Nomenclature for factors of the HLA system, 2010

Citation Report

#	ARTICLE	IF	CITATIONS
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2	Immunopathology of Organ Transplantation. , 0, , 1-24.		0
4	Nomenclature for factors of the HLA system, update February 2010. International Journal of Immunogenetics, 2010, 37, 317-321.	0.8	0
5	Nomenclature for factors of the HLA system, update January 2010. International Journal of Immunogenetics, 2010, 37, 313-316.	0.8	0
6	Sequence of a novel HLAâ€B*51 allele in a volunteer haematopoietic stem cell donor. International Journal of Immunogenetics, 2010, 37, 399-400.	0.8	3
7	Nomenclature for factors of the HLA system, update April 2010*. International Journal of Immunogenetics, 2010, 37, 519-527.	0.8	0
8	Nomenclature for factors of the HLA system, update May 2010*. International Journal of Immunogenetics, 2010, 37, 529-533.	0.8	0
9	Genomics in human renal transplantation. Current Opinion in Immunology, 2010, 22, 689-693.	2.4	1
10	Allele Name Translation Tool and Update NomenCLature: software tools for the automated translation of HLA allele names between successive nomenclatures. Tissue Antigens, 2010, 75, 457-461.	1.0	5
11	A novel HLA-B*35 variant, HLA-B* 35:02:03, identified by sequence-based typing in a bone marrow donor. Tissue Antigens, 2010, 76, no-no.	1.0	0
12	Nomenclature for factors of the HLA system, update January 2010. Tissue Antigens, 2010, 76, no-no.	1.0	4
13	Nomenclature for factors of the HLA system, update February 2010. Tissue Antigens, 2010, 76, no-no.	1.0	3
14	The novel HLA-B*15:180 allele appears to be a recombinant B*08/B*15 allele. Tissue Antigens, 2010, 76, 334-335.	1.0	3
15	Characterization of a novel HLA allele HLA-B*15:178 in a Chinese individual. Tissue Antigens, 2010, 76, 333-334.	1.0	3
16	HLA-B*18:35, a new HLA-B*18 allele identified by sequence-based typing in a Brazilian volunteer bone marrow donor. Tissue Antigens, 2010, 76, 336-337.	1.0	4
17	A new polymorphic position in exon 2 shows the novel allele HLA-DPB1*123:01. Tissue Antigens, 2010, 76, 338-339.	1.0	4
18	Identification of the novel allele HLA-B*51:84 by sequence-based typing method in a Taiwanese individual. Tissue Antigens, 2010, 76, 337-338.	1.0	3
19	Sixty-five novel alleles at the HLA-A, -B, and -DRB1 loci identified from National Marrow Donor Program volunteer donors. Tissue Antigens, 2010, 76, 319-324.	1.0	4

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20	Identification of a novel HLA-A*11:57 allele in a Chinese bone marrow donor. Tissue Antigens, 2010, 76, 332-333.	1.0	4
21	A novel HLA-A*03 allele, HLA-A*03:71. Tissue Antigens, 2010, 76, 331-331.	1.0	3
22	A new HLA-B*15 variant, B*15:144, identified by sequence-based typing in a Chinese individual. Tissue Antigens, 2010, 76, 421-422.	1.0	3
23	A novel HLA-DRB1 allele, DRB1*08:36, identified by sequence-based typing. Tissue Antigens, 2010, 76, 426-427.	1.0	7
24	Identification of a novel HLA-C allele, C*12:10:02, in a Chinese individual. Tissue Antigens, 2010, 76, 424-425.	1.0	4
25	Identification of a novel HLA-B51 allele, HLA-B*51:94. Tissue Antigens, 2010, 76, 423-423.	1.0	4
26	A novel HLA-C allele, HLA-Cw*06:20. Tissue Antigens, 2010, 76, 425-425.	1.0	3
27	A novel HLA-B allele, B*14:10, identified by haplotype-specific sequencing. Tissue Antigens, 2010, 76, 497-498.	1.0	3
28	Identification of a novel allele HLA-B*35:137 in a Chinese leukemia patient. Tissue Antigens, 2010, 76, 498-499.	1.0	6
29	A novel HLA-B allele, B*40:123, carrying a nucleotide substitution at a conserved codon. Tissue Antigens, 2010, 76, 499-500.	1.0	4
30	Identification of the novel allele HLA-B*13:01:03 by sequence-based typing method in a Taiwanese individual. Tissue Antigens, 2010, 76, 496-497.	1.0	3
31	A novel allele, HLA-A*11:01:10, identified by sequence-based typing in a Chinese individual. Tissue Antigens, 2010, 76, 495-495.	1.0	4
32	Nomenclature for factors of the HLA system, update April 2010. Tissue Antigens, 2010, 76, 501-508.	1.0	15
33	Nomenclature for factors of the HLA system, update May 2010. Tissue Antigens, 2010, 76, 509-513.	1.0	2
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42	Nomenclature for factors of the HLA system, update April 2010. Human Immunology, 2010, 71, 1043-1050.	1.2	0
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50	From Trees to the Forest: Genes to Genomics. Biology of Blood and Marrow Transplantation, 2011, 17, S52-S57.	2.0	2
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55	Nomenclature for factors of the HLA system, update November 2010. Human Immunology, 2011, 72, 370-375.	1.2	0
56	Nomenclature for factors of the HLA system, update December 2010. Human Immunology, 2011, 72, 376-377.	1.2	1
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60	Nomenclature for factors of the HLA system, update February 2011. Human Immunology, 2011, 72, 674-678.	1.2	0
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64	A â€œsilentâ€•nucleotide substitution in exon 4 is responsible for the â€œalternative expressionâ€•of HLA-A*01:01:38L through aberrant splicing. Human Immunology, 2011, 72, 717-22.	1.2	16
65	Definitions of histocompatibility typing terms: Harmonization of Histocompatibility Typing Terms Working Group. Human Immunology, 2011, 72, 1214-1216.	1.2	30
66	Nomenclature for factors of the HLA system, update April 2011. Human Immunology, 2011, 72, 979-984.	1.2	0
67	Nomenclature for factors of the HLA system, update May 2011. Human Immunology, 2011, 72, 985-989.	1.2	0
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83	Identification of a novel HLAâ€DRB1 allele, DRB1*11:95. Tissue Antigens, 2011, 77, 160-161.	1.0	4
84	Sequenceâ€based typing identification of the novel HLAâ€A*24:135 variant in a Maldivian family with a bone marrow patient. Tissue Antigens, 2011, 77, 152-153.	1.0	3
85	B*39:60, a novel HLAâ€B*39 allele identified by sequenceâ€based typing. Tissue Antigens, 2011, 77, 155-156.	1.0	8
86	Two novel HLA class I alleles, HLAâ€B*40:122 and HLAâ€B*40:127. Tissue Antigens, 2011, 77, 156-157.	1.0	4
87	Identification of a novel HLAâ€A*26 allele, A*26:36, by sequenceâ€based typing. Tissue Antigens, 2011, 77, 154-155.	1.0	0
88	The complete genomic sequence of HLAâ€DRB1*10:01:01 was identified by sequencing in a Han Chinese individual. Tissue Antigens, 2011, 77, 159-160.	1.0	5
89	Nomenclature for factors of the HLA system, update July 2010. Tissue Antigens, 2011, 77, 162-166.	1.0	0
90	Nomenclature for factors of the HLA system, update August 2010. Tissue Antigens, 2011, 77, 167-175.	1.0	3
91	Identification of nine novel HLAâ€DRB1 alleles, HLAâ€DRB1*04:91, DRB1*07:18, DRB1*11:01:12, DRB1*12:02:05, DRB1*12:22, DRB1*12:23, DRB1*13:100, DRB1*15:45, and DRB1*15:46 by polymerase chain reactionâ€"sequenceâ€based typing. Tissue Antigens, 2011, 77, 264-266.	1.0	4
92	HLAâ€C*03:93, a novel HLAâ€C*03 allele identified by sequenceâ€based typing. Tissue Antigens, 2011, 77, 266-267.	1.0	3
93	Characterization of a novel HLA allele, HLAâ€B*40:128, in a Chinese individual. Tissue Antigens, 2011, 77, 260-261.	1.0	4

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95	Identification of a novel HLA–B allele, B*56:31, by sequence–based typing in a Lithuanian individual. Tissue Antigens, 2011, 77, 262-263.	1.0	3
96	Identification of a novel HLA–B*35 variant allele, B*35:151. Tissue Antigens, 2011, 77, 258-259.	1.0	3
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98	Analysis of the complete cDNA sequences of HLA–DRB1 alleles with group–specific amplification primers in the Chinese Han population. Tissue Antigens, 2011, 77, 329-332.	1.0	10
99	HLA–DRB1*13:99, a novel HLA–DRB1*13 allele identified by sequence–based typing. Tissue Antigens, 2011, 77, 343-344.	1.0	3
100	HLA–DQB1*05:06, a novel HLA–DQB1*05 allele identified by sequence–based typing. Tissue Antigens, 2011, 77, 344-346.	1.0	4
101	Identification of a novel HLA–B*35:42:02 allele in a Chinese bone marrow donor. Tissue Antigens, 2011, 77, 341-342.	1.0	4
102	A single nomenclature and associated database for alleles at the major histocompatibility complex class II *DRB1* locus of sheep. Tissue Antigens, 2011, 77, 546-553.	1.0	16
103	*MICA*002:04*: a new allele identified by sequence–based typing in a Chinese individual. Tissue Antigens, 2011, 78, 74-76.	1.0	3
104	Identification of a novel *HLA–B*52* allele in a Brazilian individual: *B*52:21*. Tissue Antigens, 2011, 77, 342-343.	1.0	3
105	Nomenclature for factors of the HLA system, update December 2010. Tissue Antigens, 2011, 77, 362-363.	1.0	5
106	Nomenclature for factors of the HLA system, update November 2010. Tissue Antigens, 2011, 77, 355-361.	1.0	3
107	Nomenclature for factors of the HLA system, update October 2010. Tissue Antigens, 2011, 77, 349-354.	1.0	3
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110	The novel allele, *HLA–B*07:68:02*, identified in a German cord blood donor and her father. Tissue Antigens, 2011, 77, 598-599.	1.0	3
111	Two new HLA–C alleles identified in one volunteer bone marrow donor: *C*15:44* and *C*07:137:02*. Tissue Antigens, 2011, 78, 73-74.	1.0	3

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113	The new HLA‐C allele C*07:170 shows a new polymorphism at amino acid position 147. Tissue Antigens, 2011, 78, 72-73.	1.0	3
114	Identification of a novel HLA-C allele, HLA-C*04:82, by sequence-based typing in a Chinese individual. Tissue Antigens, 2011, 78, 71-72.	1.0	5
115	Eight new alleles found in Flanders (Belgium), including the <i>HLA‐DRB1*12:24N</i> allele. Tissue Antigens, 2011, 78, 45-48.	1.0	4
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117	<i>C</i>*<i>07:185</i>, a novel <i>HLA‐C</i>*<i>07</i> allele identified by sequence‐based typing. Tissue Antigens, 2011, 78, 155-156.	1.0	3
118	Identification of a novel <i>HLA‐B*40</i> null allele, <i>HLAB*40:155N</i>. Tissue Antigens, 2011, 78, 154-155.	1.0	4
119	A novel HLA‐DRB4 allele, <i>DRB4*01:08</i>, identified by sequence‐based typing. Tissue Antigens, 2011, 78, 158-159.	1.0	3
120	Characterization of three new alleles <i>HLA‐A*02:241</i>, <i>HLA‐A*02:242</i> and <i>HLA‐A*30:04:02</i>. Tissue Antigens, 2011, 78, 152-153.	1.0	3
121	The novel <i>HLA‐G*01:03:01:02</i> allele differs from <i>G*01:03:01:01</i> by a possible inversion event in intron 3. Tissue Antigens, 2011, 78, 159-160.	1.0	5
122	Identification of a new HLA-B*08 allele in a family of Iraqi origin. Tissue Antigens, 2011, 78, 153-154.	1.0	3
123	Characterization of a novel HLA allele <i>HLA‐B*15:25:03</i> in a Chinese individual. Tissue Antigens, 2011, 78, 219-220.	1.0	0
124	Characterization of new HLA‐A and ‐B alleles from Kazakhstan. Tissue Antigens, 2011, 78, 217-218.	1.0	4
125	Identification of a novel <i>HLA‐DRB1*04:94N</i> allele by polymerase chain reaction sequence‐based typing. Tissue Antigens, 2011, 78, 226-227.	1.0	5
126	Identification of the novel HLA‐B allele <i>B*57:29</i> by cloning and sequencing in a Chinese individual. Tissue Antigens, 2011, 78, 221-222.	1.0	3
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129	Nomenclature for factors of the HLA system, update February 2011. Tissue Antigens, 2011, 77, 613-617.	1.0	0

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132	Identification of a novel HLAâ€DPB1 allele, <i>DPB1*131:01</i>, by sequenceâ€based typing. Tissue Antigens, 2011, 78, 222-223.	1.0	5
133	Nomenclature for factors of the HLA system, update March 2011. Tissue Antigens, 2011, 78, 77-79.	1.0	2
134	Two novel HLA-A alleles: HLA-A*31:01:09 and HLA-A*33:30. Tissue Antigens, 2011, 78, 218-219.	1.0	4
135	Identification of a novel allele <i>HLAâ€DRB1*08:41</i> in a Chinese donor. Tissue Antigens, 2011, 78, 294-295.	1.0	4
136	Sequencingâ€based typing identifies three new HLA alleles: <i>C*02:29</i>, <i>C*06:29</i> and <i>DQB1*03:24</i>. Tissue Antigens, 2011, 78, 290-291.	1.0	4
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138	<i>HLAâ€B*51:79</i> is a novel allele associated with a conserved haplotype. Tissue Antigens, 2011, 78, 288-289.	1.0	3
139	Characterization and polymorphic analysis of 4.5 kb genomic fullâ€length HLAâ€C in the Chinese Han population. Tissue Antigens, 2011, 78, 102-114.	1.0	9
140	Identification of the new HLA-A*31:48 allele in an Italian patient. Tissue Antigens, 2011, 78, 286-287.	1.0	3
141	One amino acid change located in the conserved region of the alpha 1 domain specifies the novel <i>HLAâ€C*07:147</i> allele. Tissue Antigens, 2011, 78, 292-294.	1.0	3
142	A new <i>HLAâ€B*15</i> allele, <i>B*15:220</i>, found in three individuals sharing the <i>HLAâ€A*66:01</i>, <i>HLAâ€C*12:03</i> and <i>HLADRB1*07:01</i> alleles. Tissue Antigens, 2011, 78, 287-288.	1.0	4
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144	A novel HLA allele derived from a likely DRB1/DRB3 gene conversion event: <i>HLAâ€DRB3*01:15</i>. Tissue Antigens, 2011, 78, 405-407.	1.0	4
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146	A novel HLAâ€A allele detected by sequenceâ€based typing: <i>A*68:66</i>. Tissue Antigens, 2011, 78, 397-397.	1.0	1
147	Sequencing of a single HLAâ€B genotype including two rare alleles allows the detection of a new allele, <i>B*44:130</i>. Tissue Antigens, 2011, 78, 398-399.	1.0	3

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149	Identification of a novel HLA-B allele, B*35:01:25, by sequence-based typing in a Korean individual. Tissue Antigens, 2011, 78, 400-401.	1.0	3
150	A novel <i>HLA‑B*48</i> allele, <i>B*48:01:03</i>, identified by sequence‑based typing. Tissue Antigens, 2011, 78, 401-402.	1.0	3
151	Novel HLA class I and II alleles identified during routine registry typing in 2010. Tissue Antigens, 2011, 78, 263-266.	1.0	4
152	Sequence‑based HLA high‑resolution typing of a bone marrow donor/recipient pair shows the novel HLA allele <i>DQB1*06:43</i>. Tissue Antigens, 2011, 78, 461-462.	1.0	4
153	Characterization of 236 novel alleles at the HLA-A, -B, -C, -DRB1, -DQB1 and -DPB1 loci from China Marrow Donor Program. Tissue Antigens, 2011, 78, 267-270.	1.0	9
154	Identification of a new <i>HLA‑DRB1* 14</i> allele, <i>DRB1*14:99</i>, by HLA typing of an Asian individual. Tissue Antigens, 2011, 78, 466-467.	1.0	3
155	Identification of a novel <i>HLA‑C*02</i> allele, <i>HLA‑C*02:02:09</i>. Tissue Antigens, 2011, 78, 456-457.	1.0	3
156	Identification of a novel <i>HLA‑C*07:02:25</i> allele by polymerase chain reaction sequence‑based typing in a Chinese leukemia patient. Tissue Antigens, 2011, 78, 457-459.	1.0	4
157	Identification of a novel <i>HLA‑DRB1</i> allele, <i>HLA‑DRB1*12:27</i>, in a Chinese individual. Tissue Antigens, 2011, 78, 465-466.	1.0	6
158	Sequencing of a new <i>HLA‑DRB1*04:98</i> allele in a Spanish donor. Tissue Antigens, 2011, 78, 462-463.	1.0	3
159	Identification of a novel <i>HLA‑A</i> allele, <i>A*29:28</i>, in an East African population. Tissue Antigens, 2011, 78, 451-452.	1.0	5
160	A single nucleotide deletion in exon 2 produces a novel null allele, <i>HLA‑DRB1*01:33N</i>. Tissue Antigens, 2011, 78, 463-464.	1.0	4
161	Identification of a novel <i>HLA‑B*40</i> allele, <i>B*40:162</i>, in a Chinese individual. Tissue Antigens, 2011, 78, 452-454.	1.0	3
162	A new <i>HLA‑B*52</i> allele, <i>B*52:23</i>, detected in a patient before bone marrow transplantation. Tissue Antigens, 2011, 78, 455-456.	1.0	4
163	A novel HLA-B44 allele, B*44:127, was identified by sequencing-based typing. Tissue Antigens, 2011, 78, 454-455.	1.0	4
164	Nomenclature for factors of the HLA system, update April 2011. Tissue Antigens, 2011, 78, 299-306.	1.0	0
165	Nomenclature for factors of the HLA system, update May 2011. Tissue Antigens, 2011, 78, 307-313.	1.0	4

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167	Genetic variation and hitchhiking between structurally polymorphic Alu insertions and HLAâ€A, â€B, and â€C alleles and other retroelements within the MHC class I region. Tissue Antigens, 2011, 78, 359-377.	1.0	19
168	A community standard for immunogenomic data reporting and analysis: proposal for a STrengthening the REporting of Immunogenomic Studies statement. Tissue Antigens, 2011, 78, 333-344.	1.0	50
169	Nomenclature for factors of the HLA system, update July 2011. Tissue Antigens, 2011, 78, 468-473.	1.0	1
170	Nomenclature for factors of the HLA system, update August 2011. Tissue Antigens, 2011, 78, 474-481.	1.0	1
171	Nomenclature for factors of the HLA system, update September 2011. Tissue Antigens, 2011, 78, 482-484.	1.0	1
172	Nomenclature for factors of the HLA system, update June 2010*. International Journal of Immunogenetics, 2011, 38, 77-81.	0.8	0
173	The HLA-B*83:01 allele is generated by a gene conversion event including whole of exon 2 and partial introns 1 and 2 between B*44 and B*56 alleles. International Journal of Immunogenetics, 2011, 38, 73-75.	0.8	3
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338	Analysis of HLAâ€DM polymorphisms in the Chinese Han population. Tissue Antigens, 2012, 79, 157-164.	1.0	6
339	Nomenclature for factors of the HLA system, update October 2011. Tissue Antigens, 2012, 79, 218-221.	1.0	0
340	Nomenclature for factors of the HLA system, update November 2011. Tissue Antigens, 2012, 79, 222-229.	1.0	1
341	Nomenclature for factors of the HLA system, update December 2011. Tissue Antigens, 2012, 79, 230-234.	1.0	2
342	Characterization of a new allele: *HLAâ€A*03:134*. Tissue Antigens, 2012, 79, 309-310.	1.0	6
343	HLA-A*02:07:02, a new allele identified by sequence-based typing in a Korean individual. Tissue Antigens, 2012, 79, 384-385.	1.0	8
344	A novel *HLAâ€A*31* allele, *A*31:57*, identified by sequenceâ€based typing. Tissue Antigens, 2012, 79, 386-387.	1.0	4
345	Nomenclature for factors of the HLA system, update January 2012. Tissue Antigens, 2012, 79, 393-397.	1.0	8
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350	Characterization of the novel HLA-C*07:195 allele in an Italian hematopoietic stem cell volunteer donor, detected by sequence-based typing. Tissue Antigens, 2012, 80, 67-68.	1.0	3
351	Detection of a new *HLAâ€B*15* allele, *HLAâ€B*15:238*, in a voluntary stem cell donor. Tissue Antigens, 2012, 80, 66-67.	1.0	3
352	Description of *HLAâ€A*33:49* in a Spanish cord blood unit. Tissue Antigens, 2012, 80, 65-66.	1.0	3
353	Nomenclature for factors of the HLA system, update February 2012. Tissue Antigens, 2012, 80, 72-77.	1.0	4
354	Nomenclature for factors of the HLA system, update March 2012. Tissue Antigens, 2012, 80, 202-204.	1.0	0
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357	Identification of a novel *HLAâ€A*24:02:55* allele by sequenceâ€based typing. Tissue Antigens, 2012, 80, 193-194.	1.0	3
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359	Identification of a novel HLAâ€DPB1 allele, *DPB1*138:01*, by sequenceâ€based typing. Tissue Antigens, 2012, 80, 195-196.	1.0	5
360	Characterization of the novel *HLAâ€DPB1*139:01* allele identified by sequenceâ€based typing in a Chinese Han individual. Tissue Antigens, 2012, 80, 197-198.	1.0	4
361	Detection of a novel HLAâ€A allele, designated *A*02:334*. Tissue Antigens, 2012, 80, 191-191.	1.0	3
362	Identification of a novel HLA-DQB1*03:38 allele by polymerase chain reaction sequence-based typing in a Chinese bone marrow donor. Tissue Antigens, 2012, 80, 198-199.	1.0	8
363	Characterization of three new HLAâ€B alleles: *B*35:05:03, B*52:29* and *B*57:01:13*. Tissue Antigens, 2012, 80, 270-271.	1.0	3
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370	Identification of a novel HLA-C*01:61 allele by polymerase chain reaction sequence-based typing in a Chinese leukemia patient. Tissue Antigens, 2012, 80, 275-276.	1.0	4
371	Heterogeneity of <i><scp>HLAâ€‹DRB1</scp>*04</i> alleles and haplotypes in the Croatian population. Tissue Antigens, 2012, 80, 219-223.	1.0	2
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375	A newHLA-B*40:186allele identified by sequence-based typing in a Korean individual. Tissue Antigens, 2012, 80, 383-384.	1.0	3
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378	Discovery of the novel <scp>HLA</scp>â€‹<scp>DRB</scp>1*09:01:08 allele in a <scp>T</scp>aiwanese volunteer bone marrow donor and identification of the probable <scp>HLA</scp>â€‹<scp>A</scp>,â€‹<scp>HLA</scp>â€‹<scp>B</scp> and <scp>HLA</scp>â€‹<scp>DRB</scp>1 haplotype in association with <scp>DRB</scp>1*09:01:08. International Journal of Immunogenetics, 2013, 40, 149-150.	0.8	22
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482	Sequencing of a novelHLA-DRB1*09:01:07allele in a Chinese individual. Tissue Antigens, 2013, 82, 151-152.	1.0	4
483	The novel allele <i><scp>DRB1</scp>*07:23</i> was revealed by LABType HD DRB1 typing test and confirmed by sequenceâ€based <scp>HLA</scp> typing. Tissue Antigens, 2013, 82, 360-361.	1.0	4
484	A novelHLA-Aallele,A*31:65,was identified by sequence-based typing in a Chinese potential donor. Tissue Antigens, 2013, 81, 173-175.	1.0	3
485	Identification of a novel <scp>HLA</scp>â€A allele, <i>A*02:357</i>, in a Chinese individual. Tissue Antigens, 2013, 81, 171-172.	1.0	3
486	Sequenceâ€based typing identified a new <i><scp>HLA</scp>â€B*40</i> allele, <i><scp>HLA</scp>â€B*40:124:02</i>. Tissue Antigens, 2013, 81, 464-465.	1.0	4
487	Identification of the novel <i><scp>HLA</scp>â€A*24:233</i> allele in a Chinese individual. Tissue Antigens, 2013, 82, 424-425.	1.0	4
488	Characterization of the novel <i><scp>HLAâ€DRB1</scp>*13:26:02</i> allele. Tissue Antigens, 2013, 82, 152-153.	1.0	3
489	Advancing allele groupâ€specific amplification of the complete <i><scp>HLA</scp>â€C</i> geneâ€"isolation of novel alleles from three allele groups (<i>C*04</i>, <i>C*07</i> and <i>C*08</i>). Tissue Antigens, 2013, 82, 280-285.	1.0	2
490	The novel allele, <i><scp>HLAâ€DQB1</scp>*02:27</i>, identified in a German cord blood donor and his mother. Tissue Antigens, 2013, 82, 440-441.	1.0	4
491	Characterization of a novel variant allele <scp><i>HLAâ€DPA1*02:02:05</i></scp>, identified in a Chinese Han individual. Tissue Antigens, 2013, 81, 234-236.	1.0	5
492	Identification of a novelHLA-A*33:03:11allele by polymerase chain reaction sequence-based typing in a Chinese cord blood donor. Tissue Antigens, 2013, 82, 59-60.	1.0	8

#	ARTICLE	IF	CITATIONS
493	Nomenclature for factors of the ‹scp›HLA‹/scp› system, update November 2012. Tissue Antigens, 2013, 81, 249-252.	1.0	1
494	Identification of a new ‹scp›HLA‹/scp›â€‹ allele, ‹i›‹scp›HLA‹/scp›â€‹C*08:76‹/i› in a Caucasian individual. Tissue Antigens, 2013, 82, 69-70.	1.0	3
495	Identification of a novelHLA-B*40allele,HLA-B*40:211, in a Chinese individual. Tissue Antigens, 2013, 82, 207-207.	1.0	3
496	A novel ‹scp›HLA‹/scp› allele, ‹i›‹scp›HLA‹/scp›â€‹B*39:46‹/i›, identified by sequenceâ€‹based typing in a Chinese volunteer bone marrow donor. Tissue Antigens, 2013, 81, 462-463.	1.0	3
497	Identification of a novel ‹scp›HLA‹/scp›â€‹B allele, ‹i›B*46:39‹/i›, by sequenceâ€‹based typing. Tissue Antigens, 2013, 82, 351-352.	1.0	3
498	Identification of a new HLA allele,HLA-DRB1*03:80, in a Chinese Han ethnic individual. Tissue Antigens, 2013, 81, 236-237.	1.0	5
499	Nomenclature for factors of the ‹scp›HLA‹/scp› system, update August 2013. International Journal of Immunogenetics, 2013, 40, 533-537.	0.8	0
500	Nomenclature for factors of the ‹scp›HLA‹/scp› system, update ‹scp›S‹/scp›eptember 2012. International Journal of Immunogenetics, 2013, 40, 165-170.	0.8	1
501	A novel ‹i›HLAâ€‹DQB1*04‹/i› allele, ‹i›DQB1*04:10‹/i›, identified in a Japanese individual. Tissue Antigens, 2013, 82, 148-149.	1.0	4
502	A novel ‹scp›HLA‹/scp› allele, ‹i›‹scp›HLA‹/scp›â€‹A*80:03‹/i›, identified in a Brazilian individual. Tissue Antigens, 2013, 82, 349-350.	1.0	4
503	Nomenclature for factors of the ‹scp›HLA‹/scp› system, update ‹scp›M‹/scp›arch 2013. International Journal of Immunogenetics, 2013, 40, 343-347.	0.8	0
504	Characterization of the novel ‹scp›HLA‹/scp›â€‹‹scp›DQB‹/scp›1*06:48 allele by groupâ€‹specific sequencing. International Journal of Immunogenetics, 2013, 40, 322-323.	0.8	4
505	Nomenclature for factors of the ‹scp›HLA‹/scp› system, update April 2013. Tissue Antigens, 2013, 82, 219-223.	1.0	7
506	Characterization of the novelHLA-DPA1*02:05allele identified by sequence-based typing. Tissue Antigens, 2013, 82, n/a-n/a.	1.0	4
507	The novel ‹i›‹scp›HLA‹/scp›â€‹B*44:02:27‹/i› allele, identified by sequencingâ€‹based typing of a candidate stemâ€‹cell donor. Tissue Antigens, 2013, 81, 230-231.	1.0	4
508	Characterisation of a new ‹scp›HLAâ€‹DRB1‹/scp› allele: ‹i›‹scp›DRB1‹/scp›*03:85‹/i›. Tissue Antigens, 2013, 82, 358-359.	1.0	4
509	Nomenclature for factors of the ‹scp›HLA‹/scp› system, update July 2013. Tissue Antigens, 2013, 82, 447-451.	1.0	1
510	A novel HLA allele,HLA-B*40:227, was identified by polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2013, 82, 208-209.	1.0	4

#	Article	IF	Citations
511	A novel <scp>MICA</scp> allele, <i>MICA*069</i>, identified by sequenceâ€based typing in a Chinese individual. Tissue Antigens, 2013, 81, 240-241.	1.0	6
512	A novel HLAâ€A allele detected by sequenceâ€based typing: <i>HLAâ€A*24:223</i>. Tissue Antigens, 2013, 82, 139-140.	1.0	3
513	Identification of a novel <scp>HLA</scp>â€B allele <i><scp>HLA</scp>â€B*07:185</i> in a Japanese individual. Tissue Antigens, 2013, 82, 434-436.	1.0	3
514	Identification of a new <scp>HLA</scp>â€A null allele, <i>A*30:<scp>59N</scp></i>, with a stop codon in exon 3, by sequenceâ€based typing. Tissue Antigens, 2013, 82, 430-432.	1.0	3
515	Identification of two novelHLA-B*54alleles,B*54:01:03andB*54:01:04by polymerase chain reaction sequence-based typing. Tissue Antigens, 2013, 82, 63-65.	1.0	7
516	Characterization of a novel allele,HLA-DQB1*06:47. Tissue Antigens, 2013, 82, 74-75.	1.0	8
517	Characterization of 27 novel <scp>HLA</scp> class <scp>II</scp> alleles from China Marrow Donor Program. Tissue Antigens, 2013, 82, 201-202.	1.0	5
518	Identification of three novel human leukocyte antigen alleles,HLA-B*58:43,HLA-C*03:190, andHLA-DPA1*01:12, in an East African cohort. Tissue Antigens, 2013, 82, 131-133.	1.0	4
519	The novel allele <i><scp>HLA</scp>â€C*03:04:36</i> identified by sequenceâ€based typing. Tissue Antigens, 2013, 82, 353-354.	1.0	3
520	Nomenclature for factors of the <scp>HLA</scp> system, update <scp>N</scp>ovember 2012. International Journal of Immunogenetics, 2013, 40, 252-255.	0.8	0
521	A novel <scp>HLA</scp> allele, <i><scp>HLA</scp>â€A*02:393</i>, identified in a two generation Italian family. Tissue Antigens, 2013, 82, 203-204.	1.0	3
522	Discovery of the rare <scp>HLA</scp>â€B*39:77 allele in an unrelated <scp>T</scp>aiwanese bone marrow stem cell donor using the sequenceâ€based typing method. International Journal of Immunogenetics, 2013, 40, 331-334.	0.8	0
523	Nomenclature for factors of the <scp>HLA</scp> system, update August 2012. International Journal of Immunogenetics, 2013, 40, 161-164.	0.8	1
524	A single nucleotide insertion in exon 2 produces a novel <i><scp>HLA</scp>â€B*13</i> null allele, <i><scp>HLA</scp>â€B*13:<scp>63N</scp></i>. Tissue Antigens, 2013, 81, 459-460.	1.0	4
525	Nomenclature for factors of the <scp>HLA</scp> system, update May 2013. Tissue Antigens, 2013, 82, 224-228.	1.0	0
526	Nomenclature for factors of the <scp>HLA</scp> system, update July 2013. International Journal of Immunogenetics, 2013, 40, 528-532.	0.8	0
527	Identification of the novel <i><scp>HLA</scp>â€B*50:18</i> allele variant in an Italian unrelated bone marrow donor. Tissue Antigens, 2013, 81, 466-468.	1.0	3
528	A novel <scp>HLA</scp>â€C allele, <i><scp>HLA</scp>â€C*06:45</i>, identified by sequenceâ€based typing in a Chinese individual. Tissue Antigens, 2013, 82, 67-68.	1.0	4

#	ARTICLE	IF	CITATIONS
529	The novel <i><scp>HLA</scp>â€C*12:92</i> allele is characterized by one amino acid exchange located in the Tâ€cell receptor binding region of the alpha 2 domain. Tissue Antigens, 2013, 82, 355-356.	1.0	3
530	The novelHLA-A*24:215allele, identified by sequencing-based typing of a stem cell transplant patient and the sibling donor. Tissue Antigens, 2013, 82, 138-139.	1.0	3
531	A novel <scp>HLA</scp>â€B allele, B*58:01:12, detected in a Taiwanese volunteer bone marrow stem cell donor using sequenceâ€based typing method. International Journal of Immunogenetics, 2013, 40, 324-325.	0.8	3
532	Identification of the novelHLA-C*07:315allele. Tissue Antigens, 2013, 82, 354-355.	1.0	3
533	The new <scp>HLA</scp>â€C variant <i><scp>HLA</scp>â€C*05:26</i> is likely to be structurally identical to the <i>C*05:<scp>01P</scp></i> alleles. Tissue Antigens, 2013, 81, 179-180.	1.0	4
534	A novel <i><scp>HLA</scp>â€B*13</i> allele, <i>B*13:68</i>, was identified by sequencingâ€based typing. Tissue Antigens, 2013, 82, 204-205.	1.0	4
535	<i><scp>HLA</scp>â€B*40:06:06</i>, a novel allele, which has arisen by silent mutation in codon 45. Tissue Antigens, 2013, 82, 142-144.	1.0	3
536	Detection of a new <scp>HLA</scp>â€A allele, designated <i><scp>HLA</scp>â€A*32:53</i>. Tissue Antigens, 2013, 82, 348-348.	1.0	3
537	Identification of the novel <scp>HLA</scp>â€B*40:221 allele in a Taiwanese hematopoietic stem cell donor using a sequenceâ€based typing method. International Journal of Immunogenetics, 2013, 40, 320-321.	0.8	4
538	Nomenclature for factors of the <scp>HLA</scp> system, update March 2013. Tissue Antigens, 2013, 81, 480-484.	1.0	5
539	Identification of a novel <scp>HLAâ€DQB1</scp> allele, <i><scp>DQB1</scp>*05:19</i>, in an African American family by sequenceâ€based typing. Tissue Antigens, 2013, 82, 150-151.	1.0	4
540	Identification of the new HLA-DPB1 allele,DPB1*162:01, in a Venezuelan family. Tissue Antigens, 2013, 82, 442-443.	1.0	3
541	Nomenclature for factors of the HLA system, update June 2013. Tissue Antigens, 2013, 82, 229-233.	1.0	5
542	Detection of the newHLA-DRB1*14:129allele in a voluntary stem cell donor. Tissue Antigens, 2013, 81, 229-230.	1.0	4
543	Novel <i><scp>HLA</scp>â€C*07:314</i> allele identified by sequenceâ€based typing in a French lymphoblastic leukemia patient. Tissue Antigens, 2013, 82, 439-440.	1.0	3
544	The novel <i><scp>HLA</scp>â€A*26:89</i> allele identified by sequenceâ€based typing. Tissue Antigens, 2013, 82, 427-428.	1.0	4
545	Genomic fullâ€length sequence of two <scp>HLA</scp>â€A alleles, <i>A*30:01:01</i> and <i>A*30:04:01</i>, identified by cloning and sequencing. Tissue Antigens, 2013, 82, 295-296.	1.0	3
546	A novel HLA allele,HLA-C*01:02:18, was identified by polymerase chain reaction sequence-based typing in a Chinese leukemia patient. Tissue Antigens, 2013, 82, 65-66.	1.0	7

#	ARTICLE	IF	CITATIONS
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548	HLA-DRB1*03:49, a novel allele identified by group-specific sequence-based typing in a north European individual. Tissue Antigens, 2013, 82, 357-358.	1.0	0
549	Characterization of a novelHLA-C*07allele,HLA-C*07:56:01. Tissue Antigens, 2013, 81, 181-182.	1.0	3
550	HLA-B*58:45, a variant ofHLA-B*58, found in a Taiwanese unrelated hematopoietic stem cell donor. Tissue Antigens, 2013, 82, 438-439.	1.0	3
551	A novel allele, <i><scp>HLA</scp>â€A*03:01:09</i>, identified by sequenceâ€based typing in a Chinese individual. Tissue Antigens, 2013, 81, 452-453.	1.0	3
552	A novel HLA-B allele,B*40:01:06, identified by sequence-based typing in a Chinese individual. Tissue Antigens, 2013, 81, 176-177.	1.0	3
553	Identification of a new <i><scp>HLA</scp>â€B*46</i> allele, <i>B*46:37</i>, in a Chinese individual. Tissue Antigens, 2013, 81, 465-466.	1.0	3
554	Sequencing of a novel <scp>HLA</scp>â€B allele, <i>B*51:153</i>, in a Spanish individual. Tissue Antigens, 2013, 82, 297-297.	1.0	3
555	Two novel <scp>HLA</scp> class <scp>II</scp> alleles, <i><scp>DRB1</scp>*11:131</i> and <i><scp>DQB1</scp>*05:01:05</i>, identified by sequenceâ€based typing. Tissue Antigens, 2013, 82, 299-300.	1.0	7
556	Sequencing of the novel <i><scp>HLA</scp>â€B*49:24</i> and <i><scp>HLAâ€DRB1</scp>*03:64</i> alleles. Tissue Antigens, 2013, 81, 177-178.	1.0	5
557	Identification of a novelHLA-DRB1*11:129allele in a Chinese individual by sequence-based typing. Tissue Antigens, 2013, 81, 237-239.	1.0	5
558	Identification of a newHLA-DQB1*03variant,DQB1*03:39, by sequence-based typing in a Chinese Han individual. Tissue Antigens, 2013, 82, 213-214.	1.0	4
559	A novel allele: <i><scp>HLA</scp>â€B*13:01:06</i>, identified by sequenceâ€based typing in a Chinese individual. Tissue Antigens, 2013, 81, 457-459.	1.0	3
560	Genotype List String: a grammar for describing <scp>HLA</scp> and <scp>KIR</scp> genotyping results in a text string. Tissue Antigens, 2013, 82, 106-112.	1.0	56
561	A novel <scp>HLA</scp>â€A allele, <i>A*24:191</i>, was identified by sequenceâ€based typing in a Chinese donor. Tissue Antigens, 2013, 82, 423-424.	1.0	4
562	Nomenclature for factors of the <scp>HLA</scp> system, update <scp>M</scp>ay 2013. International Journal of Immunogenetics, 2013, 40, 429-433.	0.8	2
563	Identification of the novelHLA-B*15:257allele by polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2013, 82, 62-63.	1.0	7
564	Identification of a novel allele <i>MICA</i>*<i>008:05</i> by sequenceâ€based typing. Tissue Antigens, 2013, 82, 217-218.	1.0	3

#	Article	IF	Citations
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566	The novelHLA-Aallele,A*26:74, identified by sequence-based typing in a Chinese individual. Tissue Antigens, 2013, 82, 426-427.	1.0	3
567	Nomenclature for factors of the <scp>HLA</scp> system, update January 2013. Tissue Antigens, 2013, 81, 471-473.	1.0	0
568	Nomenclature for factors of the <scp>HLA</scp> system, update <scp>A</scp>pril 2013. International Journal of Immunogenetics, 2013, 40, 424-428.	0.8	0
569	Characterization of the novelHLA-DPA1*02:02:04allele. Tissue Antigens, 2013, 82, 77-78.	1.0	4
570	An Analysis of Characteristics of Subjects Examined for Incretin Effects on Pancreatic Pathology. Diabetes Technology and Therapeutics, 2013, 15, 609-618.	2.4	27
571	Anti–"IFN-Î³ autoantibodies in adults with disseminated nontuberculous mycobacterial infections are associated with HLA-DRB1*16:02 and HLA-DQB1*05:02 and the reactivation of latent varicella-zoster virus infection. Blood, 2013, 121, 1357-1366.	0.6	145
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577	HLA-E, HLA-F and HLA-G â€" The Non-Classical Side of the MHC Cluster. , 2014, , .		22
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580	Histocompatibility in Kidney Transplantation. , 2014, , 142-160.		0
581	The <i>HLAâ€A*02:481</i> allele was identified in unrelated Brazilians sharing <i>HLAâ€B*15:17</i>, <i>C*07:01P</i>, <i>DRB1*13:02</i> and <i>DQB1*06:04</i>. Tissue Antigens, 2014, 84, 577-578.	1.0	3
582	<i><scp>HLA</scp>â€A*02:01:79</i>, a novel allele, which has arisen by silent mutation in codon 230. Tissue Antigens, 2014, 84, 323-324.	1.0	3

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585	Description of the novel <i><scp>HLAâ€DPB1</scp>*166:01</i> allele identified by sequenceâ€based typing. Tissue Antigens, 2014, 83, 298-299.	1.0	4
586	A new <i><scp>HLA</scp>â€A*24</i> allele, <i><scp>HLA</scp>â€A*24:02:87</i>, identified by sequencingâ€based typing in a Chinese volunteer bone marrow donor. Tissue Antigens, 2014, 84, 413-414.	1.0	3
587	<i><scp>HLA</scp>â€B*40:02:18</i>, a new <i><scp>HLA</scp>â€B*40</i> allele. Tissue Antigens, 2014, 83, 429-430.	1.0	3
588	Nomenclature for factors of the <scp>HLA</scp> system, update June 2014. International Journal of Immunogenetics, 2014, 41, 452-455.	0.8	0
589	Two novel alleles <scp>HLA</scp>â€A*02:433 and <scp>HLA</scp>â€A*02:434 identified in Saudi bone marrow donors using sequenceâ€based typing. International Journal of Immunogenetics, 2014, 41, 338-339.	0.8	6
590	A new <scp>HLA</scp> allele, <i><scp>HLA</scp>â€B*08:108</i>, described in two unrelated Spanish individuals. Tissue Antigens, 2014, 83, 130-131.	1.0	3
591	Nomenclature for factors of the <scp>HLA</scp> system, update May 2014. Tissue Antigens, 2014, 84, 335-341.	1.0	0
592	Nomenclature for factors of the <scp>HLA</scp> system, update January 2014. Tissue Antigens, 2014, 83, 435-438.	1.0	0
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594	A novel <scp>HLA</scp>â€C allele, <i>C*08:01:10</i> was identified in a Chinese leukemia patient. Tissue Antigens, 2014, 84, 419-420.	1.0	4
595	External quality assessment of patient <scp>HLA</scp>â€B*57:01 testing prior to abacavir prescription. International Journal of Immunogenetics, 2014, 41, 277-280.	0.8	9
596	A new nonâ€expressed allele <i><scp>HLA</scp>â€A*03:<scp>168N</scp></i>, identified by serological and sequenceâ€based typing in a voluntary Norwegian hematopoietic stem cell donor. Tissue Antigens, 2014, 83, 195-196.	1.0	3
597	Nomenclature for factors of the <scp>HLA</scp> system, update May 2014. International Journal of Immunogenetics, 2014, 41, 445-451.	0.8	0
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599	Variation and linkage disequilibrium between a structurally polymorphic <i>Alu</i> located near the <i><scp>OR</scp>12D2</i> gene of the extended major histocompatibility complex class I region and <scp>HLA</scp>â€A alleles. International Journal of Immunogenetics, 2014, 41, 250-261.	0.8	2
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603	Nomenclature for factors of the HLA system, update September 2014. Human Immunology, 2014, 75, 1284-1287.	1.2	0
604	A novel HLA-C null allele,HLA-C*05:99N. Tissue Antigens, 2014, 84, 420-421.	1.0	3
605	Identification of a novel <i><scp>HLA</scp>â€B*15</i> variant allele, <i><scp>HLA</scp>â€B*15:276</i>, in an umbilical cord blood donor. Tissue Antigens, 2014, 83, 428-429.	1.0	3
606	Identification of a novel <scp>HLAâ€DPB1</scp> allele, <i><scp>HLAâ€DPB1</scp>*167:01</i>, in a Chinese individual. Tissue Antigens, 2014, 83, 299-300.	1.0	4
607	Three novelHLA-A*24alleles identified in Chinese Han individuals. Tissue Antigens, 2014, 83, 427-428.	1.0	4
608	Description and molecular modeling of four novel <scp>HLA</scp>â€B alleles identified in Brazilian individuals. Tissue Antigens, 2014, 83, 55-57.	1.0	5
609	Detection of a novel <scp>HLA</scp>â€B allele, <i><scp>HLA</scp>â€B*08:111</i>, in an Italian bone marrow donor. Tissue Antigens, 2014, 83, 57-59.	1.0	3
610	Sequencing of a new <scp>HLA</scp>â€A allele: <i><scp>HLA</scp>â€A*11:01:54</i>. Tissue Antigens, 2014, 83, 360-361.	1.0	3
611	The novel allele, <i>HLAâ€DPB1*04:01:15</i>, is shared by a cord blood unit and corresponding maternal sample. Tissue Antigens, 2014, 83, 201-203.	1.0	4
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770	Two novel <scp>HLA</scp> alleles, <i><scp>HLA</scp>â€A*30:02:01:03</i> and <i><scp>HLA</scp>â€C*08:113</i>, identified in a South African bone marrow donor. Tissue Antigens, 2015, 85, 291-293.	1.0	6
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772	Characterization of a new HLA allele: <i>A*02:548</i>. Tissue Antigens, 2015, 86, 55-56.	1.0	3
773	A novel HLA allele,HLA-DRB1*15:116, discovered in a Taiwanese unrelated hematopoietic stem cell donor. Tissue Antigens, 2015, 86, 149-150.	1.0	4
774	Three new HLAâ€DQB1 alleles â€" <i>DQB1</i>*<i>03</i>:<i>113</i>, <i>DQB1</i>*<i>06</i>:<i>02</i>:<i>15</i> and <i>DQB1</i>*<i>06</i>:<i>129</i>. Tissue Antigens, 2015, 86, 216-217.	1.0	4
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906	Two novel <scp>HLA</scp>â€A alleles: <i>A*24:258</i> and <i>A*24:305</i> were identified in Chinese individuals. Hla, 2016, 87, 173-174.	0.4	4
907	<i><scp>HLA</scp>â€A*11:231</i>, a novel variant of <i><scp>HLA</scp>â€A*11</i>, found in a Taiwanese hematopoietic stem cell donor. Hla, 2016, 87, 170-171.	0.4	3
908	HLA-A*24:334, a novel variant ofHLA-A*24, discovered in a Taiwanese individual. Hla, 2016, 87, 390-391.	0.4	2
909	Discovery of <i><scp>HLA</scp>â€B*58:77</i>, a novel <i><scp>HLA</scp>â€B*58</i> variant, in a Taiwanese blood donor. Hla, 2016, 87, 398-399.	0.4	2
910	HLA-B*40:167, a novel allele, identified in a Chinese individual. Hla, 2016, 87, 392-394.	0.4	2

#	ARTICLE	IF	CITATIONS
911	Identification of the novel <i><scp>HLA</scp>â€‹B*52:42</i> allele by polymerase chain reaction sequenceâ€‹based typing in a Chinese bone marrow donor. Hla, 2016, 87, 400-402.	0.4	4
912	A novel <i><scp>HLA</scp>â€‹A*02</i> allele, <i>A*02:436</i>, was identified by sequencingâ€‹based typing. Hla, 2016, 87, 453-454.	0.4	3
913	HLA-A*24:333, a novelHLA-A*24variant, discovered in a Taiwanese individual. Hla, 2016, 87, 458-459.	0.4	3
914	HLAâ€‹B allele dropout in PCR sequenceâ€‹specific oligonucleotide probe typing due to intronic polymorphism in the novel B*58:01:01:02 allele. International Journal of Immunogenetics, 2016, 43, 180-183.	0.8	5
915	Nomenclature for factors of the HLA system, update October 2015. International Journal of Immunogenetics, 2016, 43, 111-115.	0.8	0
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917	A novel <scp>HLA</scp>â€‹B allele, <i><scp>HLA</scp>â€‹B*35:269</i>. Hla, 2016, 87, 107-109.	0.4	3
918	Identification of a new HLAâ€‹B allele, <i>HLAâ€‹B*15:300</i>, in a Chinese individual. Hla, 2016, 87, 175-176.	0.4	5
919	Discovery of <i><scp>HLA</scp>â€‹A*11:01:69</i>, a novel <i><scp>HLA</scp>â€‹A*11:01</i> variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2016, 87, 166-167.	0.4	3
920	A new <scp>HLA</scp>â€‹A allele, <i><scp>HLA</scp>â€‹A*11:120</i>, sequenced in a Chinese hematopoietic stem cell donor. Hla, 2016, 87, 168-170.	0.4	4
921	Discovery ofHLA-B*55:02:10, a novelHLA-B*55:02variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2016, 87, 182-183.	0.4	5
922	<i><scp>HLA</scp>â€‹B*40:01:44</i>, a novel variant of <i><scp>HLA</scp>â€‹B*40:01</i>, discovered in a Taiwanese hematopoietic stem cell donor. Hla, 2016, 87, 179-180.	0.4	3
923	HLA-B*40:326,a novel variant ofHLA-B*40, discovered in a Taiwanese blood donor. Hla, 2016, 87, 394-395.	0.4	2
924	A novel allele <i><scp>HLA</scp>â€‹C*07:445</i> identified in a French hematopoietic stem cell donor. Hla, 2016, 87, 403-404.	0.4	2
925	<i>HLAâ€‹B*46:65</i>, a novel <i>HLAâ€‹B*46</i> variant, detected in a Taiwanese unrelated bone marrow stem cell donor. Hla, 2016, 87, 397-398.	0.4	2
926	Identification of a novel HLAâ€‹A allele, <i>HLAâ€‹A*02:590</i>, in a Korean individual. Hla, 2016, 87, 454-455.	0.4	4
927	Identification of a novel HLAâ€‹C allele, <i>HLAâ€‹C*07:477</i> in a Drung Chinese individual. Hla, 2016, 87, 468-469.	0.4	3
928	Characterization of a novel <i><scp>MICA</scp></i> allele, <i><scp>MICA</scp>*012:05</i>, by cloning and sequencing. International Journal of Immunogenetics, 2016, 43, 244-245.	0.8	3

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930	Characterization of the novel *HLA–C*02:106* allele in a hematopoietic stem cell volunteer donor. Hla, 2016, 88, 52-53.	0.4	3
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932	A novel HLA–B*14 allele – B*14:53 – genetics and serology. International Journal of Immunogenetics, 2016, 43, 236-239.	0.8	7
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934	Identification of the novel *HLA–B*07:261* allele. Hla, 2016, 87, 102-103.	0.4	3
935	Identification of a novel *HLA–B*40* allele, *HLA–B*40:324*, by sequence–based typing, in a Chinese individual. Hla, 2016, 87, 110-111.	0.4	3
936	A novel HLA–DRB1 allele, *HLA–DRB1*12:48*. Hla, 2016, 87, 190-191.	0.4	3
937	*HLA–A*24:02:96*, a novel variant of *HLA–A*24:02:01:01*, identified in a Chinese individual by sequence–based typing. Hla, 2016, 87, 171-172.	0.4	3
938	A novel allele, *HLA–B*55:70* was identified in a Chinese cord blood donor. Hla, 2016, 87, 183-185.	0.4	3
939	*HLA–A*11:235Q*, a novel *HLA–A*11* variant, detected in a Taiwanese individual. Hla, 2016, 87, 456-458.	0.4	3
940	*HLA–B*46:40:02*, a novel *HLA–B*46* allele identified in a Chinese individual by sequence–based typing. Hla, 2016, 87, 462-464.	0.4	4
941	Identification of a novel *HLA–C*03* variant, *C*03:303*, by sequence–based typing in a Dai Chinese individual. Hla, 2016, 87, 465-466.	0.4	3
942	Identification of a novel *HLA–A*02* variant, *HLA–A*02:614*, in a Taiwanese individual. Hla, 2016, 87, 455-456.	0.4	3
943	A novel HLA–B allele, *HLA–B*15:399*, identified in a Chinese individual. Hla, 2016, 88, 264-265.	0.4	3
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949	Nomenclature for factors of the HLA system, update August 2016. Human Immunology, 2016, 77, 1324-1328.	1.2	0
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963	Identification of a novelHLA-B*07variant,B*07:269, by sequence-based typing in a Taiwanese bone marrow stem cell donor. Hla, 2016, 88, 198-199.	0.4	3
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967	Identification of a novel <scp>HLA</scp>â€B null allele, <i><scp>HLA</scp>â€B*41:<scp>45N</scp></i>, by <scp>HLA</scp> typing of two related Caucasoid individuals. Hla, 2016, 88, 50-51.	0.4	3
968	Identification of a novel HLA allele, <i>HLAâ€B*08:163</i>, in a platelet donor. Hla, 2016, 88, 263-264.	0.4	4
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972	Recognition of a novel <i>HLAâ€A*11</i> variant, <i>HLAâ€A*11:196</i>, in a Taiwanese unrelated bone marrow stem cell donor. Hla, 2016, 88, 39-40.	0.4	3
973	Detection of a novelHLA-B*46:01variant,HLA-B*46:01:22, in a Taiwanese individual. Hla, 2016, 88, 205-206.	0.4	3
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980	A novelHLA-B*38variant,B*38:15, detected in a Taiwanese individual. Hla, 2016, 88, 125-126.	0.4	3
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986	Description of two new HLAâ€A alleles, <i>HLAâ€A*02:572</i> and <i>HLAâ€A*03:225</i>. Hla, 2016, 87, 39-40.	0.4	3
987	Correction of the <i><scp>HLAâ€DQB1</scp>*04:01:01</i> sequence at position 79 in exon 1. Hla, 2016, 87, 57-58.	0.4	3
988	A novel HLA-C null allele,HLA-C*08:121N. Hla, 2016, 87, 55-56.	0.4	3
989	Identification of a novelHLA-A*02variant,HLA-A*02:586, in a Taiwanese bone marrow hematopoietic stem cell donor. Hla, 2016, 87, 40-41.	0.4	3
990	A novel <scp>HLAâ€DRB1</scp> allele, <i><scp>DRB1</scp>*15:66:02</i> was identified in a Chinese potential donor by sequenceâ€based typing. Hla, 2016, 87, 61-63.	0.4	3
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992	A single nucleotide insertion in Exon 3 produces a novelHLA-B*58null allele,HLA-B*58:72N. Hla, 2016, 87, 54-55.	0.4	3
993	Identification of a novelHLA-A*02variant,HLA-A*02:621, in a Taiwanese bone marrow donor. Hla, 2016, 88, 194-195.	0.4	4
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995	Discovery ofHLA-B*58:80, a novelHLA-B*58variant, in a Taiwanese bone marrow donor. Hla, 2016, 88, 127-129.	0.4	3
996	Identification of two novel <scp>HLAâ€DQB1</scp> alleles, <i><scp>HLAâ€DQB1</scp>*03:164</i> and <i><scp>HLAâ€DQB1</scp>*03:165</i> in Chinese individuals. Hla, 2016, 88, 316-317.	0.4	4
997	Nomenclature for factors of the <scp>HLA</scp> system, update September 2016. International Journal of Immunogenetics, 2016, 43, 426-433.	0.8	1
998	Identification of a novelHLA-C*06variant allele,HLA-C*06:166, by sequence-based typing in a Chinese individual. Hla, 2016, 88, 315-316.	0.4	0
999	A novel HLAâ€B allele, <i>HLAâ€B*13:69</i>. Hla, 2016, 88, 122-123.	0.4	3
1000	Characterization of a novel HLA-DRB1 allele,HLA-DRB1*13:215, in an Italian bone marrow donor. Hla, 2016, 88, 269-269.	0.4	3
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1005	Identification of a novelHLA-B*50allele,HLA-B*50:01:08, in a Spanish hematological patient. Hla, 2016, 88, 266-267.	0.4	3
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1017	Nomenclature for factors of the HLA system, update April 2016. Hla, 2016, 88, 131-135.	0.4	0
1018	Identification of the novel *HLAâ€B*39:01:23* allele by polymerase chain reaction sequenceâ€based typing. Hla, 2016, 88, 310-311.	0.4	2
1019	Identification of a novel allele, *HLA*â€A*26:01:01:03N*, by fullâ€length genome sequencing. Hla, 2016, 88, 260-261.	0.4	4
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1024	<i>HLAâ€B*15:388</i>, a novel variant of <i>HLAâ€B*15</i>, discovered in a Taiwanese individual. Hla, 2016, 88, 204-205.	0.4	3
1025	Identification of a novelHLA-B*15allele,HLA-B*15:374, in a Chinese individual. Hla, 2016, 88, 307-308.	0.4	0
1026	Genomic fullâ€length sequence of two <scp>HLA</scp>â€A alleles, <i>A*24:08</i> and <i>A*24:10</i>, identified by cloning and sequencing. Hla, 2016, 88, 300-302.	0.4	0
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1033	Genomic fullâ€length sequence of two new HLAâ€C alleles, <i><scp>HLAâ€C</scp>*04:239</i> and <i><scp>HLAâ€C</scp>*05:137</i>. Hla, 2016, 88, 313-314.	0.4	0
1034	Nomenclature for factors of the <scp>HLA</scp> system, update May 2016. International Journal of Immunogenetics, 2016, 43, 314-319.	0.8	0
1035	<i>HLAâ€DRB1*16:39,</i> a novel <i>HLAâ€DRB1*16</i> variant, discovered in a Taiwanese bone marrow hematopoietic stem cell donor. Hla, 2016, 88, 124-125.	0.4	5
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1040	A novel allele, <i><scp>HLA</scp>â€‘B*55:77</i>, identified by sequenceâ€‘based typing in a Chinese individual. Hla, 2016, 87, 185-186.	0.4	3
1041	Detection of <i><scp>HLA</scp>â€‘C*04:212</i>, a novel <i><scp>HLA</scp>â€‘C*04</i> variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2016, 87, 187-189.	0.4	3
1042	Identification of a novel <i><scp>HLA</scp>â€‘C*16</i> variant, <i><scp>HLA</scp>â€‘C*16:90</i>, in a Chinese individual. Hla, 2016, 87, 189-190.	0.4	3
1043	Identification of a novel allele,HLA-A*24:02:50, by sequence-based typing. Hla, 2016, 87, 388-390.	0.4	2
1044	Detection of <i><scp>HLAâ€‘DRB1</scp>*04:207</i>, a novel <i><scp>HLAâ€‘DRB1</scp>*04</i> variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2016, 87, 406-407.	0.4	3
1045	<i><scp>HLA</scp>â€‘A*02:513</i>, a novel <i><scp>HLA</scp>â€‘A*02</i> allele. Hla, 2016, 87, 382-383.	0.4	2
1046	Identification of the novel <i>HLAâ€‘DRB1*12:50</i> allele by polymerase chain reaction sequenceâ€‘based typing in a Chinese individual. Hla, 2016, 87, 473-474.	0.4	6
1047	A new HLA-B allele,B*52:44, sequenced in a Chinese individual. Hla, 2016, 87, 464-465.	0.4	3
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1054	Nomenclature for factors of the HLA system, update October 2015. Hla, 2016, 87, 117-121.	0.4	2
1055	Nomenclature for factors of the HLA system, update November 2015. Hla, 2016, 87, 122-125.	0.4	1
1056	Identification of two novel <scp>HLA</scp>â€‘A alleles, <scp><i>HLA</i></scp><i>â€‘A*03:181</i> and <scp><i>HLA</i></scp><i>â€‘A*03:229</i> in Chinese individuals. Hla, 2016, 87, 165-166.	0.4	5

#	ARTICLE	IF	CITATIONS
1057	Discovery ofHLA-B*35:307, a novelHLA-B*35variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2016, 87, 178-179.	0.4	5
1058	Identification of a novel <i><scp>HLAâ€DRB1</scp>*14</i> allele, <i><scp>HLAâ€DRB1</scp>*14:150</i>, in a Chinese individual. Hla, 2016, 87, 407-408.	0.4	1
1059	<i><scp>HLA</scp>â€A*33:102</i>, a novel allele identified by sequenceâ€based typing in cord blood from a Korean woman. Hla, 2016, 87, 391-392.	0.4	2
1060	Identification of a novel <i><scp>HLA</scp>â€A*02</i> variant, <i><scp>HLA</scp>â€A*02:610,</i> in a Taiwanese individual. Hla, 2016, 87, 385-386.	0.4	2
1061	<i><scp>HLA</scp>â€B*40:327</i>, a novel <i><scp>HLA</scp>â€B*40</i> variant, discovered in a Taiwanese blood donor. Hla, 2016, 87, 395-397.	0.4	2
1062	<i>HLAâ€C*07:486</i>, a novel <i>HLAâ€C*07</i> variant, detected in a Taiwanese individual. Hla, 2016, 87, 404-405.	0.4	2
1063	<i><scp>HLA</scp>â€C*16:01:01:02</i>, a recently isolated <scp>HLA</scp> class I allele from Southern Africa. Hla, 2016, 87, 470-471.	0.4	3
1064	Isolation of fullâ€length <i><scp>HLA</scp>â€C*18:02</i> allele in an individual from Subâ€Saharan Africa. Hla, 2016, 87, 471-472.	0.4	3
1065	<i>HLAâ€A*02:06:21,</i> a novel variant of <i>HLAâ€A*02:06</i>, discovered in a Taiwanese bone marrow hematopoietic stem cell donor. Hla, 2016, 87, 36-37.	0.4	3
1066	Identification of a novel <i>HLAâ€A*02 allele</i>, <i>HLAâ€A*02:432</i>, in a Chinese individual. Hla, 2016, 87, 37-38.	0.4	3
1067	A novel allele <i><scp>HLA</scp>â€A*11:152</i> was identified by sequenceâ€based typing in a Chinese individual. Hla, 2016, 87, 41-42.	0.4	3
1068	Nomenclature for factors of the <scp>HLA</scp> system, update August 2015. International Journal of Immunogenetics, 2016, 43, 55-59.	0.8	0
1069	Nomenclature for factors of the <scp>HLA</scp> system, update September 2015. International Journal of Immunogenetics, 2016, 43, 60-64.	0.8	0
1070	Nomenclature for factors of the HLA system, update December 2015. Human Immunology, 2016, 77, 432-435.	1.2	0
1071	The Secreted Protein Rv1860 of Mycobacterium tuberculosis Stimulates Human Polyfunctional CD8⁺T Cells. Vaccine Journal, 2016, 23, 282-293.	3.2	13
1072	Allele Frequencies Net Database: Improvements for storage of individual genotypes and analysis of existing data. Human Immunology, 2016, 77, 238-248.	1.2	107
1073	Identification and characterization a novel <scp>HLA</scp>â€A allele, A*02:355, by sequenceâ€based typing in a Chinese potential donor. International Journal of Immunogenetics, 2017, 44, 35-37.	0.8	3
1074	<i><scp>HLA</scp>â€C*07:544</i>, a novel <i><scp>HLA</scp>â€C*07</i> variant, detected in a Taiwanese bone marrow donor. Hla, 2017, 89, 120-121.	0.4	3

#	ARTICLE	IF	CITATIONS
1075	Combining one‐step Sanger sequencing with phasing probe hybridization for HLA class I typing yields rapid, G‐group resolution predicting 99% of unique full length protein sequences. Hla, 2017, 89, 90-97.	0.4	4
1076	A new <i>HLA‐B*55</i> allele, <i>B*55:83N</i> with a stop codon in exon 4 generated by a point mutation<i>,</i> identified in a Chinese individual. Hla, 2017, 89, 119-120.	0.4	3
1077	Nomenclature for factors of the HLA system, update October 2016. Human Immunology, 2017, 78, 294-305.	1.2	0
1078	HLA typing in diverse populations. ISBT Science Series, 2017, 12, 107-111.	1.1	0
1079	Identification of a novel <i>HLA‐B*40</i> allele, <i>HLA‐B*40:332</i>, in a Korean individual. Hla, 2017, 89, 117-118.	0.4	3
1080	Nomenclature for factors of the HLA system, update December 2016. Human Immunology, 2017, 78, 316-321.	1.2	1
1081	<scp>RNA</scp> processing and protein expression of <i><scp>HLA</scp>‐B*07:<scp>44N</scp></i>. Hla, 2017, 89, 230-234.	0.4	5
1082	Identification of the novel <i><scp>HLA‐DRB1</scp>*14:54:06</i> allele by sequencing‐based typing in a Chinese bone marrow donor. Hla, 2017, 89, 172-173.	0.4	3
1083	Nomenclature for factors of the <scp>HLA</scp> system, update October 2016. Hla, 2017, 89, 177-189.	0.4	1
1084	<i><scp>HLA</scp>‐B*15:414</i>, a novel variant of <i><scp>HLA</scp>‐B*15</i>, discovered in a Taiwanese individual. Hla, 2017, 89, 242-243.	0.4	4
1085	Nomenclature for factors of the <scp>HLA</scp> system, update November 2016. International Journal of Immunogenetics, 2017, 44, 79-85.	0.8	1
1086	A novel HLA-E allele,HLA-E*01:01:01:06, identified in a Chinese Leukemia patient. Hla, 2017, 89, 260-262.	0.4	4
1087	Identification of an HLA class I allele closely involved in the autoantigen presentation in acquired aplastic anemia. Blood, 2017, 129, 2908-2916.	0.6	71
1088	Detection of a novel <i><scp>HLA</scp>‐C*03</i> variant, <i><scp>HLA</scp>‐C*03:227</i>, in a Taiwanese individual. Hla, 2017, 89, 253-254.	0.4	5
1089	Identification of a novel <i><scp>HLA</scp>‐A*02:01:01</i> variant, <i><scp>HLA</scp>‐A*02:01:01:09</i>, in a Taiwanese bone marrow donor. Hla, 2017, 89, 301-302.	0.4	3
1090	Full‐length sequences of 3 <scp>HLA</scp>‐B alleles, <i>B*07:05:01:01</i>, <i>B*14:01:01</i> and <i>B*18:02</i>, confirmed by cloning and sequencing. Hla, 2017, 89, 305-308.	0.4	3
1091	Full‐length sequences of 3 <scp>HLA</scp>‐B alleles, <i>B*40:01:01</i>, <i>B*40:03</i> and <i>B*40:40</i>, confirmed by cloning and sequencing. Hla, 2017, 89, 321-324.	0.4	3
1092	Full‐length sequences of <i><scp>HLA</scp>‐B*39:05:01</i> and <i>B*39:<scp>38Q</scp></i>, confirmed by cloning and sequencing. Hla, 2017, 89, 159-162.	0.4	3

#	Article	IF	Citations
1093	Full‐length sequences of 3 <i><scp>HLA</scp>‐B*56</i> alleles, <i>B*56:01:01:01</i>, <i>B*56:03</i> and <i>B*56:04</i>, confirmed by cloning and sequencing. Hla, 2017, 89, 246-250.	0.4	3
1094	Nomenclature for factors of the <scp>HLA</scp> system, update November 2016. Hla, 2017, 89, 190-197.	0.4	5
1095	Identification of a novel <scp>HLA</scp>‐B allele, <i><scp>HLA</scp>‐B*55:81</i>. Hla, 2017, 89, 166-167.	0.4	3
1096	Exon 2 sequencing of the new <scp>HLA</scp>‐<scp>DRB</scp>1 allele, <scp>DRB</scp>1*13:216. International Journal of Immunogenetics, 2017, 44, 38-39.	0.8	3
1097	A new <i>MICA</i> allele, <i>MICA*007:07</i>, characterized by cloning and sequencing. International Journal of Immunogenetics, 2017, 44, 145-147.	0.8	3
1098	Algorithm with Heuristics for Kidney Allocation in Transplant Information System. IFMBE Proceedings, 2017, , 213-218.	0.2	2
1099	Identification of a novel allele, <i><scp>HLA</scp>‐B*15:01:23</i>, in a platelet donor by sequence‐based typing. Hla, 2017, 90, 37-39.	0.4	3
1100	Characterization of the novel <i><scp>HLA</scp>‐A*32:95</i> allele, identified in the Republic of Kazakhstan. Hla, 2017, 90, 112-113.	0.4	4
1101	Full‐length sequence of 2 <scp>HLA</scp>‐B alleles, <i>B*52:01:01:01</i> and <i>B*52:01:02:01</i>, identified by cloning and sequencing. Hla, 2017, 89, 163-165.	0.4	2
1102	Confirmation of the <i><scp>HLA</scp>‐C*16:97</i> allele in multiple individuals, a new common and well‐defined allele?. Hla, 2017, 89, 170-171.	0.4	2
1103	Detection of a novelHLA-A*11variant,A*11:255, in a Taiwanese individual. Hla, 2017, 89, 238-239.	0.4	3
1104	Identification of a new <i><scp>HLA‐DQB1</scp>*06</i> allele, <i><scp>HLA‐DQB1</scp>*06:210</i>, by monoallelic Sanger sequencing. Hla, 2017, 90, 132-133.	0.4	3
1105	HLA-C*07:566, a novelHLA-C*07variant, detected in a Taiwanese hematopoietic stem cell donor. Hla, 2017, 89, 255-256.	0.4	3
1106	A novel <scp>HLA</scp>‐E allele, <i><scp>HLA</scp>‐E*01:01:01:07</i>, identified in a Chinese leukemia patient. Hla, 2017, 89, 327-330.	0.4	4
1107	Full‐length sequences of 4 <i><scp>HLA</scp>‐B*15</i> alleles, <i>B*15:03:01:01</i>, <i>B*15:13:01</i>, <i>B*15:18:01:01</i> and <i>B*15:25:01</i>, confirmed by cloning and sequencing. Hla, 2017, 89, 309-312.	0.4	3
1108	HLA-A*11:256Q, a novelHLA-A*11variant, detected in a Taiwanese individual. Hla, 2017, 89, 302-304.	0.4	4
1109	Identification of the novel <i><scp>HLA</scp>‐B*27:147</i> allele by polymerase chain reaction sequence‐based typing. Hla, 2017, 90, 115-116.	0.4	3
1110	Identification of the novel null allele, <i><scp>HLA</scp>‐C*01:<scp>109N</scp></i>, using polymerase chain reaction sequence‐based typing in a Chinese leukemia patient. Hla, 2017, 89, 252-253.	0.4	4

#	ARTICLE	IF	CITATIONS
1111	The novel <i><scp>HLAâ€DRB1</scp>*15:140</i> allele discovered in a Taiwanese unrelated hematopoietic stem cell donor. Hla, 2017, 89, 259-260.	0.4	4
1112	A novel <i><scp>HLA</scp>â€A*24</i> allele, <i><scp>A</scp>*24:02:61</i>, confirmed in a Chinese individual. Hla, 2017, 89, 52-53.	0.4	4
1113	<i>HLAâ€DQB1*03:01:34</i>, a novel allele, which has arisen by silent mutation in codon 87. Hla, 2017, 89, 62-64.	0.4	4
1114	Identification of the new <i><scp>HLAâ€DQA1</scp>*01:<scp>15N</scp></i> allele in an Italian patient. Hla, 2017, 90, 130-131.	0.4	2
1115	Nomenclature for factors of the HLA system, update March 2017. Human Immunology, 2017, 78, 461-465.	1.2	0
1116	HLA class I binding prediction via convolutional neural networks. Bioinformatics, 2017, 33, 2658-2665.	1.8	99
1117	Nomenclature for factors of the HLA system, update February 2017. Human Immunology, 2017, 78, 455-460.	1.2	0
1118	Gorilla MHC class I gene and sequence variation in a comparative context. Immunogenetics, 2017, 69, 303-323.	1.2	12
1119	Detection of a novel <scp>HLA</scp>â€B allele, <i><scp>HLA</scp>â€B*39:119</i>, in a Chinese individual. Hla, 2017, 90, 116-118.	0.4	2
1120	Identification of a novel <scp>HLA</scp>â€B allele, <i><scp>HLA</scp>â€B*08:177</i>, in a Hungarian patient and her <scp>HLA</scp> identical sibling. Hla, 2017, 90, 113-114.	0.4	2
1121	Identification of the novel <i><scp>HLAâ€DQB1</scp>*02:85</i> and <i><scp>HLAâ€DRB1</scp>*01:01:30</i> alleles in Russian individuals. Hla, 2017, 90, 135-136.	0.4	3
1122	Identification of the novel <i><scp>HLA</scp>â€B*40:01:41</i> allele by polymerase chain reaction sequenceâ€based typing in a Chinese cord blood donor. Hla, 2017, 90, 118-120.	0.4	3
1123	Identification of a novel HLA-C allele,HLA-C*15:134, in a Taiwanese hematopoietic stem cell donor. Hla, 2017, 89, 256-257.	0.4	3
1124	Fullâ€length sequences of 4 <i><scp>HLA</scp>â€B*35</i> alleles, <i>B*35:02:01:01</i>, <i>B*35:03:01:01</i>, <i>B*35:05:01:01</i> and <i>B*35:08:01:01</i>, confirmed by cloning and sequencing. Hla, 2017, 89, 317-321.	0.4	3
1125	The full length genomic sequence of a novel <i><scp>HLA</scp>â€A*24</i> allele, <i><scp>HLA</scp>â€A*24:353</i>, identified in a patient with hepatitis B infection. Hla, 2017, 89, 304-305.	0.4	3
1126	Detection of a new <i><scp>HLA</scp>â€B*44</i> allele, <i><scp>HLA</scp>â€B*44:02:45,</i> by monoallelic Sanger sequencing. Hla, 2017, 90, 124-124.	0.4	2
1127	Nomenclature for factors of the <scp>HLA</scp> system, update October 2016. International Journal of Immunogenetics, 2017, 44, 71-78.	0.8	0
1128	Fullâ€length sequences of <i><scp>HLA</scp>â€B*67:01:01</i> and <i>B*67:01:02</i>, confirmed by cloning and sequencing. Hla, 2017, 89, 324-327.	0.4	3

#	ARTICLE	IF	CITATIONS
1129	Identification of MHC Haplotypes Associated with Drug-induced Hypersensitivity Reactions in Cynomolgus Monkeys. Toxicologic Pathology, 2017, 45, 127-133.	0.9	5
1130	Identification of the novel <i>HLAâ€DRB1*08:69</i> allele by polymerase chain reaction sequenceâ€based typing in a Chinese cord blood donor. Hla, 2017, 89, 64-65.	0.4	5
1131	Detection of <i><scp>HLA</scp>â€C*04:247</i>, a novel <i><scp>HLA</scp>â€C*04</i> variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2017, 89, 60-61.	0.4	3
1132	<i><scp>HLA</scp>â€B*13:95</i>, a novel variant of <i><scp>HLA</scp>â€B*13</i>, discovered in a Taiwanese blood donor. Hla, 2017, 89, 54-55.	0.4	3
1133	Description of a new <i><scp>HLA</scp>â€A*02</i> allele, <i>A*02:658</i>, in a Russian individual. Hla, 2017, 89, 235-236.	0.4	4
1134	Fullâ€length sequence of <i><scp>HLA</scp>â€B*55:02:01:01</i>, confirmed by cloning and sequencing. Hla, 2017, 90, 45-48.	0.4	3
1135	Identification of a novel <scp>HLA</scp> allele, <i><scp>HLA</scp>â€B*41:50</i>, in a French individual. Hla, 2017, 90, 122-123.	0.4	2
1136	<scp>TypeLoader</scp>: A fast and efficient automated workflow for the annotation and submission of novel fullâ€length <scp>HLA</scp> alleles. Hla, 2017, 90, 25-31.	0.4	20
1137	Detection of the HLAâ€DQB1 allele, <i>DQB1*03:82</i>, in a Kazakh patient with acute myeloid leukemia. Hla, 2017, 90, 181-182.	0.4	6
1138	Identification of a novel <scp>HLA</scp>â€A allele, <i><scp>HLA</scp>â€A*02:505</i>, by sequenceâ€based typing in a patient with tuberculosis. Hla, 2017, 90, 106-107.	0.4	3
1139	A novel HLA allele, HLA-C*15:02:01:04 , identified in a Taiwanese individual. Hla, 2017, 90, 50-51.	0.4	3
1140	Identification of a novel <i><scp>HLA</scp>â€B*27</i> variant, <i>B*27:112</i>, by sequenceâ€based typing in a Taiwanese donor. Hla, 2017, 90, 175-176.	0.4	3
1141	The <i><scp>DRB1</scp>*15:11</i> allele discovered in a Taiwanese unrelated hematopoietic stem cell donor. Hla, 2017, 90, 184-185.	0.4	4
1142	Nomenclature for factors of the HLA system, update February 2017. Hla, 2017, 90, 62-69.	0.4	0
1143	Sequenceâ€based typing identification of a novel <scp>HLA</scp>â€C allele, <i><scp>C</scp>*12:214</i>, in a Han Chinese individual. Hla, 2017, 90, 179-180.	0.4	3
1144	Polymorphism at residue 156 of the new <i><scp>HLA</scp>â€A*02:683</i> allele suggests immunological relevance. Hla, 2017, 90, 107-109.	0.4	3
1145	The new <i>HLAâ€B*50:51</i> allele was generated by intralocus recombination between <i>B*50:01:01:02</i> and <i>B*14:02:01</i>. Hla, 2017, 90, 176-178.	0.4	5
1146	A novel <scp>HLA</scp>â€C allele, <i><scp>HLA</scp>â€C*01:01:01:07</i>, was identified in a Chinese patient with Posnerâ€Schlossman syndrome. Hla, 2017, 90, 136-140.	0.4	6

#	ARTICLE	IF	CITATIONS
1147	Identification of a novel <i><scp>HLA</scp>â€B*18</i> allele, <i>B*18:119</i>, using sequenceâ€based typing in a caucasoid individual. Hla, 2017, 89, 244-245.	0.4	3
1148	Description of the novel <scp>HLA</scp>â€A allele, <i><scp>HLA</scp>â€A*11:229</i>, identified by sequenceâ€based typing in a Chinese individual. Hla, 2017, 90, 111-112.	0.4	3
1149	Detection of a novel <i><scp>HLA</scp>â€A*30</i> variant, <i>A*30:109</i>, in a Taiwanese individual. Hla, 2017, 90, 36-37.	0.4	3
1150	Nomenclature for factors of the <scp>HLA</scp> system, update December 2016. Hla, 2017, 89, 198-205.	0.4	1
1151	Characterization of three new <scp>HLA</scp> Class I Alleles in Spanish Caucasians, <scp>HLA</scp>â€A*02:620, <scp>HLA</scp>â€B*27:150 and <scp>HLA</scp>â€B*07:05:01:02. International Journal of Immunogenetics, 2017, 44, 148-150.	0.8	4
1152	Role of Genetic Polymorphism of ALDH2 in Hematopoietic Stem Cell Transplantation. Biology of Blood and Marrow Transplantation, 2017, 23, 1374-1380.	2.0	4
1153	A new allele, <i><scp>HLA</scp>â€A*03:<scp>275N</scp></i>. Hla, 2017, 90, 109-110.	0.4	2
1154	Identification of the novel <i><scp>HLAâ€DRB1</scp>*15:127</i> allele by polymerase chain reaction sequenceâ€based typing in a Chinese bone marrow donor. Hla, 2017, 90, 133-134.	0.4	3
1155	Identification of a novel <i><scp>HLA</scp>â€B*13</i> allele, <i><scp>HLA</scp>â€B*13:99</i>, by sequenceâ€based typing in German bone marrow donor. Hla, 2017, 89, 158-159.	0.4	4
1156	Limited HLA sequence variation outside of antigen recognition domain exons of 360 10 of 10 matched unrelated hematopoietic stem cell transplant donorâ€recipient pairs. Hla, 2017, 89, 39-46.	0.4	20
1157	Identification of 2 novel <scp>HLA</scp>â€B alleles, <i><scp>HLA</scp>â€B*55:02:09</i> and <i><scp>HLA</scp>â€B*55:80</i> in Chinese individuals. Hla, 2017, 90, 48-50.	0.4	6
1158	Identification of the novel <i>HLAâ€B*39:01:01:04</i> allele in a Chinese individual by sequenceâ€based typing. Hla, 2017, 89, 115-117.	0.4	3
1159	Two new HLA class I alleles described in a Spanish individual, <i>HLAâ€A*11:01:01:04</i> and <i>HLAâ€B*35:330</i>. Hla, 2017, 89, 236-237.	0.4	3
1161	Principles and Recommendations for Standardizing the Use of the Next-Generation Sequencing Variant File in Clinical Settings. Journal of Molecular Diagnostics, 2017, 19, 417-426.	1.2	19
1162	International Conference on Advancements of Medicine and Health Care through Technology; 12th - 15th October 2016, Cluj-Napoca, Romania. IFMBE Proceedings, 2017, , .	0.2	1
1163	Nomenclature for factors of the <scp>HLA</scp> system, update December 2016. International Journal of Immunogenetics, 2017, 44, 86-92.	0.8	0
1164	Nomenclature for factors of the HLA system, update November 2016. Human Immunology, 2017, 78, 306-315.	1.2	0
1165	Identification of 3 novel HLAâ€B alleles: <i>B*08:173</i>, <i>B*18:72:03</i> and <i>B*53:05:02</i>. Hla, 2017, 89, 114-115.	0.4	3

#	ARTICLE	IF	CITATIONS
1166	A novel <scp>HLA</scp> allele, <i><scp>HLA</scp>â€B*50:48</i>, identified by sequencingâ€based typing. Hla, 2017, 89, 57-58.	0.4	5
1167	Identification of a novel HLAâ€B allele, <i>HLAâ€B*40:238</i>, in a Taiwanese individual. Hla, 2017, 90, 252-253.	0.4	2
1168	The fullâ€length sequence of <i><scp>HLAâ€B</scp>*59:01:01:01</i> confirmed by cloning and sequencing. Hla, 2017, 90, 255-258.	0.4	2
1169	Identification of a novel allele, <i>HLAâ€C*01:135</i>, by fullâ€length genomic sequencing. Hla, 2017, 90, 258-259.	0.4	2
1170	Two novel alleles, <i>HLAâ€A*02:643N</i> and <i>HLAâ€B*53:44</i>, identified in Brazilian individuals. Hla, 2017, 90, 362-364.	0.4	4
1171	<i>HLAâ€A*24:388N</i>: a novel <i>HLAâ€A*24</i> allele identified by sequenceâ€based typing. Hla, 2017, 90, 364-365.	0.4	3
1172	Nomenclature for factors of the <scp>HLA</scp> system, update May 2017. Hla, 2017, 90, 193-196.	0.4	2
1173	Identification of a novel allele, <i><scp>HLA</scp>â€C*02:02:33</i>, by fullâ€length genomic sequencing. Hla, 2017, 90, 313-314.	0.4	2
1174	Characterization of the novel <i>HLAâ€A*02:07:10</i> allele by sequencingâ€based typing. Hla, 2017, 90, 361-362.	0.4	3
1175	<i><scp>HLAâ€DPB1</scp>*519:01</i>, a new allele identified by sequenceâ€based typing in a Korean individual. Hla, 2017, 90, 318-319.	0.4	2
1176	Nomenclature for factors of the <scp>HLA</scp> system, update June 2017. International Journal of Immunogenetics, 2017, 44, 243-250.	0.8	0
1177	The novel <i>HLAâ€B*08:183</i> allele identified by sequenceâ€based typing in a Caucasian leukemia patient. Hla, 2017, 90, 367-368.	0.4	3
1178	Identification of the novel allele, <i><scp>HLA</scp>â€A*02:653</i>, in an Italian patient. Hla, 2017, 90, 300-301.	0.4	1
1179	Identification of the novel allele, <i>HLAâ€DRB1*09:30</i>, by sequenceâ€based high resolution typing. Hla, 2017, 90, 379-380.	0.4	4
1180	Detection of a novel <i><scp>HLAâ€DRB1</scp>*09</i> variant, <i><scp>HLAâ€DRB1</scp>*09:31</i>, in a College of American Pathologists <scp>HLA</scp> survey sample. Hla, 2017, 90, 320-321.	0.4	3
1181	<i><scp>HLAâ€DPB1</scp>*518:01</i>, a new allele identified by sequenceâ€based typing in a Korean individual. Hla, 2017, 90, 316-318.	0.4	3
1182	Association of the HLA-B alleles with carbamazepine-induced Stevensâ€"Johnson syndrome/toxic epidermal necrolysis in the Javanese and Sundanese population of Indonesia: the important role of the HLA-B75 serotype. Pharmacogenomics, 2017, 18, 1643-1648.	0.6	36
1183	Characterization of the novel <i><scp>HLA</scp>â€B*49:39</i> allele identified in a German leukaemia patient. Hla, 2017, 90, 43-45.	0.4	3

#	ARTICLE	IF	CITATIONS
1184	Nomenclature for factors of the <scp>HLA</scp> system, update June 2017. Hla, 2017, 90, 197-203.	0.4	2
1185	<i>HLAâ€B*40:302</i>, a novel allele identified by sequenceâ€based typing in a Korean individual. Hla, 2017, 90, 368-369.	0.4	0
1186	Identification of <i>HLAâ€B*58:01:21</i>, a novel allele in a Korean individual. Hla, 2017, 90, 371-372.	0.4	3
1187	Melanocytes: Target Cells of an HLA-C*06:02â€"Restricted Autoimmune Response in Psoriasis. Journal of Investigative Dermatology, 2017, 137, 2053-2058.	0.3	33
1188	Identification of a new <i><scp>HLAâ€B</scp>*51:226</i> allele in a Chinese bone marrowâ€related donor. Hla, 2017, 90, 370-371.	0.4	3
1189	Characterization of the novel <i>HLAâ€DRB1*13:241</i> allele by sequencingâ€based typing. Hla, 2017, 90, 380-381.	0.4	3
1190	Identification of the <i>HLAâ€DQB1*06:123</i> allele in an unrelated stem cell donor from the Saudi Registry. Hla, 2017, 90, 262-263.	0.4	3
1191	Sequenceâ€based typing of a novel <scp>HLAâ€DRB1</scp> allele, <i><scp>HLAâ€DRB1</scp>*14:32:03</i>, in a Chinese individual. Hla, 2017, 90, 325-326.	0.4	2
1192	Identification of a novel allele, <i><scp>HLA</scp>â€A*02:01:131</i>, by fullâ€length genomic sequencing. Hla, 2017, 90, 360-361.	0.4	3
1193	<i><scp>HLA</scp>â€A*33:<scp>74N</scp></i>, a novel <i><scp>HLA</scp>â€A*33</i> variant, identified by sequenceâ€based typing in a Taiwanese individual. Hla, 2017, 90, 365-366.	0.4	5
1194	A new allele, <i><scp>HLAâ€DQA1</scp>*02:01:02</i>. Hla, 2017, 90, 376-377.	0.4	4
1195	A novel allele, <i>HLAâ€C*14:02:01:03</i>, identified by fullâ€length genomic sequencing. Hla, 2017, 90, 260-261.	0.4	2
1196	Discovery of a novel <i>HLAâ€B*07</i> variant, <i>HLAâ€B*07:294</i>, in a Chinese individual. Hla, 2017, 90, 251-252.	0.4	1
1197	<i>HLAâ€A*02:687</i>, a novel allele identified by sequenceâ€based typing in cord blood from a Korean woman. Hla, 2017, 90, 246-247.	0.4	1
1198	<i>HFE</i> gene polymorphism defined by sequenceâ€based typing of the Brazilian population and a standardized nomenclature for <i>HFE</i> allele sequences. Hla, 2017, 90, 238-242.	0.4	3
1199	Identification of the novel <i>HLAâ€A*11:141</i> allele in a Chinese bone marrow donor. Hla, 2017, 90, 247-248.	0.4	3
1200	Identification of a new HLAâ€A allele, <i>HLAâ€A*02:07:09</i>, in a Chinese nonâ€Hodgkin lymphoma patient. Hla, 2017, 90, 243-244.	0.4	1
1201	Fullâ€length sequencing of the HLA region identified a novel allele, <i>HLAâ€B*52:70</i>. Hla, 2017, 90, 253-254.	0.4	2

#	ARTICLE	IF	CITATIONS
1202	Epitopes and motifs of the <scp>HLA</scp>â€B*14 allele family products and related <scp>HLA</scp>â€B14 crossâ€reactive specificities. International Journal of Immunogenetics, 2017, 44, 225-233.	0.8	2
1203	Detection of a novel <i><scp>HLA</scp>â€A*11</i> variant, <i>A*11:263</i>, in a Taiwanese individual. Hla, 2017, 90, 303-304.	0.4	2
1204	<i><scp>HLA</scp>â€B*40:01:45</i>, a novel variant of <i><scp>HLA</scp>â€B*40:01</i>, discovered in a Taiwanese hematopoitic stem cell donor. Hla, 2017, 90, 311-312.	0.4	2
1205	Detection of a novel allele, <i><scp>HLAâ€A</scp>*24:198</i>, by sequenceâ€based typing in a Chinese individual. Hla, 2017, 90, 306-307.	0.4	1
1206	Confirmed the fullâ€length sequence of <i><scp>HLA</scp>â€B*44:03:02</i> by cloning and sequencing. Hla, 2017, 90, 125-127.	0.4	2
1207	Nomenclature for factors of the HLA system, update May 2017. Human Immunology, 2017, 78, 586-588.	1.2	0
1208	Nomenclature for factors of the HLA system, update January 2017. Human Immunology, 2017, 78, 451-454.	1.2	0
1209	Nomenclature for factors of the HLA system, update April 2017. Human Immunology, 2017, 78, 582-585.	1.2	0
1210	Nomenclature for factors of the HLA system, update June 2017. Human Immunology, 2017, 78, 589-593.	1.2	0
1211	Discovery of a novel <i><scp>HLA</scp>â€A*02</i> variant, <i><scp>HLA</scp>â€A*02:684,</i> in a Taiwanese individual. Hla, 2017, 90, 301-302.	0.4	1
1212	<i><scp>HLA</scp>â€A*24:02:01:09</i>, a new allele identified by sequenceâ€based typing in a Korean individual. Hla, 2017, 90, 304-305.	0.4	1
1213	Myocardial Tissue Engineering for Regenerative Applications. Current Cardiology Reports, 2017, 19, 78.	1.3	29
1214	Nomenclature for factors of the <scp>HLA</scp> system, updated on January 2017. International Journal of Immunogenetics, 2017, 44, 171-176.	0.8	0
1215	A novel allele <i><scp>HLA</scp>â€B*27:149</i> identified by sequenceâ€based typing in a Chinese individual. Hla, 2017, 90, 309-310.	0.4	1
1216	A novel <scp>HLA</scp>â€B*18 allele, <scp>HLA</scp>â€B*18:124, identified in a German volunteer bone marrow donor. Hla, 2017, 89, 56-57.	0.4	4
1217	Identification of <i>DPB1*581:01</i>, a novel HLAâ€DPB1 allele in a Chinese Mongolian individual. Hla, 2017, 89, 61-62.	0.4	4
1218	Nomenclature and Serology of HLA Class I and Class II Alleles. Current Protocols in Immunology, 2017, 118, A.1S.1-A.1S.6.	3.6	14
1219	<i><scp>HLAâ€DRB3</scp>*02:<scp>61Q</scp></i>, a novel <scp>HLAâ€DRB3</scp> allele identified in a volunteer bone marrow donor. Hla, 2017, 90, 186-187.	0.4	3

#	Article	IF	Citations
1220	*HLA–DQB1*05:144*, a novel allele, discovered in a Southeast Asian donor for stem cell transplantation. Hla, 2017, 90, 182-183.	0.4	4
1221	Nomenclature for factors of the <scp>HLA</scp> system, update March 2017. International Journal of Immunogenetics, 2017, 44, 187-193.	0.8	0
1222	Full–length sequences of 4 *<scp>HLA</scp>–B*15* alleles, *B*15:07:01:01*, *B*15:27:01*, *B*15:32:01* and *B*15:58*, confirmed by cloning and sequencing. Hla, 2017, 89, 313-317.	0.4	3
1223	Four new HLA class I alleles, *HLA–A*02:681*, *HLA–A*30:111*, *HLA–A*68:164* and *HLA–B*35:01:46*. Hla, 2017, 90, 174-175.	0.4	3
1224	Nomenclature for factors of the HLA system, update March 2017. Hla, 2017, 90, 70-76.	0.4	0
1225	Characterization of a novel *<scp>HLA</scp>–C*04* allele, *<scp>HLA</scp>–C*04:277*. Hla, 2017, 90, 315-316.	0.4	2
1226	A variant of *<scp>HLA–DRB1</scp>*11*, *<scp>HLA–DRB1</scp>*11:143*, discovered in a Chinese hematopoietic stem cell donor. Hla, 2017, 90, 321-323.	0.4	3
1227	Detection of a novel *<scp>HLA–DRB1</scp>*12* variant, *<scp>HLA–DRB1</scp>*12:67*, in a Taiwanese individual. Hla, 2017, 90, 323-324.	0.4	2
1228	Full–length sequences of 3 <scp>HLA</scp>–B alleles, *<scp>HLA–B</scp>*27:04:01*, *<scp>B</scp>*27:07:01* and *<scp>B</scp>*27:25*, confirmed by cloning and sequencing. Hla, 2017, 90, 40-43.	0.4	3
1229	Nomenclature for factors of the <scp>HLA</scp> system, update April 2017. Hla, 2017, 90, 188-192.	0.4	4
1230	Identification of a novel <scp>HLA</scp> allele, *<scp>HLA</scp>–B*15:416*, in a bone marrow hematopoietic stem cell donor. Hla, 2017, 90, 39-40.	0.4	3
1231	Full–length sequences of 2 <scp>HLA</scp>–B alleles, *B*48:03:01* and *B*48:04:01*, confirmed by cloning and sequencing. Hla, 2017, 90, 128-130.	0.4	2
1232	Identification of the new *<scp>HLA–DRB1</scp>*14:186* allele in an Italian bone marrow donor. Hla, 2017, 90, 183-184.	0.4	3
1233	Characterization of the novel *HLA–DQB1*05:155* allele by sequencing–based typing. Hla, 2017, 90, 377-378.	0.4	4
1234	Nomenclature for factors of the HLA system, update September 2017. Human Immunology, 2017, 78, 765-767.	1.2	0
1235	HLA class Ib in pregnancy and pregnancy-related disorders. Immunogenetics, 2017, 69, 581-595.	1.2	49
1236	Rheumatoid arthritis: Recent advances on its etiology, role of cytokines and pharmacotherapy. Biomedicine and Pharmacotherapy, 2017, 92, 615-633.	2.5	227
1237	Nomenclature for factors of the HLA system, update January 2017. Hla, 2017, 90, 56-61.	0.4	1

#	Article	IF	Citations
1238	Hapl-o-Mat: open-source software for HLA haplotype frequency estimation from ambiguous and heterogeneous data. BMC Bioinformatics, 2017, 18, 284.	1.2	36
1239	Genomic and genetic studies of systemic sclerosis: A systematic review. Human Immunology, 2017, 78, 153-165.	1.2	43
1240	HLA-DRB1 the notorious gene in the mosaic of autoimmunity. Immunologic Research, 2017, 65, 82-98.	1.3	101
1241	Comparison of exome-based HLA class I genotyping tools: identification of platform-specific genotyping errors. Journal of Human Genetics, 2017, 62, 397-405.	1.1	55
1242	*HLAâ€A*30:99* shows a new amino acid position 156 within *HLAâ€A*30* subtypes. Hla, 2017, 89, 53-54.	0.4	4
1243	Discovery of *HLAâ€B*51:209*, a novel *HLAâ€B*51* variant, in a Taiwanese individual. Hla, 2017, 89, 58-59.	0.4	3
1244	A novel allele, *HLAâ€DRB1*12:60N*, detected in a Taiwanese unrelated hematopoietic stem cell donor. Hla, 2017, 89, 65-66.	0.4	3
1245	The investigation of the origin of Southern Tunisians using HLA genes. Journal of Human Genetics, 2017, 62, 419-429.	1.1	15
1246	Nomenclature for factors of the <scp>HLA</scp> system, update May 2017. International Journal of Immunogenetics, 2017, 44, 239-242.	0.8	0
1247	Characterization of a novel *<scp>HLA</scp>â€B*39:01:01â€* related allele, *<scp>HLA</scp>â€B*39:130*, by cloning and phasing. International Journal of Immunogenetics, 2017, 44, 356-358.	0.8	3
1248	Nomenclature for factors of the HLA system, update August 2017. Human Immunology, 2017, 78, 762-764.	1.2	0
1249	Nomenclature for factors of the HLA system, update July 2017. Human Immunology, 2017, 78, 758-761.	1.2	0
1250	Nomenclature for factors of the HLA system, update September 2017. Hla, 2017, 90, 391-395.	0.4	6
1251	*<scp>HLAâ€B</scp>*67:01:03*, a <scp>HLA</scp> novel allele, identified in a Chinese platelet donor. Hla, 2017, 90, 373-374.	0.4	3
1252	A novel HLAâ€DRB1 allele, *HLAâ€DRB1*14:127:02*, detected in a Chinese hematopoietic stem cell donor. Hla, 2017, 90, 382-383.	0.4	3
1253	Nomenclature for factors of the HLA system, update August 2017. Hla, 2017, 90, 388-390.	0.4	2
1254	Nomenclature for factors of the <scp>HLA</scp> system, update April 2017. International Journal of Immunogenetics, 2017, 44, 234-238.	0.8	0
1255	Sequence identification, serological reactivity and family genetics of a novel <scp>HLA</scp> allele, *<scp>HLAâ€A</scp>*26:82*. Hla, 2017, 90, 308-309.	0.4	1

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1256	Nomenclature for factors of the HLA system, update July 2017. Hla, 2017, 90, 384-387.	0.4	1
1257	Genomic sequences of two novel <scp>HLAâ€C</scp> alleles, <i><scp>HLAâ€C</scp>*15:143</i> and <i><scp>HLAâ€C</scp>*07:109:02</i>. Hla, 2017, 90, 374-376.	0.4	3
1258	The Immune Epitope Database and Analysis Resource in Epitope Discovery and Synthetic Vaccine Design. Frontiers in Immunology, 2017, 8, 278.	2.2	369
1259	HLA-check: evaluating HLA data from SNP information. BMC Bioinformatics, 2017, 18, 334.	1.2	8
1260	Future Perspectives in HLA Typing Technologies. , 0, , .		3
1261	Characterization of two new HLAâ€B alleles, <i>HLAâ€B*07:299</i> and <i>HLAâ€B*35:350</i>. Hla, 2018, 91, 132-133.	0.4	2
1262	The novel allele, HLA-A*11:264 , was identified in a Chinese individual by sequence-based typing. Hla, 2018, 91, 533-534.	0.4	4
1263	Nomenclature for factors of the <scp>HLA</scp> system, update December 2017. International Journal of Immunogenetics, 2018, 45, 88-93.	0.8	1
1264	Characterization of the novel <i>HLAâ€B*48:43</i> allele by sequencingâ€based typing. Hla, 2018, 91, 139-140.	0.4	2
1265	A novel <i>HLAâ€B</i> allele: <i>HLAâ€B*40:241</i>. Hla, 2018, 91, 138-139.	0.4	2
1266	Nomenclature for factors of the HLA system, update December 2017. Human Immunology, 2018, 79, 251-254.	1.2	0
1267	The novel <i>HLAâ€DQB1*03:01:01:12</i> allele, identified by nextâ€generation sequencing in a Chinese family. Hla, 2018, 91, 221-222.	0.4	7
1268	A new <i>HLAâ€DQB1*04</i> allele, <i>HLAâ€DQB1*04:01:05,</i> identified in a Korean individual. Hla, 2018, 91, 312-313.	0.4	4
1269	Identification of the novel <i>HLAâ€C*07:530</i> allele by polymerase chain reaction sequenceâ€based typing. Hla, 2018, 91, 213-215.	0.4	5
1270	Identification of a novel HLAâ€C allele, <i>HLAâ€C*15:151</i>, in a Taiwanese individual. Hla, 2018, 91, 308-309.	0.4	3
1271	Characterization of the novel <i>HLAâ€C*16:116</i> allele by sequencingâ€based typing. Hla, 2018, 91, 309-311.	0.4	3
1272	Identification of the novel <i>HLAâ€C*03</i> allele, <i>HLAâ€C*03:03:35</i>. Hla, 2018, 91, 306-307.	0.4	3
1273	<i>HLAâ€A*30:125</i> shows a new antigen binding domain created from <i>HLAâ€A*30:01</i> and <i>HLAâ€A*30:02</i>. Hla, 2018, 91, 128-130.	0.4	2

#	ARTICLE	IF	CITATIONS
1274	A novel *HLA‐DQB1*03* variant, *HLA‐DQB1*03:23:03*, identified in a Chinese individual. Hla, 2018, 91, 541-542.	0.4	5
1275	Nomenclature for factors of the HLA system, update August 2017. International Journal of Immunogenetics, 2018, 45, 31-34.	0.8	0
1276	Nomenclature for factors of the HLA system, update February 2018. Human Immunology, 2018, 79, 511-515.	1.2	0
1277	Characterization of a novel allelic variant in HLA-B*40 lineage, HLA-B*40:298:02, by cloning and sequencing. International Journal of Immunogenetics, 2018, 45, 143-145.	0.8	3
1278	Nomenclature for factors of the HLA system, update March 2018. Human Immunology, 2018, 79, 516-525.	1.2	2
1279	Genetic Mechanisms Involved in the Generation of HLA Alleles in Brazilians: Description and Comparison of HLA Alleles. Transplantation Proceedings, 2018, 50, 835-840.	0.3	5
1280	Genomic sequences of <scp>HLA</scp>‐A*68:169, <scp>HLA</scp>‐B*07:298 and <scp>HLA</scp>‐B*39:129. International Journal of Immunogenetics, 2018, 45, 140-142.	0.8	3
1281	Detection of a novel *HLA‐DRB1*12* variant, *HLA‐DRB1*12:68*, in a Taiwanese individual. Hla, 2018, 91, 145-146.	0.4	2
1282	Characterization of a novel allele, HLA-C*02:135N, by full-length gene sequencing in a bone marrow donor. Hla, 2018, 91, 538-539.	0.4	3
1283	Identification of the novel HLA-B*51:230 allele in a Saudi individual. Hla, 2018, 92, 49-50.	0.4	3
1284	Identification of the novel *HLA‐B*08:01:01:02* allele in a Saudi individual. Hla, 2018, 92, 173-173.	0.4	3
1285	A novel null allele, HLA-DRB1*15:148N , identified in a Chinese family. Hla, 2018, 91, 544-545.	0.4	4
1286	Nomenclature for factors of the HLA system, update January 2018. Human Immunology, 2018, 79, 506-510.	1.2	0
1287	The full‐length genomic sequence of a novel *HLA‐C*07* allele, *HLA‐C*07:63*, identified in a Chinese individual. Hla, 2018, 91, 539-541.	0.4	3
1288	Immunopharmacogenomics towards personalized cancer immunotherapy targeting neoantigens. Cancer Science, 2018, 109, 542-549.	1.7	45
1289	Identification of the novel allele, HLA‐C*01:136 in a German cord blood donor originating from Azerbaijan. Hla, 2018, 91, 305-306.	0.4	3
1290	Identification of the new HLA‐DPB1 allele, DPB1*647:01, in an Italian patient with leukemia. Hla, 2018, 91, 311-312.	0.4	4
1291	Next‐generation sequencing‐based typing of a new allele *HLA‐DQB1*03:01:01:20* in a Chinese family. Hla, 2018, 91, 223-224.	0.4	6

#	Article	IF	Citations
1292	The novel null allele, *HLAâ€B*40:338N*, was identified in a Chinese leukemia patient. Hla, 2018, 91, 303-305.	0.4	4
1293	Characterization of the novel *HLAâ€B*40:01:31* allele. Hla, 2018, 91, 200-201.	0.4	3
1294	A new *HLAâ€C*05* allele, *HLAâ€C*05:156*, characterized by fullâ€length hemizygous sequencing. Hla, 2018, 91, 212-213.	0.4	3
1295	A novel allele, *HLAâ€B*51:220*, identified in an individual from south of Brazil. Hla, 2018, 91, 202-204.	0.4	4
1296	Comprehensive human leukocyte antigen genotyping of patients with type 1 diabetes mellitus in Taiwan. Pediatric Diabetes, 2018, 19, 699-706.	1.2	7
1297	Identification of a novel allele, *HLAâ€A*03:289*, by sequenceâ€based typing in Lithuanian patient. Hla, 2018, 91, 197-198.	0.4	4
1298	Risk of HLA Homozygous Cord Blood Transplantation: Implications for Induced Pluripotent Stem Cell Banking and Transplantation. Stem Cells Translational Medicine, 2018, 7, 173-179.	1.6	9
1299	A novel *HLAâ€A*02* allele, *A*02:692*, was identified by sequencingâ€based typing. Hla, 2018, 91, 59-60.	0.4	3
1300	Collection and storage of HLA NGS genotyping data for the 17th International HLA and Immunogenetics Workshop. Human Immunology, 2018, 79, 77-86.	1.2	16
1301	Identification of the novel *HLAâ€B*13:98* allele in a Chinese individual. Hla, 2018, 91, 133-134.	0.4	3
1302	Characterisation of the novel HLAâ€C null allele, *HLAâ€C*01:21N*, in Finnish individuals. Hla, 2018, 91, 146-147.	0.4	9
1303	The fullâ€length sequence of the HLAâ€C allele, *HLAâ€C*15:13:01:01*. Hla, 2018, 91, 216-220.	0.4	3
1304	Identification of the novel allele, *HLAâ€B*14:56*, in a Brazilian individual. Hla, 2018, 91, 199-200.	0.4	3
1305	Next-generation sequencing characterization of HLA in multi-generation families of Kuwaiti descent. Human Immunology, 2018, 79, 137-142.	1.2	5
1306	Nomenclature for factors of the <scp>HLA</scp> system, update September 2017. International Journal of Immunogenetics, 2018, 45, 35-39.	0.8	1
1307	More than 150 novel variants of HLA class I genes detected in German Stem Cell Donor Registry and UCLA International Cell Exchange samples. Hla, 2018, 91, 187-194.	0.4	15
1308	Sequencing of the novel *HLAâ€A*02:01:72* allele in a Chinese hematopoietic stem cell donor. Hla, 2018, 91, 56-57.	0.4	3
1309	The novel allele, *HLAâ€A*24:383*, was identified in a Chinese individual by sequenceâ€based typing. Hla, 2018, 91, 65-66.	0.4	3

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1310	Identification of the novel <i>HLAâ€B*27:103</i> allele in a Chinese bone marrow donor. Hla, 2018, 92, 175-176.	0.4	0
1311	Ancient DNA study reveals HLA susceptibility locus for leprosy in medieval Europeans. Nature Communications, 2018, 9, 1569.	5.8	67
1312	A novel HLA-A*31 allele, A*31:126N , was identified by sequence-based typing. Hla, 2018, 91, 534-535.	0.4	3
1313	Nomenclature for factors of the <scp>HLA</scp> system, update October 2017. International Journal of Immunogenetics, 2018, 45, 65-74.	0.8	0
1314	Nomenclature for factors of the <scp>HLA</scp> system, update January 2018. International Journal of Immunogenetics, 2018, 45, 152-159.	0.8	0
1315	Nomenclature for factors of the <scp>HLA</scp> system, update February 2018. International Journal of Immunogenetics, 2018, 45, 160-167.	0.8	1
1316	Discovery of <i>HLAâ€C*14:87</i>, a novel <i>HLAâ€C*14</i> variant, in a Taiwanese individual. Hla, 2018, 91, 72-73.	0.4	3
1317	Characterization of the novel HLAâ€B*07:305 allele by sequencingâ€based typing. Hla, 2018, 91, 296-297.	0.4	3
1318	The fullâ€length sequence of the HLAâ€C allele, <i>HLAâ€C*03:02:01</i>. Hla, 2018, 91, 204-207.	0.4	3
1319	Four novel alleles in Korean individuals, <i>HLAâ€B*40:323</i>, <i>DRB1*14:177</i>, <i>DQB1*03:200</i>, and <i>DQB1*06:205</i>. Hla, 2018, 91, 300-301.	0.4	4
1320	Identification of the novel <i>HLAâ€B*18:37:02</i> allele in a Croatian individual. Hla, 2018, 91, 299-300.	0.4	4
1321	Nomenclature for factors of the HLA system, update December 2017. Hla, 2018, 91, 248-252.	0.4	0
1322	Three novel HLA alleles discovered in Koreans, <i>HLAâ€A*26:118</i>, <i>DQB1*02:65</i> and <i>DPB1*05:01:07</i>. Hla, 2018, 91, 293-294.	0.4	4
1323	Nomenclature for factors of the <scp>HLA</scp> system, update November 2017. International Journal of Immunogenetics, 2018, 45, 75-87.	0.8	0
1324	The fullâ€length sequence of the HLAâ€C allele, <i>HLAâ€C*03:40:01</i>. Hla, 2018, 91, 208-211.	0.4	3
1325	Identification of the novel <i>HLAâ€DQB1*06:209</i> allele in a Chinese individual. Hla, 2018, 91, 543-544.	0.4	5
1326	Identification of a novel <i>HLAâ€C*12</i> allele, <i>HLAâ€C*12:221</i>. Hla, 2018, 91, 215-216.	0.4	3
1327	Nomenclature for factors of the HLA system, update November 2017. Hla, 2018, 91, 237-247.	0.4	0

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1328	A novel HLA class I allele: HLA-A*11:01:49. Hla, 2018, 91, 531-532.	0.4	3
1329	A novel HLA‐B allele, *HLA‐B*40:245* was identified in a patient with hepatitis B virus infection. Hla, 2018, 92, 52-53.	0.4	3
1330	*HLA‐A*02:01:01:28*, a novel HLA allele identified by next‐generation sequencing in a Chinese family. Hla, 2018, 91, 195-196.	0.4	5
1331	Identification of the novel allele, *HLA‐A*01:234*, in the mother of a German cord blood donor. Hla, 2018, 91, 530-531.	0.4	3
1332	Transcriptome sequencing of the long-nosed bandicoot (Perameles nasuta) reveals conservation and innovation of immune genes in the marsupial order Peramelemorphia. Immunogenetics, 2018, 70, 327-336.	1.2	3
1333	Inference from the stationary distribution of allele frequencies in a family of Wright‐Fisher models with two levels of genetic variability. Theoretical Population Biology, 2018, 122, 78-87.	0.5	1
1334	Characterization of the novel HLA‐A*24:391 allele by sequencing‐based typing. Hla, 2018, 91, 292-293.	0.4	3
1335	Identification of the novel *HLA‐B*40:333* allele by polymerase chain reaction sequence‐based typing. Hla, 2018, 91, 302-303.	0.4	3
1336	Detection of an *HLA‐B*15* variant, *HLA‐B*15:192*, in a Taiwanese individual. Hla, 2018, 91, 536-537.	0.4	3
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1342	Identification of the novel *HLA‐B*57:91* allele, by next‐generation sequencing in a Spanish individual. Hla, 2018, 91, 69-70.	0.4	3
1343	A novel HLA‐B allele, *HLA‐B*35:348*, was identified by sequencing‐based typing. Hla, 2018, 91, 68-69.	0.4	3
1344	Identification of a novel allele, *HLA‐A*02:189*, in a Chinese Han individual. Hla, 2018, 91, 58-59.	0.4	3
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1351	Identification of the novel *HLAâ€DQB1*03:181* allele in a Chinese leukemia patient. Hla, 2018, 91, 142-143.	0.4	3
1352	Identification of two new HLA class I alleles in Italian bone marrow donors: *A*31:125* and *B*44:269*. Hla, 2018, 91, 130-131.	0.4	2
1353	Sequenceâ€based typing identification of the novel *HLAâ€C*07:565* variant in a Chinese family. Hla, 2018, 91, 71-72.	0.4	3
1354	Identification of the novel *HLAâ€B*27:04:06* allele in a Chinese bone marrow donor. Hla, 2018, 91, 136-137.	0.4	3
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1356	A novel *HLAâ€DRB1*07* allele, *HLAâ€DRB1*07:01:22*, identified in a Chinese individual. Hla, 2018, 91, 143-144.	0.4	3
1357	*HLAâ€DQB1*02:102*, a novel allele identified by nextâ€generation sequencing in a Spanish individual. Hla, 2018, 91, 140-141.	0.4	3
1358	The novel *HLAâ€A*24:02* variant, *HLAâ€A*24:02:56*, identified by sequencing in a Chinese individual. Hla, 2018, 91, 63-65.	0.4	3
1359	Characterization of the novel *HLAâ€DPB1*763:01* allele by sequencingâ€based typing. Hla, 2018, 92, 429-431.	0.4	5
1360	The novel HLAâ€B allele, *HLA*â€*B*50:31*, was identified by sequencing genomic DNA. Hla, 2018, 92, 415-417.	0.4	4
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1365	Identification of a novel HLAâ€DPB1 allele, <i>HLAâ€DPB1*612:01</i>, in a Chinese individual. Hla, 2018, 92, 428-429.	0.4	5
1366	The HLA-C*06:66 allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2018, 92, 318-319.	0.4	2
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1372	<i>HLAâ€B*40:55</i>, an <i>HLAâ€B*40</i> variant, identified in Taiwanese individuals. Hla, 2018, 92, 50-51.	0.4	4
1373	Nomenclature for factors of the HLA system, update September 2018. Hla, 2018, 92, 354-371.	0.4	5
1374	Characterization of the novel HLAâ€DQA1*04:05 allele by sequencingâ€based typing. Hla, 2018, 93, 59-60.	0.4	4
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1378	The HLA-C*03:272 allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2018, 92, 316-318.	0.4	2
1379	Detection of an <i>HLAâ€B*58</i> variant, <i>HLAâ€B*58:19</i>, by sequenceâ€based typing in a Taiwanese individual. Hla, 2018, 92, 420-421.	0.4	3
1380	Nomenclature for factors of the HLA system, update July 2018. Hla, 2018, 92, 328-340.	0.4	1
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1393	HLA-B*40:330 allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2018, 92, 248-250.	0.4	3
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1395	The HLA-DRB1*09:29 allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2018, 92, 261-262.	0.4	4
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1402	Characterization of the novel <i>HLAâ€C*07:639</i> allele by sequencingâ€based typing. Hla, 2018, 92, 422-423.	0.4	4
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1404	The novel HLA-DRB1*15:158 allele discovered in a Taiwanese individual. Hla, 2018, 92, 264-265.	0.4	3
1405	The HLA-A*33:110 allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2018, 92, 242-243.	0.4	3
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1422	A novel HLA‐B variant, <i>HLA‐B*14:02:16</i>, discovered by amplicon‐based next‐generation sequencing. Hla, 2018, 92, 323-324.	0.4	4
1423	A novel <i>HLA‐A*24</i> allele, <i>A*24:231</i>, was identified by sequence‐based typing. Hla, 2018, 92, 96-97.	0.4	3
1424	Characterization of the novel <i>HLA‐DQB1*03:279</i> allele by sequencing‐based typing. Hla, 2018, 92, 63-64.	0.4	3
1425	Identification of a novel HLA-C allele, C*02:10:03 , in a Venezuelan patient. Hla, 2018, 92, 53-54.	0.4	3
1426	Identification of an HLA-B*27 variant, B*27:120 , by sequence-based typing in a Taiwanese bone marrow stem cell donor. Hla, 2018, 92, 47-49.	0.4	3
1427	The novel <i>HLA‐A*02:625</i> allele was identified in a Chinese bone marrow donor. Hla, 2018, 92, 94-95.	0.4	3
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1433	HLA-B*15:436 , a novel variant of HLA-B*15 , discovered in a Taiwanese individual. Hla, 2018, 92, 178-179.	0.4	1
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1444	A novel HLAâ€A null allele, <i>HLAâ€A*66:28N</i>. Hla, 2018, 92, 169-170.	0.4	2
1445	Identification of an HLAâ€A null allele, <i>HLAâ€A*33:129N</i>. Hla, 2018, 92, 243-244.	0.4	4
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1452	Discovery of HLA-B*58:92 , a novel HLA-B*58 variant, in a Taiwanese individual. Hla, 2018, 92, 182-183.	0.4	2
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1473	Nomenclature for factors of the <scp>HLA</scp> system, update May 2018. International Journal of Immunogenetics, 2018, 45, 290-296.	0.8	1
1474	Identification of a novel HLA allele, HLA-A*01:127 , in a donor from China. Hla, 2018, 92, 235-236.	0.4	4
1475	The new allele <i>HLAâ€DRB1*14:54:01:04</i> in a Chinese family identified using nextâ€generation sequencing. Hla, 2018, 92, 320-321.	0.4	2
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1482	Genetic Association in the HLA Region. Methods in Molecular Biology, 2018, 1793, 111-134.	0.4	9
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1486	The <i>HLAâ€DRB1*14:208</i> allele detected in a Taiwanese individual. Hla, 2019, 94, 460-461.	0.4	2
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1493	Detection of the novel <i>HLA‑A</i> allele, <i>HLA‑A*02:01:01:58</i>, in a Japanese individual. Hla, 2019, 94, 435-436.	0.4	2
1494	Discovery of the novel <i>HLA‑B</i> allele, <i>HLA‑B*51:01:01:36</i> in a Japanese individual. Hla, 2019, 94, 445-447.	0.4	2
1495	<i>HLA‑DQB1*03:01:01:21Q</i>, a novel HLA‑DQB1 allele, detected in a Spanish volunteer bone marrow donor. Hla, 2019, 94, 393-394.	0.4	4
1496	A novel HLA class I allele, <i>HLA‑B*46:01:21</i>, identified by sequence‑based typing in a Chinese individual. Hla, 2019, 94, 444-445.	0.4	2
1497	<i>HLA‑B*13:64</i>, a novel allele, identified in a Chinese individual. Hla, 2019, 94, 376-378.	0.4	3
1498	Characterization of the novel <i>HLA‑DPA1*01:20</i> allele by sequencing‑based typing. Hla, 2019, 94, 396-397.	0.4	2
1499	Identification of a novel <i>HLA‑A*11</i> allele, <i>HLA‑A*11:333</i>, in a Chinese individual. Hla, 2019, 94, 438-440.	0.4	2
1500	Identification of the novel <i>HLA‑C*07:02:107</i> allele in a volunteer donor from the China Marrow Donor Program. Hla, 2019, 94, 388-389.	0.4	4
1501	Characterization of the novel HLA‑C*07:724 allele by sequencing‑based typing. Hla, 2019, 94, 77-78.	0.4	4
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1504	Identification of the novel HLA‑B allele, <i>HLA‑B*15:539</i>, in a South‑Sudanese individual. Hla, 2019, 94, 380-381.	0.4	2
1505	Characterization of the novel <i>HLA‑DQA1*01:22</i> allele by sequencing‑based typing. Hla, 2019, 94, 333-334.	0.4	2
1506	Description of two new HLA alleles: <i>HLA‑A*30:118</i> and <i>HLA‑C*03:02:17</i>. Hla, 2019, 94, 371-373.	0.4	2
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1510	Nomenclature for factors of the HLA system, update April, May and June 2019. Hla, 2019, 94, 188-271.	0.4	5
1511	In a healthy Chinese blood donor, a novel allele, <i>HLAâ€Ɛ*01:03:07</i> was detected. Hla, 2019, 94, 389-392.	0.4	2
1512	A novel HLA allele, <i>HLAâ€B*56:67</i>, identified in a Melanesian individual from New Caledonia. Hla, 2019, 94, 384-386.	0.4	2
1513	Identification of a novel allele, <i>MICA*088N</i>, in a Chinese Kazakh individual. Hla, 2019, 94, 400-401.	0.4	2
1514	A novel HLAâ€B allele, <i>HLAâ€B*44:03:01:19</i>, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2019, 94, 381-382.	0.4	2
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1516	Identification of the novel <i>HLAâ€A*02:839</i> allele in a Brazilian candidate for bone marrow donation. Hla, 2019, 94, 365-366.	0.4	8
1517	Characterization of the novel HLAâ€DQB1*03:353 allele by sequencingâ€based typing. Hla, 2019, 94, 86-87.	0.4	4
1518	A novel <i>HLAâ€A*24:460</i> allele identified in a hematopoietic stem cell transplantation donor and recipient. Hla, 2019, 94, 368-369.	0.4	2
1519	A novel HLAâ€B allele, <i>HLAâ€B*08:242</i>, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2019, 94, 375-376.	0.4	2
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1521	Characterization of the novel <i>HLAâ€DPA1*02:15</i> allele by sequencingâ€based typing. Hla, 2019, 94, 179-180.	0.4	2
1522	The novel <i>HLAâ€C*04:387</i> allele is likely generated by intraâ€locus recombination event. Hla, 2019, 94, 535-537.	0.4	2
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1524	Characterization of an <i>HLAâ€B*55:01:01</i> variant, <i>HLAâ€B*55:01:01:14</i>, identified in a Brazilian individual. Hla, 2019, 94, 449-450.	0.4	2
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1526	Three new HLA alleles, <i>HLAâ€A*02:01:01:126</i>, â€<i>B*35:01:01:40</i>, and <i>â€B*44:02:01:37</i> detected in Spanish donors. Hla, 2019, 94, 514-515.	0.4	2

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1528	Characterization of four novel HLA‑DPA1*01:03:01 variants, identified in Brazilian individuals. Hla, 2019, 94, 546-547.	0.4	2
1529	Recognition of the novel <i>HLA‑A*11:01:01:25</i> allele in the tribal population of Nagaland, North‑East India. Hla, 2019, 94, 515-517.	0.4	2
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1540	Identification of <i>HLA‑A*24:02:78</i> by next‑generation sequencing in a Chinese Han individual. Hla, 2019, 94, 519-521.	0.4	2
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1544	Description of a novel HLA null allele, <i>DRB1*15:176N</i>. Hla, 2019, 94, 462-463.	0.4	2
1545	Identification of a novel recombinant allele, <i>HLA‑DPB1*835:01:01:02</i>, in Black South African individuals. Hla, 2019, 94, 549-551.	0.4	3

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1547	<i>HLAâ€DRB1*11:259</i> allele detected in a Taiwanese individual. Hla, 2019, 94, 540-542.	0.4	2
1548	Characterization of three novel HLAâ€DQA1*03:03:01 variants, identified in Brazilian individuals. Hla, 2019, 94, 542-543.	0.4	2
1549	Characterization of the novel <i>HLAâ€DQB1*06:01:22</i> allele by nextâ€generation sequencing. Hla, 2019, 94, 543-545.	0.4	2
1550	Characterization of an <i>HLAâ€B*15:10:01</i> variant, <i>HLAâ€B*15:10:01:05</i>, identified in a Brazilian individual. Hla, 2019, 94, 528-529.	0.4	2
1551	Identification of the novel HLAâ€B allele, <i>HLAâ€B*27:198</i>, from a Korean individual. Hla, 2019, 94, 530-531.	0.4	2
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1568	Identification of a novel HLAâ€C allele, HLAâ€C*16:123N , in an Emirati blood donor. Hla, 2019, 93, 505-506.	0.4	5
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1576	Sequenceâ€based typing identification of a novel allele <i>HLAâ€A*30:115</i>. Hla, 2019, 94, 314-315.	0.4	2
1577	<i>HLAâ€C*08:01:25</i>, a novel HLA allele, which has arisen by a silent mutation in codon 271. Hla, 2019, 94, 162-163.	0.4	2
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1590	Detection of an ‹i›HLAâ€‹A*33‹/i› variant, ‹i›HLAâ€‹A*33:176N‹/i›, in a Taiwanese individual. Hla, 2019, 94, 316-317.	0.4	2
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1592	Characterization of two new HLAâ€‹C alleles, ‹i›HLAâ€‹C*05:01:01:17‹/i› and â€‹‹i›C*16:152‹/i›. Hla, 2019, 94, 159-160.	0.4	2
1593	Characterization of the novel ‹i›HLAâ€‹DQA1*01:25‹/i› allele by sequencingâ€‹based typing. Hla, 2019, 94, 174-175.	0.4	2
1594	The novel HLAâ€‹DRB1*04:01:10 allele detected in a Taiwanese individual. Hla, 2019, 94, 83-84.	0.4	4
1595	The ‹i›HLAâ€‹DRB1*04:90‹/i› allele recognized in a Taiwanese individual. Hla, 2019, 94, 167-168.	0.4	2
1596	Identification of the new ‹i›HLAâ€‹C*16:02:17‹/i› allele in a Brazilian candidate donor for bone marrow donation. Hla, 2019, 94, 332-333.	0.4	2
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1610	Identification of the novel HLAâ€C*04:286 allele by nextâ€generation sequencing in a Chinese cord blood donor. Hla, 2019, 94, 73-74.	0.4	4
1611	Detection of an HLAâ€A*02 variant, HLAâ€A*02:474 , in a Taiwanese individual. Hla, 2019, 94, 57-58.	0.4	4
1612	Characterization of the novel <i>HLAâ€A*11:324</i> allele by sequencingâ€based typing. Hla, 2019, 94, 155-156.	0.4	2
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1628	Characterization of the novel <i>HLAâ€DRB3*03:01:07</i> allele by sequencingâ€based typing. Hla, 2019, 93, 240-241.	0.4	2
1629	Identification of three new HLA alleles in the Spanish population: <i>HLAâ€C*05:203</i>, <i>C*15:10:04</i> and <i>DRB1*01:99</i>. Hla, 2019, 93, 234-235.	0.4	2
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1631	Identification of a novel <i>HLAâ€B*40</i> variant allele, <i>B*40:405</i> by nextâ€generation sequencing. Hla, 2019, 93, 491-492.	0.4	5
1632	A novel HLAâ€B allele: HLAâ€B*58:01:13. Hla, 2019, 93, 492-493.	0.4	5
1633	Characterization of the novel HLAâ€C*08:159 allele by nextâ€generation sequencing. Hla, 2019, 93, 502-503.	0.4	6
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1635	Detection of an HLAâ€C*03 variant, HLAâ€C*03:258 , in a Taiwanese individual. Hla, 2019, 93, 125-127.	0.4	3
1636	Identification of a novel allele, HLAâ€DPB1*835:01 , in an African American renal transplant candidate. Hla, 2019, 93, 245-245.	0.4	4
1637	Identification of the novel HLAâ€B allele, HLAâ€B*52:56 by sequenceâ€based typing in a Chinese individual. Hla, 2019, 93, 226-227.	0.4	2

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1639	Characterization of the novel HLAâ€C*07:708 allele by sequencingâ€based typing. Hla, 2019, 93, 235-236.	0.4	2
1640	Detection of <i>HLAâ€C*08:147</i>, an <i>HLAâ€C*08</i> variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2019, 93, 237-238.	0.4	2
1641	Characterization of the novel <i>HLAâ€DQA1*05:14</i> allele by sequencingâ€based typing. Hla, 2019, 93, 241-243.	0.4	2
1642	Characterization of the novel <i>HLAâ€DPB1*896:01</i> allele by sequencingâ€based typing. Hla, 2019, 93, 246-247.	0.4	2
1643	Detection of an <i>HLAâ€C*03:04</i> variant, <i>HLAâ€C*03:04:71</i>, in a Taiwanese individual. Hla, 2019, 93, 229-230.	0.4	3
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1647	Characterization of the novel HLAâ€DRB1*14:207 allele by sequencingâ€based typing. Hla, 2019, 94, 85-86.	0.4	4
1648	The novel HLAâ€A*03 variant, HLAâ€A*03:08:01:02 , detected by sequencingâ€based typing. Hla, 2019, 94, 60-61.	0.4	4
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1652	Characterization of the novel <i>HLAâ€C*03:372</i> allele by nextâ€generation sequencing. Hla, 2019, 94, 71-73.	0.4	5
1653	Characterization of the novel HLAâ€B*07:02:73 allele by sequencingâ€based typing. Hla, 2019, 94, 65-66.	0.4	4
1654	Identification of the novel <i>HLAâ€C*12:220</i> allele in a Chinese individual. Hla, 2019, 94, 80-81.	0.4	5
1655	Identification of two novel HLA class II alleles, HLAâ€DQB1*02:139 and HLAâ€DQB1*02:140 , in northern Chinese individuals. Hla, 2019, 93, 508-510.	0.4	5

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1657	Characterization of the novel HLAâ€B*51:228 allele in a Chinese individual. Hla, 2019, 93, 120-122.	0.4	2
1658	Tools for building, analyzing and evaluating HLA haplotypes from families. Human Immunology, 2019, 80, 633-643.	1.2	11
1659	Characterization of the novel HLAâ€B*46:01:23 allele in a Chinese bone marrow donor. Hla, 2019, 93, 224-226.	0.4	2
1660	Detection of a novel <i>HLAâ€A*11</i> variant, <i>A*11:292</i>, in a Taiwanese individual. Hla, 2019, 93, 40-41.	0.4	2
1661	Identification of the novel HLAâ€C*03:365 allele in a Chinese bone marrow donor. Hla, 2019, 93, 231-232.	0.4	4
1662	Description of four new HLA alleles: <i>HLAâ€A*01:288</i>, â€<i>A*02:06:23</i>, â€<i>A*32:121</i> and â€<i>DRB1*07:100</i>. Hla, 2019, 93, 220-221.	0.4	3
1663	Identification of the novel HLAâ€A*01:289 allele in an Italian patient. Hla, 2019, 93, 484-485.	0.4	5
1664	Characterization of the novel HLAâ€C*08:154 allele by sequencingâ€based typing. Hla, 2019, 93, 238-240.	0.4	5
1665	Discovery of HLAâ€B*58:74 , a novel HLAâ€B*58 variant, in Singaporean Chinese individuals. Hla, 2019, 93, 496-497.	0.4	4
1666	Two novel HLA alleles, HLAâ€C*02:163 and HLAâ€C*04:348 , identified in Russian individuals. Hla, 2019, 93, 228-229.	0.4	2
1667	Characterization of the novel HLAâ€A*33:170 allele by sequencingâ€based typing. Hla, 2019, 93, 221-223.	0.4	2
1668	The novel HLAâ€C*08:80 allele identified by fullâ€length sequencing of the HLA region. Hla, 2019, 93, 500-501.	0.4	5
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1670	Characterization of the novel HLAâ€B*15:271 allele by sequencingâ€based typing in a Chinese bone marrow donor. Hla, 2019, 93, 106-107.	0.4	3
1671	Identification of the novel HLAâ€A*29:02:01:06 allele in a Saudi individual. Hla, 2019, 93, 102-103.	0.4	2
1672	Identification of the novel HLAâ€B*35:01:01:16 allele in a Saudi individual. Hla, 2019, 93, 111-111.	0.4	2
1673	HLAâ€DQB1*06:276 , a novel HLA allele fund in a patient from Guineaâ€Bissau. Hla, 2019, 93, 243-244.	0.4	3

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1676	Characterization of seven new HLA alleles, <i>A*24:14:01:04</i>, <i>A*29:02:01:07</i>, <i>C*06:02:01:37</i>, <i>C*07:830</i>, <i>C*16:162</i>, <i>C*16:01:01:07</i> and <i>DQA1*01:02:05</i>. Hla, 2019, 94, 521-522.	0.4	12
1677	A novel allele, <i>HLAâ€C*07:01:01:30</i> identified using thirdâ€generation sequencing. Hla, 2019, 94, 455-456.	0.4	2
1678	<i>HLAâ€DQB1*05:217</i>, an <i>HLAâ€DQB1*05:01:01:01</i> variant, identified in a Taiwanese individual. Hla, 2019, 94, 178-179.	0.4	7
1679	<i>HLAâ€A*33:03:01:06</i> allele identified in an individual from the Kashmiri Brahmin population of North India. Hla, 2019, 94, 524-526.	0.4	4
1680	Histocompatibility in Renal Transplantation. , 2019, , 139-156.		0
1681	Nomenclature for factors of the HLA system, update July, August and September 2019. Hla, 2019, 94, 552-624.	0.4	11
1682	Identification of a novel HLAâ€C allele, <i>HLAâ€C*07:778</i>, in an unrelated hematopoietic stem cell donor. Hla, 2019, 94, 161-162.	0.4	3
1683	<i>HLAâ€A*11:01:46</i>, a novel <i>HLAâ€A*11</i> variant, detected in a Chinese individual. Hla, 2019, 94, 517-519.	0.4	3
1684	Characterization of the first <i>HLAâ€B*15:31</i> variant, <i>B*15:31:01:02</i>, found in a Brazilian individual. Hla, 2019, 94, 529-530.	0.4	3
1685	Validation of new HLAâ€F alleles assigned by nextâ€generation sequencing. Hla, 2019, 93, 131-132.	0.4	20
1686	Direct HLA Genetic Comparisons Identify Highly Matched Unrelated Donor-Recipient Pairs with Improved Transplantation Outcome. Biology of Blood and Marrow Transplantation, 2019, 25, 921-931.	2.0	21
1687	Characterization of the novel HLA-DRB5*02:21 allele by sequencing-based typing. Hla, 2019, 93, 58-59.	0.4	3
1688	Characterization of a novel <i>MICB</i> variant in an individual from the Chinese Uyghur population, <i>MICB*005:09</i>, by cloning and sequencing. International Journal of Immunogenetics, 2019, 46, 17-19.	0.8	2
1689	Identification of the new <i>HLAâ€DRB5*01:01:01:02</i> allele in a Chinese individual. Hla, 2019, 93, 56-57.	0.4	3
1690	Allelic nomenclature for the duplicated MHC class II DQ genes in sheep. Immunogenetics, 2019, 71, 347-351.	1.2	4
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1693	Identification of the novel HLAâ€A*30:02:01:04 allele in a Saudi individual. Hla, 2019, 93, 103-104.	0.4	3
1694	Characterization of the novel <i>HLAâ€A*30:135</i> allele by sequencingâ€based typing. Hla, 2019, 93, 46-47.	0.4	3
1695	Characterization of the novel HLA-A*03:315 allele by sequencing-based typing. Hla, 2019, 93, 39-40.	0.4	3
1696	Assessing the utilization of high-resolution 2-field HLA typing in solid organ transplantation. American Journal of Transplantation, 2019, 19, 1955-1963.	2.6	39
1697	Identification of the novel HLA allele, HLAâ€B*35:03:23 , by nextâ€generation sequencing. Hla, 2019, 93, 112-113.	0.4	3
1698	Characterization of the novel HLAâ€DRB1*11:245 allele by sequencingâ€based typing. Hla, 2019, 93, 133-134.	0.4	2
1699	Identification of the novel HLAâ€C*04:01:01:31 allele in a Saudi individual. Hla, 2019, 93, 127-128.	0.4	2
1700	The new HLAâ€C*05:199 was generated by intralocus recombination involving C*05:01:01:01 and C*16:01:01:01 alleles. Hla, 2019, 93, 128-130.	0.4	3
1701	The novel HLAâ€DRB1*15:01:37 allele discovered in a Taiwanese individual. Hla, 2019, 93, 134-135.	0.4	3
1702	Identification of the novel HLAâ€C*02:151 allele in Russian bone marrow donors. Hla, 2019, 93, 124-125.	0.4	3
1703	Identification of the novel HLAâ€A*31:124 allele by sequenceâ€based typing in a Chinese cord blood donor. Hla, 2019, 93, 104-105.	0.4	2
1704	Human leukocyte antigen polymorphisms and Kaposi's sarcomaâ€associated herpesvirus infection outcomes: A call for deeper exploration. Journal of Medical Virology, 2019, 91, 541-548.	2.5	7
1705	Identification of the novel HLAâ€DQB1*05:01:01:05 allele in a Saudi individual. Hla, 2019, 93, 137-138.	0.4	3
1706	Identification of the novel HLAâ€B*49:01:01:04 allele in a Saudi individual. Hla, 2019, 93, 119-120.	0.4	2
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1713	Recognition of <i>HLAâ€C*03:36</i> allele in a Taiwanese individual. Hla, 2020, 95, 56-57.	0.4	2
1714	<i>HLAâ€A*11:01:01:16, â€B*15:548, â€DPA1*02:02:02:06Q, â€DPA1*02:02:08</i>, and <i>â€DPA1*02:25</i>, identified in Chinese Mongolians. Hla, 2020, 95, 45-47.	0.4	4
1715	A novel <i>HLAâ€A*33</i> allele, <i>HLAâ€A*33:131</i>, identified by sequencingâ€based typing. Hla, 2020, 95, 133-134.	0.4	2
1716	<i>HLAâ€DQB1*03:13</i> identified in two unrelated Taiwanese individuals. Hla, 2020, 95, 76-78.	0.4	2
1717	Injectable Ovalbuminâ€Based Composite Implant for Photothermal Tumor Therapy. ChemBioChem, 2020, 21, 865-873.	1.3	3
1718	Characterization of the new <i>HLAâ€DQB1*06:344</i> allele by nextâ€generation sequencing in a Korean individual. Hla, 2020, 95, 155-156.	0.4	2
1719	The novel <i>HLAâ€B*58:118</i> allele was identified during highâ€resolution HLA typing. Hla, 2020, 95, 214-215.	0.4	2
1720	The novel <i>HLAâ€DQB1*06:358</i> allele, identified by Nextâ€Generation Sequencing in a Saudi individual. Hla, 2020, 95, 157-158.	0.4	2
1721	Identification of the novel <i>HLAâ€B*46:78</i> allele by next generation sequencing in a Chinese individual. Hla, 2020, 95, 137-138.	0.4	2
1722	Characterization of the novel <i>HLAâ€DRB3*03:37</i> allele by sequencingâ€based typing. Hla, 2020, 95, 152-153.	0.4	2
1723	Description of the novel null allele, <i>HLAâ€DPA1*01:29N</i> in a Lotha individual from Nagaland, Northâ€East India. Hla, 2020, 95, 82-83.	0.4	2
1724	An Illumina approach to MHC typing of Atlantic salmon. Immunogenetics, 2020, 72, 89-100.	1.2	7
1725	A novel HLAâ€E allele, <i>HLAâ€E*01:12:01:01</i>, identified in a healthy Chinese blood donor. Hla, 2020, 95, 144-146.	0.4	5
1726	<i>HLAâ€C*06:287</i>, a novel <i>HLAâ€C*06</i> allele identified by sequenceâ€based typing. Hla, 2020, 95, 64-65.	0.4	2
1727	Characterization of an <i>HLAâ€C*03:04:01</i> variant, <i>HLAâ€C*03:04:01:39</i>, identified in a Brazilian individual. Hla, 2020, 95, 55-56.	0.4	2

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1729	Characterization of the novel allele <i>HLAâ€DPA1*02:01:01:12</i> from an Ao tribal individual from Nagaland, Northâ€East India. Hla, 2020, 95, 86-87.	0.4	2
1730	Characterization of two novel HLAâ€DQA1*05:05:01 variants, identified in Brazilian individuals. Hla, 2020, 95, 75-76.	0.4	1
1731	<i>HLAâ€C*07:06:01:06</i>â€"A new HLAâ€C allele identified in an Ao tribal individual from Nagaland, <i>Northâ€East India</i>. Hla, 2020, 95, 65-66.	0.4	2
1732	Identification of the novel HLA allele, <i>HLAâ€C*07:780</i>, identified in a Danish woman. Hla, 2020, 95, 69-71.	0.4	2
1733	Characterization of the first <i>HLAâ€DPA1*04:02</i> variant, <i>DPA1*04:02:01:02</i>, found in a Brazilian individual. Hla, 2020, 95, 90-91.	0.4	2
1734	Identification of the novel allele, <i>HLAâ€A*26:193</i>, in a patient with Myelodysplastic Syndrome. Hla, 2020, 95, 48-49.	0.4	2
1735	<i>HLAâ€C*04:01:01:78</i>, a novel variant of <i>HLAâ€C*04</i>, detected in a Sumi individual from Nagaland, Northâ€East India. Hla, 2020, 95, 58-59.	0.4	4
1736	Simultaneous confidence corridors for mean functions in functional data analysis of imaging data. Biometrics, 2020, 76, 427-437.	0.8	10
1737	Identification of <i>HLAâ€C*16:46</i> allele in a Kuwaiti individual. Hla, 2020, 95, 71-72.	0.4	3
1738	A novel allele <i>HLAâ€B*15:02:01:04</i>, recognized in an individual from the Angami tribe, Nagaland, Northâ€East India. Hla, 2020, 95, 49-50.	0.4	2
1739	Discovery of the novel allele, <i>HLAâ€C*03:04:01:38</i> in two individuals from the Ao tribe of Nagaland, <i>Northâ€East India</i>. Hla, 2020, 95, 53-54.	0.4	4
1740	The IPD Project: a centralised resource for the study of polymorphism in genes of the immune system. Immunogenetics, 2020, 72, 49-55.	1.2	27
1741	Peptideâ€MHC class I and class II tetramers: From flow to mass cytometry. Hla, 2020, 95, 169-178.	0.4	17
1742	Characterization of the novel <i>HLAâ€DPA1*01:03:16</i> allele by sequencingâ€based typing. Hla, 2020, 95, 158-159.	0.4	2
1743	Detection of an <i>HLAâ€C*03</i> variant, <i>HLAâ€C*03:85</i>, in a Taiwanese individual. Hla, 2020, 95, 216-217.	0.4	2
1744	Characterization of the novel <i>HLAâ€DRB1*15:178</i> allele by sequencingâ€based typing. Hla, 2020, 95, 149-150.	0.4	2
1745	Detection of an <i>HLAâ€DRB1*14</i> variant, <i>HLAâ€DRB1*14:141</i>, in a Taiwanese individual. Hla, 2020, 95, 579-580.	0.4	2

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1746	Nomenclature for factors of the HLA system, update July, August and September 2019. International Journal of Immunogenetics, 2020, 47, 67-138.	0.8	0
1747	The novel <i>HLAâ€A*26:01:64</i> allele, identified by nextâ€generation sequencing in a bone marrow donor of Bashkir origin. Hla, 2020, 95, 484-485.	0.4	2
1748	Discovery of the novel HLA allele, <i>HLAâ€A*30:72</i>, in a Chinese platelet donor by sequenceâ€based typing. Hla, 2020, 95, 131-133.	0.4	3
1749	A novel HLAâ€E allele, <i>HLAâ€E*01:03:01:05</i>, identified in a healthy Chinese blood donor. Hla, 2020, 95, 222-224.	0.4	2
1750	Revealing factors determining immunodominant responses against dominant epitopes. Immunogenetics, 2020, 72, 109-118.	1.2	20
1751	Characterization of the novel <i>HLAâ€DPB1*04:01:42</i> allele by sequencingâ€based typing. Hla, 2020, 95, 161-163.	0.4	2
1752	Characterization of an <i>HLAâ€C*04:01:01</i> variant, <i>HLAâ€C*04:01:01:76</i>, identified in a Brazilian individual. Hla, 2020, 95, 141-142.	0.4	2
1753	Identification of a novel <i>HLAâ€C*07</i> allele in a Russian individual: <i>HLAâ€C*07:839N</i>. Hla, 2020, 95, 142-143.	0.4	2
1754	Characterization of the novel <i>HLAâ€DRB1*12:82</i> allele by sequencingâ€based typing. Hla, 2020, 95, 147-148.	0.4	2
1755	Characterization and confirmation of the <i>HLAâ€DQA1*04:01:01:09</i> allele, identified in a Brazilian individual. Hla, 2020, 95, 153-154.	0.4	2
1756	Characterization of the novel <i>HLAâ€DRB3*02:02:23</i> allele by sequencingâ€based typing. Hla, 2020, 95, 150-151.	0.4	2
1757	Identification of a novel allele, <i>HLAâ€B*52:01:24</i>, by sequenceâ€based typing in a Chinese individual. Hla, 2020, 95, 139-141.	0.4	2
1758	Identification of the novel <i>HLAâ€A*30:162</i> allele in a Brazilian candidate donor for bone marrow donation. Hla, 2020, 95, 208-209.	0.4	2
1759	Identification of the novel <i>HLAâ€DQA1*02:11</i> allele in a Brazilian candidate donor for bone marrow donation. Hla, 2020, 95, 228-229.	0.4	2
1760	A novel allele <i>HLAâ€B*38:02:01:02</i>, identified in an Angami individual from Nagaland, Northâ€East India. Hla, 2020, 95, 51-52.	0.4	2
1761	<i>HLAâ€DQB1*04:02:13</i>, an <i>HLAâ€DQB1*04:02</i> variant, identified in a Kuwaiti individual. Hla, 2020, 95, 80-81.	0.4	2
1762	<i>HLAâ€DPA1*02:01:01:13</i>, a novel HLAâ€DPA1 allele, detected in an Ao tribal individual from Nagaland, Northâ€East India. Hla, 2020, 95, 88-90.	0.4	4
1763	The novel <i>HLAâ€C*04:400</i> allele, identified in a Konyakâ€Naga tribal individual from Northâ€East India. Hla, 2020, 95, 62-63.	0.4	2

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1764	Detection of <i>HLA‑C*07:27:01</i> in a Taiwanese individual. Hla, 2020, 95, 67-68.	0.4	3
1765	Detection of an <i>HLA‑B*15</i> variant, <i>HLA‑B*15:109</i>, in a Taiwanese individual. Hla, 2020, 95, 135-136.	0.4	2
1766	Characterization of the novel <i>HLA‑DPA1*02:26</i> allele by sequencing‑based typing. Hla, 2020, 95, 160-161.	0.4	2
1767	Detection of an <i>HLA‑A*30</i> variant, <i>HLA‑A*30:114</i>, in a Taiwanese individual. Hla, 2020, 95, 206-207.	0.4	2
1768	Characterization of two novel <i>HLA‑DQA1*05:05:01</i> variants, identified in Brazilian individuals. Hla, 2020, 95, 230-231.	0.4	2
1769	<i>HLA‑C*07:154</i>, an <i>HLA‑C*07</i> variant, detected in a Taiwanese blood donor. Hla, 2020, 95, 218-219.	0.4	3
1770	Characterization of the novel HLA‑A*26:199 allele by sequencing‑based typing. Hla, 2020, 96, 499-500.	0.4	6
1771	Nomenclature for factors of the <scp>HLA</scp> system, update April, May, and June 2020. Hla, 2020, 96, 384-412.	0.4	16
1772	HLA‑A *02:411 identified in a platelet donor from China. Hla, 2020, 96, 491-493.	0.4	7
1773	Novel <scp><i>HLA‑B*81:02:02</i></scp> allele identified in a Saudi individual. Hla, 2020, 96, 644-645.	0.4	3
1774	Identification of <i><scp>HLA‑A</scp>*31:73</i> in a platelet donor from China by sequence‑based typing. Hla, 2020, 96, 628-631.	0.4	3
1775	The novel <scp><i>HLA‑B</i></scp><i>*57:135</i> allele was identified during high‑resolution <scp>HLA</scp> typing. Hla, 2020, 96, 642-644.	0.4	4
1776	Nomenclature for factors of the HLA system, update April, May and June 2020. International Journal of Immunogenetics, 2020, 47, 456-483.	0.8	6
1777	Detection of the novel <scp>HLA‑B</scp> allele, <scp><i>HLA‑B*27:199</i></scp>, in a Korean individual. Hla, 2020, 96, 345-347.	0.4	6
1778	A novel <i><scp>HLA‑B</scp>*52</i> allele, <i>B*52:100</i>, was identified by sequencing‑based typing. Hla, 2020, 96, 522-523.	0.4	7
1779	Characterization of a novel <scp>HLA</scp> allele, <i><scp>DPB1</scp>*1049:01</i>, in a pediatric patient with severe aplastic anemia. Hla, 2020, 96, 758-759.	0.4	3
1780	An immunoinformatics study on the spike protein of SARS-CoV-2 revealing potential epitopes as vaccine candidates. Heliyon, 2020, 6, e04865.	1.4	8
1781	The <scp><i>HLA‑A*33:03:42</i></scp> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 96, 334-335.	0.4	6

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1782	The novel <scp><i>HLAâ€C*07:02:01:97</i></scp> allele identified in a Chang tribal individual from <scp>Nagaland</scp>, <scp>Northâ€East India</scp>. Hla, 2020, 96, 530-531.	0.4	2
1783	The novel <scp><i>HLAâ€DQB1*06:03:01:06</i></scp> allele identified in a <scp>Saudi</scp> individual. Hla, 2020, 96, 661-662.	0.4	3
1784	Identification of a novel <scp>HLAâ€DQB1</scp> allele, <scp><i>HLAâ€DQB1*03:168</i></scp>, by sequenceâ€based typing in a Taiwanese individual. Hla, 2020, 96, 546-547.	0.4	6
1785	The discovery of the first <scp><i>HLAâ€DQA1*01:01:02</i></scp> variant, <scp><i>DQA1*01:01:02:02</i></scp>, found in a Brazilian individual. Hla, 2020, 96, 115-116.	0.4	3
1786	Identification of the novel <i>HLAâ€A*02</i> allele, <i>HLAâ€A*02:725</i>. Hla, 2020, 95, 476-478.	0.4	2
1787	The discovery of two <scp><i>HLAâ€DPA1</i></scp><i>*02:01:01</i> variants, found in Brazilian individuals. Hla, 2020, 96, 555-556.	0.4	6
1788	Characterization of the novel <i><scp>HLAâ€C</scp>*07:862</i> allele in a North Indian hematopoietic stem cell donor. Hla, 2020, 96, 532-533.	0.4	6
1789	Novel and extended HLA class I and II alleles encountered in Kashmiri Brahmin population from North India. Hla, 2020, 96, 487-489.	0.4	9
1790	Detection of the <i><scp>HLAâ€DRB4</scp>*01:44</i> allele in a Kuwaiti individual. Hla, 2020, 96, 537-539.	0.4	7
1791	Critical Review of Existing MHC I Immunopeptidome Isolation Methods. Molecules, 2020, 25, 5409.	1.7	15
1792	Characterization of the novel <scp><i>HLAâ€B*07:381</i></scp> allele by nextâ€generation sequencing. Hla, 2020, 96, 726-727.	0.4	3
1793	Characterization of the novel <scp><i>HLAâ€A*11:349</i></scp> allele by nextâ€generation sequencing. Hla, 2020, 96, 714-715.	0.4	3
1794	Characterization of the novel <scp><i>HLAâ€B*07:385</i></scp> allele by nextâ€generation sequencing. Hla, 2020, 96, 727-728.	0.4	3
1795	Detection of the novel HLAâ€B *51:232:02 variant in an inhabitant from the island of Crete. Hla, 2020, 97, 364-366.	0.4	4
1796	Identification of three new <scp>HLAâ€DQB1</scp> alleles: <i><scp>HLAâ€DQB1</scp>*03:02:33, â€<scp>DQB1</scp>*04:79</i> and <i>â€<scp>DQB1</scp>*05:249</i>. Hla, 2021, 97, 378-380.	0.4	3
1797	Characterization of the novel <scp><i>HLAâ€C*14:115</i></scp> allele by nextâ€generation sequencing. Hla, 2020, 96, 737-739.	0.4	3
1798	Two novel HLAâ€A alleles, <i>HLAâ€A*03:399</i> and <i>â€A*24:17:01:02</i>, detected in inhabitants from the island of Crete. Hla, 2021, 97, 353-356.	0.4	4
1799	Characterization of the novel <scp><i>HLAâ€C*03:489</i></scp> allele by nextâ€generation sequencing. Hla, 2020, 96, 732-733.	0.4	3

#	ARTICLE	IF	CITATIONS
1800	Characterization of the novel <i><scp>HLAâ€C</scp>*06:283</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 734-735.	0.4	3
1801	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*03:<scp>400N</scp></i> allele by nextâ€generation sequencing. Hla, 2020, 96, 749-750.	0.4	3
1802	Characterization of the novel <i><scp>HLAâ€B</scp>*15:474</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 729-730.	0.4	3
1803	Two novel <scp><i>HLAâ€DQA1</i></scp><i>*03:03:01</i> variants identified in Brazilian individuals: <scp><i>DQA1</i></scp><i>*03:03:01:13</i> and <i><scp>DQA1</scp>*03:03:01:14</i>. Hla, 2020, 96, 747-748.	0.4	3
1804	<i><scp>HLAâ€DQB1</scp>*05:02:12</i>, an <i><scp>HLAâ€DQB1</scp>*05:02:01:01</i> variant, identified in a Taiwanese individual. Hla, 2020, 96, 551-552.	0.4	6
1805	The novel <scp>HLAâ€DQB1</scp> allele, <i><scp>DQB1</scp>*05:02:23</i>. Hla, 2020, 96, 552-553.	0.4	6
1806	Terasaki Institute: Innovating Personalized Health through Convergent Science and Bioengineering. Matter, 2020, 3, 324-326.	5.0	0
1807	Characterization of the novel <scp><i>HLAâ€DQB1*02:141</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 369-370.	0.4	6
1808	Characterization of five novel <scp>HLA</scp> alleles: <scp><i>HLAâ€A</i></scp><i>*01:217</i>, â€<i>A*24:314</i>, â€<i>A*26:106</i>, â€<i>B*57:78</i> and <i>â€C*05:145</i>. Hla, 2020, 96, 490-491.	0.4	6
1809	Genomic fullâ€length sequence of the <i><scp>HLAâ€A</scp>*11:334</i> allele, identified by cloning and sequencing. Hla, 2020, 96, 495-496.	0.4	6
1810	Three novel HLAâ€DQB1 *05 variants identified in Brazilian individuals. Hla, 2020, 96, 549-551.	0.4	6
1811	Intron 2 deletion generated the <scp><i>HLAâ€C*08:01:01:05</i></scp> allele in a <scp>Sumi</scp> individual from <scp>Nagaland</scp>, <scp>Northâ€East India</scp>. Hla, 2020, 96, 534-535.	0.4	3
1812	A simple electronic tool for assessing amino acid sequence polymorphisms within exon-2 of HLA-DPB1 alleles. Human Immunology, 2020, 81, 430-436.	1.2	0
1813	Identification of Novel Candidate Epitopes on SARS-CoV-2 Proteins for South America: A Review of HLA Frequencies by Country. Frontiers in Immunology, 2020, 11, 2008.	2.2	23
1814	Frequencies and haplotype associations of non-expressed HLA alleles in ethnically diverse populations on the National Marrow Donor Programâ€™s Be The Match Registry. Human Immunology, 2020, 81, 580-587.	1.2	7
1815	Intron 2 substitution resulted in <scp><i>HLAâ€DQB1*02:01:01:02</i></scp> variant in a Kashmiri Brahmin individual from North India. Hla, 2020, 96, 539-541.	0.4	6
1816	The novel <scp><i>HLAâ€C*03:04:01:47</i></scp> allele sequence identified using Pacific biosciences <scp>SMRT</scp> sequencing. Hla, 2020, 96, 525-526.	0.4	6
1817	Characterization of the novel <scp><i>HLAâ€B*07:355</i></scp> allele by nextâ€generation sequencing. Hla, 2020, 96, 724-725.	0.4	3

#	ARTICLE	IF	CITATIONS
1818	Specific amino acid patterns define split specificities of HLA-B15 antigens enabling conversion from DNA-based typing to serological equivalents. Immunogenetics, 2020, 72, 339-346.	1.2	2
1819	Enabling Routine MHC-II-Associated Peptide Proteomics for Risk Assessment of Drug-Induced Immunogenicity. Journal of Proteome Research, 2020, 19, 3792-3806.	1.8	11
1820	Characterization of a new <scp>HLAâ€A</scp> allele, <i><scp>HLAâ€A</scp>*29:128</i>, in a Spanish individual. Hla, 2020, 96, 501-502.	0.4	7
1821	Genomic fullâ€length sequence of the <i><scp>HLAâ€B</scp>*13:68</i> allele, identified by fullâ€length groupâ€specific sequencing. Hla, 2020, 96, 631-633.	0.4	3
1822	Identification of the novel <scp><i>HLAâ€A*26:206N</i></scp> allele by next generation sequencing in an <scp>Emirati</scp> bone marrow donor. Hla, 2020, 96, 717-718.	0.4	3
1823	<i>HLAâ€B*40:462</i> was likely generated by a recombination event between <i>B*40:01:02</i> and <i>B*13:02:01</i>. Hla, 2020, 96, 518-519.	0.4	6
1824	Identification of the novel <scp><i>HLAâ€C*15:219</i></scp> allele in a volunteer donor from the China Marrow Donor Program. Hla, 2020, 96, 741-742.	0.4	3
1825	Characterization of the novel <scp><i>HLAâ€DQB1*02:162N</i></scp> allele by nextâ€generation sequencing. Hla, 2021, 98, 244-246.	0.4	3
1826	Characterization of the novel <scp><i>HLAâ€DQB1*03:01:46</i></scp> variant allele in a French hematopoietic stem cell donor. Hla, 2020, 96, 370-371.	0.4	6
1827	HLAâ€B*51:315 , identified using nextâ€generation sequencingâ€based typing in a Korean individual. Hla, 2020, 96, 349-350.	0.4	6
1828	Characterization of the novel <scp>HLA</scp> allele: <scp><i>HLAâ€B*15:437</i></scp> in a Chinese bone marrow donor. Hla, 2020, 96, 511-513.	0.4	7
1829	Characterization of the novel <scp><i>HLAâ€C*03:517</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 527-528.	0.4	6
1830	Two novel alleles <scp><i>HLAâ€DQB1*03:01:01:23</i></scp> and <scp><i>DQB1*03:01:01:24</i></scp> in a <scp>Konyak Naga</scp> individual from <scp>Northâ€East India</scp>. Hla, 2020, 96, 541-543.	0.4	6
1831	Five novel <i><scp>HLAâ€DQA1</scp>*01</i> variants identified in Brazilian individuals. Hla, 2020, 96, 361-362.	0.4	6
1832	Identification of the novel <scp><i>HLAâ€A*30:171</i></scp> allele in a volunteer donor from the China Marrow Donor Program. Hla, 2020, 96, 721-722.	0.4	3
1833	Characterization of the novel HLAâ€DRB3*01:86 allele by sequencingâ€based typing. Hla, 2020, 96, 535-537.	0.4	6
1834	Characterization of the novel <i><scp>HLAâ€A</scp>*29:141</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 719-720.	0.4	3
1835	Characterization of the novel <i><scp>HLAâ€B</scp>*53:62</i> allele by sequencingâ€based typing. Hla, 2020, 96, 640-642.	0.4	3

#	ARTICLE	IF	CITATIONS
1836	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*04:78</i> allele by sequencingâ€based typing. Hla, 2020, 96, 547-549.	0.4	6
1837	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*05:176</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 750-752.	0.4	3
1838	Fullâ€length sequence of the novel <scp><i>HLAâ€C*12:02:36</i></scp> allele by next generation sequencing in a Chinese individual. Hla, 2020, 96, 650-651.	0.4	3
1839	Detection of an <i><scp>HLAâ€C</scp>*01:02:01:01</i> variant, <i><scp>HLAâ€C</scp>*01:02:43</i>, in a Taiwanese individual. Hla, 2020, 96, 645-646.	0.4	3
1840	Detection of a novel allele, <scp><i>HLAâ€B*15:01:39</i></scp>, by sequenceâ€based typing in a platelet donor from China. Hla, 2020, 96, 633-635.	0.4	3
1841	Novel <scp><i>HLAâ€DPB1*14:01:11</i></scp> allele, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2020, 96, 245-246.	0.4	6
1842	Characterization of the novel <scp><i>HLAâ€DPB1*1089:01</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 247-248.	0.4	6
1843	Novel <scp><i>HLAâ€C*06:284</i></scp> allele, identified by <scp>nextâ€generation</scp> sequencing in a Saudi individual. Hla, 2020, 96, 224-225.	0.4	6
1844	The novel <i><scp>HLAâ€C</scp>*07:93:02</i> allele identified in a healthy individual from Brazil. Hla, 2020, 96, 648-649.	0.4	3
1845	The novel <scp><i>HLAâ€C*14:02:34</i></scp> allele identified in a healthy individual from Brazil. Hla, 2020, 96, 652-653.	0.4	3
1846	The novel <i><scp>HLAâ€B</scp>*42:02:02</i> allele identified in a Brazilian family. Hla, 2020, 96, 638-640.	0.4	3
1847	Identification of the novel <i><scp>HLAâ€C</scp>*05:230</i> allele in a Brazilian individual. Hla, 2020, 96, 647-648.	0.4	3
1848	A novel <i><scp>HLAâ€C</scp>*15:02</i> variant, <i><scp>HLAâ€C</scp>*15:02:43</i>, identified in a healthy individual from Brazil. Hla, 2020, 96, 653-654.	0.4	3
1849	Characterization of the novel <scp><i>HLAâ€DQA1*03:15</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 236-237.	0.4	6
1850	A novel <scp><i>HLAâ€B*08</i></scp> allele, <scp><i>HLAâ€B*08:253</i></scp>, was identified by next generation sequencing in two Russian individuals. Hla, 2020, 96, 341-342.	0.4	7
1851	Characterization of the novel <scp><i>HLAâ€DRB3*02:02:25</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 359-360.	0.4	6
1852	The novel <i><scp>HLAâ€DPA1</scp>*01:03:01:21</i> allele in an individual of the Ao tribe from Nagaland, <scp>Northâ€East</scp><i>India</i>. Hla, 2020, 96, 376-377.	0.4	2
1853	Recognition of the novel <scp><i>HLAâ€DQB1*05:03:01:04</i></scp> allele in an <scp>Angami</scp> individual from <scp>Nagaland</scp>, <scp>Northâ€East India</scp>. Hla, 2020, 96, 554-555.	0.4	2

#	ARTICLE	IF	CITATIONS
1854	<scp><i>HLAâ€B*38:64</i></scp>, an <scp><i>HLAâ€B*38</i></scp> variant, detected in a Singaporean Malay unrelated hematopoietic stem cell donor. Hla, 2020, 96, 217-218.	0.4	6
1855	Novel <scp><i>HLAâ€B*50:66</i></scp> allele, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2020, 96, 222-223.	0.4	6
1856	Characterization of the novel <scp><i>HLAâ€DQA1*03:01:06</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 234-235.	0.4	6
1857	Detection of the <i><scp>HLAâ€A</scp>*02:935</i> allele in a Taiwanese individual. Hla, 2020, 96, 623-624.	0.4	3
1858	HLAâ€DQB1*05:66:01 , a novel variant of HLAâ€DQB1*05 , identified in a Singaporean Malay bone marrow donor. Hla, 2020, 96, 240-241.	0.4	6
1859	Characterization of the novel <scp><i>HLAâ€DPB1*1098:01N</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 249-251.	0.4	6
1860	<scp><i>HLAâ€DQB1*06:132</i></scp>, an <scp><i>HLAâ€DQB1*06</i></scp> variant, discovered in a Singaporean Malay bone marrow donor. Hla, 2020, 96, 243-244.	0.4	6
1861	Identification of four new HLA alleles, <scp><i>HLAâ€B*40:455</i></scp>, â€‹<scp><i>C*03:521</i></scp>, â€‹<scp><i>C*03:04:81</i></scp> and <i>â€‹<scp>DQB1*03:431</scp></i>. Hla, 2020, 96, 219-220.	0.4	6
1862	Characterization of the novel <scp><i>HLAâ€A*31:01:34</i></scp> allele by polymerase chain reaction sequencingâ€based typing. Hla, 2020, 96, 502-504.	0.4	7
1863	Similar patterns of genetic diversity and linkage disequilibrium in Western chimpanzees (Pan) Tj ETQq1 1 0.784314 rgBT /Overlock 10 BMC Evolutionary Biology, 2020, 20, 119.	3.2	2
1864	<scp><i>HLAâ€DQB1*05:116</i></scp>, an <scp><i>HLAâ€DQB1*05</i></scp> variant, detected in a Singaporean Chinese individual. Hla, 2020, 96, 238-239.	0.4	6
1865	Identification of an <i><scp>HLAâ€B</scp>*35:01:01:01</i> variant, <i><scp>HLAâ€B</scp>*35:01:23</i>, in a Taiwanese individual. Hla, 2020, 96, 635-637.	0.4	3
1866	Characterization of the novel HLAâ€DRB1*15:170 allele in a French hematopoietic stem cell donor. Hla, 2020, 96, 358-359.	0.4	6
1867	Characterization of the novel <i><scp>HLAâ€DRB1</scp>*15:175</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 746-747.	0.4	3
1868	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*05:237</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 752-753.	0.4	4
1869	A novel <scp>HLAâ€B</scp> allele, <i><scp>HLAâ€B</scp>*44:493</i>, detected in a potential hematopoietic stem cell donor. Hla, 2020, 96, 730-732.	0.4	3
1870	Two novel <scp><i>HLAâ€DPA1</i></scp><i>*01:03:01</i> variants identified in Brazilian individuals: <scp><i>DPA1</i></scp><i>*01:03:01:38</i> and <i><scp>DPA1</scp>*01:03:01:39</i>. Hla, 2020, 96, 755-756.	0.4	3
1871	Nomenclature for factors of the <scp>HLA</scp> system, update July, August, and September 2020. Hla, 2020, 96, 760-797.	0.4	6

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1872	Identification of the novel <i><scp>HLAâ€A</scp>*02:837</i> and <i>â€A*02:888</i> alleles by nextâ€generation sequencing in two Chinese individuals. Hla, 2021, 97, 345-349.	0.4	3
1873	The discovery of the <i><scp>HLAâ€DQB1</scp>*02:02:13</i> allele, found in a Brazilian individual. Hla, 2020, 96, 656-657.	0.4	4
1874	Characterization of the novel <i><scp>HLAâ€C</scp>*07:841</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 736-737.	0.4	3
1875	<scp><i>HLAâ€C*03:03:01:32</i></scp> shows an alternative splicing producing a functional protein with an extended cytoplasmic tail. Hla, 2020, 96, 523-525.	0.4	6
1876	The novel <scp><i>HLAâ€DQB1*05:254</i></scp> allele identified in a Taiwanese individual. Hla, 2020, 96, 659-660.	0.4	3
1877	Characterization of the new <scp>HLA</scp> allele <i><scp>HLAâ€A</scp>*25:68</i> by nextâ€generation sequencing. Hla, 2020, 96, 627-628.	0.4	3
1878	Novel HLAâ€DPB1 alleles in Spanish individuals: <i>DPB1*02:01:57</i>, <i>DPB1*17:01:04</i>, <i>DPB1*1117:01</i> and <i>DPB1*1124:01</i>. Hla, 2020, 96, 757-758.	0.4	3
1879	Two novel HLAâ€B variants identified in Russian individuals, HLAâ€B *56:74 and â€B*58:124. Hla, 2020, 97, 369-371.	0.4	3
1880	Genomic fullâ€length sequence of the HLAâ€A *24:233 allele, identified by fullâ€length groupâ€specific sequencing. Hla, 2020, 97, 356-358.	0.4	3
1881	Characterization of the novel <i><scp>HLAâ€B</scp>*15:547</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 637-638.	0.4	3
1882	Characterization of the novel <i><scp>HLAâ€A</scp>*32:134</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 723-724.	0.4	3
1883	Characterization of the novel <i><scp>HLAâ€DRB1</scp>*01:106</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 742-744.	0.4	3
1884	Characterization of three new <i><scp>HLAâ€A</scp>*11</i> alleles in Russian individuals by nextâ€generation sequencing. Hla, 2021, 97, 351-353.	0.4	3
1885	Identification of a novel <i><scp>HLAâ€B</scp>*40</i> allele, <i><scp>HLAâ€B</scp>*40:468</i>, by nextâ€generation sequencing in a Chinese individual. Hla, 2021, 97, 363-364.	0.4	3
1886	Characterization of the novel <i><scp>HLAâ€C</scp>*15:203</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 739-740.	0.4	4
1887	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*06:352</i> allele by nextâ€generation sequencing. Hla, 2020, 96, 754-755.	0.4	3
1888	A novel HLAâ€A allele, HLAâ€A*31:72 , detected in a Chinese hematopoietic stem cell donor and platelet donor. Hla, 2020, 96, 504-507.	0.4	6
1889	Characterization of the novel <scp><i>HLAâ€A*11:361</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 497-498.	0.4	6

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1890	Two novel <scp>HLAâ€DRB1</scp> alleles, <i><scp>DRB1</scp>*11:261</i> and <i><scp>DRB1</scp>*13:286</i> identified by sequencing in Brazilian individuals. Hla, 2020, 96, 744-745.	0.4	3
1891	Characterization of the novel <scp><i>HLAâ€A*24:470</i></scp> allele by nextâ€generation sequencing. Hla, 2020, 96, 716-717.	0.4	3
1892	Characterization of the novel <i><scp>HLAâ€B</scp>*40:450</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 160-162.	0.4	3
1893	Characterization of the novel <scp><i>HLAâ€B*18:161</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 513-514.	0.4	7
1894	The novel <scp>HLAâ€B</scp> allele, <scp><i>HLAâ€B*44:345N</i></scp>, discovered in a Korean family. Hla, 2020, 96, 220-221.	0.4	8
1895	Recognition of an <scp><i>HLAâ€A*24:02</i></scp> variant, <scp><i>HLAâ€A*24:02:31</i></scp>, in a Taiwanese individual. Hla, 2020, 96, 331-332.	0.4	6
1896	Characterization of two novel <scp>HLA</scp> alleles, <scp><i>C*03:03:58</i></scp> and <scp><i>DQB1*06:288</i></scp>, in a French hematopoietic stem cell donor. Hla, 2020, 96, 353-355.	0.4	6
1897	Characterization of the novel <scp><i>HLAâ€C*02:185</i></scp> allele in a kidney transplant recipient. Hla, 2020, 96, 352-353.	0.4	6
1898	A new HLAâ€C allele, <scp><i>HLAâ€C*07:02:83</i></scp>, identified in a Chinese family. Hla, 2020, 96, 103-104.	0.4	2
1899	Identification of a novel HLAâ€A*01 variant, HLAâ€A*01:211 , in a Singaporean Malay bone marrow donor. Hla, 2020, 96, 329-330.	0.4	6
1900	Identification of the novel HLAâ€B*13:109 allele by polymerase chain reaction sequenceâ€based typing. Hla, 2020, 96, 342-343.	0.4	7
1901	Characterization of the novel HLAâ€DQB1*05:236N null allele in a French hematopoietic stem cell donor. Hla, 2020, 96, 373-375.	0.4	6
1902	Rapid, highly accurate and costâ€effective openâ€source simultaneous complete <scp>HLA</scp> typing and phasing of class I and <scp>II</scp> alleles using nanopore sequencing. Hla, 2020, 96, 163-178.	0.4	20
1903	Extracting Structured Genotype Information from Free-Text HLA Reports Using a Rule-Based Approach. Journal of Korean Medical Science, 2020, 35, e78.	1.1	6
1904	Detection of <scp><i>HLAâ€B*48:50</i></scp> in a Chinese individual. Hla, 2020, 96, 347-349.	0.4	6
1905	The novel HLAâ€DRB3*03:39 allele, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2020, 96, 114-115.	0.4	2
1906	Nomenclature for factors of the <scp>HLA</scp> system, update January, February and March 2020. Hla, 2020, 95, 599-637.	0.4	8
1907	Characterization of the novel <scp><i>HLAâ€DQA1*01:49</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 233-234.	0.4	6

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1908	Characterization of the novel <scp><i>HLAâ€B*27:198</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 515-516.	0.4	6
1909	Usefulness of the Hematopoietic Stem Cell Donor Pool as a Source of HLA-Homozygous Induced Pluripotent Stem Cells for Haplobanking: Combined Analysis of the Cord Blood Inventory and Bone Marrow Donor Registry. Biology of Blood and Marrow Transplantation, 2020, 26, e202-e208.	2.0	4
1910	Characterization of the novel variant allele, <i><scp>HLAâ€B</scp>*13:82</i>, identified in a Chinese Han individual. Hla, 2020, 96, 510-511.	0.4	6
1911	The discovery of the first <scp><i>HLAâ€DQB1*03:04:01</i></scp> variant, <scp><i>DQB1*03:04:01:02</i></scp>, found in a Brazilian individual. Hla, 2020, 96, 543-544.	0.4	6
1912	Whole-genome sequencing of Chinese centenarians reveals important genetic variants in aging WGS of centenarian for genetic analysis of aging. Human Genomics, 2020, 14, 23.	1.4	6
1913	Characterization of the novel HLAâ€DRB1*04:275 allele by sequencingâ€based typing. Hla, 2020, 96, 356-357.	0.4	6
1914	The novel HLAâ€DRB1 allele, HLAâ€DRB1*01:108 , identified in a Korean individual. Hla, 2020, 96, 364-366.	0.4	6
1915	Specific donor HLA allotypes as predictors of cytomegalovirus disease risk in acute myeloid leukemia. Hla, 2020, 96, 445-455.	0.4	5
1916	Characterization of the novel <scp><i>HLAâ€C*01:195</i></scp> allele. Hla, 2020, 96, 350-351.	0.4	6
1917	Identification of the novel <scp><i>HLAâ€B*46:74</i></scp> allele by polymerase chain reaction sequenceâ€based typing. Hla, 2020, 96, 520-521.	0.4	6
1918	Identifying the HLA DRB1-DQB1 molecules and predicting epitopes associated with high-risk HPV infection clearance and redetection. Scientific Reports, 2020, 10, 7306.	1.6	9
1919	Discovery of <scp><i>HLAâ€B*35:368</i></scp>, a novel <scp><i>HLAâ€B*35</i></scp> variant, in a Singaporean Malay hematopoietic stem cell donor. Hla, 2020, 96, 94-95.	0.4	2
1920	Two novel <scp>HLA</scp> alleles, <scp><i>HLAâ€C*07:04:20</i></scp> and <scp><i>HLAâ€DRB1*07:34:02</i></scp>, detected in Russian individuals from Irkutsk. Hla, 2020, 96, 226-227.	0.4	6
1921	The novel <scp><i>HLAâ€DRB1*14:221</i></scp> allele was identified during highâ€resolution <scp>HLA</scp> typing. Hla, 2020, 96, 231-232.	0.4	7
1922	Characterization of the novel <i>HLAâ€DPA1*01:03:19</i> allele by sequencingâ€based typing. Hla, 2020, 96, 129-130.	0.4	2
1923	The <scp><i>HLAâ€DRB1*04:05:21</i></scp> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 96, 110-111.	0.4	2
1924	The <scp><i>HLAâ€A*24:480</i></scp> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 96, 332-334.	0.4	6
1925	The novel HLAâ€DRB1*13:290 allele, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2020, 96, 229-230.	0.4	6

#	Article	IF	Citations
1926	Detection of an <i>HLAâ€DRB1*14</i> variant, <i>HLAâ€DRB1*14:44:01</i>, in a Taiwanese individual. Hla, 2020, 95, 577-578.	0.4	2
1927	<i>HLAâ€A*02:877</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 480-482.	0.4	2
1928	Characterization of the novel <scp><i>HLAâ€DRB3*02:142</i></scp> allele by sequencingâ€based typing. Hla, 2020, 95, 581-582.	0.4	2
1929	Characterization of the novel HLAâ€C*01:154 allele by polymerase chain reaction sequencingâ€based typing. Hla, 2020, 95, 498-499.	0.4	2
1930	Nomenclature for factors of the HLA system, update October, November and December 2019. Hla, 2020, 95, 232-264.	0.4	4
1931	Detection of an <i>HLAâ€A*02</i> variant, <i>HLAâ€A*02:346</i>, in a Taiwanese individual. Hla, 2020, 96, 203-204.	0.4	6
1932	Twelve new HLA class I alleles described in the Spanish population. Hla, 2020, 96, 85-86.	0.4	2
1933	Identification of the novel <scp><i>HLAâ€DQB1*03:280</i></scp> allele by polymerase chain reaction sequenceâ€based typing. Hla, 2020, 96, 122-123.	0.4	2
1934	The <i>HLAâ€B*54:41</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 496-497.	0.4	2
1935	The novel <scp><i>HLAâ€B*07:387</i></scp> allele, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2020, 96, 213-214.	0.4	6
1936	A new HLAâ€DQA1 allele, <i>HLAâ€DQA1*01:26</i>. Hla, 2020, 96, 119-120.	0.4	2
1937	Confirmation of the recently described <scp>HLAâ€DPA1</scp> allele, <scp><i>HLAâ€DPA1*02:02:08</i></scp>. Hla, 2020, 96, 130-131.	0.4	2
1938	The novel <scp><i>HLAâ€A*68:227</i></scp> allele, identified by <scp>Nextâ€Generation Sequencing</scp> in a <scp>Saudi</scp> individual. Hla, 2020, 96, 337-339.	0.4	6
1939	<scp><i>HLAâ€B*15:349:02</i></scp>, a novel variant of <scp><i>HLAâ€B*15</i></scp>, discovered in a Singaporean Malay bone marrow donor. Hla, 2020, 96, 344-345.	0.4	6
1940	Discovery of a novel <scp><i>HLAâ€A*02</i></scp> variant, <scp><i>HLAâ€A*02:402</i></scp>, in a Singaporean cord blood unit. Hla, 2020, 96, 205-206.	0.4	6
1941	Characterization of the novel HLAâ€C*15:160N allele. Hla, 2020, 96, 227-229.	0.4	7
1942	Detection of an <scp><i>HLAâ€A*02</i></scp> variant, <scp><i>HLAâ€A*02:611</i></scp>, in a Singaporean Chinese individual. Hla, 2020, 95, 204-205.	0.4	2
1943	The <scp><i>HLAâ€A*02:842</i></scp> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 478-480.	0.4	0

#	ARTICLE	IF	CITATIONS
1944	Recognition of <scp><i>HLAâ€B*15:56</i></scp>, a variant of <scp><i>HLAâ€B*15</i></scp>, in a Taiwanese individual. Hla, 2020, 96, 215-216.	0.4	6
1945	Detection of the novel <i>HLAâ€DQB1*03:404</i> allele in an Ao individual from Nagaland, Northâ€East India. Hla, 2020, 95, 78-79.	0.4	2
1946	Detection of <i>HLAâ€C*12:109</i>, an <i>HLAâ€C*12</i> variant, in a Taiwanese hematopoietic stem cell donor. Hla, 2020, 95, 220-221.	0.4	3
1947	The <i>HLAâ€B*15:529</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 490-491.	0.4	2
1948	The <i>HLAâ€B*07:367</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 488-489.	0.4	2
1949	Detection of <i>HLAâ€C*14:20</i> in a Taiwanese individual. Hla, 2020, 95, 499-501.	0.4	2
1950	The <i>HLAâ€A*31:154</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 485-486.	0.4	2
1951	Identification of a novel allele, <i>HLAâ€DPB1*18:01:01:04</i>, in an African American renal transplant candidate. Hla, 2020, 95, 591-592.	0.4	2
1952	Characterization of <scp><i>HLAâ€A*01:302</i></scp> in two Spanish individuals by sequencingâ€based typing. Hla, 2020, 95, 474-476.	0.4	2
1953	Characterization of the novel <scp><i>HLAâ€B*44:192:04</i></scp> allele by sequencingâ€based typing. Hla, 2020, 95, 573-574.	0.4	2
1954	Characterization of the novel HLAâ€DQA1*05:23 allele by sequencingâ€based typing. Hla, 2020, 96, 120-121.	0.4	2
1955	Characterization of the novel HLAâ€DQB1*06:361 allele by sequencingâ€based typing. Hla, 2020, 96, 125-127.	0.4	2
1956	Detection of an HLAâ€DQB1*06 variant, HLAâ€DQB1*06:364, in a Russian individual. Hla, 2020, 96, 127-128.	0.4	2
1957	The novel HLAâ€A*33 variant, HLAâ€A*33:03:43, detected by next generation sequencing. Hla, 2020, 96, 210-211.	0.4	6
1958	<i>HLAâ€B*38:165N</i> and <i>HLAâ€B*51:309</i> alleles identified in two Chinese cervical intraepithelial neoplasia patients. Hla, 2020, 96, 96-97.	0.4	2
1959	Identification of the novel <i>HLAâ€DRB1*09:40</i> allele in a Chinese individual. Hla, 2020, 96, 111-113.	0.4	2
1960	Characterization of the novel <scp><i>HLAâ€DQA1*05:05:05</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 372-373.	0.4	6
1961	<scp><i>HLAâ€DPB1*526:01</i></scp> detected in two Singaporean Indian individuals. Hla, 2020, 96, 132-133.	0.4	2

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1962	Characterization of three new alleles, <i><scp>HLAâ€B</scp>*08:254</i>, â€<i><scp>DQB1</scp>*02:172</i> and <i>â€<scp>DPB1</scp>*1101:01</i> in Spanish individuals. Hla, 2020, 96, 508-509.	0.4	6
1963	Access to stem cell data and registration of pluripotent cell lines: The Human Pluripotent Stem Cell Registry (hPSCreg). Stem Cell Research, 2020, 47, 101887.	0.3	14
1964	Characterization of the novel <scp><i>HLAâ€DQB1*03:01:46</i></scp> allele by sequencingâ€based typing. Hla, 2020, 96, 544-545.	0.4	6
1965	Two novel <scp>HLA</scp> alleles, <scp><i>HLAâ€DRB1*14:223</i></scp> and <scp><i>HLAâ€DQB1*03:01:49</i></scp>, detected in a Buryat individual. Hla, 2020, 96, 375-376.	0.4	6
1966	Discovery of an <scp><i>HLAâ€A*33</i></scp> variant, <scp><i>HLAâ€A*33:200</i></scp>, in a Taiwanese individual. Hla, 2020, 96, 507-508.	0.4	6
1967	Nomenclature for factors of the HLA system, update January, February and March 2020. International Journal of Immunogenetics, 2020, 47, 359-395.	0.8	0
1968	Chechens from Chechen Republic, Russia. Hla, 2020, 96, 83-84.	0.4	0
1969	Umbilical Cord Blood Units Cryopreserved in the Public Cord Blood Bank: A Breakthrough in iPSC Haplobanking?. Cell Transplantation, 2020, 29, 096368972092615.	1.2	4
1970	Characterization of the novel <i>HLAâ€B*27:13:02</i> allele by sequencingâ€based typing. Hla, 2020, 96, 92-93.	0.4	2
1971	Nomenclature for factors of the HLA system, update October, November and December 2019. International Journal of Immunogenetics, 2020, 47, 190-225.	0.8	0
1972	Characterization of the novel <i>HLAâ€B*46:41N</i> allele. Hla, 2020, 95, 212-213.	0.4	2
1973	Complete genetic sequence of 15 novel HLAâ€H alleles. Hla, 2020, 96, 133-135.	0.4	4
1974	Identification of two novel HLAâ€C alleles, <i>HLAâ€C*07:02:92</i> and <i>HLAâ€C*07:828</i> in Chinese individuals. Hla, 2020, 96, 104-106.	0.4	4
1975	The novel <scp><i>HLAâ€A*02:787</i></scp> allele was identified by polymerase chain reaction sequenceâ€based typing. Hla, 2020, 96, 211-213.	0.4	8
1976	Characterization of a novel <scp><i>HLAâ€DRB1*03</i></scp> allele, <scp><i>DRB1*03:171</i></scp>. Hla, 2020, 96, 108-109.	0.4	2
1977	A novel HLAâ€DQA1 allele, DQA1*01:05:03, identified in a patient with suspected celiac disease. Hla, 2020, 96, 117-118.	0.4	2
1978	Characterization of the novel <i>HLAâ€C*04:408</i> allele by sequencingâ€based typing. Hla, 2020, 96, 101-102.	0.4	2
1979	The <i>HLAâ€B*46:01:26</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 492-493.	0.4	2

#	ARTICLE	IF	CITATIONS
1980	HLA-E expression in diffuse glioma: relationship with clinicopathological features and patient survival. BMC Neurology, 2020, 20, 59.	0.8	17
1981	The <i>HLAâ€B*44:454</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 210-212.	0.4	2
1982	The <i>HLAâ€B*51:284</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 494-495.	0.4	2
1983	The <i>HLAâ€DRB1*04:277</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2020, 95, 225-226.	0.4	3
1984	Detection of <i>HLAâ€DRB1*16:35</i> in a Taiwanese unrelated bone marrow stem cell donor. Hla, 2020, 95, 227-228.	0.4	2
1985	Characterization of the novel <scp><i>HLAâ€DQA1*04:08</i></scp> allele by sequencingâ€based typing. Hla, 2020, 95, 584-585.	0.4	2
1986	Identification of the novel allele, <i>HLAâ€C*15:02:32</i>, in a Chinese individual. Hla, 2020, 96, 106-108.	0.4	3
1987	<i>HLAâ€DQB1*05:05:02</i>, an <i>HLAâ€DQB1*05:05</i> variant, identified in a Taiwanese individual. Hla, 2020, 96, 124-125.	0.4	4
1988	<scp><i>HLAâ€A*02:05:01:10</i></scp>, <scp><i>â€A*26:01:01:30</i></scp>, <scp><i>â€C*07:01:01:75</i></scp>, and <scp><i>â€C*07:04:01:12</i></scp> identified in Spanish donors. Hla, 2020, 96, 207-208.	0.4	7
1989	Characterization of three novel HLAâ€DPA1*02:01:01 variants, identified in Brazilian individuals. Hla, 2020, 95, 84-85.	0.4	3
1990	Characterization of two novel HLAâ€DQB1*05:01:01 variants, identified in Brazilian individuals. Hla, 2020, 95, 586-587.	0.4	2
1991	Characterization of two novel HLAâ€DQA1*03:03:01 variants, identified in Brazilian individuals. Hla, 2020, 95, 583-584.	0.4	2
1992	The novel <i>HLAâ€A*02:916</i> allele identified in a Brazilian candidate donor for bone marrow donation. Hla, 2020, 96, 89-90.	0.4	2
1993	Identification of a novel <i>HLAâ€B*54</i> allele, <i>HLAâ€B*54:01:08</i>, in a Korean individual. Hla, 2020, 96, 98-99.	0.4	2
1994	The novel HLAâ€A*31:177 allele identified in a Brazilian candidate donor for bone marrow donation. Hla, 2020, 96, 90-91.	0.4	2
1995	Four new HLA alleles, <i>HLAâ€A*02:912</i>, <i>â€B*15:557</i>, <i>â€B*15:558</i>, and <i>â€B*40:97</i>, identified in the Chinese Han population. Hla, 2020, 96, 87-89.	0.4	2
1996	Description of two new HLA alleles: <i>HLAâ€DRB1*07:112</i> and <i>HLAâ€DQB1*02:169</i>. Hla, 2020, 95, 576-577.	0.4	2
1997	HLA Allele and Haplotype Frequencies in Three Urban Mexican Populations: Genetic Diversity for the Approach of Genomic Medicine. Diagnostics, 2020, 10, 47.	1.3	5

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1998	Common, intermediate and wellâ€documented HLA alleles in world populations: CIWD version 3.0.0. Hla, 2020, 95, 516-531.	0.4	93
1999	Characterization of two novel <scp><i>HLAâ€DQB1*06:02:01</i></scp> variants, identified in Brazilian individuals. Hla, 2020, 95, 587-588.	0.4	2
2000	Characterization of the novel <i>HLAâ€A*11:280</i> allele by nextâ€generation sequencing in a Chinese cord blood donor. Hla, 2020, 95, 482-483.	0.4	2
2001	Identification of the novel allele, <scp><i>HLAâ€DPB1*906:01</i></scp>, in a Caucasian individual by nextâ€generation sequencing. Hla, 2020, 95, 592-594.	0.4	2
2002	The <i>HLAâ€DRB1*04:68</i> allele recognized in a Taiwanese individual. Hla, 2020, 95, 575-576.	0.4	2
2003	HLA-A, -B, -C, -DRB1 and -DQB1 allele and haplotype frequencies and phylogenetic analysis of Bahraini population. Gene, 2020, 735, 144399.	1.0	3
2004	Characterization of two new HLA alleles: <i>HLAâ€A*33:199</i> and <i>HLAâ€B*13:140</i>. Hla, 2020, 95, 487-488.	0.4	2
2005	Characterization of the novel HLAâ€B*07:398 allele in a French hematopoietic stem cell donor. Hla, 2020, 96, 339-340.	0.4	6
2006	Nextâ€generation sequencing of <scp>HLA</scp>: validation and identification of new polymorphisms in a Brazilian population. Hla, 2020, 96, 13-23.	0.4	8
2007	Single molecule realâ€time DNA sequencing of the full HLAâ€E gene for 212 reference cell lines. Hla, 2020, 95, 561-572.	0.4	5
2008	The discovery of a <scp><i>HLAâ€C*17:51</i></scp> variant, <scp><i>C*17:51:02</i></scp>, found in a Brazilian individual. Hla, 2020, 96, 355-356.	0.4	6
2009	Identification of seven novel <scp>HLAâ€C</scp> alleles. Hla, 2020, 96, 99-101.	0.4	2
2010	Identification of a novel <scp><i>HLAâ€DQB1*05</i></scp> variant allele, <scp><i>HLAâ€DQB1*05:247</i></scp> by nextâ€generation sequencing. Hla, 2020, 96, 242-243.	0.4	6
2011	Identification of four novel <scp>HLAâ€A</scp> alleles. Hla, 2020, 96, 202-203.	0.4	7
2012	Novel <scp><i>HLAâ€DPB1*10:01:05</i></scp> allele, identified by nextâ€generation sequencing in a Saudi individual. Hla, 2020, 96, 379-381.	0.4	6
2013	Two novel <scp>HLAâ€DRB1</scp> alleles detected in inhabitants from the island of Crete. Hla, 2021, 97, 163-166.	0.4	4
2014	Characterization of 15 novel <scp>HLA</scp> alleles by next generation sequencing in Brazilian individuals. Hla, 2021, 97, 60-62.	0.4	3
2015	Detection of the novel <i><scp>HLAâ€DQB1</scp>*03:439</i> variant in an inhabitant from the island of Crete. Hla, 2021, 97, 88-90.	0.4	4

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2016	Characterization of the novel <scp><i>HLAâ€DPA1*01:42</i></scp> allele by sequencingâ€based typing. Hla, 2021, 97, 93-94.	0.4	3
2017	The novel <i><scp>HLAâ€B</scp>*15:554</i> allele identified in four Brazilian individuals. Hla, 2021, 97, 145-146.	0.4	3
2018	The novel allele <i><scp>HLAâ€A</scp>*11:362</i> identified in two unrelated bone marrow donors from Russia. Hla, 2021, 97, 218-219.	0.4	3
2019	<i><scp>HLAâ€A</scp>*11:263</i> is found on the haplotype: <i><scp>HLAâ€A</scp>*11:<scp>263â€C</scp>*03:04:<scp>01â€B</scp>*13:01</i>. Hla, 2021, 97, 138-139.	0.4	3
2020	Individual HLAs influence immunological events in allogeneic stem cell transplantation from HLA-identical sibling donors. Bone Marrow Transplantation, 2021, 56, 646-654.	1.3	0
2021	The novel <i><scp>HLAâ€B</scp>*15:555</i> allele identified in a healthy Brazilian individual. Hla, 2021, 97, 73-74.	0.4	3
2022	Detection of the <i><scp>HLAâ€B</scp>*51:322</i> allele in a Russian individual. Hla, 2021, 97, 155-156.	0.4	3
2023	Characterization of the novel HLAâ€B*35:460Q allele by nextâ€generation sequencing. Hla, 2021, 97, 361-362.	0.4	3
2024	Two novel <scp>HLAâ€C</scp> alleles, <i><scp>HLAâ€C</scp>*15:228</i> and <i>â€C*04:434</i>, detected in inhabitants from the island of Crete. Hla, 2021, 97, 243-245.	0.4	4
2025	<i><scp>HLAâ€DRB1</scp>*14:54:09</i> and <i>â€‹<scp>DRB1</scp>*14:54:10</i>, were identified by nextâ€generation sequencing in Chinese cord blood donors. Hla, 2021, 97, 166-169.	0.4	6
2026	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*06:371</i> allele by sequencingâ€based typing. Hla, 2021, 97, 175-176.	0.4	3
2027	Characterization of the novel <i><scp>HLAâ€DRB1</scp>*01:107</i> allele by nextâ€generation sequencing. Hla, 2021, 97, 83-85.	0.4	3
2028	Characterization of the novel <i><scp>HLAâ€B</scp>*44:476</i> allele by nextâ€generation sequencing. Hla, 2021, 97, 554-555.	0.4	3
2029	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*03:417</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 246-247.	0.4	3
2030	Detection of two novel alleles, <i>HLAâ€A*02:943</i> and <i>â€B*51:104:02</i>, in Greek cord blood units. Hla, 2021, 97, 214-215.	0.4	3
2031	A snapshot of human leukocyte antigen (HLA) diversity using data from the Allele Frequency Net Database. Human Immunology, 2021, 82, 496-504.	1.2	13
2032	Genomic fullâ€length sequence of the <i><scp>HLAâ€A</scp>*11:264</i> allele was identified in a Chinese bone marrow donor. Hla, 2021, 97, 67-69.	0.4	3
2033	Characterization of two new <scp>HLA</scp> alleles, <scp><i>HLAâ€A</i></scp><i>*02:942</i> and <i><scp>HLAâ€DQB1</scp>*06:02:47</i>. Hla, 2021, 97, 66-67.	0.4	4

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2034	Genomic fullâ€length sequence of the <i><scp>HLAâ€B</scp>*40:01:43</i> allele was identified by fullâ€length groupâ€specific sequencing. Hla, 2021, 97, 76-78.	0.4	3
2035	Identification of a novel <scp>HLAâ€B</scp> null allele, <i><scp>HLAâ€B</scp>*27:<scp>225N</scp></i>. Hla, 2021, 97, 232-233.	0.4	3
2036	A novel <i><scp>HLAâ€A</scp>*24</i> allele, <i><scp>HLAâ€A</scp>*24:512</i>, identified by nextâ€generation sequencing in a Chinese individual. Hla, 2021, 97, 220-221.	0.4	3
2037	Characterization of the new <i><scp>HLAâ€B</scp>*35:482</i>, detected in a potential hematopoietic stem cell donor. Hla, 2021, 97, 233-235.	0.4	3
2038	Characterization of the novel HLAâ€DQB1 *05: 235N allele by nextâ€generation sequencing. Hla, 2021, 97, 254-255.	0.4	3
2039	Characterization of the novel <i><scp>HLAâ€DRB1</scp>*08:97</i> allele by nextâ€generation sequencing. Hla, 2021, 97, 248-250.	0.4	3
2040	Identification of the novel <scp><i>HLAâ€A*02:406</i></scp> allele in a Chinese individual. Hla, 2021, 97, 64-65.	0.4	5
2041	Identification of the novel <scp>HLAâ€DQB1</scp>, allele <i><scp>DQB1</scp>*06:02:44</i> by nextâ€generation sequencing. Hla, 2021, 97, 91-92.	0.4	3
2042	Hemizygous amplification and complete Sanger sequencing of <i><scp>HLAâ€C</scp>*07:37:01:02</i> from a South European Caucasoid. Hla, 2021, 97, 159-161.	0.4	4
2043	Identification of two novel <scp>HLAâ€DQB1</scp> alleles by nextâ€generation sequencing <scp><i>DQB1*04:73N</i></scp> and <scp><i>DQB1*03:19:01:02Q</i></scp>. Hla, 2021, 97, 171-172.	0.4	4
2044	The <i><scp>HLAâ€DRB1</scp>*11:23:02</i> allele confirmed in a Chinese individual by nextâ€generation sequencing. Hla, 2021, 97, 85-86.	0.4	3
2045	The <i><scp>HLAâ€C</scp>*03:539</i> allele identified in a kidney transplantation recipient. Hla, 2021, 97, 158-159.	0.4	3
2046	Discovery of the <i><scp>HLAâ€DRB1</scp>*14:227</i> allele, a variant of <i><scp>HLAâ€DRB1</scp>*14</i>, in a Taiwanese bone marrow donor. Hla, 2021, 97, 169-171.	0.4	3
2047	Characterization of the novel <i><scp>HLAâ€A</scp>*30:154</i> allele by nextâ€generation sequencing. Hla, 2021, 97, 224-226.	0.4	3
2048	Characterization of the novel <scp><i>HLAâ€A*02:944</i></scp> allele by sequencingâ€based typing. Hla, 2021, 97, 216-217.	0.4	3
2049	A novel variant of <i><scp>HLAâ€C</scp>*12</i>, <i><scp>HLAâ€C</scp>*12:130</i>, detected in a Taiwanese individual. Hla, 2021, 97, 161-163.	0.4	3
2050	Characterization of the novel <i><scp>HLAâ€DQA1</scp>*01:39</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 240-241.	0.4	3
2051	Identification of three new HLA class I alleles: <i>HLAâ€B*50:73</i>, <i>â€C*08:218</i> and <i>â€C*15:229</i>. Hla, 2021, 97, 235-236.	0.4	3

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2052	The genomic full‐length characterization of <i>HLA‐C*07:166</i> shows it was likely generated by a recombination event. Hla, 2021, 97, 239-240.	0.4	3
2053	Identification of 11 novel <scp><i>HLA‐A</i></scp>, <i>‐B</i>, <i>‐C</i>, <i>‐</i><scp><i>DRB1</i>,</scp> and <i>‐<scp>DQB1</scp></i> alleles in the 1000 Genomes Project panel. Hla, 2021, 97, 246-248.	0.4	4
2054	Characterization of the novel <i><scp>HLA‐C</scp>*14:114</i> allele by next‐generation sequencing. Hla, 2021, 97, 373-374.	0.4	3
2055	The novel <scp><i>HLA‐A*02:941</i></scp> allele was identified during high‐resolution <scp>HLA</scp> typing. Hla, 2021, 97, 136-138.	0.4	4
2056	The novel <scp>HLA‐A</scp> allele, <i><scp>HLA‐A</scp>*01:353</i>. Hla, 2021, 97, 134-136.	0.4	4
2057	A novel <i><scp>HLA‐B</scp>*13</i> variant, <i><scp>HLA‐B</scp>*13:146</i>, identified by next‐generation sequencing in a Chinese individual. Hla, 2021, 97, 226-227.	0.4	3
2058	Characterization of the novel <i><scp>HLA‐DRB1</scp>*11:260</i> allele by next‐generation sequencing. Hla, 2021, 97, 87-88.	0.4	3
2059	The <i><scp>HLA‐A</scp>*02:01:175</i> allele newly identified in a Korean hematopoietic stem cell donor by next‐generation sequencing. Hla, 2021, 97, 62-64.	0.4	3
2060	A new <scp><i>HLA‐B*39</i></scp> allele, <scp><i>B*39:168</i></scp>, closely related to <scp><i>B*39:05:01:02</i></scp>. Hla, 2021, 97, 75-76.	0.4	3
2061	Characterization of the novel <i><scp>HLA‐B</scp>*18:181</i> allele by next‐generation sequencing. Hla, 2021, 97, 230-231.	0.4	3
2062	The <i><scp>HLA‐B</scp>*35:01:64</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2021, 97, 147-148.	0.4	3
2063	Characterization of the novel <i><scp>HLA‐B</scp>*15:514</i> allele in a French hematopoietic stem cell donor. Hla, 2021, 97, 143-145.	0.4	3
2064	Identification of a novel <i><scp>HLA‐A</scp>*31</i> variant, <i><scp>HLA‐A</scp>*31:01:02:31</i>, in a Saudi individual. Hla, 2021, 97, 358-359.	0.4	3
2065	Identification of the novel <i><scp>HLA‐C</scp>*15:210</i> allele by polymerase chain reaction sequence‐based typing. Hla, 2021, 97, 241-243.	0.4	3
2066	Two new <scp>HLA</scp> alleles, <scp><i>HLA‐B</i></scp><i>*18:200</i> and <scp><i>HLA‐C</i></scp><i>*04:435</i> detected in Russian donors. Hla, 2021, 97, 459-460.	0.4	4
2067	Detection of the <i><scp>HLA‐A</scp>*33:146</i> allele in a Taiwanese individual. Hla, 2021, 97, 142-143.	0.4	3
2068	Identification of an <i><scp>HLA‐B</scp>*40:01:01:01</i> variant, <i><scp>HLA‐B</scp>*40:33</i>, in a Taiwanese individual. Hla, 2021, 97, 149-150.	0.4	3
2069	Characterization of the novel <i><scp>HLA‐DQA1</scp>*01:19</i> allele by next‐generation sequencing. Hla, 2021, 97, 250-251.	0.4	3

#	ARTICLE	IF	CITATIONS
2070	Characterization of the novel <i><scp>HLAâ€DQA1</scp>*01:38:01:01</i> allele by nextâ€generation sequencing. Hla, 2021, 97, 252-253.	0.4	3
2071	Identification of a novel <i><scp>HLAâ€C</scp>*03:04</i> allele, <i><scp>HLAâ€C</scp>*03:04:84</i>, in a Korean individual. Hla, 2021, 97, 156-158.	0.4	3
2072	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*05:272</i> allele in a South African patient by nextâ€generation sequencing. Hla, 2021, 97, 173-174.	0.4	3
2073	Characterization of the novel <i><scp>HLAâ€B</scp>*44:452</i> allele by nextâ€generation sequencing. Hla, 2021, 97, 153-154.	0.4	3
2074	Recognition of an <i><scp>HLAâ€C</scp>*03:04:01:01</i> variant, <i><scp>HLAâ€C</scp>*03:04:37</i>, in a Taiwanese individual. Hla, 2021, 97, 78-80.	0.4	3
2075	The fullâ€length sequence of the novel <i><scp>HLAâ€C</scp>*08:190</i> allele, identified by cloning and sequencing. Hla, 2021, 97, 80-81.	0.4	3
2076	Identification of the novel <i><scp>HLAâ€A</scp>*01:01:01:53</i> allele generated by recombination in intron 1. Hla, 2021, 97, 133-134.	0.4	3
2077	A new <scp>HLAâ€B</scp> allele, <i><scp>HLAâ€B</scp>*35:481</i>. Hla, 2021, 98, 159-160.	0.4	3
2078	Characterization of the novel <scp><i>HLAâ€B*07:384</i></scp> allele by nextâ€generation sequencing. Hla, 2021, 97, 71-73.	0.4	3
2079	Introduction and some history. , 2021, , 1-4.		0
2080	Allogeneic hematopoietic stem cell transplant recipients in Spain: Human leukocyte antigen characteristics and diversity by highâ€resolution analysis. Hla, 2021, 97, 198-213.	0.4	2
2081	Distributions of HLAâ€A , â€B, and â€DRB1 alleles typed by ampliconâ€based next generation sequencing in Korean volunteer donors for unrelated hematopoietic stem cell transplantation. Hla, 2021, 97, 112-126.	0.4	7
2082	Two new <scp>HLA</scp> alleles, <scp><i>HLAâ€B</i></scp><i>*15:583</i> and <scp><i>DRB1</i></scp><i>*11:279</i>, detected in individuals from the Irkutsk region. Hla, 2021, 97, 458-459.	0.4	3
2083	HLA in Hematopoietic Stem Cell Transplantation. Organ and Tissue Transplantation, 2021, , 43-54.	0.0	0
2084	HLA Methods. , 2021, , .		0
2085	Nomenclature and naming conventions for HLA. , 2021, , 23-31.		0
2086	<i><scp>HLAâ€A</scp>*11:<scp>382N</scp></i>, a novel <scp>HLAâ€A</scp> null allele identified by nextâ€generation sequencing. Hla, 2021, 97, 448-449.	0.4	4
2087	<i><scp>HLAâ€A</scp>*11:77</i>, a variant of <i><scp>HLAâ€A</scp>*11</i>, detected in a Taiwanese patient. Hla, 2021, 97, 445-446.	0.4	3

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2088	Characterization of the novel HLAâ€A*02:939 allele by sequencingâ€based typing. Hla, 2021, 97, 436-437.	0.4	3
2089	Description of two new <scp>HLA</scp> alleles: <i><scp>HLAâ€A</scp>*24:<scp>517N</scp></i> and <i><scp>HLAâ€B</scp>*46:86</i>. Hla, 2021, 97, 451-452.	0.4	3
2091	Considerations in using human pluripotent stem cellâ€"derived pancreatic beta cells to treat type 1 diabetes. , 2021, , 173-203.		0
2092	Kalmyks from Republic of Kalmykia, Russia. Hla, 2021, 97, 177-179.	0.4	1
2093	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*06:374</i> allele by sequencingâ€based typing. Hla, 2021, 97, 382-383.	0.4	3
2094	The Genetics of Multiple Sclerosis. , 2021, , 155-172.		0
2095	Identification of a novel <i><scp>HLAâ€B</scp>*18</i> variant, <i><scp>HLAâ€B</scp>*18:01:01:52</i>, in a Saudi individual. Hla, 2021, 97, 359-360.	0.4	4
2096	Three novel <scp>HLAâ€C</scp> alleles identified in Russian individuals: <i>C*04:01:124</i>, <i>C*12:02:38</i>, and <i>C*12:03:64</i>. Hla, 2021, 97, 237-239.	0.4	4
2097	Pharmacogenetics and personalized medicine. , 2021, , 193-219.		0
2098	Detection of an HLAâ€C*08 variant, HLAâ€C*08:01:31 , in a Chinese individual. Hla, 2021, 97, 465-466.	0.4	3
2099	The novel HLAâ€DQB1 *05:02:24 allele, identified in a Russian bone marrow donor. Hla, 2021, 97, 380-381.	0.4	3
2100	A novel allele, <i><scp>HLAâ€C</scp>*15:227</i>, identified when typing <scp>COVID</scp>â€19 patients. Hla, 2021, 97, 377-378.	0.4	6
2101	BSHI guideline: HLA matching and donor selection for haematopoietic progenitor cell transplantation. International Journal of Immunogenetics, 2021, 48, 75-109.	0.8	22
2102	Major histocompatibility complex (MHC) associations with diseases in ethnic groups of the Arabian Peninsula. Immunogenetics, 2021, 73, 131-152.	1.2	10
2103	Complex Linkage Disequilibrium Effects in HLA-DPB1 Expression and Molecular Mismatch Analyses of Transplantation Outcomes. Transplantation, 2021, 105, 637-647.	0.5	7
2104	Two novel HLAâ€DRB3 alleles, DRB3*02:151 and DRB3*03:48. Hla, 2021, 97, 383-385.	0.4	3
2105	Identification of the novel <scp><i>HLAâ€C*03:04:79</i></scp> allele in a <scp><i>Chinese</i></scp> bone marrow donor. Hla, 2021, 97, 371-373.	0.4	3
2106	Characterization of the novel <i><scp>HLAâ€DRB3</scp>*03:49</i> allele by sequencingâ€based typing. Hla, 2021, 97, 477-478.	0.4	3

#	Article	IF	Citations
2107	Characterization of the novel HLAâ€DPB1*1151:01 allele by sequencingâ€based typing. Hla, 2021, 97, 470-471.	0.4	3
2108	The <i><scp>HLAâ€A</scp>*02:954</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2021, 97, 439-441.	0.4	3
2109	The novel <scp>HLAâ€A</scp> allele, <i><scp>HLAâ€A</scp>*02:952</i>, first described in a Brazilian individual. Hla, 2021, 97, 438-439.	0.4	3
2110	Discovery of the novel <i><scp>HLAâ€B</scp>*46:87</i> allele in a Taiwanese patient. Hla, 2021, 97, 461-462.	0.4	3
2111	Recognition of <i><scp>HLAâ€B</scp>*15:35</i> and its associated <scp>HLA</scp> haplotype. Hla, 2021, 97, 456-458.	0.4	3
2112	Characterization of the novel <i><scp>HLAâ€DPA1</scp>*01:44</i> allele by sequencingâ€based typing. Hla, 2021, 97, 466-468.	0.4	3
2113	Nomenclature for factors of the HLA system, update October, November and December 2020. International Journal of Immunogenetics, 2021, 48, 266-298.	0.8	0
2114	Nomenclature for factors of the <scp>HLA</scp> system, update <scp>October, November and December</scp> 2020. Hla, 2021, 97, 256-295.	0.4	4
2115	Identification of the novel <i><scp>HLAâ€DPB1</scp>*1149:01</i>. Hla, 2021, 97, 468-469.	0.4	3
2116	Description of two new <scp>HLA</scp> alleles: <i><scp>HLAâ€DRB1</scp>*11:262</i> and <i><scp>HLAâ€DRB1</scp>*11:268</i>. Hla, 2021, 97, 474-477.	0.4	3
2117	A novel <scp>HLAâ€C</scp> allele, <i><scp>HLAâ€C</scp>*14:125</i>. Hla, 2021, 97, 375-377.	0.4	3
2118	Three <scp>HLAâ€A</scp> alleles, <i>A*11:01:89</i>, <i>A*11:01:96</i> and <i>A*11:01:01:14</i> were identified in Chinese individuals. Hla, 2021, 97, 442-444.	0.4	3
2119	The <i><scp>HLAâ€B</scp>*13:144</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2021, 97, 548-550.	0.4	3
2120	Three novel HLAâ€DQB1 alleles identified in Brazilian individuals by nextâ€generation sequencing. Hla, 2021, 97, 472-473.	0.4	3
2121	Three novel HLAâ€DQA1 alleles identified in Brazilian individuals by nextâ€generation sequencing. Hla, 2021, 98, 74-75.	0.4	3
2122	Recognition of the <i><scp>HLAâ€A</scp>*24:353</i> allele and its associated <scp>HLA</scp> haplotype in a Taiwanese patient. Hla, 2021, 97, 529-530.	0.4	3
2123	Characterization of the novel <i><scp>HLAâ€DQA1</scp>*03:11</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 78-79.	0.4	3
2124	Recognition of the HLAâ€B *15:86 allele and its associated HLA haplotype in a Taiwanese individual. Hla, 2021, 97, 552-554.	0.4	3

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2125	The <i><scp>HLAâ€DRB1</scp>*09:45</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2021, 98, 238-239.	0.4	3
2126	Nomenclature for factors of the HLA system, update October, November and December 2020. Human Immunology, 2021, 82, 193-213.	1.2	0
2127	Standard reference sequences for submission of <scp>HLA</scp> genotyping for the 18th International HLA and Immunogenetics Workshop. Hla, 2021, 97, 512-519.	0.4	6
2128	Incidence and impact of allele-specific anti-HLA antibodies and high-resolution HLA genotyping on assessing immunologic compatibility. Human Immunology, 2021, 82, 147-154.	1.2	11
2129	Characterization of the novel <i><scp>DQA1</scp>*01:01:09</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 403-404.	0.4	3
2130	The <i><scp>HLAâ€A</scp>*24:<scp>514N</scp></i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2021, 97, 527-529.	0.4	3
2131	Identification of the novel <i><scp>HLAâ€A</scp>*68:250</i> allele in a volunteer bone marrow donor from Sao Paulo, Brazil. Hla, 2021, 97, 541-543.	0.4	3
2132	Two novel <scp>HLA</scp> alleles identified in Russian bone marrow donors: <i><scp>HLAâ€A</scp>*02:957</i> and â€<i>C*12:03:67</i>. Hla, 2021, 97, 523-524.	0.4	3
2133	Characterization of the novel <i><scp>HLAâ€DQA1</scp>*05:13</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 241-242.	0.4	3
2134	Characterization of the novel <i><scp>HLAâ€DQA1</scp>*05:19</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 243-244.	0.4	3
2135	A new <scp>HLAâ€DQB1</scp> allele, <i><scp>HLAâ€DQB1</scp>*04:62</i>, identified in a Chinese family. Hla, 2021, 98, 248-249.	0.4	3
2136	<i><scp>HLAâ€A</scp>*24:521</i> and <i><scp>HLAâ€B</scp>*13:152</i> identified by nextâ€generation sequencing in Russian bone marrow donors. Hla, 2021, 97, 531-532.	0.4	3
2137	Characterization of the novel <i><scp>HLAâ€C</scp>*14:124</i> allele by next generation sequencing. Hla, 2021, 97, 562-563.	0.4	5
2138	Identification of the novel <i><scp>HLAâ€C</scp>*01:202</i> allele in a Korean individual. Hla, 2021, 97, 560-561.	0.4	3
2139	Characterization of the novel <i><scp>HLAâ€B</scp>*56:76</i> allele by sequencingâ€based typing. Hla, 2021, 98, 66-67.	0.4	3
2140	A novel <i>B*07</i> variant allele, <i>B*07:416</i>, identified by nextâ€generation sequencing. Hla, 2021, 97, 545-546.	0.4	4
2141	Identification of <i><scp>HLAâ€A</scp>*01:01:01:76</i>, <i>â€B*15:03:12</i>, <i>â€B*49:01:01:14</i>, and <i>â€C*04:01:01:115</i> by <scp>nextâ€generation</scp> sequencing. Hla, 2021, 97, 520-521.	0.4	3
2142	Characterization of the novel <i><scp>HLAâ€B</scp>*08:67:<scp>02N</scp></i> allele by nextâ€generation sequencing. Hla, 2021, 98, 55-56.	0.4	3

#	ARTICLE	IF	CITATIONS
2143	Identification of the novel <scp>HLA</scp> allele, <i><scp>HLAâ€DPA1</scp>*01:46</i>, identified in a man of Serbian origin. Hla, 2021, 98, 79-81.	0.4	3
2145	Characterization of the novel <i><scp>HLAâ€A</scp>*26:01:66</i> allele by sequencingâ€based typing. Hla, 2021, 97, 532-533.	0.4	3
2146	Characterization of the novel <i><scp>HLAâ€B</scp>*35:29:03</i> allele by nextâ€generation sequencing. Hla, 2021, 97, 550-552.	0.4	3
2147	The novel allele, <i><scp>HLAâ€A</scp>*32:148</i>, identified by next generation sequencing in a Saudi individual. Hla, 2021, 97, 537-538.	0.4	3
2148	Identification of the novel <i><scp>HLAâ€A</scp>*68:100</i> allele by <scp>PCRâ€SBT</scp> in a Chinese bone marrow donor. Hla, 2021, 97, 539-540.	0.4	4
2149	The novel <scp>HLAâ€DPB1</scp> allele, <scp><i>HLAâ€DPB1*04:01:51</i></scp>, first described in a Brazilian individual. Hla, 2021, 98, 85-86.	0.4	3
2150	Characterization of the novel <i><scp>HLAâ€C</scp>*01:02:56</i> and <i>HLAâ€C*01:02:57</i> alleles by sequencingâ€based typing. Hla, 2021, 97, 557-560.	0.4	3
2151	Characterization of the novel <i><scp>HLAâ€C</scp>*06:311</i> allele by sequencingâ€based typing. Hla, 2021, 97, 565-566.	0.4	3
2152	Recognition of the <i><scp>HLAâ€C</scp>*01:22</i> allele in a Taiwanese individual. Hla, 2021, 97, 555-557.	0.4	3
2153	The <i><scp>HLAâ€A</scp>*11:384</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2021, 97, 525-526.	0.4	3
2154	The novel <scp>HLAâ€DPA1</scp> allele, <i><scp>HLAâ€DPA1</scp>*04:03</i>, first described in a Brazilian individual. Hla, 2021, 98, 81-82.	0.4	3
2155	Identification of two novel alleles, <i><scp>HLAâ€A</scp>*02:01:193</i> and <i>â€<scp>DQA1</scp>*02:17</i>, in Greek individuals. Hla, 2021, 97, 521-523.	0.4	3
2156	On Path to Informing Hierarchy of Eplet Mismatches as Determinants of Kidney Transplant Loss. Kidney International Reports, 2021, 6, 1567-1579.	0.4	24
2157	Characterization of the novel allele <i><scp>HLAâ€DRB1</scp>*01:01:36</i> in an Emirati donor. Hla, 2021, 98, 176-177.	0.4	3
2158	Characterization of the novel <i><scp>HLAâ€DQA1</scp>*05:18</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 494-496.	0.4	3
2159	Characterization of the novel <i>HLAâ€DQB1*03:362</i> allele in a Chinese family. Hla, 2021, 98, 410-412.	0.4	3
2160	Recognition of a novel <scp><i>HLAâ€B</i></scp><i>*13</i> allele, <scp><i>HLAâ€B</i></scp><i>*13:153</i>, in a Russian individual. Hla, 2021, 97, 547-548.	0.4	3
2161	The novel <scp><i>HLAâ€DPB1</i></scp><i>*571:01</i> allele characterized by <scp>SMRT DNA</scp> sequencing in an African Caribbean individual. Hla, 2021, 98, 87-89.	0.4	3

#	ARTICLE	IF	CITATIONS
2162	Characterization of two new HLA alleles: <scp><i>HLAâ€A</i></scp><i>*02:982</i> and <i>HLAâ€C*04:441</i>. Hla, 2021, 98, 47-48.	0.4	4
2163	Discovery of the novel <scp><i>HLAâ€C*06:195</i></scp> allele in a Singaporean unrelated hematopoietic stem cell donor. Hla, 2021, 97, 563-564.	0.4	3
2164	The novel <scp><i>HLAâ€DQB1</i></scp><i>*06:03:27</i> allele characterised by sequenceâ€based typing in a European bone marrow donor. Hla, 2021, 98, 498-500.	0.4	3
2165	Detection of the <scp><i>HLAâ€DQB1*06:02:43</i></scp> allele in a <scp>Taiwanese</scp> leukemic patient. Hla, 2021, 98, 412-414.	0.4	3
2166	Three novel <scp>HLA</scp> alleles detected in individuals from Russia: <scp><i>HLAâ€A</i></scp><i>*26:209</i>, â€<scp><i>DRB1</i></scp><i>*03:01:33</i>, and â€<i><scp>DQB1</scp>*03:447</i>. Hla, 2021, 97, 535-536.	0.4	3
2167	The novel <scp><i>HLAâ€C*04:407</i></scp> allele was identified in a Chinese individual. Hla, 2021, 98, 68-69.	0.4	3
2168	Sequencing of the new HLA class I alleles, <i>HLAâ€A*68:02:01:14</i>, â€<i>B*35:510</i>, and â€<i>C*07:907</i>. Hla, 2021, 97, 543-544.	0.4	3
2169	Characterization of the novel <scp><i>HLAâ€A</i></scp>*<i>26:208</i> allele by sequencingâ€based typing. Hla, 2021, 97, 534-535.	0.4	3
2170	Characterization of the novel <scp><i>HLAâ€C</i></scp><i>*07:944</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 73-74.	0.4	3
2171	Current Status of Korean Red Cross HLA-Matched Platelet Donor Registry. The Korean Journal of Blood Transfusion, 2021, 32, 1-10.	0.1	2
2172	Cost-effectiveness analysis of genotyping for HLA-B*15:02 in Indonesian patients with epilepsy using a generic model. Pharmacogenomics Journal, 2021, 21, 476-483.	0.9	9
2173	Associations of highâ€resolutionâ€typingâ€defined <i>MICA</i> and <i>MICB</i> polymorphisms, and the levels of soluble MICA and MICB with Oral Squamous Cell Carcinoma in Bulgarian patients. Journal of Oral Pathology and Medicine, 2021, 50, 758-765.	1.4	6
2174	Characterization of the novel <i><scp>HLAâ€C</scp>*06:314</i> allele by sequencingâ€based typing. Hla, 2021, 98, 70-71.	0.4	3
2175	High-resolution HLA typing by long reads from the R10.3 Oxford nanopore flow cells. Human Immunology, 2021, 82, 288-295.	1.2	28
2176	A novel <scp><i>MICB</i></scp> allele, <i><scp>MICB</scp>*004:02</i>, identified in a western china Uyghur individual. Hla, 2021, 98, 418-419.	0.4	3
2177	The novel HLAâ€DQB1 *03: 282N allele was identified in a Chinese individual. Hla, 2021, 98, 408-410.	0.4	3
2178	Detection of the <i><scp>HLAâ€A</scp>*02:191</i> allele in a Taiwanese individual. Hla, 2021, 98, 45-46.	0.4	3
2179	Validation of new <scp><i>HLAâ€J</i></scp> alleles assigned by next generation sequencing. Hla, 2021, 98, 173-175.	0.4	5

#	Article	IF	Citations
2180	Characterization of the novel <scp><i>HLAâ€C</i></scp><i>*01:214</i> allele by sequencingâ€based typing. Hla, 2021, 98, 167-168.	0.4	3
2181	Buryats from Republic of Buryatia and Irkutsk Region, Russia. Hla, 2021, 98, 262-264.	0.4	0
2182	Characterization of the novel <i><scp>HLAâ€A</scp>*36:12</i> allele by sequencingâ€based typing. Hla, 2021, 98, 51-53.	0.4	3
2183	Systematic comparative study of computational methods for HLA typing from nextâ€generation sequencing. Hla, 2021, 97, 481-492.	0.4	5
2184	Nomenclature for factors of the HLA system, update January, February and March 2021. International Journal of Immunogenetics, 2021, 48, 340-384.	0.8	2
2185	Recognition of the <i><scp>HLAâ€B</scp>*38:35</i> allele in a Filipino kidney donor. Hla, 2021, 98, 60-61.	0.4	3
2186	Discovery of the novel <i><scp>HLAâ€A</scp>*31:195</i> allele in a Taiwanese individual. Hla, 2021, 98, 50-51.	0.4	3
2187	Identification of a novel allele <i>HLAâ€C*03:560N</i> by nextâ€generation sequencing in a hematopoietic stem cell recipient. Hla, 2022, 99, 392-394.	0.4	3
2188	Recognition of the <i><scp>HLAâ€C</scp>*03:88</i> allele in a Singaporean bone marrow donor. Hla, 2021, 98, 168-170.	0.4	3
2189	Characterization of the novel HLAâ€DRB1*11:282 allele by sequencingâ€based typing. Hla, 2021, 98, 182-184.	0.4	3
2190	Characterization of the novel <i><scp>HLAâ€C</scp>*06:317</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 71-73.	0.4	3
2191	The novel <scp><i>HLAâ€B</i></scp> allele, <i><scp>HLAâ€B</scp>*55:01:27</i>. Hla, 2021, 98, 64-65.	0.4	3
2192	Discovery of the novel <i><scp>HLAâ€B</scp>*35:518</i> allele in a Taiwanese individual. Hla, 2021, 98, 56-58.	0.4	3
2193	Characterization of the novel <i><scp>HLAâ€DPA1</scp>*01:57</i> allele by sequencingâ€based typing. Hla, 2021, 98, 83-84.	0.4	3
2194	Characterization of the novel <i><scp>HLAâ€DQB1</scp>*06:385</i> allele by sequencingâ€based typing. Hla, 2021, 98, 573-574.	0.4	3
2195	Two novel <scp>HLA</scp> alleles, <scp><i>HLAâ€DRB1</i></scp><i>*12:90</i> and <i>HLAâ€<scp>DQB1</scp>*03:458</i>, identified by nextâ€generation sequencing. Hla, 2021, 98, 187-188.	0.4	4
2196	Description of two new <scp>HLA</scp> alleles: <i><scp>HLAâ€A</scp>*24:02:129</i> and <i><scp>HLAâ€A</scp>*24:02:135</i>. Hla, 2021, 98, 146-148.	0.4	3
2197	Discovery of the novel <i><scp>HLAâ€A</scp>*11:01:111</i> allele in a Taiwanese individual. Hla, 2021, 98, 48-49.	0.4	3

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2199	The novel <i><scp>HLAâ€ÐRB1</scp>*03:01:32</i> allele identified using nextâ€generation sequencing. Hla, 2021, 98, 177-178.	0.4	4
2200	Altered Immune Phenotypes and HLA-DQB1 Gene Variation in Multiple Sclerosis Patients Failing Interferon Î² Treatment. Frontiers in Immunology, 2021, 12, 628375.	2.2	0
2201	Identification of the novel <i><scp>HLAâ€B</scp>*40:125:03</i> allele in a Chinese bone marrow donor. Hla, 2021, 98, 62-64.	0.4	3
2202	Characterization of the novel <i><scp>HLAâ€A</scp>*01:367</i> allele by sequencingâ€based typing. Hla, 2021, 98, 43-44.	0.4	3
2203	Identification of three novel <scp>HLA</scp> alleles: <i><scp>HLAâ€A</scp>*68:01:58</i>, <i>â€B*27:05:52</i> and <i>â€<scp>DRB1</scp>*14:04:09</i>. Hla, 2021, 98, 53-54.	0.4	4
2204	Detection of the <i><scp>HLAâ€B</scp>*18:116</i> allele in a Singaporean bone marrow donor. Hla, 2021, 98, 58-59.	0.4	3
2205	HLAâ€associated adverse drug reactions â€scoping review. Clinical and Translational Science, 2021, 14, 1648-1658.	1.5	10
2206	Discovery of the novel <i><scp>HLAâ€C</scp>*01:179</i> allele in a southern Chinese patient. Hla, 2021, 98, 165-166.	0.4	3
2207	Unique Pathogen Peptidomes Facilitate Pathogen-Specific Selection and Specialization of MHC Alleles. Molecular Biology and Evolution, 2021, 38, 4376-4387.	3.5	7
2208	High-resolution characterization of the structural features and genetic variation of six feline leukocyte antigen class I loci via single molecule, real-time (SMRT) sequencing. Immunogenetics, 2021, 73, 381-393.	1.2	3
2209	<i><scp>HLAâ€ÐQB1</scp>*05:239</i> and â€<i><scp>DQB1</scp>*05:250</i>, were identified by sequencing in <scp>Chinese</scp> bone marrow donors. Hla, 2021, 98, 496-498.	0.4	3
2210	Characterization of the novel <i><scp>HLAâ€B</scp>*18:204</i> allele by sequencingâ€based typing. Hla, 2021, 98, 157-158.	0.4	3
2211	Detection of the <i><scp>HLAâ€A</scp>*11:01:06</i> allele in a Taiwanese individual. Hla, 2021, 98, 140-141.	0.4	3
2212	The novel <scp>HLAâ€A</scp> allele, <i><scp>HLAâ€A</scp>*68:272</i>, first described in a Brazilian individual. Hla, 2021, 98, 228-229.	0.4	3
2213	Recognition of the new <i><scp>HLAâ€ÐQB1</scp>*06</i> variant, <i><scp>HLAâ€ÐQB1</scp>*06:03:41</i>, in a Russian individual. Hla, 2021, 98, 501-502.	0.4	3
2214	Identification of the novel <i><scp>HLAâ€ÐPA1</scp>*01:49</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 251-252.	0.4	3
2215	HLA diversity in the Russian population assessed by next generation sequencing. Medical Immunology (Russia), 2021, 23, 509-522.	0.1	5
2216	Characterization of the novel allele, <i><scp>HLAâ€ÐQB1</scp>*04:74</i>, by sequencingâ€based typing in a Chinese Han individual. Hla, 2021, 98, 249-250.	0.4	3

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2217	Characterization of the novel <scp><i>HLAâ€B*51:296</i></scp> allele by nextâ€generation sequencing. Hla, 2021, 98, 163-164.	0.4	3
2218	Identification of the novel <i><scp>HLAâ€C</scp>*07:02:01:115</i> allele in a volunteer bone marrow donor. Hla, 2021, 98, 172-173.	0.4	4
2219	Characterization of the novel <i><scp>HLAâ€C</scp>*05:01:58</i> allele by sequencingâ€based typing. Hla, 2021, 98, 170-171.	0.4	3
2220	Identification of the novel <i><scp>HLAâ€A</scp>*11:361:02</i> allele in a Chinese patient. Hla, 2021, 98, 141-143.	0.4	3
2221	Identification of <i><scp>HLAâ€DPB1</scp>*1104:01</i> by <scp>nextâ€generation</scp> sequencing in a Chinese individual. Hla, 2021, 98, 414-416.	0.4	3
2222	Identification of the novel HLAâ€DRB1*03:01:32 in an Italian patient. Hla, 2021, 98, 179-180.	0.4	4
2223	A substitution in exon 2 resulted in the novel <i><scp>HLAâ€A</scp>*30:140</i> variant identified in a Chinese individual. Hla, 2021, 98, 226-228.	0.4	3
2224	Identification of the novel <i><scp>HLAâ€DRB1</scp>*04:305</i> allele in a Chinese leukemia patient. Hla, 2021, 98, 180-182.	0.4	4
2225	<i><scp>HLAâ€A</scp>*11:396</i>, <i>â€B*55:112</i>, and <i>â€<scp>DQA1</scp>*01:01:08</i> identified in individuals from Zhuang population of China. Hla, 2021, 98, 148-150.	0.4	3
2226	Discovery of the novel <scp><i>HLAâ€A</i></scp><i>*11:396</i> allele in a Chinese Han individual. Hla, 2021, 98, 144-145.	0.4	3
2227	Immunoinformatics based prediction of recombinant multi-epitope vaccine for the control and prevention of SARS-CoV-2. AEJ - Alexandria Engineering Journal, 2021, 60, 3087-3097.	3.4	9
2228	Detection of the <i><scp>HLAâ€DRB1</scp>*14:22</i> allele in a Taiwanese individual. Hla, 2021, 98, 185-186.	0.4	3
2229	Sequenceâ€based typing identification of the novel allele <i><scp>HLAâ€A</scp>*30:170</i>. Hla, 2021, 98, 153-155.	0.4	3
2230	The novel <i><scp>HLAâ€A</scp>*26:174</i> allele was identified in a Chinese individual. Hla, 2021, 98, 151-153.	0.4	3
2231	Identification of the novel <i><scp>HLAâ€DQB1</scp>*05:275</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 571-572.	0.4	3
2232	Characterization of novel HLA class I alleles: <i>HLAâ€A*02:984</i>, <i>â€B*18:205</i>, <i>â€B*57:142N</i>, <i>â€C*02:204</i>, and <i>â€C*16:185</i>. Hla, 2021, 98, 380-381.	0.4	3
2233	Identification of a new <i><scp>HLAâ€B</scp>*44</i> allele, <i><scp>HLAâ€B</scp>*44:02:68</i>, by next generation sequencing. Hla, 2021, 98, 162-163.	0.4	3
2234	Characterization of the novel <i><scp>HLAâ€B</scp>*14:01:13</i> allele by sequencingâ€based typing. Hla, 2021, 98, 155-156.	0.4	3

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2235	Description of four new <scp>HLA</scp> alleles: <i><scp>HLAâ€B</scp>*40:02:01:26</i>, <i>HLAâ€<scp>DQB1</scp>*03:01:29</i>, <i>HLAâ€<scp>DQB1</scp>*04:02:01:12</i>, and <i>HLAâ€<scp>DPB1</scp>*1195:01</i>. Hla, 2021, 98, 229-230.	0.4	3
2236	Characterization of the novel <i><scp>HLAâ€DPB1</scp>*1139:01</i> allele by sequencingâ€based typing. Hla, 2021, 98, 254-256.	0.4	3
2237	Identification of the <scp>HLAâ€DPA1</scp>*02:33 allele by nextâ€generation sequencing in a Chinese individual. Hla, 2021, 98, 252-254.	0.4	3
2238	Characterization of the novel <i><scp>HLAâ€DQA1</scp>*01:01:10</i> allele by sequencingâ€based typing. Hla, 2021, 98, 405-406.	0.4	3
2239	The changing landscape of HLA typing: Understanding how and when HLA typing data can be used with confidence from bench to bedside. Human Immunology, 2021, 82, 466-477.	1.2	4
2240	Discovery of the novel <i><scp>HLAâ€B</scp>*52:<scp>49N</scp></i> allele in a Chinese individual. Hla, 2021, 98, 553-555.	0.4	3
2241	Identification of a novel <scp>HLAâ€C</scp>*07 allele, <scp>HLAâ€C</scp>*07:943, in a Spanish individual. Hla, 2021, 98, 558-560.	0.4	3
2242	Characterization of the novel <i><scp>HLAâ€B</scp>*15:598</i> allele identified in a Korean individual by sequenceâ€based typing. Hla, 2021, 98, 387-389.	0.4	3
2243	Naming HLA diversity: A review of HLA nomenclature. Human Immunology, 2021, 82, 457-465.	1.2	25
2244	Characterization of the novel <scp><i>HLAâ€C</i></scp><i>*04:450</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 231-232.	0.4	3
2245	Identification of the novel <scp><i>HLAâ€B</i></scp><i>*35:03:01:22</i> allele in an individual of Nepalese origin. Hla, 2021, 98, 389-390.	0.4	3
2246	The novel <scp>HLAâ€DQA1</scp> allele, <scp><i>HLAâ€DQA1</i></scp><i>*03:03:05</i>, first described in a Brazilian individual. Hla, 2021, 98, 407-408.	0.4	3
2247	Characterization of the novel <scp><i>HLAâ€A</i></scp><i>*03:01:102</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 382-383.	0.4	3
2248	MiDASâ€"Meaningful Immunogenetic Data at Scale. PLoS Computational Biology, 2021, 17, e1009131.	1.5	12
2249	Characterization of the novel <i><scp>HLAâ€DPB1</scp>*1219:01</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 256-257.	0.4	3
2250	Recognition of the <i><scp>HLAâ€C</scp>*07:199:01</i> allele in a Singaporean unrelated hematopoietic stem cell donor. Hla, 2021, 98, 395-396.	0.4	3
2251	Characterization of a novel <i><scp>HLAâ€C</scp>*04:01:01:102</i> allele in an individual from West Bengal. Hla, 2021, 98, 393-394.	0.4	3
2252	Identification of the novel <i><scp>HLAâ€DRB1</scp>*11:271</i> allele by nextâ€generation sequencing. Hla, 2021, 98, 401-403.	0.4	3

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2253	Detection of the <i><scp>HLA‑DRB1</scp>*04:80</i> allele, a variant of <i><scp>HLA‑DRB1</scp>*04</i>, in a Taiwanese individual. Hla, 2021, 98, 235-236.	0.4	3
2254	Characterization of the novel <scp><i>HLA‑B</i></scp><i>*52:02:02</i> allele. Hla, 2021, 98, 476-477.	0.4	3
2255	Characterization of the novel <scp><i>HLA‑DRB1</i></scp><i>*04:326</i> allele by next generation sequencing. Hla, 2021, 98, 236-237.	0.4	3
2256	Description of two new <scp>HLA</scp> alleles: <i><scp>HLA‑B</scp>*46:80</i> and <i><scp>HLA‑B</scp>*46:81</i> identified in Chinese individuals. Hla, 2021, 98, 391-393.	0.4	6
2257	Nomenclature for factors of the HLA system, update April, May and June 2021. International Journal of Immunogenetics, 2021, 48, 443-484.	0.8	0
2258	Characterization of the novel <i><scp>HLA‑DPA1</scp>*01:61</i> allele by sequencing‑based typing. Hla, 2021, 98, 577-578.	0.4	3
2259	Identification of the novel <i><scp>HLA‑C</scp>*14:02:01:22</i> allele in an Indian individual. Hla, 2021, 98, 232-233.	0.4	3
2260	Identification of a novel <i><scp>HLA‑A</scp>*11</i> allele, <i><scp>HLA‑A</scp>*11:399</i>, in a Chinese Individual. Hla, 2021, 98, 223-224.	0.4	3
2261	The novel <scp><i>HLA‑DQB1</i></scp> allele, <i><scp>HLA‑DQB1</scp>*03:457</i>, identified in a Korean kidney transplant recipient. Hla, 2021, 98, 570-571.	0.4	3
2262	Characterization of the novel <scp>HLA</scp> allele, <i><scp>HLA‑DRB1</scp>*08:76</i>. Hla, 2021, 98, 399-401.	0.4	3
2263	Recognition of the <i><scp>HLA‑A</scp>*24:141</i> allele in a Taiwanese individual. Hla, 2021, 98, 386-387.	0.4	3
2264	Characterization of the novel HLA‑B*57:02:01:03 allele by sequencing‑based typing. Hla, 2021, , .	0.4	3
2265	Identification of a novel <scp>HLA‑A</scp> allele, <scp><i>HLA‑A</i></scp>*<i>31:97</i>, by sequence‑based typing. Hla, 2021, 98, 383-385.	0.4	3
2266	Identification of the novel <i><scp>HLA‑DRB3</scp>*02:02:19</i> allele. Hla, 2021, 98, 488-490.	0.4	3
2267	A novel <scp>HLA</scp> allele, <scp><i>HLA‑A*32:74</i></scp>, detected by sequencing in a Chinese individual. Hla, 2021, 98, 541-543.	0.4	3
2268	Identification of the novel <i>HLA‑DQB1*03:454</i> allele using next‑generation sequencing. Hla, 2021, 98, 568-569.	0.4	4
2269	A novel <scp>HLA‑B</scp> allele, <scp><i>HLA‑B*50:04:02</i></scp>, detected in a Russian hematopoietic stem cell donor. Hla, 2021, 98, 551-552.	0.4	3
2270	Identification of three novel <scp>HLA‑DRA</scp> alleles by next‑generation sequencing. Hla, 2021, 98, 560-562.	0.4	3

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2271	A novel <i><scp>HLAâ€C</scp>*15</i> allele, <i><scp>HLAâ€C</scp>*15:192</i>, identified by next generation sequencing in a Chinese individual. Hla, 2021, 98, 485-486.	0.4	3
2272	Discovery of the novel <i>HLA</i> allele, <i>HLAâ€DRB1</i>*<i>15:123</i>, in a hematopoietic stem cell donor from China. Hla, 2021, 98, 564-566.	0.4	3
2273	Identification of the novel <scp>HLAâ€DPA1</scp>, <i><scp>DPA1</scp>*01:56</i> by nextâ€generation sequencing. Hla, 2021, 98, 502-503.	0.4	3
2274	Characterization of the novel <i><scp>HLAâ€DPA1</scp>*01:60</i> allele by sequencingâ€based typing. Hla, 2021, 98, 575-576.	0.4	3
2275	Characterization of the novel <i><scp>HLAâ€C</scp>*04:451</i> allele by sequencingâ€based typing. Hla, 2021, 98, 483-485.	0.4	3
2276	Nomenclature for factors of the HLA system, update April, May and June 2021. Human Immunology, 2021, 82, 985-1013.	1.2	0
2277	Detection of the novel HLA allele, <i>HLAâ€B*46:64</i>, in a Chinese platelet donor by sequenceâ€based typing. Hla, 2021, 98, 548-551.	0.4	3
2278	Characterization of the novel HLAâ€B*44:02:73 allele by sequencingâ€based typing. Hla, 2021, 98, 474-476.	0.4	3
2279	Characterization of the novel <i><scp>HLAâ€C</scp>*07:01:101</i> allele by sequencingâ€based typing. Hla, 2021, 98, 556-557.	0.4	3
2280	Identification of novel HLA alleles discovered in 2019â€"2021. Human Immunology, 2021, 82, 982-984.	1.2	4
2281	Characterization of the novel <scp><i>HLAâ€C</i></scp>*<i>01:214</i> allele by sequencingâ€based typing. Hla, 2021, 98, 481-483.	0.4	3
2282	Characterization of the novel <i>HLAâ€A*31:199</i> allele by sequencingâ€based typing. Hla, 2021, 98, 540-541.	0.4	3
2283	Full genomic sequence of the <scp><i>HLAâ€DQB1*04:51</i></scp> allele identified by nextâ€generation sequencing. Hla, 2022, 99, 142-144.	0.4	3
2284	Characterization of the novel <i>HLAâ€A</i>*<i>68:277</i> allele by sequencingâ€based typing. Hla, 2021, 98, 544-545.	0.4	3
2285	Novel <scp><i>HLAâ€A</i></scp>, <scp><i>HLAâ€B</i></scp>, and <scp><i>HLAâ€DRB1</i></scp> alleles identified in Brazilian individuals. Hla, 2022, 99, 31-32.	0.4	3
2286	Characterization of the novel <scp><i>HLAâ€C</i></scp>*<i>15:241</i> allele by sequencingâ€based typing. Hla, 2021, 98, 397-399.	0.4	3
2287	A novel <scp>HLA</scp> class I allele: <scp><i>HLAâ€A</i></scp><i>*24:241</i>. Hla, 2021, 98, 471-472.	0.4	3
2288	Characterization of the novel <i><scp>HLAâ€A</scp>*24:538</i> allele by sequencingâ€based typing. Hla, 2021, 98, 473-474.	0.4	3

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2289	Identification of the novel <scp><i>HLAâ€DQB1*06:386</i></scp> allele by nextâ€generation sequencing. Hla, 2022, 99, 146-147.	0.4	3
2290	Nomenclature for factors of the <scp>HLA</scp> system, update January, February and March 2021. Hla, 2021, 98, 265-312.	0.4	1
2291	Nomenclature for factors of the <scp>HLA</scp> system, update April, May and June 2021. Hla, 2021, 98, 313-353.	0.4	4
2292	The novel <scp>HLAâ€DRB1</scp>*15:01:42 allele was identified by nextâ€generation sequencing. Hla, 2021, 98, 487-488.	0.4	3
2293	The novel <scp>HLAâ€DQB1</scp>*05:240 allele was likely generated by recombination between <scp>DQB1</scp>*05:01 and <scp>DQB1</scp>*03:02. Hla, 2022, 99, 144-145.	0.4	3
2294	Peptides of H. sapiens and P. falciparum that are predicted to bind strongly to HLA-A*24:02 and homologous to a SARS-CoV-2 peptide. Acta Tropica, 2021, 221, 106013.	0.9	5
2295	The novel <scp>HLAâ€A</scp>*31:191 allele was identified in a Chinese platelet donor. Hla, 2022, 99, 38-40.	0.4	3
2296	Identification of the novel <scp>HLAâ€B</scp>*55:107 allele in a Chinese bone marrow donor. Hla, 2021, 98, 478-479.	0.4	3
2297	Characterization of the novel <scp><i>HLAâ€C</i></scp><i>*04:438</i> allele by next generation sequencing. Hla, 2022, 99, 53-54.	0.4	3
2298	A new allele, <scp><i>HLAâ€C</i></scp><i>*08:228</i>, identified by sequenceâ€based typing in a Korean individual. Hla, 2022, 99, 59-60.	0.4	3
2299	Report of 13 new HLA alleles found in Spanish individuals. Hla, 2021, 98, 467-469.	0.4	3
2300	The novel <scp>HLAâ€DRB1</scp>*14:<scp>222N</scp> allele was identified by nextâ€generation sequencing. Hla, 2021, 98, 562-564.	0.4	3
2301	Discovery of the <scp>HLAâ€C</scp>*03:561 allele, a variant of <scp>HLAâ€C</scp>*03, in a Chinese individual. Hla, 2022, 99, 127-129.	0.4	3
2302	Challenges for the standardized reporting of NGS HLA genotyping: Surveying gaps between clinical and research laboratories. Human Immunology, 2021, 82, 820-828.	1.2	4
2303	The genomic fullâ€length sequence of the <i><scp>HLAâ€A</scp>*24:257</i> allele, identified by fullâ€length groupâ€specific sequencing. Hla, 2021, 98, 536-538.	0.4	3
2304	Sequenceâ€based typing identification of the novel <scp><i>HLAâ€B*27:226</i></scp> variant allele in a Chinese family. Hla, 2021, 98, 545-546.	0.4	3
2305	Detection of the novel allele, A*01:380:01:02, in a Bahraini individual by next generation sequencing. Hla, 2022, 99, 33-34.	0.4	3
2306	Characterization of the novel <scp><i>HLAâ€DQA1</i></scp><i>*01:80</i> allele by next generation sequencing. Hla, 2022, 99, 68-69.	0.4	3

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2307	The novel <i>HLAâ€A*24</i> allele, <i>HLAâ€A*24:487</i>, identified in three unrelated bone marrow donors in Colombia. Hla, 2021, 98, 538-540.	0.4	8
2308	Nomenclature for factors of the HLA system, update January, February and March 2021. Human Immunology, 2021, 82, 679-711.	1.2	2
2309	Efficacy of guselkumab on axial involvement in patients with active psoriatic arthritis and sacroiliitis: a post-hoc analysis of the phase 3 DISCOVER-1 and DISCOVER-2 studies. Lancet Rheumatology, The, 2021, 3, e715-e723.	2.2	53
2310	The discovery of the <i><scp>HLAâ€B</scp>*56:75</i> allele shows that <scp>HLA</scp> sequencing is important in all transplant settings. Hla, 2021, 97, 463-464.	0.4	3
2311	Characterization of two novel <scp>HLA</scp> alleles, <scp><i>HLAâ€A</i></scp><i>*25:71</i> and <scp><i>HLAâ€C</i></scp><i>*07:899</i> in Russian individuals. Hla, 2021, 97, 453-454.	0.4	3
2312	Characterization of the novel <i>HLAâ€A*11:376</i> allele by sequencingâ€based typing. Hla, 2021, 97, 447-448.	0.4	3
2313	High-Throughput MHC I Ligand Prediction Using MHCflurry. Methods in Molecular Biology, 2020, 2120, 113-127.	0.4	4
2314	Prospects for Designing â€˜Universalâ€™ Stem Cell Lines. , 2013, , 147-173.		2
2315	IMGT/HLA and the Immuno Polymorphism Database. Methods in Molecular Biology, 2014, 1184, 109-121.	0.4	18
2316	The Role of Adaptive Immunity in Idiopathic Pulmonary Fibrosis: Hiding in Plain Sight. , 2014, , 123-159.		3
2317	Transfusionsmedizin und ImmunhÃ¤matologie. , 2011, , .		12
2318	Polymorphic KIR-HLA System Regulates Natural Killer Cell Response. , 2016, , 369-380.		1
2319	Genome-wide Study Identifies Association between HLA-Bâˆ—55:01 and Self-Reported Penicillin Allergy. American Journal of Human Genetics, 2020, 107, 612-621.	2.6	34
2320	Like Wings of a Bird: Functional Divergence and Complementarity between HLA-A and HLA-B Molecules. Molecular Biology and Evolution, 2021, 38, 1580-1594.	3.5	7
2322	<i><scp>HLA</scp>â€B*40:356</i>, identified by nextâ€generation sequence based typing in a Chinese tuberculosis patient. Hla, 2017, 90, 312-313.	0.4	2
2323	Characterization of the novel <i><scp>HLAâ€C</scp>*16:173</i> allele by sequencingâ€based typing. Hla, 2021, 97, 82-83.	0.4	3
2324	<i><scp>HLAâ€A</scp>*02:934</i>, identified in National bone marrow donor registry named after Vasya Perevoshchikov, Russian Federation. Hla, 2021, 97, 349-351.	0.4	3
2325	Detection of a novel <scp>HLAâ€B</scp> allele, <i><scp>HLAâ€B</scp>*55:71</i>, in a Chinese hematopoietic stem cell donor and platelet donor. Hla, 2021, 97, 366-368.	0.4	3

#	Article	IF	Citations
2326	The novel <scp>HLAâ€A</scp> allele, <i>HLAâ€A*01:354</i>, identified in a Buryat individual. Hla, 2021, 97, 435-436.	0.4	4
2327	Three novel <scp>HLA</scp> alleles detected in Spanish donors, <i><scp>HLAâ€A</scp>*03:01:93</i>, â€‹<i>C*07:18:08</i>, and â€‹<i>C*03:03:01:41</i>. Hla, 2021, 97, 441-442.	0.4	3
2328	The novel <scp>HLAâ€A</scp> allele, <i><scp>HLAâ€A</scp>*24:516</i>, first described in two unrelated Brazilian individuals. Hla, 2021, 97, 450-451.	0.4	4
2329	Cross-presentation and genome-wide screening reveal candidate T cells antigens for a herpes simplex virus type 1 vaccine. Journal of Clinical Investigation, 2012, 122, 654-673.	3.9	83
2330	Genetic Differentiation in a Sample from Northern Mexico City Detected by HLA System Analysis: Impact in the Study of Population Immunogenetics. Human Biology, 2017, 89, 181.	0.4	5
2331	Distinguishing functional polymorphism from random variation in the sequences of >10,000 HLA-A, -B and -C alleles. PLoS Genetics, 2017, 13, e1006862.	1.5	129
2332	Molecular Characterization of HIV-1 CRF01_AE in Mekong Delta, Vietnam, and Impact of T-Cell Epitope Mutations on HLA Recognition (ANRS 12159). PLoS ONE, 2011, 6, e26244.	1.1	9
2333	TEPITOPEpan: Extending TEPITOPE for Peptide Binding Prediction Covering over 700 HLA-DR Molecules. PLoS ONE, 2012, 7, e30483.	1.1	100
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2335	Unraveling the Genetic Basis of Aspirin Hypersensitivity in Asthma Beyond Arachidonate Pathways. Allergy, Asthma and Immunology Research, 2013, 5, 258.	1.1	50
2336	Allogeneic amniotic membrane-derived mesenchymal stromal cell transplantation in a porcine model of chronic myocardial ischemia. Journal of Stem Cells and Regenerative Medicine, 2012, 8, 171-180.	2.2	14
2337	Genetic barriers in transplantation medicine. World Journal of Transplantation, 2016, 6, 532.	0.6	6
2338	Characterization of the novel <i>HLAâ€A*03:425</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 193-195.	0.4	3
2339	Characterization of the novel <scp><i>HLAâ€B</i></scp><i>*51:01:01:83</i> allele in three individuals from North India. Hla, 2022, 99, 132-133.	0.4	3
2340	Identification of the novel <i><scp>HLAâ€C</scp>*03:03:01:<scp>52N</scp></i> allele, a spliceâ€site variant at the boundary of intron1 and exon2. Hla, 2022, 99, 50-51.	0.4	5
2341	Characterization of the novel <i><scp>HLAâ€B</scp>*13:02:01:18</i> allele in a North Indian individual. Hla, 2022, 99, 40-41.	0.4	3
2342	Characterization of the novel <scp><i>HLAâ€B*52:01:01:18</i></scp> allele in a <scp>West Indian</scp> individual. Hla, 2022, 99, 124-125.	0.4	3
2343	Detection of a novel allele, <i><scp>HLAâ€DPA1</scp>*01:58</i>, originating from a recombination within exon 2 of the <scp>DPA1</scp> gene. Hla, 2022, 99, 72-74.	0.4	3

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2344	Characterization of the novel <scp><i>HLA‑C</i></scp>*<i>06:312</i> allele by two next‑generation sequencing methods. Hla, 2022, 99, 130-131.	0.4	0
2345	Identification of the novel allele, HLA‑B *44:256 , in an individual from West Bengal in Eastern India. Hla, 2021, , .	0.4	3
2346	A Comprehensive Review of HLA and Severe Cutaneous Adverse Drug Reactions: Implication for Clinical Pharmacogenomics and Precision Medicine. Pharmaceuticals, 2021, 14, 1077.	1.7	27
2347	The novel <i><scp>HLA‑A</scp></i>*01:390</i> allele was identified in a Russian individual. Hla, 2022, 99, 34-36.	0.4	3
2348	Identification of a novel HLA‑DPB1*02:01:61 allele in an Indian individual. Hla, 2021, , .	0.4	3
2349	Detection of the novel <scp>HLA</scp> allele, <i><scp>HLA‑DQA1</scp>*01:65</i>, identified in a Danish donor. Hla, 2022, 99, 66-67.	0.4	3
2350	Detection of an <scp><i>HLA‑DRB1*07</i></scp> variant, <scp><i>HLA‑DRB1*07:130</i></scp>, in a <scp>Russian Kalmyk</scp> individual. Hla, 2022, 99, 63-64.	0.4	3
2351	Detection of the new <i>HLA‑DQB1*06:384</i> allele in a Russian individual. Hla, 2022, 99, 71-72.	0.4	3
2352	The four novel <scp>HLA‑B</scp> alleles in unrelated Russian individuals: <i><scp>HLA‑B</scp></i>*38:01:18</i>, <i>‑B*39:01:28</i>, <i>‑B*40:484</i>, <i>‑B*44:498</i>. Hla, 2022, 99, 42-44.	0.4	3
2353	Characterization of the novel <i><scp>HLA‑C</scp>*05:255</i> allele by next‑generation sequencing. Hla, 2022, 99, 55-56.	0.4	3
2354	Characterization of the novel <i><scp>HLA‑A</scp>*68:280</i> allele by sequencing‑based typing. Hla, 2022, 99, 200-201.	0.4	3
2355	A novel <i><scp>HLA‑A</scp>*68</i> allele, <i><scp>HLA‑A</scp>*68:253</i>, identified by next generation sequencing in a Russian individual. Hla, 2022, 99, 116-118.	0.4	3
2356	Characterization of the novel <i><scp>HLA‑C</scp>*05:01:68</i> allele by two next‑generation sequencing methods. Hla, 2022, 99, 216-217.	0.4	3
2357	Identification of the novel <i><scp>HLA‑A</scp>*30:181</i> allele by next‑generation sequencing. Hla, 2022, 99, 115-116.	0.4	3
2358	Characterization of the novel <i><scp>HLA‑B</scp>*42:01:05</i> allele by two next‑generation sequencing methods. Hla, 2022, 99, 119-121.	0.4	3
2359	Characterization of the novel <scp><i>HLA‑C*01:213</i></scp> allele by next‑generation sequencing in a Chinese family. Hla, 2022, 99, 125-127.	0.4	3
2360	Characterization of the novel <i><scp>HLA‑C</scp>*07:977</i> allele by two next‑generation sequencing methods. Hla, 2022, 99, 121-122.	0.4	3
2361	The novel <i><scp>HLA‑DQB1</scp>*03:445</i> allele was identified in two unrelated bone marrow donors from Russia. Hla, 2022, 99, 69-70.	0.4	3

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2362	Identification of the novel <scp>HLAâ€DPA1</scp> allele, <scp><i>DPA1</i></scp>*<i>02:53</i> by nextâ€generation sequencing. Hla, 2022, 99, 149-150.	0.4	3
2363	Two novel <scp>HLAâ€C</scp> alleles, <i><scp>HLAâ€C</scp>*04:450</i> and <i><scp>HLAâ€C</scp>*15:02:51</i>, detected in individuals from Russia. Hla, 2022, 99, 129-130.	0.4	3
2364	Description of a novel allele <i><scp>HLAâ€DRB1</scp>*16:02:10</i>, identified in a bone marrow donor. Hla, 2022, 99, 135-136.	0.4	3
2365	Identification of the new <i><scp>HLAâ€C</scp>*03</i> variant, <i><scp>HLAâ€C</scp>*03:563</i>, by next generation sequencing. Hla, 2022, 99, 52-53.	0.4	3
2366	A novel allele, <scp>HLAâ€DPA1</scp>*01:<scp>87Q</scp> identified by next generation sequencing. Hla, 2022, 99, 147-148.	0.4	3
2367	A single nucleotide mutation in exon 2 produces a novel <i><scp>HLAâ€DRB1</scp>*07</i> null allele, <i><scp>HLAâ€DRB1</scp>*07:<scp>129N</scp></i>. Hla, 2022, 99, 133-134.	0.4	3
2368	Characterization of the novel <i><scp>HLAâ€C</scp>*02:207</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 213-214.	0.4	3
2369	Recognition of the <i>HLAâ€B*51:185</i> allele in a Taiwanese individual. Hla, 2022, 99, 122-124.	0.4	3
2370	The novel HLAâ€C *07:975 allele identified in an Italian bone marrow donor. Hla, 2021, , .	0.4	3
2371	<scp><i>HLAâ€B</i></scp>*<i>48:53</i>, a novel <scp><i>HLAâ€B</i></scp> allele with two exonic mutations. Hla, 2022, 99, 212-213.	0.4	3
2372	Das HLA-System. , 2010, , 189-203.		0
2377	From SNP Genotyping to Improved Pediatric Healthcare. Translational Bioinformatics, 2012, , 359-378.	0.0	0
2379	HLA Typing in Support of Hematopoietic Cell Transplantation from Unrelated Donors. , 2013, , 1307-1320.		0
2382	Tissue Typing, Crossmatch and Antibody Incompatibility in Kidney Transplantation. , 2014, , 767-781.		0
2384	PATHWAYS OF RECOGNITION OF FOREIGN ANTIGENS IN THE ADAPTIVE IMMUNE RESPONSE TO ALLOGENEIC ORGAN TRANSPLANTATION. Vestnik Transplantologii I Iskusstvennykh Organov, 2016, 17, 104-117.	0.1	1
2385	HLA System and Giant Cell Arteritis. , 2016, , 97-108.		0
2387	Compatibilidad HLA donante-receptor y probabilidades de trasplante renal en una poblaciÃ³n colombiana. Revista Ciencias De La Salud, 2016, 14, 147-160.	0.1	1
2389	Nomenclature for factors of the HLA system, update February 2017*. International Journal of Immunogenetics, 2017, 44, 177-186.	0.8	0

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2394	Peculiarities of phenotypes of patients with pyelo- and glomerulonephritis by HLA distribution analysis. Ukrainian Journal of Nephrology and Dialysis, 2018, , 11-18.	0.0	0
2397	Annual Report of Korean Association of External Quality Assessment Service on Histocompatibility Testing (2018). Journal of Laboratory Medicine and Quality Assurance, 2019, 41, 130-152.	0.1	2
2398	Characterization of the novel <i><scp>HLAâ€B</scp>*15:10:08</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 201-202.	0.4	3
2399	<i><scp>HLAâ€A</scp>*02:294</i>, a variant of <i><scp>HLAâ€A</scp>*02:01:01:01</i>, detected in a Taiwanese individual. Hla, 2022, 99, 36-38.	0.4	3
2400	Recognition of the <i><scp>HLAâ€DRB1</scp>*14:119</i> allele in a Singaporean bone marrow donor. Hla, 2022, 99, 222-224.	0.4	3
2401	Characterization of the novel <i><scp>HLAâ€C</scp>*07:01:105</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 218-219.	0.4	3
2402	Characterization of the novel <i><scp>HLAâ€B</scp>*40:02:35</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 209-210.	0.4	3
2406	Characterization of the novel <i><scp>HLAâ€C</scp>*03:294</i> allele by sequencingâ€based typing in a Taiwanese individual. Hla, 2022, 99, 215-216.	0.4	3
2407	Characterization of the novel <i><scp>HLAâ€DPA1</scp>*01:03:34</i> allele by sequencingâ€based typing. Hla, 2022, 99, 227-228.	0.4	3
2408	Identification of the novel <scp>HLAâ€B</scp> allele, <scp><i>HLAâ€B*44:532</i></scp> by nextâ€generation sequencing. Hla, 2022, 99, 210-211.	0.4	3
2409	Detection of the <i><scp>HLAâ€B</scp>*15:404</i> allele in a Singaporean bone marrow donor. Hla, 2022, 99, 205-206.	0.4	3
2410	Recognition of the <i><scp>HLAâ€A</scp>*11:85</i> allele in a Taiwanese bone marrow donor. Hla, 2022, 99, 195-196.	0.4	3
2411	A new <scp>HLAâ€B</scp> allele, <i>HLAâ€B*07:422</i>. Hla, 2022, 99, 118-119.	0.4	3
2412	Nomenclature for factors of the HLA system, update July, August and September 2020*. International Journal of Immunogenetics, 2021, 48, 36-74.	0.8	1
2413	HLA in Hematopoietic Stem Cell Transplantation. Organ and Tissue Transplantation, 2021, , 1-13.	0.0	0
2414	Analysis of variants associated with abnormal drug responses, genetics, and genomics in drug design. , 2022, , 209-235.		0
2415	Immunogenetic surveillance to histocompatibility. , 2022, , 85-126.		0
2416	Immunogenetics: the developmental course. , 2022, , 21-40.		0

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2417	Nomenclature for factors of the HLA system, update July, August and September 2021. Human Immunology, 2021, 83, 169-169.	1.2	0
2418	Characterization of the novel <i><scp>HLAâ€A</scp>*29:02:38</i> allele by sequencingâ€based typing. Hla, 2022, 99, 198-200.	0.4	3
2419	Graphical user interface for the haplotype frequency estimation software Hapl-o-Mat. Human Immunology, 2022, 83, 107-112.	1.2	3
2420	Characterization of the novel <scp>HLAâ€DQA1</scp>*05:01:07 allele by sequencingâ€based typing. Hla, 2022, 99, 138-139.	0.4	3
2421	Detection of the <i><scp>HLAâ€A</scp>*02:56:02</i> allele in a Taiwanese individual. Hla, 2022, 99, 192-193.	0.4	3
2422	A single nucleotide mutation in exon 3 produces the novel <i><scp>HLAâ€DPB1</scp>*700:<scp>01N</scp></i> allele. Hla, 2022, 99, 152-153.	0.4	3
2423	Recognition of the <i><scp>HLAâ€B</scp>*39:36</i> allele in a Taiwanese bone marrow donor. Hla, 2022, 99, 207-208.	0.4	3
2424	Value and Use of Genetic Test of Celiac Disease. , 2022, , 99-119.		0
2425	Identification of the novel <scp>HLAâ€B</scp>*38:97 allele in a Chinese individual. Hla, 2022, 99, 382-384.	0.4	3
2426	Characterization of the novel <scp><i>HLAâ€A</i></scp>*<i>23:118</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 197-198.	0.4	3
2427	The <i>HLAâ€B*51:353</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2022, 99, 385-387.	0.4	3
2428	Characterization of the novel <scp><i>HLAâ€DPB1*665:01:02</i></scp> allele by sequencingâ€based typing. Hla, 2022, 99, 150-152.	0.4	3
2429	Characterization of the novel <i>HLAâ€DQA1*01:76</i> allele by sequencingâ€based typing. Hla, 2022, 99, 136-137.	0.4	3
2430	Approaching Genetics Through the MHC Lens: Tools and Methods for HLA Research. Frontiers in Genetics, 2021, 12, 774916.	1.1	12
2431	Characterization of the novel alleles: <i><scp>HLAâ€A</scp>*01:305</i>, <i><scp>HLA</scp>â€B*58:120</i>, <i><scp>HLA</scp>â€C*04:<scp>428Q</scp></i> and <i><scp>HLA</scp>â€C*05:01:56</i>. Hla, 2022, 99, 370-371.	0.4	3
2432	Characterization of the novel <i><scp>HLAâ€A</scp>*01:405</i> allele by two next generation sequencing methods. Hla, 2022, 99, 371-372.	0.4	3
2433	Characterization of the novel <i><scp>HLAâ€A</scp>*01:406</i> allele by two next generation sequencing methods. Hla, 2022, 99, 617-618.	0.4	3
2434	Identification of the novel <i><scp>HLAâ€B</scp>*51:01:01:78</i> allele and confirmation of extended sequence of <i>B*18:01:26</i>. Hla, 2022, 99, 637-638.	0.4	4

#	ARTICLE	IF	CITATIONS
2435	Characterization of the novel <i><scp>HLAâ€DRB1</scp>*11:296</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 657-659.	0.4	3
2436	The novel <i>HLAâ€B*46:85</i> allele identified by sequencingâ€based typing in a Chinese individual. Hla, 2022, 99, 634-635.	0.4	3
2437	Identification of the novel <scp>HLAâ€DRB3</scp>*02:174 allele by nextâ€generation sequencing. Hla, 2022, 99, 224-225.	0.4	3
2438	The novel allele <i><scp>HLAâ€DQA1</scp>*05:05:10</i> was detected in a Nigerian who also lacks the <scp>DRB5</scp> gene in the <i><scp>DRB1</scp>*16</i> haplotype. Hla, 2022, 99, 667-668.	0.4	3
2439	Identification of a novel <scp>HLAâ€DRB1</scp> null allele, <scp><i>HLAâ€DRB1</i></scp><i>*13</i>:<scp><i>298N</i></scp>, in two siblings from Colombia. Hla, 2022, 99, 659-660.	0.4	4
2440	Characterization of the novel <i><scp>HLAâ€C</scp>*06:346</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 641-642.	0.4	3
2441	Transmembrane Helices Are an Over-Presented and Evolutionarily Conserved Source of Major Histocompatibility Complex Class I and II Epitopes. Frontiers in Immunology, 2021, 12, 763044.	2.2	2
2442	The novel <i>HLAâ€DRB1*04:05:19</i> allele identified by sequencingâ€based typing. Hla, 2022, 99, 652-654.	0.4	3
2443	The novel <scp><i>HLAâ€DRB1</i></scp><i>*13:306</i> allele, identified by sequencingâ€based typing in a Chinese individual. Hla, 2022, 99, 661-662.	0.4	3
2444	Detection of the <i><scp>HLAâ€A</scp>*68:99:02</i> allele in a Russian unrelated hematopoietic cell donor. Hla, 2022, 99, 627-628.	0.4	0
2445	Characterization of the novel <i><scp>HLAâ€C</scp>*16:184</i> allele by nextâ€generation sequencing. Hla, 2022, 99, 649-650.	0.4	3
2446	Identification of the novel <i><scp>HLAâ€A</scp>*01:348</i> allele in a Chinese individual. Hla, 2022, 99, 615-617.	0.4	3
2447	New HLAâ€DQB1 intronic variants detected by nextâ€generation sequencing. Hla, 2022, , .	0.4	3
2448	Characterization of the novel <i><scp>HLAâ€B</scp>*55:122</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 387-388.	0.4	3
2449	Characterization of the novel <i>HLAâ€DRB1*13:320</i> allele by nextâ€generation sequencing. Hla, 2022, 99, 404-405.	0.4	3
2450	Characterization of the novel <i><scp>HLAâ€A</scp>*03:436</i> allele by sequencingâ€based typing. Hla, 2022, 99, 621-623.	0.4	3
2451	Identification of the novel <scp><i>HLAâ€DPB1</i>*<i>1273:01</i></scp> allele by polymerase chain reaction sequenceâ€based typing. Hla, 2022, 99, 672-673.	0.4	3
2452	The novel <scp><i>HLAâ€B</i></scp><i>*35:501</i> allele, identified by sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2022, 100, 367-368.	0.4	3

#	ARTICLE	IF	CITATIONS
2453	Identification of the novel <scp>HLAâ€QA1</scp>*01:82 allele by nextâ€generation sequencing. Hla, 2022, 100, 183-184.	0.4	3
2454	Nomenclature for factors of the HLA system, update October, November and December 2021. Human Immunology, 2022, , .	1.2	0
2455	Characterization of seven new <scp>HLA</scp> alleles, <i><scp>HLAâ€A</scp>*01:407</i>, â€<i>A*01:408</i>, â€<i>A*03:434</i>, â€<i>B*40:<scp>508N</scp></i>, â€<i>B*40:<scp>511N</scp></i>, â€<i><scp>DRB1</scp>*04:336</i>, and â€<i><scp>DRB1</scp>*11:<scp>297Q</scp></i>. Hla, 2022, 99, 619-621.	0.4	3
2456	The novel <i>HLAâ€B*51:328</i> allele identified by sequencingâ€based typing in a Chinese individual. Hla, 2022, 100, 75-76.	0.4	3
2457	Identification of the novel <i><scp>HLAâ€B</scp>*42:28</i> allele by <scp>nextâ€generation</scp> sequencing. Hla, 2022, 99, 384-385.	0.4	3
2458	Identification of the novel <scp>HLAâ€DRB1</scp>*04:335 allele by nextâ€generation sequencing. Hla, 2022, 99, 656-657.	0.4	3
2459	Identification of the novel <scp>HLAâ€C</scp>*07:976 allele by polymerase chain reaction sequenceâ€based typing. Hla, 2022, 99, 643-644.	0.4	4
2460	<i><scp>HLAâ€DRB1</scp>*15:01:43</i> and <i><scp>HLAâ€DRB1</scp>*15:01:44</i> alleles were identified by nextâ€generation sequencing. Hla, 2022, 99, 664-666.	0.4	4
2461	Characterization of the novel <i><scp>HLAâ€A</scp>*02:01:01:206</i> allele in a Northern European individual. Hla, 2022, 100, 59-60.	0.4	3
2462	Identification of the novel <scp>HLAâ€DQB1</scp>*03:471 allele by nextâ€generation sequencing. Hla, 2022, 99, 670-671.	0.4	3
2463	The novel <i><scp>HLAâ€DRB1</scp>*03:01:01:05</i> and <i>â€<scp>DPB1</scp>*04:02:01:21</i> alleles identified in patients with acute leukemia. Hla, 2022, 99, 650-652.	0.4	3
2464	<i>HLAâ€C*01:224N</i>, a novel null <scp>HLAâ€C</scp> allele characterized by two nextâ€generation sequencing methods. Hla, 2022, 99, 640-641.	0.4	3
2465	The novel <i>HLAâ€A*69:01:04</i> allele identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2022, 99, 628-629.	0.4	3
2466	Nomenclature for factors of the HLA system, update July, August and September 2021. International Journal of Immunogenetics, 2022, 49, 90-122.	0.8	0
2467	Identification of the novel <i><scp>HLAâ€C</scp>*01</i>:<i>03</i>:<i>02</i>:<i>25</i> allele in an acute lymphoblastic leukemia patient. Hla, 2022, 100, 90-92.	0.4	3
2468	Identification of the novel allele, <i>HLAâ€A*01:01:01:92</i>, in an individual from North India. Hla, 2022, 99, 614-615.	0.4	3
2469	Identification of the novel <scp><i>HLAâ€DRB1*15:184</i></scp> allele by <scp>nextâ€generation</scp> sequencing. Hla, 2022, 99, 666-667.	0.4	3
2470	Detection of the novel <i>HLAâ€B*55:123</i> allele in a Taiwanese bone marrow donor. Hla, 2022, 99, 638-639.	0.4	3

#	Article	IF	Citations
2471	The novel <i>HLAâ€C*07:1000</i> allele was likely generated by an intragenic recombination event. Hla, 2022, 99, 645-646.	0.4	3
2472	Description of two new alleles: <i><scp>HLAâ€B</scp>*50:79</i> and <i><scp>HLAâ€DRB1</scp>*04:332</i>. Hla, 2022, 99, 635-637.	0.4	3
2473	Full genomic sequence of the <i><scp>HLAâ€C</scp>*15:02:03</i> allele identified by next generation sequencing. Hla, 2022, 99, 646-648.	0.4	3
2474	Nomenclature for factors of the HLA System, update October, November and December 2021. International Journal of Immunogenetics, 2022, 49, 123-167.	0.8	1
2475	The novel HLAâ€DRB1*12:02:11 allele identified by nextâ€generation sequencing in a Chinese bone marrow donor. Hla, 2021, , .	0.4	3
2476	Nomenclature for factors of the <scp>HLA</scp> system, update July, August and September 2021. Hla, 2021, 98, 579-618.	0.4	3
2477	Immunogenetics and its utility in therapeutics. , 2022, , 1-34.		0
2478	The Concept of Immunogenetics. Advances in Experimental Medicine and Biology, 2022, 1367, 1-17.	0.8	0
2479	Discovery of the novel <i>HLAâ€A*11:417N</i> allele in a Taiwanese individual. Hla, 2022, 100, 61-62.	0.4	3
2480	Identification of the novel allele, <i><scp>HLAâ€DRB1</scp>*08:03:12</i>, in a Chinese cord blood donor. Hla, 2022, 100, 94-95.	0.4	3
2481	Identification of the novel <scp>HLA</scp> allele, <scp><i>HLAâ€C*04:01:01:129</i></scp>, in an individual from northeast <scp>India</scp>. Hla, 2022, 99, 396-397.	0.4	3
2482	Nomenclature for factors of the <scp>HLA</scp> system, update October, November and December 2021. Hla, 2022, 99, 231-278.	0.4	4
2483	Genome Editing of Pluripotent Stem Cells for Adoptive and Regenerative Cell Therapies. , 2022, 1, 77-90.		0
2484	<i>HLAâ€A*32:34</i>, a variant of <i>HLAâ€A*32</i>, detected in a Taiwanese individual. Hla, 2022, 99, 625-626.	0.4	3
2485	Full genomic sequence of the <i><scp>HLAâ€C</scp>*07:308</i> allele identified by next generation sequencing. Hla, 2022, 100, 86-88.	0.4	3
2486	Discovery of the novel <i><scp>HLAâ€B</scp>*40:<scp>483N</scp></i> Allele in a Taiwanese individual. Hla, 2022, 99, 630-631.	0.4	3
2487	Detection of an <i><scp>HLAâ€A</scp>*24</i> variant, <i><scp>HLAâ€A</scp>*24</i>:<i>255</i>, in a Taiwanese individual. Hla, 2022, 100, 66-68.	0.4	3
2488	Human leukocyte antigen B polymorphism and association between HLAâ€B27 and endoplasmic reticulum aminopeptidase 1 rs30187 SNP in patients with ankylosing spondylitis in Bangladesh. Rheumatology & Autoimmunity, 2022, 2, 15-21.	0.3	0

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2489	Concurrent use of two independent methods prevents erroneous HLA typing of deceased organ donors â€" An important strategy for patient safety and accurate virtual crossmatching for broader sharing. Human Immunology, 2022, 83, 458-466.	1.2	2
2490	Three <scp>HLAâ€DQB1</scp> alleles, <i><scp>DQB1</scp>*03:432</i>, <i><scp>DQB1</scp>*03:454</i> and <i><scp>DQB1</scp>*03:465</i> were identified in Chinese individuals. Hla, 2022, 100, 97-99.	0.4	3
2491	Identification of the novel <i><scp>HLAâ€DPB1</scp>*03:01:14</i> allele by nextâ€generation sequencing in a Chinese cord blood donor. Hla, 2022, 100, 101-103.	0.4	3
2492	Characterization of the novel <i><scp>HLAâ€B</scp>*46:01:28</i> allele. Hla, 2022, 100, 73-74.	0.4	3
2493	Characterization of the novel <i><scp>HLAâ€C</scp>*03:04:94</i> allele by nextâ€generation sequencing. Hla, 2022, 100, 78-79.	0.4	3
2494	A clinicianâ€™s guide to HLA matching in allogeneic hematopoietic stem cell transplant. Human Immunology, 2022, 83, 687-694.	1.2	9
2495	Characterization of a novel HLA-A*11:335 allele resulting from a rare interlocus recombination involving HLA-A*11:01:01:01/126 and HLA-H*02:07/14/18 alleles with nanopore sequencing, in a volunteer from the China Marrow Donor Program. BMC Medical Genomics, 2022, 15, 58.	0.7	0
2497	The MHC class I MICA gene is a histocompatibility antigen in kidney transplantation. Nature Medicine, 2022, 28, 989-998.	15.2	20
2498	Characterization of a novel variant allele, <i>HLAâ€C*03:587</i>, identified in a Chinese Han individual. Hla, 2022, 100, 82-83.	0.4	3
2499	Immunogenetic profile of MIC (A, B) HLA loci linked to MHC antigenic complex in Russians of the Chelyabinsk Region. Medical Immunology (Russia), 2022, 24, 41-52.	0.1	0
2500	<i>HLAâ€B*58:41</i>, a variant of <i>HLAâ€B*58</i>, detected in a Taiwanese individual. Hla, 2022, 100, 76-78.	0.4	3
2501	The novel HLA class II allele, <i>DPB1*1284:01</i>, identified using nextâ€generation sequencing. Hla, 2022, 100, 103-104.	0.4	3
2502	UM171 Expansion of Cord Blood Improves Donor Availability and HLA Matching For All Patients, Including Minorities. Transplantation and Cellular Therapy, 2022, 28, 410.e1-410.e5.	0.6	6
2503	Characterization of six novel HLA alleles, <i>HLAâ€A*24:565</i>, <i>â€A*29:160</i>, <i>â€B*07:444</i>, <i>â€B*57:156</i>, <i>â€C*18:15</i> and <i>â€DPB1*795:01:02</i>. Hla, 2022, 100, 68-69.	0.4	3
2504	Identification of the novel <i><scp>HLAâ€C</scp>*04:440</i> allele using nextâ€generation sequencing. Hla, 2022, 100, 84-86.	0.4	3
2505	Characterization of the novel <i>HLAâ€DRB1*04:05:23</i> allele by polymerase chain reaction sequenceâ€based typing. Hla, 2022, , .	0.4	3
2506	Full genomic sequence of the <scp><i>HLAâ€DPA1*02:46</i></scp> allele identified by next generation sequencing. Hla, 2022, 100, 99-101.	0.4	4
2507	Identification of the novel <i>HLAâ€A*24:02:138</i> allele in a Chinese individual. Hla, 2022, 100, 64-66.	0.4	3

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2508	Principles of Virtual Crossmatch Testing for Kidney Transplantation. Kidney International Reports, 2022, 7, 1179-1188.	0.4	9
2509	Characterization of the novel <scp>HLAâ€DRB1</scp>*13:03:12 allele by two nextâ€generation sequencing methods. Hla, 2022, 100, 96-97.	0.4	3
2510	A novel <scp>HLAâ€A</scp> null allele, <i><scp>HLAâ€A</scp>*31:<scp>188N</scp></i>, identified by nextâ€generation sequencing in a Chinese individual. Hla, 2022, 100, 70-71.	0.4	3
2511	<scp><i>HLAâ€B*40:36</i></scp>, a variant of <scp><i>HLAâ€B*40</i></scp>, recognized in a <scp>Taiwanese individual</scp>. Hla, 2022, 100, 71-73.	0.4	3
2512	Identification of the novel <i>HLAâ€B*46:83</i> allele by sequencingâ€based typing in a Chinese individual. Hla, 2022, 100, 161-163.	0.4	3
2513	Characterization of the novel <scp><i>HLAâ€DQA1*01:81</i></scp> allele by sequencingâ€based typing. Hla, 2022, 100, 181-182.	0.4	3
2514	Characterization of the novel <i>HLAâ€A*31:11:02</i> allele by next generation sequencing. Hla, 2022, 100, 157-158.	0.4	3
2515	Recognition of the <i>HLAâ€A*24:516</i> allele in a Chinese patient and his daughter. Hla, 2022, 100, 151-153.	0.4	3
2516	Nextâ€generation sequencing identifies two novel HLA class II alleles, <i>HLAâ€DRB1*01:115</i> and <i>HLAâ€DRB1*14:224</i>. Hla, 2022, 100, 176-177.	0.4	3
2517	Association of Human Leukocyte Antigen Genotypes with Severe Acute Respiratory Syndrome Coronavirus 2 Vaccine-Induced Subacute Thyroiditis. Thyroid, 2022, 32, 640-647.	2.4	15
2518	Characterization of the novel <scp><i>HLAâ€DRB1*11:04:21</i></scp> allele by sequencingâ€based typing. Hla, 2022, 100, 179-180.	0.4	4
2519	Characterization of two new alleles: <i><scp>HLAâ€B</scp>*51:363</i> and <i><scp>HLAâ€DRB1</scp>*13:<scp>322N</scp></i>. Hla, 2022, 100, 165-166.	0.4	3
2520	Genomic fullâ€length sequence of the <i><scp>HLAâ€C</scp>*03:227</i> allele was identified by full length groupâ€specific sequencing. Hla, 2022, 100, 79-81.	0.4	3
2521	Identification of the novel <i>HLAâ€A*02:981</i> allele by sequencingâ€based typing. Hla, 2022, 100, 142-143.	0.4	3
2522	The novel <scp>HLAâ€C</scp> *06:325 allele identified in a Korean individual awaiting kidney transplantation. Hla, 2022, , .	0.4	3
2523	Genomic fullâ€length sequence of the <i>HLAâ€B*44:348</i> allele was identified by next generation sequencing. Hla, 2022, 100, 160-161.	0.4	4
2524	Identification of the novel <i>HLAâ€DPB1*1289:01</i> allele by <scp>nextâ€generation</scp> sequencing. Hla, 2022, 100, 188-189.	0.4	3
2525	<i><scp>HLAâ€DRB1</scp>*03:190</i>, a novel <scp>HLAâ€DRB1</scp> allele suspected with microlymphocytoxicity and confirmed by sequencing. Hla, 2022, 99, 219-220.	0.4	3

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2526	The <i><scp>HLAâ€B</scp>*58:01:42</i> allele identified in a volunteer bone marrow donor. Hla, 2022, 99, 391-392.	0.4	3
2527	Identification of novel <i><scp>HLAâ€DQA1</scp>*03:02:03</i> allele using nextâ€generation sequencing. Hla, 2022, 99, 225-226.	0.4	3
2528	Genetic markers for psoriatic arthritis among patients with psoriasis. Part II: HLA genes. Vestnik Dermatologii I Venerologii, 2021, 97, 6-17.	0.2	2
2529	Characterization of the novel <i>HLAâ€DRB1*04:334</i> allele by two nextâ€generation sequencing methods. Hla, 2022, 99, 401-402.	0.4	3
2530	Description of two new <scp>HLAâ€C</scp> alleles: <scp><i>HLAâ€C</i></scp><i>*07:900</i> and <i><scp>HLAâ€C</scp>*07:906</i>. Hla, 2022, 99, 399-400.	0.4	3
2531	The novel <scp>HLAâ€A</scp> allele, <scp>HLAâ€A</scp>*02:01:202, first described in a Brazilian individual. Hla, 2022, 99, 373-374.	0.4	3
2532	The novel <scp>HLAâ€A</scp> allele, <i><scp>HLAâ€A</scp>*29:158</i>, first described in two Brazilian individuals. Hla, 2022, 99, 376-377.	0.4	3
2533	The novel <scp>HLAâ€A</scp> allele, <i><scp>HLAâ€A</scp>*33:221</i>, first described in a Brazilian individual. Hla, 2022, 99, 379-380.	0.4	3
2534	Characterization of the novel <scp><i>HLAâ€A*30:02:28</i></scp> allele by sequencingâ€based typing. Hla, 2022, 99, 377-378.	0.4	3
2535	Characterization of the novel <i>HLAâ€A*11:383N</i> and <i>HLAâ€A*11:388N</i> alleles by nextâ€generation sequencing. Hla, 2022, 99, 374-375.	0.4	4
2536	Identification of the novel <i><scp>HLAâ€C</scp>*07:446</i> allele in a volunteer bone marrow donor. Hla, 2022, 99, 397-399.	0.4	3
2537	Characterization of the novel <i><scp>HLAâ€DRB4</scp>*01:151</i> allele by sequencingâ€based typing. Hla, 2022, 99, 64-66.	0.4	3
2538	Identification of two <scp>HLAâ€C</scp> alleles with new amino acid residues in the Î±â€3 domain, <scp><i>HLAâ€C*03:581</i></scp> and â€<scp><i>C*05:267</i></scp>. Hla, 2022, 99, 394-395.	0.4	3
2539	The novel <i><scp>HLAâ€DRB1</scp>*03:178</i>, â€<i><scp>DRB1</scp>*03:179</i>, and â€<i><scp>DRB1</scp>*11:276</i> alleles identified in a healthy Brazilian individuals. Hla, 2022, 99, 61-62.	0.4	3
2540	Detection of the <i><scp>HLAâ€B</scp>*15:360</i> allele in a Taiwanese bone marrow donor. Hla, 2022, 99, 203-204.	0.4	3
2541	Five novel <scp>HLAâ€A</scp> , â€B, and â€C alleles identified in Brazilian individuals by nextâ€generation sequencing. Hla, 2022, 99, 368-369.	0.4	3
2542	Two novel <scp>HLAâ€DRB1</scp> alleles, <i>HLAâ€DRB1*04:333</i> and <i>â€DRB1*15:01:48</i>, identified by sequencing in Russian individuals. Hla, 2022, 99, 221-222.	0.4	3
2543	Genomic fullâ€length sequence of the <i>HLAâ€B*52:100</i> allele, identified by fullâ€length groupâ€specific sequencing. Hla, 2022, 100, 167-169.	0.4	3

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2544	The <i>HLAâ€C*15:250</i> allele identified in a volunteer bone marrow donor. Hla, 2022, 100, 174-176.	0.4	3
2545	Detection of an <i><scp>HLAâ€A</scp>*02</i> variant, <i><scp>HLAâ€A</scp>*02:99:01</i>, in a Taiwanese individual. Hla, 2022, 100, 144-145.	0.4	3
2546	The novel <i><scp>HLAâ€DRB1</scp>*01:129</i> allele identified in a kidney transplant recipient. Hla, 2022, 100, 178-179.	0.4	3
2547	Genomic sequence of the <scp><i>HLAâ€A</i></scp><i>*11:01:54</i> allele identified by <scp>nextâ€generation</scp> sequencing in a transplant donor. Hla, 2022, 100, 146-148.	0.4	4
2548	Identification of a novel <i><scp>HLAâ€C</scp>*08</i> allele, <i><scp>HLAâ€C</scp>*08:242</i> in a Chinese individual. Hla, 2022, 100, 172-174.	0.4	3
2549	<i><scp>HLAâ€A</scp>*26:35</i>, a variant of <i><scp>HLAâ€A</scp>*26</i>, detected in a Taiwanese individual. Hla, 2022, 100, 153-155.	0.4	3
2550	The novel <i>HLAâ€A*24:49:02</i> and <i>HLAâ€DQB1*06:428</i> alleles identified in Indian bone marrow donors. Hla, 2022, 100, 150-151.	0.4	3
2551	The novel <i><scp>HLAâ€B</scp>*51:01:83</i> allele was identified by nextâ€generation sequencing. Hla, 2022, 100, 163-165.	0.4	3
2564	Human Leucocyte Antigens. , 0, , 34-49.		0
2565	Recognition of an <scp><i>HLAâ€DQB1*06:319</i></scp> variant, <scp><i>HLAâ€DQB1*06:319:02</i></scp>, in an hematopoietic stem cell donor. Hla, 2022, 100, 297-298.	0.4	4
2566	Sequenceâ€based typing identification of the novel allele <i><scp>HLAâ€B</scp>*40:482</i>. Hla, 2022, 100, 270-271.	0.4	3
2567	Genomic sequence of the <i>HLA</i>â€<i>B</i>*<i>15</i>:<i>01</i>:<i>02</i> and <i>HLA</i>â€<i>C</i>*<i>08</i>:<i>72</i>:<i>01</i> alleles identified in Gujarati individuals. Hla, 2022, 100, 267-268.	0.4	3
2568	Four novel <scp>HLAâ€Class II</scp> alleles: <scp><i>DQA1*03:31</i></scp>, <scp><i>DQB1*05:01:01:19</i></scp>, <scp><i>DPA1*02:01:01:20</i></scp> and <scp><i>DPA1*02:01:01:28</i></scp>, characterized in <scp>Spanish</scp> individuals. Hla, 2022, 100, 292-293.	0.4	3
2569	Characterization of the novel allele <scp>HLAâ€B</scp> *35:251:02. Hla, 2022, , .	0.4	3
2570	A novel <scp>HLAâ€C</scp> allele, <scp><i>HLAâ€C*15:244</i></scp>. Hla, 2022, 100, 283-284.	0.4	3
2571	Identification of the novel <i><scp>HLAâ€DQB1</scp>*04:85</i> allele by nextâ€generation sequencing. Hla, 2022, 100, 295-296.	0.4	3
2572	Identification of the novel allele <scp><i>HLAâ€B*13:157</i></scp> by sequenceâ€based typing. Hla, 2022, 100, 265-266.	0.4	3
2573	Identification of the novel <i>HLAâ€B*56:01:18</i> allele using nextâ€generation sequencing. Hla, 2022, 100, 274-275.	0.4	3

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2574	Nomenclature for factors of the HLA system, update January, February and March 2022. International Journal of Immunogenetics, 2022, 49, 215-241.	0.8	0
2575	Recognition of the <scp><i>HLAâ€A*11:398</i></scp> allele in a Chinese patient and his sister. Hla, 2022, 100, 258-260.	0.4	3
2576	The novel <i><scp>HLAâ€DRB1</scp>*12:96</i> allele was likely generated by recombination between <i><scp>DRB1</scp>*12:01</i> and <i><scp>DRB1</scp>*14:01</i>. Hla, 2022, 100, 290-291.	0.4	3
2577	Discovery of the novel <i>HLA</i>â€<i>C</i>*<i>08:22:02</i> allele in a Taiwanese individual. Hla, 2022, 100, 171-172.	0.4	3
2578	Nomenclature for factors of the HLA system, update January, February and March 2022. Human Immunology, 2022, 83, 589-589.	1.2	0
2579	Identification of the novel <i>HLAâ€A*26:01:75</i> allele using nextâ€generation sequencing. Hla, 2022, 100, 260-261.	0.4	4
2580	Discovery of the novel <scp><i>HLAâ€DRB1*09:31:02</i></scp> allele in a College of American Pathologists <scp>HLA</scp> Survey specimen. Hla, 2022, 100, 288-289.	0.4	3
2581	A new strategy for systematically classifying <scp>HLA</scp> alleles into serological specificities. Hla, 2022, 100, 193-231.	0.4	3
2582	Detection of an <scp><i>HLAâ€C*01</i></scp> variant, <scp><i>HLAâ€C*01:212</i></scp>, in a <scp>Chinese</scp> individual. Hla, 2022, 100, 275-277.	0.4	3
2583	Genomic fullâ€length sequence of the <i>HLA</i>â€<i>A</i>*<i>02</i>:<i>406</i> allele was identified by fullâ€length groupâ€specific sequencing. Hla, 2022, 100, 256-258.	0.4	3
2584	Identification of the novel <i><scp>HLAâ€C</scp>*08:243</i> allele in a Spanish bone marrow donor.. Hla, 2022, , .	0.4	3
2585	The novel <scp>HLAâ€DQB1</scp> allele, <i><scp>HLAâ€DQB1</scp>*04:72</i>, detected in a potential hematopoietic stem cell donor. Hla, 2022, , .	0.4	3
2586	The genomic fullâ€length sequence of the <i>HLA</i>â€<i>A</i>*<i>02</i>:<i>344</i> allele, identified by fullâ€length groupâ€specific sequencing. Hla, 2022, 100, 254-256.	0.4	3
2587	Detection of a novel <scp>HLAâ€B</scp> allele, <scp><i>HLAâ€B*48:54</i></scp> in a <scp>Chinese</scp> individual. Hla, 2022, 100, 272-273.	0.4	3
2588	Characterization of the novel <i>HLA</i>â€<i>C</i>*<i>07</i>:<i>1001N</i> allele by nextâ€generation sequencing. Hla, 2022, 100, 384-385.	0.4	3
2589	Human Leukocyte Antigen (HLA) System: Genetics and Association with Bacterial and Viral Infections. Journal of Immunology Research, 2022, 2022, 1-15.	0.9	21
2590	Characterization of the novel <i>HLAâ€C*03:598</i> allele. Hla, 2022, 100, 277-278.	0.4	3
2591	Nomenclature for factors of the <scp>HLA</scp> system, update January, February, and March 2022. Hla, 2022, 99, 674-701.	0.4	4

#	ARTICLE	IF	CITATIONS
2592	Identification of the Novel HLA-A*24:518N Null Allele and Evaluation of its Cell Surface Expression on Lymphocytes. Transplantation, 2022, 106, e312-e313.	0.5	1
2593	Discovery of the novel <i><scp>HLAâ€DRB1</scp>*07:136</i> allele in a Taiwanese patient. Hla, 2022, 100, 285-286.	0.4	3
2594	Discovery of the novel <i>HLAâ€C*08:03:05</i> allele in a Taiwanese individual. Hla, 2022, 100, 386-387.	0.4	3
2596	The novel <i>HLA</i><scp>â€</scp><i>B</i>*<i>07</i>:<i>461</i> allele and confirmation of the <i>HLA</i>â€<i>C</i>*<i>15</i>:<i>193</i> allele in individuals from <scp>E</scp>astern <scp>I</scp>ndia. Hla, 2022, 100, 262-263.	0.4	3
2597	Discovery of the <i>HLAâ€C*08:99</i> allele in a Chinese individual. Hla, 2022, 100, 278-280.	0.4	3
2598	The novel <scp>HLAâ€DRB1</scp>*12:01:10 allele was identified by nextâ€generation sequencing. Hla, 2022, 100, 389-390.	0.4	3
2599	Full genomic sequence of the <i><scp>HLAâ€DRB3</scp>*02:22:01</i> allele by single molecule realâ€time sequencing technology. Hla, 2022, 100, 394-396.	0.4	3
2600	Discovery of the novel <i>HLAâ€DRB1*08:113</i> allele in a Taiwanese bone marrow donor. Hla, 2022, 100, 287-288.	0.4	3
2601	Identification of the novel <i><scp>HLAâ€DQB1</scp>*03:483</i> allele by sequencingâ€based typing. Hla, 2022, 100, 400-401.	0.4	3
2602	Sequencing of a novel <scp>HLAâ€DQB1</scp> allele, <i><scp>DQB1</scp>*04:02:01:<scp>16Q</scp></i>, with a mutation in the intron 3 donor splicing site. Hla, 2022, 100, 401-402.	0.4	3
2603	Discovery of the novel <i>HLAâ€B*13:173</i> allele in a Taiwanese individual. Hla, 2022, 100, 364-365.	0.4	3
2605	Super high-resolution single-molecule sequence-based typing of HLA class I alleles in HIV-1 infected individuals in Ghana. PLoS ONE, 2022, 17, e0269390.	1.1	2
2606	Identification of the novel HLAâ€B*51:367 allele in a Cord Blood Donor by next generation sequencing. Hla, 0, , .	0.4	3
2607	Identification of the novel <scp><i>HLAâ€C*06:318</i></scp> allele by nextâ€generation sequencing in a Chinese individual. Hla, 2022, 100, 381-382.	0.4	3
2609	Identification of the novel HLAâ€DPA1*01:88 allele by nextâ€generation sequencing. Hla, 0, , .	0.4	4
2610	Identification of the novel <i>HLAâ€A*74:03:03</i> allele by sequencingâ€based typing. Hla, 2022, 100, 361-362.	0.4	3
2611	The novel <scp>HLAâ€C</scp> allele, <scp><i>HLAâ€C*03:537</i></scp> in a <scp>Chinese</scp> individual. Hla, 2022, 100, 376-377.	0.4	4
2612	The <i><scp>HLAâ€B</scp>*07:457</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2022, 100, 362-364.	0.4	3

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2613	The novel <scp>HLA</scp> class I allele, <scp>HLAâ€B</scp> *14:110, identified by <scp>Nextâ€Generation</scp> Sequencing. Hla, 0, , .	0.4	4
2614	Sequence Diversity and Differences at the Highly Duplicated MHC-I Gene Reflect Viral Susceptibility in Sympatric Pinniped Species. Journal of Heredity, 2022, 113, 525-537.	1.0	1
2615	The <i><scp>HLAâ€DRB1</scp>*12:97</i> allele identified in a volunteer donor for hematopoietic stem cell transplant. Hla, 2022, 100, 391-392.	0.4	3
2616	Identification of the novel <scp>HLA</scp> allele, <i>HLAâ€DRB1*14:50:02</i>, in an individual from southern India. Hla, 2022, 100, 392-393.	0.4	3
2617	The novel <scp>HLAâ€C</scp> allele, C*03:538 was identified by nextâ€generation sequencing. Hla, 0, , .	0.4	4
2618	Discovery of the novel <i><scp>HLAâ€DRB1</scp>*11:01:47</i> allele in a Taiwanese individual. Hla, 2022, 100, 387-388.	0.4	3
2619	Peptidome Surveillance Across Evolving SARS-CoV-2 Lineages Reveals HLA Binding Conservation in Nucleocapsid Among Variants With Most Potential for T-Cell Epitope Loss in Spike. Frontiers in Immunology, 0, 13, .	2.2	1
2620	Characterization of two new HLA alleles, <i>HLAâ€A*02:01:01:208</i> and <i>HLAâ€DQB1*06:03:44</i>. Hla, 2022, 100, 505-506.	0.4	3
2621	Characterization of four new <scp>HLA</scp> alleles: <i><scp>HLAâ€A</scp>*68:288</i>, <i>â€C*07:1012</i>, <i>â€C*13:364</i>, and <i>â€<scp>DQA1</scp>*05:51</i>. Hla, 2022, 100, 519-520.	0.4	3
2622	Identification and characterization of the novel <i>HLAâ€B*49:78</i> allele by nextâ€generation sequencing. Hla, 2022, 100, 370-371.	0.4	7
2623	Four new HLA class I alleles: <i>HLAâ€A*03:445</i>, <i>â€A*66:44</i>, <i>â€B*35:01:70</i>, and <i>â€C*07:02:01:161</i> identified in Russians. Hla, 2022, 100, 512-513.	0.4	3
2625	Description of the <i>HLAâ€B*41:01:08</i> allele: First identified in a Brazilian individual. Hla, 2022, 100, 529-530.	0.4	3
2626	Identification of the novel <i>HLAâ€DPA1</i> allele, <i>HLAâ€DPA1*01:03:34</i> in a Kuwaiti family. Hla, 2022, 100, 549-550.	0.4	3
2627	Description of two novel <scp>HLA</scp> alleles: <i>HLAâ€C*01:02:73</i> and <i>HLAâ€C*01:02:75</i>. Hla, 0, , .	0.4	3
2628	The novel <scp><i>HLAâ€DRB1</i></scp><i>*12:69</i> allele was identified in a Gujarati individual from North Gujarat, India. Hla, 2022, 100, 544-546.	0.4	3
2629	Identification of the novel <i>HLAâ€B*15:633</i> allele by nextâ€generation sequencing in an Indian individual. Hla, 2022, 100, 523-524.	0.4	3
2630	Identification of a novel <scp>HLAâ€DRB4</scp> allele, <scp><i>HLAâ€DRB4*01:152</i></scp> in a Kuwaiti family. Hla, 2022, 100, 546-547.	0.4	3
2631	Description of the novel <i>HLAâ€A</i> allele, <i>HLAâ€A*02:937</i> in a Chinese individual. Hla, 2022, 100, 508-510.	0.4	3

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2632	The genomic full‐length sequence of the <i>HLA‐A*02:304</i> allele, identified by full‐length group‐specific sequencing. Hla, 2022, 100, 506-508.	0.4	3
2633	The novel <scp><i>HLA‐C*03</i></scp> allele, <scp><i>HLA‐C*03:597</i></scp>, identified in a Chinese patient. Hla, 2022, 100, 534-536.	0.4	3
2634	Description of the <i>HLA‐DRB1*03:196</i> allele, first identified in a Brazilian individual. Hla, 2022, 100, 542-543.	0.4	3
2635	Detection of the <i><scp>HLA‐B</scp>*40:416</i> allele in a Taiwanese individual. Hla, 2022, 100, 528-529.	0.4	3
2636	Characterization of the novel <scp><i>HLA‐B</i></scp> allele, <i><scp>HLA‐B</scp>*39:01:32</i>. Hla, 2022, 100, 526-527.	0.4	3
2637	The novel <i>HLA‐A*24:520</i> allele was identified in a Chinese individual. Hla, 2022, 100, 515-517.	0.4	3
2638	Identification of the <scp>HLA‐A</scp> allele, <i>HLA‐A*11:03:01:02</i>, in two individuals in a Western Indian family. Hla, 2022, 100, 622-623.	0.4	3
2639	Identification of the novel <i>HLA‐DRB4*01:162N</i> allele using next‐generation sequencing. Hla, 2022, 100, 659-660.	0.4	3
2640	The novel <i><scp>HLA‐DQB1</scp>*03:493</i> allele, the first with glutamic acid at position‐10 in the leader peptide. Hla, 2022, 100, 665-667.	0.4	5
2641	Three novel alleles, <i>HLA‐A</i>*<i>03:446</i>, <i>HLA‐C</i>*<i>02:213</i>, and <i>HLA‐C</i>*<i>02:214</i>, identified by next‐generation sequencing. Hla, 2022, 100, 514-515.	0.4	3
2642	The novel <i>HLA‐C*06:02:102</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2022, 100, 641-643.	0.4	3
2643	Detection of the <i><scp>HLA‐A</scp>*02:759</i> allele, a variant of <i>A*02:06:01:01</i>, in a Taiwanese individual. Hla, 2022, 100, 620-622.	0.4	3
2644	Three novel HLA‐C alleles identified in Russian individuals: <i>HLA‐C*12:346, ‐C*05:01:66</i>, and <i>‐C*07:955</i>. Hla, 2022, 100, 539-541.	0.4	3
2645	Four new <scp>HLA‐DPA1</scp> intronic variants detected by next‐generation sequencing. Hla, 0, , .	0.4	3
2646	Characterization of the novel <i>HLA‐DQA1*01:89</i> allele by sequencing‐based typing. Hla, 2022, 100, 661-662.	0.4	4
2647	Identification of the novel <scp><i>HLA‐A</i></scp><i>*68:175:02</i> and ‐<i>A*68:287</i> alleles in Indian individuals. Hla, 2022, 100, 517-518.	0.4	3
2648	Characterization of the novel <scp><i>HLA‐B*53:64</i></scp> allele by next‐generation sequencing. Hla, 2022, 100, 635-636.	0.4	3
2649	Discovery of the novel <i><scp>HLA‐B</scp>*52:109</i> allele in a Taiwanese individual. Hla, 2022, 100, 530-532.	0.4	3

#	Article	IF	Citations
2650	<scp><i>HLAâ€DRB1*14:239</i></scp>, a novel <scp>HLAâ€DRB1</scp> allele with one exonic mutation. Hla, 2022, 100, 651-653.	0.4	3
2651	Identification of the novel <scp><i>HLAâ€B*15:625</i></scp> allele in a Chinese marrow donor. Hla, 2022, 100, 632-633.	0.4	3
2652	The novel <i><scp>HLAâ€C</scp>*06:327</i> allele was identified in three unrelated bone marrow donors. Hla, 2022, 100, 537-539.	0.4	3
2653	Characterization of the novel <i><scp>HLAâ€DRB3</scp>*02:<scp>179N</scp></i> allele by sequencingâ€based typing. Hla, 2022, 100, 658-659.	0.4	4
2654	Description of the <i><scp>HLAâ€C</scp>*04:01:145</i> allele, first identified in a Brazilian individual. Hla, 2022, 100, 536-537.	0.4	3
2655	Description of the <i>HLAâ€DQB1</i>*<i>03</i>:<i>491</i> allele, first identified in a <scp>B</scp>razilian individual. Hla, 2022, 100, 664-665.	0.4	3
2656	Identification of the novel <scp>HLA</scp> allele, <i>HLAâ€DRB1*14:245</i>, in an individual from Northern India. Hla, 2022, 100, 653-654.	0.4	3
2658	Characterization of the novel <i>HLAâ€DPA1*02:72</i> allele by next generation sequencing. Hla, 2022, 100, 671-672.	0.4	3
2659	Extended genomic sequence of the <i>HLAâ€DRB1*14:04:07</i> and <i>â€DRB1*15:53</i> alleles from Gujarati individuals. Hla, 2022, 100, 650-651.	0.4	4
2660	Identification of the novel <scp><i>HLAâ€DQB1*04:02:01:18</i></scp> allele in a <scp>Maharashtrian</scp> individual from <scp>India</scp>. Hla, 2022, 100, 667-668.	0.4	3
2661	Identification of the novel <i> <scp>HLAâ€C</scp> *14:02:38 </i> allele in a Chinese individual. Hla, 0, , .	0.4	3
2662	Detection of the novel allele, <i>HLAâ€A*32:165</i>, in a French individual by nextâ€generation sequencing. Hla, 2022, 100, 631-632.	0.4	3
2663	A novel <scp><i>HLAâ€DQA1*01</i></scp> allele, <scp><i>HLAâ€DQA1*01:99</i></scp>, identified by nextâ€generation sequencing. Hla, 2022, 100, 662-664.	0.4	4
2664	Description of the novel <i>HLAâ€A</i>*<i>23</i>:<i>125</i> allele in an individual from <scp>Equatorial Guinea</scp>. Hla, 2022, 100, 626-627.	0.4	3
2665	<i>HLAâ€A*24:244</i>, a variant of <i>HLAâ€A*24:02:01:01</i>, detected in a Taiwanese individual. Hla, 2022, 100, 627-629.	0.4	3
2666	Characterization of the novel <i><scp>HLAâ€C</scp>*17:01:18</i> allele by sequencingâ€based typing. Hla, 2022, 100, 646-648.	0.4	4
2667	Characterization of the novel <i>HLAâ€A*11:422</i> allele by sequencingâ€based typing. Hla, 2022, 100, 624-625.	0.4	4
2668	Sequencing of two novel <scp>HLA</scp> class I null alleles, <i>A</i>*<i>32:<scp>160N</scp></i> and <i>B</i>*<i>14:<scp>113N</scp></i>, produced by singleâ€nucleotide mutations. Hla, 2022, 100, 629-630.	0.4	3

#	ARTICLE	IF	CITATIONS
2669	Identification of the novel <scp>HLA</scp> allele, <i><scp>DQB1</scp>*06:427</i> by next‐generation sequencing method. Hla, 2022, 100, 669-670.	0.4	3
2670	The novel <i>HLA‐C*03:04:99</i> allele, identified by sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2022, 100, 637-638.	0.4	3
2671	The novel <scp>HLA‐DPB1</scp> allele, <scp><i>HLA</i>‐<i>DPB1</i></scp><i>*1344:01</i>, first identified in Korean individuals by next‐generation sequencing. Hla, 2023, 101, 96-97.	0.4	3
2672	Identification of the novel <i>HLA‐DQA1*02:01:09:01</i> allele by two different next‐generation sequencing platforms. Hla, 2023, 101, 80-82.	0.4	6
2673	Human Leukocyte Antigen (HLA) Testing in Pharmacogenomics. Methods in Molecular Biology, 2022, , 21-45.	0.4	3
2674	Characterization of the novel <scp><i>HLA‐DQB1*05:277</i></scp> allele, detected in two unrelated bone marrow donors. Hla, 2023, 101, 88-89.	0.4	2
2675	Identification of the novel <scp><i>HLA‐DRB1</i></scp>*<i>04:316</i> allele by next‐generation sequencing in a Chinese bone marrow donor. Hla, 2023, 101, 75-77.	0.4	3
2676	The novel <scp>HLA‐DPB1</scp> *1352:01 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2677	A novel <scp>HLA</scp> allele, <i><scp>HLA‐B</scp>*07:248</i>, detected in a Chinese hematopoietic stem cell donor and platelet donor. Hla, 2023, 101, 49-51.	0.4	3
2678	The novel <i>HLA‐B*35:563</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 55-56.	0.4	3
2679	The detection of three novel <scp>HLA‐DQB1</scp> alleles: <scp><i>HLA‐DQB1*02:186</i></scp>, ‐<scp><i>DQB1*06:02:49</i></scp> and ‐<scp><i>DQB1*06:391</i></scp>. Hla, 2023, 101, 82-84.	0.4	3
2680	Extended genomic sequence of the <scp><i>HLA‐DRB1</i></scp><i>*13:129</i> allele identified in a donor from the Gujarati population. Hla, 2023, 101, 77-78.	0.4	3
2681	Identification of the novel <scp><i>HLA‐DPA1*02:82</i></scp> allele in a Punjabi individual. Hla, 2023, 101, 94-95.	0.4	3
2682	The novel <i><scp>HLA‐A</scp>*02:1009</i> allele was identified in four unrelated bone marrow donors. Hla, 2023, 101, 44-45.	0.4	3
2683	The detection of three novel <scp>HLA‐A</scp> alleles: <i><scp>HLA‐A</scp>*02:1037</i>, <i>‐A*02:1038</i> and <i>‐A*02:1039</i>. Hla, 2023, 101, 146-148.	0.4	3
2684	Identification of <i>HLA‐C*03:599</i> novel allele variant in two South Indian individuals in a Brahmin family. Hla, 2023, 101, 66-67.	0.4	3
2685	Identification of the novel allele, <i>HLA‐C*06:02:38:02</i>, in a Gujarati individual from India. Hla, 2023, 101, 69-70.	0.4	3
2686	The identification of three novel <scp>HLA‐C</scp> alleles: <i>HLA‐C*01:218</i>, <i>HLA‐C*03:550</i> and <i>HLA‐C*05:01:60</i>. Hla, 2023, 101, 61-62.	0.4	3

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2687	The novel <i>HLAâ€C*03:04:74</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 63-64.	0.4	3
2688	The novel <i>HLAâ€B*52:110N</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 58-59.	0.4	3
2689	The novel <scp>HLAâ€B</scp>*52:01:01:23 allele in an individual from Uttar Pradesh, North India. Hla, 2023, 101, 56-57.	0.4	3
2690	Nomenclature for factors of the HLA system, update April, May and June 2022. Hla, 2022, 100, 409-454.	0.4	5
2691	The novel <scp><i>HLAâ€B*55:130</i></scp> allele, identified by <scp>Sanger</scp> dideoxy nucleotide sequencing in a <scp>Chinese</scp> individual. Hla, 2023, 101, 59-60.	0.4	3
2692	Genomic sequence of the <scp><i>HLAâ€A*01:109</i></scp> allele identified in a solid organ donor. Hla, 2023, 101, 42-43.	0.4	3
2693	Characterization of two new <scp>HLA</scp> class I alleles: <scp><i>HLAâ€A</i></scp>*<i>02:1065</i> and <scp><i>HLAâ€C</i></scp>*<i>02:<scp>216N</scp></i>. Hla, 2023, 101, 46-47.	0.4	3
2694	The novel <i>HLAâ€DRB1*14:246</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 78-80.	0.4	3
2695	A novel <i><scp>HLAâ€C</scp>*07</i> allele, <i><scp>HLAâ€C</scp>*07:1024</i>, was identified in a Chinese individual. Hla, 2023, 101, 72-74.	0.4	3
2696	The novel <scp><i>HLAâ€B*15:638</i></scp> allele, identified by <scp>Sanger</scp> dideoxy nucleotide sequencing in a <scp>Chinese</scp> individual. Hla, 2023, 101, 51-53.	0.4	3
2697	Identification of the novel allele, <i>HLAâ€A*02:11:01:06</i>, in two North Indian individuals in a Kshatriya family. Hla, 2023, 101, 43-44.	0.4	3
2698	Extended genomic sequence of the <i>HLA</i>â€<i>DRB1</i>*<i>15</i>:<i>68</i> and <i>HLAâ€DRB1</i>*<i>16</i>:<i>10</i>:<i>01</i> alleles in solid organ donors. Hla, 2022, 100, 655-656.	0.4	3
2699	The novel <scp>HLAâ€DQB1</scp> allele, <scp><i>DQB1*06:443</i></scp>. Hla, 2023, 101, 91-92.	0.4	3
2700	Identification of the novel <scp><i>HLAâ€A</i></scp>*24:02:01:136 allele in a Northern Indian individual from a Muslim family. Hla, 2023, 101, 158-159.	0.4	3
2701	The detection of the novel <i><scp>HLAâ€C</scp>*04:469</i> allele identified in a potential hematopoietic stem cell donor. Hla, 2023, 101, 67-69.	0.4	3
2702	The detection of four novel <scp>HLAâ€A</scp> alleles: <i>HLAâ€A*31:01:48</i>, <i>â€A*31:210</i>, <i>â€A*31:211</i> and <i>â€A*33:03:54</i>. Hla, 2023, 101, 159-161.	0.4	3
2703	Characterization of the novel <scp><i>HLAâ€B*35:547</i></scp> allele detected in a potential hematopoietic stem cell donor. Hla, 2023, 101, 53-54.	0.4	3
2704	Identification of the novel <i><scp>HLAâ€A</scp>*11:01:115</i> allele, detected in two unrelated bone marrow donors. Hla, 2023, 101, 47-48.	0.4	3

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2705	Identification of the novel <scp>HLAâ€DPB1</scp> allele, <i>HLA</i>â€‹<i>DPB1</i>*<i>1326</i>:<i>01</i>, in an <scp>I</scp>talian bone marrow donor. Hla, 2023, 101, 201-202.	0.4	3
2706	Characterization of the novel <scp>HLA</scp> allele, <i>HLA</i> â€¢<i>C</i> * <i>07</i> : <i>04</i> : <i>01</i> : <i>19</i> , in an individual from <scp>M</scp>aharashtra, <scp>I</scp>ndia. Hla, 0, , .	0.4	3
2707	A novel <i>HLAâ€DQB1*04</i> variant, <i>HLAâ€DQB1*04:90</i>, identified in a Chinese Han individual. Hla, 2023, 101, 86-88.	0.4	3
2708	Fullâ€length sequence of the novel <scp><i>HLAâ€C</i></scp>*03:566 allele by nextâ€generation sequencing in a Chinese individual. Hla, 2023, 101, 64-66.	0.4	3
2709	Fullâ€length sequence of the novel <scp><i>HLAâ€C*03:04:74</i></scp> allele by next generation sequencing in a Chinese individual. Hla, 2023, 101, 176-177.	0.4	3
2710	Identification of three novel <scp>HLA</scp> class I alleles, <i>A*11:428</i>, <i>B*37:105</i>, and <i>C*08:249</i>, found in Chinese cord bloods. Hla, 2023, 101, 156-157.	0.4	3
2711	A novel <scp><i>HLAâ€B*57</i></scp> allele, <scp><i>HLAâ€B*57:163</i></scp>, was identified by next generation sequencing typing. Hla, 2023, 101, 171-172.	0.4	3
2712	Genomic fullâ€length confirmatory sequence of <i>HLAâ€DQB1*04:59N</i> allele in three Colombian individuals. Hla, 2023, 101, 195-196.	0.4	3
2713	Identification of the novel <scp><i>HLAâ€B</i></scp>*50:01:17 allele by polymerase chain reaction sequenceâ€based typing. Hla, 2023, 101, 168-169.	0.4	3
2714	Two novel HLAâ€DQB1 alleles identified in potential hematopoietic stem cell donors: <i>DQB1*03:01:50</i> and <i>â€DQB1*03:453</i>. Hla, 2023, 101, 84-86.	0.4	3
2715	Identification of the novel <i>HLA</i> â€¢<i>DPA1</i> * <i>01</i> : <i>106</i> allele by nextâ€generation sequencing in a <scp>K</scp>orean cord blood donor. Hla, 0, , .	0.4	3
2716	Pharmacogenetics of Cutaneous Adverse Drug Reactions. Updates in Clinical Dermatology, 2022, , 3-34.	0.1	0
2717	The frequency of HLA A, B, C, DP, DQ, DR allele in patients of Turkish and Syrian nationals with allogeneic stem cell transplantation. Iraqi Journal of Hematology, 2022, .	0.0	0
2718	Identification of the novel <scp>HLA</scp> allele, <i>HLAâ€A*11:01:01:86</i>, in an individual from north India. Hla, 2023, 101, 153-154.	0.4	3
2719	The novel <i>HLAâ€C*01:02:89</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 174-175.	0.4	3
2720	Nomenclature for factors of the HLA system, update July, August and September 2022. Human Immunology, 2022, , .	1.2	0
2721	Two novel <scp>HLAâ€A</scp> alleles detected in a potential hematopoietic stem cell donors: <i>HLAâ€A*03:418</i> and <i>â€A*26:220</i>. Hla, 2023, 101, 151-153.	0.4	3
2722	Identification of the novel <i>HLAâ€B*08:79:02</i> allele by next generation sequencing. Hla, 2023, 101, 161-162.	0.4	3

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2723	The novel <i>HLAâ€B*40:02:01:39</i> allele in an individual from a Kshatriya family in North India. Hla, 0, , .	0.4	3
2724	Full genomic sequence of the <i>HLAâ€DRB3*02:32</i> allele by Single Molecule Realâ€time Sequencing Technology. Hla, 2023, 101, 188-190.	0.4	3
2725	Identification of three novel <scp>HLA</scp> class <scp>II</scp> alleles: <i>HLAâ€DQA1*02:28</i>, â€<i>DQA1*05:01:08:02</i> and â€<i>DQB1*03:02:01:13</i>. Hla, 2023, 101, 192-193.	0.4	3
2726	The identification of two novel HLA alleles, <i>HLAâ€C*06:02:103</i> and <i>HLAâ€DRB1*01:139</i> in two Russian individuals. Hla, 2023, 101, 181-182.	0.4	3
2727	Identification of a new <scp>HLAâ€B</scp> allele, <i>HLAâ€B*51:371</i>. Hla, 2023, 101, 170-171.	0.4	3
2728	Detection of the novel <scp>HLA</scp> allele, <i>HLAâ€DRB1*08:112</i>, identified in a Danish family. Hla, 2023, 101, 186-187.	0.4	4
2729	86 novel <scp> <i>HLAâ€E</i> </scp> alleles discovered through fullâ€gene sequencing of 6,227 haematopoietic cell transplant patients and unrelated donors. Hla, 0, , .	0.4	3
2730	A novel <scp>HLAâ€C</scp> null allele, <i>HLAâ€C*08:236N</i>, identified by nextâ€generation sequencing in a Chinese individual. Hla, 2023, 101, 184-185.	0.4	3
2731	Identification of the novel <i>HLAâ€B*46:01:33</i> allele by next generation sequencing in a Chinese individual. Hla, 2023, 101, 166-167.	0.4	3
2732	Identification of the novel <scp>HLA</scp> allele, <i>HLAâ€B*37:01:01:19</i>, in an individual from northern India. Hla, 2023, 101, 163-164.	0.4	3
2733	Characterization of the novel <i><scp>HLAâ€C</scp>*05:01:72</i> allele by next generation sequencing. Hla, 2023, 101, 179-181.	0.4	3
2734	The identification of three novel HLAâ€A alleles: <i>HLAâ€A*03:390:02</i>, â€<i>A*26:112:02</i> and â€<i>A*26:221</i>. Hla, 2023, 101, 148-149.	0.4	3
2735	A novel <scp>HLAâ€C</scp> allele, <scp><i>HLAâ€C</i></scp><i>*03:614</i>, identified by nextâ€generation sequencing in a Chinese individual. Hla, 2023, 101, 177-178.	0.4	3
2736	Identification of three novel <scp>HLA</scp> class I alleles: <i>HLAâ€A*02:1008</i>, <i>â€A*24:553</i> and <i>â€C*12:350</i>. Hla, 2023, 101, 145-146.	0.4	3
2737	Characterization of the novel <scp><i>HLAâ€B</i></scp>*<i>58:01:01:19</i> allele by nextâ€generation sequencing. Hla, 2023, 101, 173-174.	0.4	3
2738	Detection of the <i>HLAâ€DQB1*03:151</i> allele in a Taiwanese umbilical cord blood unit. Hla, 2023, 101, 193-194.	0.4	3
2739	Identification of the novel <i>HLAâ€C*07:02:01:184</i> allele in an Indian individual from a Kshatriya family. Hla, 0, , .	0.4	3
2740	The novel <i>HLAâ€A*11:416</i> allele was identified during highâ€resolution <scp>HLA</scp> typing. Hla, 2023, 101, 155-156.	0.4	3

#	ARTICLE	IF	CITATIONS
2741	Histocompatibility in Live Donor Kidney Transplantation. , 2022, , 69-86.		0
2742	Genomic sequence of the <scp><i>HLAâ€A*02:01:209</i></scp> allele identified in a solid organ recipient. Hla, 2023, 101, 270-271.	0.4	3
2743	Identification of the novel HLAâ€DPA1*01:03:43 allele resulting from an intralocus recombination involving the DPA1*04:01:01:03 and DPA1*01:03:01:27 alleles sequenced by Next Generation Sequencing (NGS). International Journal of Immunogenetics, 0, , .	0.8	3
2744	The novel <i>HLAâ€DQB1*04:93</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 305-307.	0.4	3
2745	The novel allele <i>HLAâ€B*35:564</i>, identified by nextâ€generation sequencing in a Chinese individual. Hla, 2023, 101, 283-284.	0.4	3
2746	Discovery of the novel <i>HLA</i>â€<i>B</i>*<i>57</i>:<i>164</i> allele, a variant of <i>HLA</i>â€<i>B</i>*<i>57</i>:<i>01</i>:<i>01</i>:<i>01</i>, in a <scp>T</scp>aiwanese individual. Hla, 2023, 101, 288-289.	0.4	3
2747	The novel <scp><i>HLAâ€B*15:644</i></scp> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 281-283.	0.4	3
2748	The IPD-IMGT/HLA Database. Nucleic Acids Research, 2023, 51, D1053-D1060.	6.5	405
2749	Characterization of the novel <i>HLAâ€DPB1*1348:01</i> allele by sequencingâ€based typing. Hla, 2023, 101, 313-314.	0.4	4
2750	Identification of the novel class <scp>II</scp> allele, <scp><i>HLAâ€DPA1*01:127</i></scp>. Hla, 2023, 101, 199-200.	0.4	3
2751	Nomenclature for factors of the <scp>HLA</scp> system, update July, August and September 2022. Hla, 2022, 100, 673-710.	0.4	3
2752	Identification of the novel <i>HLAâ€DQA1*01:01:01:11</i> and â€‹<i>DQA1*01:03:01:13</i> alleles in individuals from India. Hla, 2023, 101, 296-297.	0.4	3
2753	The novel <i>HLAâ€C*03:605</i> allele, identified using nextâ€generation sequencing in a Chinese individual. Hla, 2023, 101, 291-292.	0.4	3
2754	The novel <scp><i>HLAâ€B*40:01:78</i></scp> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 285-286.	0.4	3
2755	The novel <scp>HLAâ€A</scp> *24:02:159 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2756	Discovery of a novel <i>HLAâ€C*04</i> null allele, <i>HLAâ€C*04:279N</i>, in a Singaporean individual. Hla, 2023, 101, 293-294.	0.4	3
2757	The novel <i>HLAâ€A*02:1068Q</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 271-272.	0.4	3
2758	The novel <scp><i>HLAâ€B*55:131</i></scp> allele, identified by <scp>Sanger</scp> dideoxy nucleotide sequencing in a <scp>Chinese</scp> individual. Hla, 2023, 101, 286-287.	0.4	3

#	Article	IF	Citations
2759	The novel <i>HLAâ€DQB1*03:499N</i> allele identified by <scp>Sanger</scp> dideoxy nucleotide sequencing in a <scp>Chinese</scp> individual. Hla, 2023, 101, 304-305.	0.4	3
2760	The novel <i>HLAâ€DRB1*14:07:03</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 294-296.	0.4	3
2761	The novel <i>HLAâ€DQB1*02:01:44</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 301-302.	0.4	3
2762	Characterization of the novel <i>HLAâ€DPA1*02:01:21</i> allele by sequencingâ€based typing. Hla, 2023, 101, 309-311.	0.4	4
2763	Characterization of the novel <i>HLAâ€DQA1*02:01:14</i> allele by sequencingâ€based typing. Hla, 2023, 101, 298-299.	0.4	4
2764	Characterization of the novel <i>HLAâ€DPB1</i> * <i>11:01:06</i> allele by sequencingâ€based typing. Hla, 0, , .	0.4	1
2765	Characterization of four novel <scp>HLAâ€B</scp> alleles: <i>HLAâ€B*07:02:92</i>, <i>â€B*40:02:36</i>, â€<i>B*18:220</i>, <i>â€B*18:221</i>. Hla, 2023, 101, 274-276.	0.4	3
2766	Identification of the novel <scp> <i>HLAâ€B*07:458</i> </scp> allele, detected in two unrelated bone marrow donors. Hla, 0, , .	0.4	3
2767	Characterization of the novel HLAâ€B*08:302 allele by sequencingâ€based typing. Hla, 0, , .	0.4	4
2768	Characterization of the novel <i>HLA</i> â€•<i>DQA1</i> * <i>05</i> : <i>05</i> : <i>14</i> allele by sequencingâ€based typing. Hla, 0, , .	0.4	4
2769	Identification of the novel <scp> <i>HLAâ€DRB1*11:308</i> </scp> allele in a Greek individual. Hla, 0, , .	0.4	3
2770	Identification of <scp> <i>HLAâ€DPA1</i> </scp> * <i>01:03:01:57</i> and <scp> <i>HLAâ€DPA1</i> </scp> * <i>02:01:01:29</i> from a caseâ€"control study of atopic dermatitis. Hla, 0, , .	0.4	3
2771	The novel <i>HLAâ€C*01:239</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 289-290.	0.4	3
2772	The novel <i>HLAâ€DQB1*03:03:29</i> allele, identified by sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 302-304.	0.4	3
2773	The novel <scp>HLAâ€A</scp> *24:02:160 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2774	The novel <scp><i>HLAâ€C*12:365</i></scp> allele identified in a Gujarati individual from India. Hla, 2023, 101, 554-555.	0.4	3
2775	Identification of the novel <scp>HLAâ€C</scp> *05:269 allele by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2776	Characterization of two new <scp>HLAâ€DPB1</scp> alleles: <i>HLAâ€DPB1*1443:01</i> and â€• <i>DPB1*11:01:07</i>. Hla, 0, , .	0.4	3

#	Article	IF	Citations
2777	Identification and characterization of the novel <scp> <i>HLAâ€A*23:122</i> </scp> allele by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2778	The novel <scp> <i>HLAâ€C*07:1029</i> </scp> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2779	The novel <scp>HLAâ€DRB1</scp> *12:01:12 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2780	The novel <scp>HLAâ€B</scp> *48:01:11 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2781	The novel <i>HLAâ€C*04:490</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 546-548.	0.4	3
2782	The novel <scp>HLAâ€B</scp> *35:568 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2783	Detection of novel <scp><i>HLAâ€A*33:03:62</i></scp>, <scp><i>HLAâ€B*52:01:01:25</i></scp> alleles and confirmation of <scp><i>HLAâ€A*02:01:209</i></scp>. Hla, 2023, 101, 273-274.	0.4	3
2784	IDENTIFICATION OF THE NOVEL <scp>HLAâ€DQA1</scp> *01:04:07 ALLELE WITH A SYNONYMOUS SUBSTITUTION AND AN INTRONIC INSERTION. Hla, 0, , .	0.4	4
2785	The novel <i>HLAâ€C*04:441:01:02</i> allele was identified in three unrelated bone marrow donors. Hla, 0, , .	0.4	3
2786	Recognition of the <scp>HLAâ€C</scp> *07:1047 allele in a Russian bone marrow donor. Hla, 0, , .	0.4	3
2787	Characterization of the novel <scp> <i>HLAâ€B*15:504:02</i> </scp> allele by sequencingâ€based typing. Hla, 0, , .	0.4	3
2788	Identification of the novel <scp>HLAâ€DPB1</scp> *1328:01 allele by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2789	The novel <i>HLAâ€A*24:582</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 529-530.	0.4	3
2790	The novel <i>HLA</i> â€•<i>A</i> * <i>02</i> : <i>07</i> : <i>22</i> allele, identified by <scp>S</scp> anger dideoxy nucleotide sequencing in a <scp>C</scp> hinese individual. Hla, 0, , .	0.4	3
2791	Identification of the novel <scp><i>HLAâ€A*02:01:204</i></scp> allele by nextâ€generation sequencing. Hla, 2023, 101, 513-514.	0.4	3
2792	The novel <scp>HLAâ€C</scp> *12:368 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	0
2793	The novel <scp>HLAâ€B</scp> *13:01:21 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2794	The novel <scp>HLAâ€DQB1</scp> *06:208:02 allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3

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2795	Identification of the novel <scp>HLA</scp> alleles: <i><scp>HLAâ€A</scp>*30:201</i>, <i><scp>HLAâ€B</scp>*51:364</i> and <i><scp>HLAâ€C</scp>*08:241</i>. Hla, 2023, 101, 530-531.	0.4	3
2796	The novel <i>HLAâ€A*11</i> allele, <i>HLAâ€A*11:01:109:02</i>, identified by next generation sequencing. Hla, 2023, 101, 520-522.	0.4	3
2797	Characterization of the novel alleles: <i>HLAâ€A*31:212</i>, <i>HLAâ€B*50:01:20</i> and <i>HLAâ€C*03:593</i>. Hla, 2023, 101, 532-533.	0.4	3
2798	Identification of two novel <scp>HLA</scp> alleles, <i>HLAâ€A*03:344:02</i> and â€<i>DQB1*04:02:24</i> in Russian individuals. Hla, 2023, 101, 517-519.	0.4	3
2799	The novel <i>HLAâ€A*03</i> allele, <i>HLAâ€A*03:440</i>, identified in a potential hematopoietic stem cell donor. Hla, 2023, 101, 519-520.	0.4	3
2800	Identification of the novel <i>HLAâ€C*02:212</i> allele detected in a potential hematopoietic stem cell donor. Hla, 0, , .	0.4	3
2801	Recognition of the novel <i>HLA</i>â€<i>C</i>*<i>07</i>:<i>1002</i> allele identified in a potential hematopoietic stem cell donor. Hla, 2023, 101, 549-551.	0.4	3
2802	The novel <i>HLAâ€C*15:02:58</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2803	Two new <scp>HLA</scp> alleles, <i>HLA</i>â€<i>B</i>*<i>58</i>:<i>140</i> and â€<i>DRB1</i>*<i>07</i>:<i>145</i>, detected in inhabitants from <scp>R</scp>ussia. Hla, 2023, 101, 542-543.	0.4	3
2804	<i>HLAâ€DQA1*05:01:11</i> , a novel <scp>HLAâ€DQA1</scp> allele with an ochre stop codon. Hla, 0, , .	0.4	3
2805	Identification of the novel <scp>HLAâ€A</scp> *11:423 allele by sequencingâ€based typing. Hla, 0, , .	0.4	3
2806	The novel <i>HLAâ€A*24:561:02</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2807	Fullâ€length characterization of the <scp>HLAâ€DQB1</scp> *03:25:01 allele in two Amerindian individuals.. Hla, 0, , .	0.4	3
2808	The novel <i>HLAâ€A*02:1075</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 2023, 101, 516-517.	0.4	3
2809	Identification of a novel allele with a frameshift mutation, <i>HLAâ€DRB4*01:165N</i>, using nextâ€generation sequencing. Hla, 2023, 101, 699-700.	0.4	3
2810	Recognition of the <i>HLAâ€C*14:24:02</i> allele in a Singaporean individual. Hla, 0, , .	0.4	3
2811	Description of the novel allele <i>HLAâ€B*07:473</i> , identified in a bone marrow donor. Hla, 0, , .	0.4	3
2812	Characterization of the novel <i>HLA</i> â€<i>DQB1</i> * <i>02</i> : <i>02</i> : <i>23</i> allele in a Russian hematopoietic stem cell donor. Hla, 0, , .	0.4	3

#	Article	IF	Citations
2813	Identification of three new <scp>HLA</scp> alleles, <i><scp>HLA</scp>â€A*68:298</i>, <i><scp>HLA</scp>â€C*07:1054</i>N and <i><scp>HLA</scp>â€DRB1*15:216</i>. Hla, 2023, 101, 668-670.	0.4	3
2814	Characterization of two new <scp>HLA</scp> class I alleles: <i>HLAâ€B*35:572</i> and <i>HLAâ€C*04:491</i>. Hla, 0, , .	0.4	3
2815	Identification of the novel <scp> <i>HLAâ€DPA1*02:89</i> </scp> allele using nextâ€generation sequencing. Hla, 0, , .	0.4	3
2816	Nomenclature for factors of the HLA system, update April, May and June 2022. Human Immunology, 2023, , .	1.2	0
2817	Recognition of the novel <i>HLAâ€DQB1*05:02:01:15</i> allele in a <scp>Russian</scp> bone marrow donor. Hla, 0, , .	0.4	3
2818	Identification of the novel <i>HLA</i> â€¢<i>DQA1</i> * <i>01</i> : <i>03</i> : <i>07</i> allele using nextâ€generation sequencing. Hla, 0, , .	0.4	3
2819	A novel <scp> <i>HLAâ€DQB1*03</i> </scp> allele, <scp> <i>HLAâ€DQB1*03:509N</i> </scp> , was identified by next generation sequencing. Hla, 0, , .	0.4	3
2820	Description of the novel <scp>HLAâ€C</scp> allele, <i>HLAâ€C*12:376</i> , identified in a deceased <scp>COVID</scp> â€19 patient. Hla, 0, , .	0.4	3
2821	The <scp>HLAâ€DRB1</scp> *04:357 allele first identified in a Chinese transplant recipient. Hla, 0, , .	0.4	3
2822	HLA Class I Supertype Classification Based on Structural Similarity. Journal of Immunology, 2023, 210, 103-114.	0.4	3
2823	Detection of the <i>HLAâ€B*40:02:03</i> allele, a variant of <i>HLAâ€B*40:02:01:01</i> , in a Taiwanese individual. Hla, 0, , .	0.4	3
2824	Recognition of the <i>HLAâ€C*03:04:20</i> allele, a variant of <i>HLAâ€C*03:04:01:01</i> , in a Taiwanese individual. Hla, 0, , .	0.4	3
2825	Characterization of the novel <scp> <i>HLAâ€A*01:383</i> </scp> allele by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2826	Identification of three novel <scp>HLA</scp> class I alleles: <i>HLAâ€B*15:640</i>, â€<i>B*18:01:01:71</i> and â€<i>C*05:275</i>. Hla, 2023, 102, 77-78.	0.4	3
2827	Characterization of the novel HLAâ€DRB1 allele, <i>HLAâ€DRB1*04:328</i> in a Chinese individual. Hla, 2023, 102, 104-106.	0.4	4
2828	Characterization of the novel <i>HLAâ€A*33</i> allele, <i>HLAâ€A*33:03:55</i> in a Chinese individual. Hla, 2023, 102, 224-226.	0.4	3
2829	Identification of the novel <i><scp>HLAâ€C</scp>*12:349</i> allele in a potential hematopoietic stem cell donor. Hla, 2023, 102, 247-248.	0.4	3
2830	Identification of five new <scp>HLAâ€DQB1</scp> intronic variants by nextâ€generation sequencing. Hla, 2023, 101, 702-703.	0.4	3

#	Article	IF	Citations
2831	Identification of the novel <i>HLAâ€B*46:95N</i> allele in a Chinese blood donor. Hla, 0, , .	0.4	3
2833	Two novel <scp>HLA</scp> class <scp>I</scp> alleles, <i>HLAâ€C*04:493</i> and <i>â€A*26:01:78</i>, identified using nextâ€generation sequencing. Hla, 2023, 101, 686-687.	0.4	3
2834	Characterization of two novel <scp>HLA</scp> alleles: <i><scp>HLAâ€A</scp>*26:01:01:53</i> and <i>â€‹<scp>DRB1</scp>*01:141</i>. Hla, 2023, 101, 667-668.	0.4	3
2835	Nomenclature for factors of the <scp>HLA</scp> system, update October, November and December 2022. Hla, 2023, 101, 569-595.	0.4	4
2836	Characterization of the novel HLAâ€DRB1*11:283 allele. Hla, 2023, 102, 110-112.	0.4	3
2837	Discovery of the novel <i> <scp>HLAâ€DRB1</scp> *04:358 </i> allele, a variant of <i> <scp>HLAâ€DRB1</scp> *04 </i> , in a Taiwanese individual. Hla, 0, , .	0.4	3
2838	Detection and characterization of the novel HLA-DPA1*02:66:02N allele, with a premature stop codon in exon 2. Human Immunology, 2023, 84, 296-300.	1.2	5
2839	Identification of the novel <i>HLAâ€C</i> * <i>03:632</i> allele by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2840	Characterization of two new <scp>HLA</scp> alleles, <i>HLAâ€B*18:01:01:74</i> and <i>HLAâ€C*06:02:01:93</i> in Russian individuals. Hla, 2023, 102, 78-79.	0.4	3
2841	Identification of the novel <scp>HLAâ€C</scp> allele <i>HLAâ€C*12:02:02:22</i> , in two individuals from Central India. Hla, 0, , .	0.4	3
2842	Validation and application of new <scp>NGS</scp>â€based <scp>HLA</scp> genotyping to clinical diagnostic practice. Hla, 2023, 101, 496-506.	0.4	4
2843	Distributions of <scp>HLAâ€A</scp>, â€B, â€C, â€‹<scp>DRB1</scp> and â€‹<scp>DQB1</scp> alleles typed by next generation sequencing in Russian volunteer donors. Hla, 2023, 101, 623-633.	0.4	1
2844	The novel <i>HLAâ€B*13:176</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2845	A novel <scp> <i>HLAâ€C*06</i> </scp> allele, <scp> <i>HLAâ€C*06:02:96</i> </scp> , identified by nextâ€generation sequencing in a Chinese family. Hla, 0, , .	0.4	3
2846	Characterization of the novel <i>HLAâ€B*40:495</i> and <i>HLAâ€B*40:512</i> alleles by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2847	Identification of the novel <scp> <i>HLAâ€A*24:589</i> </scp> allele in a <scp>Korean</scp> deceased donor. Hla, 0, , .	0.4	3
2848	The novel <i>HLAâ€C*03:621</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2849	Identification of a new <scp>HLAâ€DRB1</scp> allele, <i> <scp>HLAâ€DRB1</scp> *04:361 </i>. Hla, 0, , .	0.4	3

#	Article	IF	Citations
2850	Identification of six novel HLA alleles, <i>HLAâ€A*31:208</i>, <i>â€B*08:306</i>, <i>â€C*03:582</i>, <i>â€C*04:494</i>, <i>â€C*18:18</i> and <i>â€DRB1*07:133</i>. Hla, 2023, 102, 72-74.	0.4	3
2851	MICA and MICB allele assortment in Finland. Hla, 2023, 102, 52-61.	0.4	6
2852	Identification of the <i>HLAâ€B*40:01:02:59</i> and <i>â€C*05:01:73</i> alleles in individuals from the northern regions of India. Hla, 0, , .	0.4	3
2853	The novel <scp> <i>HLAâ€DPB1*05:01:18</i> </scp> allele, identified by <scp>Sanger</scp> dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2854	Genomic sequence of the <i>HLAâ€B*13:177</i> , <i>HLAâ€C*06:02:104,</i> and <i>HLAâ€C*07:1062N</i> alleles identified in bone marrow donors. Hla, 0, , .	0.4	3
2855	Nextâ€generation sequencing reveals a novel <scp>HLAâ€C</scp> allele, <i>HLAâ€C*15:255</i>. Hla, 2023, 102, 100-102.	0.4	3
2856	Identification of the novel <i>HLAâ€C*01:02:86</i> allele in a Chinese individual. Hla, 0, , .	0.4	3
2857	Nextâ€generation sequencing identifies two novel <scp>HLAâ€DRB1</scp> alleles, <scp> <i>HLAâ€DRB1*04:362</i> </scp> and <scp> <i>HLAâ€DRB1*07:148</i> </scp>. Hla, 0, , .	0.4	3
2858	Discovery of the novel <i> <scp>HLAâ€DQB1</scp> *06:02:01:32 </i> allele, a variant of <i> <scp>HLAâ€DQB1</scp> *06:02:01:01 </i> , in a Russian individual. Hla, 0, , .	0.4	3
2859	Characterization of three novel <scp>HLA</scp> alleles: <i>HLAâ€B*44:481:02</i>, <i>HLAâ€DQB1*03:338N</i>, and <i>HLAâ€DQB1*06:467</i>. Hla, 2023, 102, 83-84.	0.4	3
2860	Characterization of the novel <i> <scp>HLAâ€DRB1</scp> *14:252 </i> allele in a Korean individual by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2861	Identification of the novel <i>HLA</i> â€•<i>A</i> * <i>02</i> : <i>974</i> allele by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2862	The novel <i> <scp>HLAâ€DQB1</scp> *06:01:33 </i> allele was identified by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2863	Description of two new <scp>HLA</scp> alleles, <scp><i>HLAâ€A*26:01:70</i></scp> and <scp><i>HLAâ€A*26:01:74</i></scp> in Chinese individuals. Hla, 2023, 102, 218-221.	0.4	4
2864	Dystrophin Expressing Chimeric (DEC) Cell Therapy for Duchenne Muscular Dystrophy: A First-in-Human Study with Minimum 6 Months Follow-up. Stem Cell Reviews and Reports, 2023, 19, 1340-1359.	1.7	5
2865	Exploration of the role of <scp>NKG2D</scp> ligands <scp>MICA</scp> and <scp>MICB</scp> in <scp>JAK2 V617F</scp> â€positive myeloproliferative neoplasms. Hla, 0, , .	0.4	0
2866	Identification of the novel <scp><i>HLAâ€C*01:242</i></scp> allele using nextâ€generation sequencing. Hla, 2023, 102, 236-237.	0.4	3
2867	Characterization of the novel <scp><i>HLAâ€DPA1*01:144</i></scp> allele by next generation sequencing. Hla, 2023, 102, 264-266.	0.4	3

#	Article	IF	Citations
2868	The novel <i>HLAâ€A*29:172</i> allele identified in a patient indicated for hematopoietic stem cell transplantation. Hla, 2023, 102, 221-223.	0.4	3
2869	Characterization of the novel <i>HLAâ€C*03:552</i> allele by sequencingâ€based typing. Hla, 2023, 102, 237-238.	0.4	3
2870	Identification of the novel <i>HLAâ€C*04:495</i> allele by sequencingâ€based typing. Hla, 2023, 102, 238-239.	0.4	3
2871	The novel <scp>HLAâ€C</scp> allele, <i>HLAâ€C*07:02:141</i> was identified in a Chinese individual. Hla, 2023, 102, 241-243.	0.4	3
2872	Characterization of the novel <i><scp>HLAâ€C</scp>*08:232</i> allele by nextâ€generation sequencing. Hla, 2023, 102, 245-247.	0.4	3
2873	World Marrow Donor Association guidelines for the reporting of novel <scp>HLA</scp> alleles. Hla, 2023, 102, 62-64.	0.4	2
2874	A novel <scp>HLAâ€DPB1</scp> allele, <i>HLAâ€DPB1*1447:01</i> , identified by nextâ€generation sequencing. Hla, 0, , .	0.4	3
2875	Identification of the novel <i>HLAâ€DQA1*01:04:04</i> allele by nextâ€generation sequencing. Hla, 2023, 102, 252-254.	0.4	3
2876	The novel <i>HLAâ€C*12:368</i> allele, identified by Sanger dideoxy nucleotide sequencing in a Chinese individual. Hla, 0, , .	0.4	3
2877	Trasplante renal con HLA idÃ©ntico de donante vivo y cadavÃ©rico: experiencia de la FundaciÃ³n Valle de Lili, Cali, Colombia. Revista Colombiana De Cirugia, 2016, 31, 170-177.	0.2	2
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2930	Tutorial: a statistical genetics guide to identifying HLA alleles driving complex disease. Nature Protocols, 2023, 18, 2625-2641.	5.5	5
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