

# F-M Zhu

## List of Publications by Year in descending order

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241  
papers

1,842  
citations

430442

18  
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454577

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256  
docs citations

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times ranked

1089  
citing authors

#	ARTICLE	IF	CITATIONS
1	Distribution of HLA allele frequencies in 82 Chinese individuals with coronavirus disease-2019 (COVID-19). <i>Hla</i> , 2020, 96, 194-196.	0.4	152
2	Distribution of killer cell immunoglobulin-like receptor genes in the Chinese Han population. <i>Tissue Antigens</i> , 2005, 65, 556-563.	1.0	107
3	HLA common and well-documented alleles in China. <i>Hla</i> , 2018, 92, 199-205.	0.4	72
4	Mutations in 3 <sup>rd</sup> -long terminal repeat of HERV-W family in chromosome 7 upregulate syncytin-1 expression in urothelial cell carcinoma of the bladder through interacting with c-Myb. <i>Oncogene</i> , 2014, 33, 3947-3958.	2.6	67
5	Distribution of ABO blood group allele and identification of three novel alleles in the Chinese Han population. <i>Vox Sanguinis</i> , 2010, 98, 554-559.	0.7	58
6	The polymorphism of HLA-A, -B, -C, -DRB3/4/5, -DRB1, -DQB1 loci in Zhejiang Han population, China using NGS technology. <i>International Journal of Immunogenetics</i> , 2021, 48, 485-489.	0.8	49
7	The distributions of HLA-A, HLA-B, HLA-C, HLA-DRB1 and HLA-DQB1 allele and haplotype at high-resolution level in Zhejiang Han population of China. <i>International Journal of Immunogenetics</i> , 2019, 46, 7-16.	0.8	46
8	A population-based study comparing multiple sclerosis clinic users and non-users in British Columbia, Canada. <i>European Journal of Neurology</i> , 2016, 23, 1093-1100.	1.7	29
9	A novel mutation +5904 C>T of RUNX1 site in the erythroid cell-specific regulatory element decreases the ABO antigen expression in Chinese population. <i>Vox Sanguinis</i> , 2018, 113, 594-600.	0.7	28
10	Molecular basis for para-Bombay phenotypes in Chinese persons, including a novel nonfunctional FUT1 allele. <i>Transfusion</i> , 2005, 45, 725-730.	0.8	25
11	Analysis for complete genomic sequence of HLA-B and HLA-C alleles in the Chinese Han population. <i>International Journal of Immunogenetics</i> , 2011, 38, 281-284.	0.8	22
12	Serological characteristic and molecular basis of A2 subgroup in the Chinese population. <i>Transfusion and Apheresis Science</i> , 2013, 48, 67-74.	0.5	21
13	HLA-A, -B and -DRB1 allele and haplotype frequencies of 8333 Chinese Han from the Zhejiang province, China. <i>International Journal of Immunogenetics</i> , 2016, 43, 86-95.	0.8	21
14	KIR3DL1 genetic diversity and phenotypic variation in the Chinese Han population. <i>Genes and Immunity</i> , 2014, 15, 8-15.	2.2	20
15	Molecular basis and zygosity determination of D variants including identification of four novel alleles in Chinese individuals. <i>Transfusion</i> , 2015, 55, 137-143.	0.8	20
16	Identification of a novel B allele with missense mutation (c.98G>C) in the ABO gene. <i>Transfusion</i> , 2017, 57, 219-220.	0.8	20
17	Birth outcomes of pregnancies fathered by men with multiple sclerosis. <i>Multiple Sclerosis Journal</i> , 2014, 20, 1260-1264.	1.4	19
18	Distribution of MICA diversity in the Chinese Han population by polymerase chain reaction sequence-based typing for exons 2-6. <i>Tissue Antigens</i> , 2009, 73, 358-363.	1.0	18

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19	MICA, MICB Polymorphisms and Linkage Disequilibrium with HLA-B in a Chinese Mongolian Population. <i>Scandinavian Journal of Immunology</i> , 2016, 83, 456-462.	1.3	17
20	Five novel HLA-A alleles, HLA-A*030108, A*2491, A*2498, A*330303, A*3317 were identified by polymerase chain reaction sequence based typing. <i>Tissue Antigens</i> , 2009, 74, 432-434.	1.0	16
21	Identification of a new HLA-DRB1 allele, HLA-DRB1*1212, and confirmation of HLA-B*1586*. <i>Tissue Antigens</i> , 2005, 65, 582-583.	1.0	15
22	Investigation of killer cell immunoglobulin-like receptor KIR2DL4 diversity by sequence-based typing in Chinese population. <i>Tissue Antigens</i> , 2006, 67, 214-221.	1.0	14
23	A dispermic chimera was identified in a healthy man with mixed field agglutination reaction in ABO blood grouping and mosaic 46, XY/46, XX karyotype. <i>Transfusion and Apheresis Science</i> , 2013, 48, 223-228.	0.5	14
24	Distribution of MICB diversity in the Zhejiang Han population: PCR sequence-based typing for exons 2-6 and identification of five novel MICB alleles. <i>Immunogenetics</i> , 2013, 65, 485-492.	1.2	14
25	Analysis of the complete genomic sequence of HLA-A alleles in the Chinese Han population. <i>International Journal of Immunogenetics</i> , 2009, 36, 351-360.	0.8	13
26	Detection of bacterial contamination of apheresis platelets in a Chinese Blood Center. <i>Transfusion Medicine</i> , 2009, 19, 357-362.	0.5	13
27	HLA-A, HLA-B, HLA-C, DRB1 allele and haplotype frequencies in 6384 umbilical cord blood units and transplantation matching and engraftment statistics in the Zhejiang cord blood bank of China. <i>International Journal of Immunogenetics</i> , 2014, 41, 13-19.	0.8	13
28	The combinatorial diversity of KIR and HLA class I allotypes in Peninsular Malaysia. <i>Immunology</i> , 2021, 162, 389-404.	2.0	12
29	HLA-A*31:56 and HLA-A*31:59 were identified by polymerase chain reaction sequence-based typing in Chinese individuals. <i>Tissue Antigens</i> , 2012, 79, 388-389.	1.0	11
30	Two novel alleles, HLA-B*46:01:11 and HLA-B*51:01:39 were identified in Chinese bone marrow donors. <i>Tissue Antigens</i> , 2015, 86, 144-145.	1.0	11
31	Diversity of the killer cell immunoglobulin-like receptor gene KIR2DS4 in the Chinese population. <i>Tissue Antigens</i> , 2007, 69, 133-138.	1.0	10
32	Analysis of the complete cDNA sequences of HLA-DRB1 alleles with group-specific amplification primers in the Chinese Han population. <i>Tissue Antigens</i> , 2011, 77, 329-332.	1.0	10
33	Human platelet antigen allele frequencies and new mutations on platelet glycoprotein genes in the Chinese Han population. <i>Transfusion Medicine</i> , 2011, 21, 330-337.	0.5	10
34	A novel FUT1 allele was identified in a Chinese individual with para-Bombay phenotype. <i>Transfusion Medicine</i> , 2011, 21, 385-393.	0.5	10
35	Identification of two novel alleles HLA-B*070209 and HLA-B*130205 by polymerase chain reaction sequence-based typing. <i>Tissue Antigens</i> , 2010, 75, 174-175.	1.0	9
36	Characterization of a novel allele, HLA-DQB1*05:03:05. <i>Tissue Antigens</i> , 2012, 79, 311-312.	1.0	9

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37	A novel allele <i>HLA*DRB1*11:119</i> was identified by polymerase chain reaction sequence-based typing in a Chinese individual. <i>Tissue Antigens</i> , 2012, 80, 68-70.	1.0	9
38	<i>HLA-A</i> locus allelic dropout in <i>Sanger</i> sequence-based typing due to the single nucleotide polymorphism of exon 1. <i>International Journal of Immunogenetics</i> , 2015, 42, 457-460.	0.8	9
39	Identification of a new allele, <i>HLA-A*3308</i> , in the Chinese population. <i>Tissue Antigens</i> , 2006, 67, 168-169.	1.0	8
40	Identification of a novel <i>HLA*DRB1*03:03:04</i> allele by polymerase chain reaction sequence-based typing in a Chinese leukemia patient. <i>Tissue Antigens</i> , 2012, 79, 214-215.	1.0	8
41	Identification of a novel <i>HLA-DQB1*03:38</i> allele by polymerase chain reaction sequence-based typing in a Chinese bone marrow donor. <i>Tissue Antigens</i> , 2012, 80, 198-199.	1.0	8
42	Identification of a novel <i>HLA-A*33:03:11</i> allele by polymerase chain reaction sequence-based typing in a Chinese cord blood donor. <i>Tissue Antigens</i> , 2013, 82, 59-60.	1.0	8
43	Characterization of a novel allele, <i>HLA-DQB1*06:47</i> . <i>Tissue Antigens</i> , 2013, 82, 74-75.	1.0	8
44	Two novel <i>HLA*DRB1*03:03</i> alleles, <i>HLA*DRB1*03:03:08</i> and <i>HLA*DRB1*03:03:13</i> , were identified in Chinese individuals. <i>Tissue Antigens</i> , 2015, 86, 66-68.	1.0	8
45	A novel <i>HLA-C</i> allele, <i>HLA-C*08:128</i> , was identified in a leukemia patient by polymerase chain reaction sequence-based typing. <i>Hla</i> , 2017, 89, 168-170.	0.4	8
46	Associations of killer cell immunoglobulin-like receptors with acute myeloid leukemia in Chinese populations. <i>Human Immunology</i> , 2017, 78, 269-273.	1.2	8
47	The novel <i>HLA*02:787</i> allele was identified by polymerase chain reaction sequence-based typing. <i>Hla</i> , 2020, 96, 211-213.	0.4	8
48	Characterization of three novel HLA alleles, <i>HLA-B*4613</i> , <i>HLA-B*4614</i> and <i>HLA-B*4618</i> , in Chinese individuals. <i>Tissue Antigens</i> , 2009, 73, 609-611.	1.0	7
49	Identification of the novel allele <i>HLA*DRB1*150205</i> by polymerase chain reaction sequence-based typing in a Chinese individual. <i>Tissue Antigens</i> , 2009, 74, 173-175.	1.0	7
50	Identification of two novel <i>HLA-B*54</i> alleles, <i>B*54:01:03</i> and <i>B*54:01:04</i> by polymerase chain reaction sequence-based typing. <i>Tissue Antigens</i> , 2013, 82, 63-65.	1.0	7
51	A novel HLA allele, <i>HLA-C*01:02:18</i> , was identified by polymerase chain reaction sequence-based typing in a Chinese leukemia patient. <i>Tissue Antigens</i> , 2013, 82, 65-66.	1.0	7
52	Identification of the novel <i>HLA-B*15:257</i> allele by polymerase chain reaction sequence-based typing in a Chinese individual. <i>Tissue Antigens</i> , 2013, 82, 62-63.	1.0	7
53	Identification of two novel alleles <i>HLA-A*11:133</i> and <i>A*11:02:05</i> by polymerase chain reaction sequence-based typing. <i>Tissue Antigens</i> , 2014, 84, 409-412.	1.0	7
54	Characterization of the novel <i>HLA-B</i> allele: <i>HLA-B*15:437</i> in a Chinese bone marrow donor. <i>Hla</i> , 2020, 96, 511-513.	0.4	7

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55	Characterization of the novel <sc><i>HLA*31:01:34</i></sc> allele by polymerase chain reaction sequencing-based typing. Hla, 2020, 96, 502-504.	0.4	7
56	Identification of the novel HLA*13:109 allele by polymerase chain reaction sequence-based typing. Hla, 2020, 96, 342-343.	0.4	7
57	Characterization of the novel HLA*15:160N allele. Hla, 2020, 96, 227-229.	0.4	7
58	A novel HLA-B*15 allele, B*9524, identified by sequence-based typing in the Chinese population. Tissue Antigens, 2007, 70, 521-522.	1.0	6
59	Full-length sequence of a novel null allele, HLA*2486N. Tissue Antigens, 2009, 73, 63-65.	1.0	6
60	Identification of a novel allele HLA-B*35:137 in a Chinese leukemia patient. Tissue Antigens, 2010, 76, 498-499.	1.0	6
61	Identification of a new HLA*11:78N allele by polymerase chain reaction sequence-based typing. Tissue Antigens, 2011, 77, 257-258.	1.0	6
62	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
63	Identification of a novel HLA-B*46:29 allele by polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2013, 81, 231-233.	1.0	6
64	A novel <sc><i>HLA*DRB1</i></sc>*05:15</i> allele was identified in a Chinese individual. Tissue Antigens, 2014, 84, 246-248.	1.0	6
65	Allelic polymorphism, mRNA and antigen expression of KIR2DL1 in the Chinese Han population. Human Immunology, 2014, 75, 245-249.	1.2	6
66	Comparison of the KIR3DS1/Bw4 distribution in Chinese healthy and acute myeloid leukemia individuals. Human Immunology, 2015, 76, 79-82.	1.2	6
67	Two novel alleles, <sc><i>HLA</i></sc>*A*02:07:06</i> and <sc><i>HLA</i></sc>*A*02:426</i>, were identified in Chinese individuals. Tissue Antigens, 2015, 85, 499-501.	1.0	6
68	Characterization of a novel HLA-B*40 allele, HLA-B*40:186:02, by cloning and sequencing. International Journal of Immunogenetics, 2016, 43, 240-241.	0.8	6
69	Identification of the novel <i>HLA*DRB1*12:50</i> allele by polymerase chain reaction sequence-based typing in a Chinese individual. Hla, 2016, 87, 473-474.	0.4	6
70	Simultaneous genotyping of human platelet alloantigen 1 to 28bw systems by multiplex polymerase chain reaction sequence-based typing. Vox Sanguinis, 2017, 112, 360-366.	0.7	6
71	Identification of 2 novel <sc>HLA</sc>*B alleles, <sc>HLA</sc>*B*55:02:09</i> and <sc>HLA</sc>*B*55:80</i> in Chinese individuals. Hla, 2017, 90, 48-50.	0.4	6
72	Characterization of the novel HLA*08:159 allele by next-generation sequencing. Hla, 2019, 93, 502-503.	0.4	6

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73	Identification of the novel <sc><i>HLA-B*46:74</i></sc> allele by polymerase chain reaction sequence-based typing. Hla, 2020, 96, 520-521.	0.4	6
74	<i><sc>HLA-DRB1</sc>*14:54:09</i> and <i><sc>DRB1</sc>*14:54:10</i>, were identified by next-generation sequencing in Chinese cord blood donors. Hla, 2021, 97, 166-169.	0.4	6
75	Description of two new <sc>HLA</sc> alleles: <i><sc>HLA-B</sc>*46:80</i> and <i><sc>HLA-B</sc>*46:81</i> identified in Chinese individuals. Hla, 2021, 98, 391-393.	0.4	6
76	Characteristic of HBV nucleic acid amplification testing yields from blood donors in China. BMC Infectious Diseases, 2021, 21, 714.	1.3	6
77	The impact of nucleic acid testing to detect human immunodeficiency virus, hepatitis C virus, and hepatitis B virus yields from a single blood center in China with 10-years review. BMC Infectious Diseases, 2022, 22, 279.	1.3	6
78	Sequence analysis of the novel allele HLA-B*5408 N in the Chinese population. Tissue Antigens, 2006, 68, 182-182.	1.0	5
79	Identification of a novel HLA-B*15 allele, B*9529, in the Chinese population. Tissue Antigens, 2008, 71, 254-255.	1.0	5
80	A novel HLA-DPA1*0204 allele was identified in a Chinese individual. Tissue Antigens, 2008, 71, 577-578.	1.0	5
81	Sequences variations in 5'-flanking region of ABO gene and correlation with ABO alleles in the indigenous Chinese. Vox Sanguinis, 2008, 94, 227-233.	0.7	5
82	Identification of the novel allele HLA-B*580102 by sequence-based typing in a Chinese individual. Tissue Antigens, 2009, 73, 74-75.	1.0	5
83	Identification of a new HLA-A*29 allele, HLA-A*290102 by sequence-based typing in a Chinese individual. Tissue Antigens, 2009, 74, 168-169.	1.0	5
84	A novel allele HLA-B*550203 was identified by polymerase chain reaction sequence-based typing. Tissue Antigens, 2009, 74, 254-255.	1.0	5
85	A novel allele HLA-B*550205 was identified by polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2010, 76, 155-6.	1.0	5
86	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
87	HLA-A*02:335 and HLA-A*02:370 were identified by polymerase chain reaction sequence-based typing in Chinese individuals. Tissue Antigens, 2012, 80, 537-539.	1.0	5
88	Identification of the novel HLA-A*26:79 allele by polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2013, 82, n/a-n/a.	1.0	5
89	Identification by sequence-based high-resolution typing of a novel HLA-C allele, <i>C*14:52</i>, in a bone marrow donor. Tissue Antigens, 2014, 83, 366-367.	1.0	5
90	A novel allele, <i>HLA-DRB1*10:07</i> was identified in a Chinese individual. Tissue Antigens, 2015, 86, 68-69.	1.0	5

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91	Identification a novel <i>HLA-B*27:105</i> allele in a Chinese bone marrow donor by polymerase chain reaction sequence-based typing. <i>Tissue Antigens</i> , 2015, 85, 212-213.	1.0	5
92	A novel <i>HLA-A*32</i> allele, <i>A*32:67</i> was identified by polymerase chain reaction sequence-based typing in a Chinese individual. <i>Tissue Antigens</i> , 2015, 85, 507-508.	1.0	5
93	Identification of a novel <i>HLA-B*54:34</i> allele by polymerase chain reaction sequence-based typing in a Chinese leukemia patient. <i>Hla</i> , 2016, 87, 180-182.	0.4	5
94	A novel <i>HLA-A*02</i> allele, <i>A*02:543</i> was identified by polymerase chain reaction sequence-based typing in a Chinese cord blood donor. <i>Hla</i> , 2016, 87, 384-385.	0.4	5
95	HLA-B allele dropout in PCR sequence-specific oligonucleotide probe typing due to intronic polymorphism in the novel <i>B*58:01:01:02</i> allele. <i>International Journal of Immunogenetics</i> , 2016, 43, 180-183.	0.8	5
96	Identification of two novel <i>HLA-A</i> alleles, <i>A*03:181</i> and <i>A*03:229</i> in Chinese individuals. <i>Hla</i> , 2016, 87, 165-166.	0.4	5
97	Identification of the novel <i>HLA-B*13:83</i> allele by polymerase chain reaction sequence-based typing in a Chinese cord blood donor. <i>Hla</i> , 2017, 89, 241-242.	0.4	5
98	Identification of the novel <i>HLA-DRB1*08:69</i> allele by polymerase chain reaction sequence-based typing in a Chinese cord blood donor. <i>Hla</i> , 2017, 89, 64-65.	0.4	5
99	Identification of the novel <i>HLA-C*07:530</i> allele by polymerase chain reaction sequence-based typing. <i>Hla</i> , 2018, 91, 213-215.	0.4	5
100	Identification of the novel <i>HLA-DQB1*06:209</i> allele in a Chinese individual. <i>Hla</i> , 2018, 91, 543-544.	0.4	5
101	Novel method for simultaneously detecting HPA and HLA antibodies using Luminex microbeads. <i>Journal of Translational Medicine</i> , 2019, 17, 249.	1.8	5
102	Identification of the novel <i>HLA-B*40:357</i> allele by polymerase chain reaction sequence-based typing. <i>Hla</i> , 2019, 94, 68-70.	0.4	5
103	Characterization of the novel <i>HLA-C*03:372</i> allele by next-generation sequencing. <i>Hla</i> , 2019, 94, 71-73.	0.4	5
104	Identification of the novel <i>HLA-C*12:220</i> allele in a Chinese individual. <i>Hla</i> , 2019, 94, 80-81.	0.4	5
105	Characterization of the novel <i>HLA-C*08:154</i> allele by sequencing-based typing. <i>Hla</i> , 2019, 93, 238-240.	0.4	5
106	Identification of a novel <i>HLA-B*4061</i> allele in the Chinese population+. <i>Tissue Antigens</i> , 2005, 66, 705-706.	1.0	4
107	A new <i>HLA-A</i> allele, <i>A*3113</i> , identified by sequence-based typing in the Chinese population+. <i>Tissue Antigens</i> , 2006, 67, 250-251.	1.0	4
108	Identification of a novel <i>HLA-B*4608</i> allele in the Chinese population+. <i>Tissue Antigens</i> , 2006, 67, 253-254.	1.0	4

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109	Identification of a novel HLA-B*39 allele, B*3936, by sequencing-based typing+. Tissue Antigens, 2006, 67, 341-342.	1.0	4
110	Identification of a novel HLA-A*2459 allele in the Chinese population+. Tissue Antigens, 2006, 68, 177-177.	1.0	4
111	Sequence analysis of the novel allele HLA-B*5522 in the Chinese population. Tissue Antigens, 2007, 69, 202-203.	1.0	4
112	Sequence analysis of the novel allele HLA-A*110106 in the Chinese population. Tissue Antigens, 2007, 70, 70-71.	1.0	4
113	HLA-A*2468, a new allele identified by sequence-based typing in the Chinese population. Tissue Antigens, 2007, 70, 256-257.	1.0	4
114	Identification of a novel allele HLA-A*3117 in the Chinese cord blood donor. Tissue Antigens, 2007, 70, 517-518.	1.0	4
115	A novel HLA-B*15 allele, B*9534, identified by sequence-based typing in the Chinese population. Tissue Antigens, 2008, 71, 256-257.	1.0	4
116	Identification of a novel HLA-B*510203 allele in a Chinese individual. Tissue Antigens, 2008, 71, 572-573.	1.0	4
117	A novel HLA-DRB1*135002 allele was identified in a cord blood donor. Tissue Antigens, 2008, 72, 78-79.	1.0	4
118	HLA-B*5530, a new allele was identified by sequence-based typing in a Chinese individual. Tissue Antigens, 2008, 72, 602-603.	1.0	4
119	HLA-A*1136 and HLA-A*1138 were identified by polymerase chain reaction sequence-based typing in Chinese individuals. Tissue Antigens, 2009, 73, 604-606.	1.0	4
120	HLA-B*4088, a new allele was identified by polymerase chain reaction sequence-based typing in a Chinese cord blood donor. Tissue Antigens, 2010, 75, 79-80.	1.0	4
121	Identification of a novel allele HLA-A*240220 by polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2010, 76, 150-1.	1.0	4
122	Identification of a novel allele HLA-A*1149 in a Chinese leukemia patient. Tissue Antigens, 2010, 76, 250-251.	1.0	4
123	Identification of a novel HLA-A*11:57 allele in a Chinese bone marrow donor. Tissue Antigens, 2010, 76, 332-333.	1.0	4
124	Identification of a new HLA-A*02 allele, HLA-A*02:230, using polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2011, 77, 150-151.	1.0	4
125	Characterization of a novel HLA allele, HLA-B*40:128, in a Chinese individual. Tissue Antigens, 2011, 77, 260-261.	1.0	4
126	Identification of a novel HLA-B*13:41 allele in a Chinese bone marrow donor. Tissue Antigens, 2011, 78, 399-400.	1.0	4



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127	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
128	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
129	Sequence-based HLA high-resolution typing of a bone marrow donor/recipient pair reveals the novel HLA allele <i>HLA-C*07:208</i> . Tissue Antigens, 2012, 80, 276-278.	1.0	4
130	Identification of a novel <i>HLA-C*01:61</i> allele by polymerase chain reaction sequence-based typing in a Chinese leukemia patient. Tissue Antigens, 2012, 80, 275-276.	1.0	4
131	Prevalence of the HPA-18w to -21w alleles in the Chinese Han population. International Journal of Immunogenetics, 2013, 40, 99-103.	0.8	4
132	Molecular basis for the p and P <sup>k</sup> phenotypes in three Chinese individuals. Transfusion Medicine, 2013, 23, 132-133.	0.5	4
133	A novel HLA allele, <i>HLA-B*40:227</i> , was identified by polymerase chain reaction sequence-based typing in a Chinese individual. Tissue Antigens, 2013, 82, 208-209.	1.0	4
134	A novel <i>HLA-C</i> allele, <i>C*08:01:10</i> was identified in a Chinese leukemia patient. Tissue Antigens, 2014, 84, 419-420.	1.0	4
135	Identification of a novel <i>HLA-DPB1</i> allele, <i>HLA-DPB1*167:01</i> , in a Chinese individual. Tissue Antigens, 2014, 83, 299-300.	1.0	4
136	A novel allele, <i>HLA-B*54:29</i> , identified by sequence-based typing in a Chinese bone marrow donor. Tissue Antigens, 2014, 83, 430-432.	1.0	4
137	Genomic full-length sequence of a novel <i>HLA-B*39:01:01:03</i> allele was identified in a Chinese individual. Tissue Antigens, 2014, 83, 132-134.	1.0	4
138	Identification of a novel <i>HLA-C*04:144</i> allele by polymerase chain reaction sequence-based typing. Tissue Antigens, 2014, 84, 245-246.	1.0	4
139	Identification of a novel <i>HLA-B*35:227</i> allele by polymerase chain reaction sequence-based typing in a Chinese bone marrow donor. Tissue Antigens, 2014, 84, 240-242.	1.0	4
140	Identification of a novel <i>HLA-A*02:06:14</i> allele by polymerase chain reaction sequence-based typing in a Chinese bone marrow donor. Tissue Antigens, 2015, 85, 287-288.	1.0	4
141	<i>HLA-C*06:103</i> , a novel allele was identified in a Chinese patient awaiting hematopoietic stem cell transplantation. Tissue Antigens, 2015, 85, 510-511.	1.0	4
142	Genomic full-length sequence of a novel <i>HLA-A*11:01:01:02</i> allele was identified in a Chinese bone marrow donor. International Journal of Immunogenetics, 2016, 43, 184-186.	0.8	4
143	A novel <i>HLA-B*15</i> allele, <i>HLA-B*15:326</i> , was identified in a Chinese bone marrow donor. Hla, 2016, 87, 176-177.	0.4	4
144	Two novel <i>HLA-A</i> 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

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145	Identification of the novel <i>&lt;sc&gt;HLA&lt;/sc&gt;â€B*52:42&lt;/i&gt; allele by polymerase chain reaction sequence-based typing in a Chinese bone marrow donor. Hla, 2016, 87, 400-402.</i>	0.4	4
146	Identification of two novel <i>&lt;sc&gt;HLAâ€DQB1&lt;/sc&gt;</i> alleles, <i>&lt;sc&gt;HLAâ€DQB1&lt;/sc&gt;*03:164&lt;/i&gt; and <i>&lt;sc&gt;HLAâ€DQB1&lt;/sc&gt;*03:165&lt;/i&gt; in Chinese individuals. Hla, 2016, 88, 316-317.</i></i>	0.4	4
147	A frame shift due to a two-nucleotide insertion results in the an <i>&lt;sc&gt;HLA&lt;/sc&gt;â€B</i> null allele, <i>&lt;sc&gt;B*39:97N&lt;/sc&gt;</i> . Hla, 2016, 88, 312-313.	0.4	4
148	Identification of the novel <i>&lt;sc&gt;HLA&lt;/sc&gt;â€B*52:01:27&lt;/i&gt; allele by polymerase chain reaction sequence-based typing. Hla, 2017, 89, 250-251.</i>	0.4	4
149	Identification of the novel null allele, <i>&lt;sc&gt;HLA&lt;/sc&gt;â€C*01:109N&lt;/sc&gt;</i> , using polymerase chain reaction sequence-based typing in a Chinese leukemia patient. Hla, 2017, 89, 252-253.	0.4	4
150	A novel mutation in <i>&lt;sc&gt;A4GALT&lt;/sc&gt;</i> was identified in a Chinese individual with p phenotype. Transfusion, 2017, 57, 215-216.	0.8	4
151	The novel null allele, <i>&lt;sc&gt;HLAâ€B*40:338N&lt;/sc&gt;</i> , was identified in a Chinese leukemia patient. Hla, 2018, 91, 303-305.	0.4	4
152	Characterization of three new HLA Class I Alleles in Chinese individuals, <i>HLAâ€B*46:68,â€B*46:71,â€B*46:72</i> . International Journal of Immunogenetics, 2018, 45, 351-353.	0.8	4
153	Identification of the novel <i>HLAâ€C*04:286</i> allele by next-generation sequencing in a Chinese cord blood donor. Hla, 2019, 94, 73-74.	0.4	4
154	Identification of the novel <i>HLAâ€C*03:365</i> allele in a Chinese bone marrow donor. Hla, 2019, 93, 231-232.	0.4	4
155	Identification of two novel <i>HLAâ€C</i> alleles, <i>&lt;sc&gt;HLAâ€C*07:02:92&lt;/sc&gt;</i> and <i>&lt;sc&gt;HLAâ€C*07:828&lt;/sc&gt;</i> in Chinese individuals. Hla, 2020, 96, 104-106.	0.4	4
156	Identification of the novel <i>&lt;sc&gt;HLAâ€DRB1&lt;/sc&gt;*04:305&lt;/i&gt; allele in a Chinese leukemia patient. Hla, 2021, 98, 180-182.</i>	0.4	4
157	Six splice site variations, three of them novel, in the ABO gene occurring in nine individuals with ABO subtypes. Journal of Translational Medicine, 2021, 19, 470.	1.8	4
158	Identification of the novel <i>&lt;sc&gt;HLAâ€C&lt;/sc&gt;*07:976</i> allele by polymerase chain reaction sequence-based typing. Hla, 2022, 99, 643-644.	0.4	4
159	<i>&lt;sc&gt;HLAâ€DRB1&lt;/sc&gt;*15:01:43&lt;/i&gt; and <i>&lt;sc&gt;HLAâ€DRB1&lt;/sc&gt;*15:01:44&lt;/i&gt; alleles were identified by next-generation sequencing. Hla, 2022, 99, 664-666.</i></i>	0.4	4
160	Characterization of the novel <i>&lt;sc&gt;HLAâ€A*11:383N&lt;/sc&gt;</i> and <i>&lt;sc&gt;HLAâ€A*11:388N&lt;/sc&gt;</i> alleles by next-generation sequencing. Hla, 2022, 99, 374-375.	0.4	4
161	The novel <i>&lt;sc&gt;HLAâ€C&lt;/sc&gt;</i> allele, <i>&lt;sc&gt;HLAâ€C*03:537&lt;/sc&gt;</i> in a Chinese individual. Hla, 2022, 100, 376-377.	0.4	4
162	The novel <i>&lt;sc&gt;HLAâ€C&lt;/sc&gt;</i> allele, <i>C*03:538</i> was identified by next-generation sequencing. Hla, 0, , .	0.4	4

#	ARTICLE	IF	CITATIONS
163	Identification of a novel allele HLA-A*9206 by sequence-based typing in the Chinese population. Tissue Antigens, 2007, 70, 257-257.	1.0	3
164	Two novel alleles HLA-B*9536 and B*4612 were identified in a healthy Chinese individual. Tissue Antigens, 2008, 71, 573-575.	1.0	3
165	Characterization of a novel HLA allele HLA-B*15:178 in a Chinese individual. Tissue Antigens, 2010, 76, 333-334.	1.0	3
166	A novel allele, <sc><i>HLA</i></sc><i>B*55:70</i> was identified in a Chinese cord blood donor. Hla, 2016, 87, 183-185.	0.4	3
167	Identification of the novel <i><sc>HLA</sc>B*27:147</i> allele by polymerase chain reaction sequence-based typing. Hla, 2017, 90, 115-116.	0.4	3
168	Identification of the novel <i><sc>HLA</sc>B*40:01:41</i> allele by polymerase chain reaction sequence-based typing in a Chinese cord blood donor. Hla, 2017, 90, 118-120.	0.4	3
169	Identification of the novel <i><sc>HLA</sc>DRB1*15:127</i> allele by polymerase chain reaction sequence-based typing in a Chinese bone marrow donor. Hla, 2017, 90, 133-134.	0.4	3
170	Identification of the novel <i>HLA-B*13:98</i> allele in a Chinese individual. Hla, 2018, 91, 133-134.	0.4	3
171	Identification of the novel <i>HLA-B*40:333</i> allele by polymerase chain reaction sequence-based typing. Hla, 2018, 91, 302-303.	0.4	3
172	Identification of the novel <i>HLA-DQB1*03:181</i> allele in a Chinese leukemia patient. Hla, 2018, 91, 142-143.	0.4	3
173	Identification of the novel <i>HLA-B*27:04:06</i> allele in a Chinese bone marrow donor. Hla, 2018, 91, 136-137.	0.4	3
174	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
175	Identification of the novel <i><sc>HLA</sc>A*02:837</i> and <i>A*02:888</i> alleles by next-generation sequencing in two Chinese individuals. Hla, 2021, 97, 345-349.	0.4	3
176	Identification of the novel allele, <i>HLA-C*15:02:32</i>, in a Chinese individual. Hla, 2020, 96, 106-108.	0.4	3
177	Identification of the novel <i><sc>HLA</sc>C*15:210</i> allele by polymerase chain reaction sequence-based typing. Hla, 2021, 97, 241-243.	0.4	3
178	Identification of the novel <sc><i>HLA</i>C*03:04:79</sc> allele in a <sc><i>Chinese</i></sc> bone marrow donor. Hla, 2021, 97, 371-373.	0.4	3
179	Description of two new <sc>HLA</sc> alleles: <i><sc>HLA</sc>DRB1*11:262</i> and <i><sc>HLA</sc>DRB1*11:268</i>. Hla, 2021, 97, 474-477.	0.4	3
180	Three <sc>HLA</sc>A 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

#	ARTICLE	IF	CITATIONS
181	Characterization of the novel <i>HLA-C*01:02:56</i> and <i>HLA-C*01:02:57</i> alleles by sequencing-based typing. Hla, 2021, 97, 557-560.	0.4	3
182	The novel <i>HLA-C*04:407</i> allele was identified in a Chinese individual. Hla, 2021, 98, 68-69.	0.4	3
183	The novel <i>HLA-DQB1*03:282N</i> allele was identified in a Chinese individual. Hla, 2021, 98, 408-410.	0.4	3
184	Description of two new <i>HLA</i> alleles: <i>HLA-A*24:02:129</i> and <i>HLA-A*24:02:135</i> . Hla, 2021, 98, 146-148.	0.4	3
185	Identification of the novel <i>HLA-B*40:125:03</i> allele in a Chinese bone marrow donor. Hla, 2021, 98, 62-64.	0.4	3
186	<i>HLA-DQB1*05:239</i> and <i>HLA-DQB1*05:250</i> , were identified by sequencing in Chinese bone marrow donors. Hla, 2021, 98, 496-498.	0.4	3
187	The novel <i>HLA-A*26:174</i> allele was identified in a Chinese individual. Hla, 2021, 98, 151-153.	0.4	3
188	Identification of the novel <i>HLA-DRB1*11:271</i> allele by next-generation sequencing. Hla, 2021, 98, 401-403.	0.4	3
189	The novel <i>HLA-DRB1*15:01:42</i> allele was identified by next-generation sequencing. Hla, 2021, 98, 487-488.	0.4	3
190	The novel <i>HLA-A*31:191</i> allele was identified in a Chinese platelet donor. Hla, 2022, 99, 38-40.	0.4	3
191	Identification of the novel <i>HLA-B*55:107</i> allele in a Chinese bone marrow donor. Hla, 2021, 98, 478-479.	0.4	3
192	The novel <i>HLA-DRB1*14:222N</i> allele was identified by next-generation sequencing. Hla, 2021, 98, 562-564.	0.4	3
193	Identification of the novel <i>HLA-A*01:348</i> allele in a Chinese individual. Hla, 2022, 99, 615-617.	0.4	3
194	The novel <i>HLA-DRB1*12:02:11</i> allele identified by next-generation sequencing in a Chinese bone marrow donor. Hla, 2021, , .	0.4	3
195	Identification of the novel allele, <i>HLA-DRB1*08:03:12</i> , in a Chinese cord blood donor. Hla, 2022, 100, 94-95.	0.4	3
196	Three <i>HLA-DQB1</i> alleles, <i>DQB1*03:432</i> , <i>DQB1*03:454</i> and <i>DQB1*03:465</i> were identified in Chinese individuals. Hla, 2022, 100, 97-99.	0.4	3
197	Identification of the novel <i>HLA-DPB1*03:01:14</i> allele by next-generation sequencing in a Chinese cord blood donor. Hla, 2022, 100, 101-103.	0.4	3
198	Characterization of the novel <i>HLA-B*46:01:28</i> allele. Hla, 2022, 100, 73-74.	0.4	3

#	ARTICLE	IF	CITATIONS
199	Characterization of the novel <i>HLA*DRB1*04:05:23</i> allele by polymerase chain reaction sequence-based typing. Hla, 2022, , .	0.4	3
200	Identification of the novel <i>HLA*EA*24:02:138</i> allele in a Chinese individual. Hla, 2022, 100, 64-66.	0.4	3
201	Identification of the novel <i>HLA*EB*46:83</i> allele by sequencing-based typing in a Chinese individual. Hla, 2022, 100, 161-163.	0.4	3
202	Description of two new <i>HLA*EC</i> alleles: <i>HLA*EC</i> and <i>HLA*EC*07:900</i> and <i>HLA*EC*07:906</i> . Hla, 2022, 99, 399-400.	0.4	3
203	The novel <i>HLA*EB*51:01:83</i> allele was identified by next-generation sequencing. Hla, 2022, 100, 163-165.	0.4	3
204	Identification of the novel <i>HLA*EQB1*04:85</i> allele by next-generation sequencing. Hla, 2022, 100, 295-296.	0.4	3
205	The novel <i>HLA*DRB1*12:01:10</i> allele was identified by next-generation sequencing. Hla, 2022, 100, 389-390.	0.4	3
206	Identification of the novel <i>HLA*EC*06:318</i> allele by next-generation sequencing in a Chinese individual. Hla, 2022, 100, 381-382.	0.4	3
207	Description of two novel <i>HLA</i> alleles: <i>HLA*EC*01:02:73</i> and <i>HLA*EC*01:02:75</i> . Hla, 0, , .	0.4	3
208	Characterization of the novel <i>HLA*EB</i> allele, <i>HLA*EB*39:01:32</i> . Hla, 2022, 100, 526-527.	0.4	3
209	The novel <i>HLA*EA*24:520</i> allele was identified in a Chinese individual. Hla, 2022, 100, 515-517.	0.4	3
210	Identification of the novel <i>HLA*EB*39:01:23</i> allele by polymerase chain reaction sequence-based typing. Hla, 2016, 88, 310-311.	0.4	2
211	Characterization of a novel allelic variant in <i>HLA*EB*46:01</i> lineage, <i>HLA*EB*46:01:25</i> , by cloning, phasing and sequencing. International Journal of Immunogenetics, 2018, 45, 347-350.	0.8	2
212	Description of two new HLA alleles: <i>HLA*EA*30:118</i> and <i>HLA*EC*03:02:17</i> . Hla, 2019, 94, 371-373.	0.4	2
213	c.830T>C mutation on the <i>ABO*A1.02</i> allele responsible for Aw phenotype. Transfusion, 2019, 59, E11-E12.	0.8	2
214	Characterization of the novel <i>HLA*EB*51:228</i> allele in a Chinese individual. Hla, 2019, 93, 120-122.	0.4	2
215	Characterization of the novel <i>HLA*EB*46:01:23</i> allele in a Chinese bone marrow donor. Hla, 2019, 93, 224-226.	0.4	2
216	Identification of the novel <i>HLA*EA*31:124</i> allele by sequence-based typing in a Chinese cord blood donor. Hla, 2019, 93, 104-105.	0.4	2

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217	Identification of the novel <i>HLA*02</i> allele, <i>HLA*02:725</i> . Hla, 2020, 95, 476-478.	0.4	2
218	Characterization of the novel <i>HLA*01:154</i> allele by polymerase chain reaction sequencing-based typing. Hla, 2020, 95, 498-499.	0.4	2
219	Identification of the novel <i>HLA*03:280</i> allele by polymerase chain reaction sequencing-based typing. Hla, 2020, 96, 122-123.	0.4	2
220	Identification of the novel <i>HLA*DRB1*09:40</i> allele in a Chinese individual. Hla, 2020, 96, 111-113.	0.4	2
221	Characterization of the novel <i>HLA*11:280</i> allele by next-generation sequencing in a Chinese cord blood donor. Hla, 2020, 95, 482-483.	0.4	2
222	Mechanism evaluation for an amino acid substitution p.Y246C of $\beta$ -glucosyltransferase enzyme with Bweak phenotype. Vox Sanguinis, 2021, 116, 464-470.	0.7	2
223	Analysis of the Genomic Sequence of ABO Allele Using Next-Generation Sequencing Method. Frontiers in Immunology, 0, 13, .	2.2	2
224	Investigation of killer cell immunoglobulin-like receptors gene KIR3DL2 diversity and confirmation of KIR3DL2*015 in a Chinese population. Tissue Antigens, 2006, 68, 220-224.	1.0	1
225	A novel HLA allele, <i>HLA*02:57</i> , was identified by polymerase chain reaction sequencing-based typing in a Chinese individual. Tissue Antigens, 2015, 86, 215-216.	1.0	1
226	Investigation of Killer Cell Immunoglobulin-Like Receptors KIR2DL2 and KIR2DL3 Diversity and Identification of Ten Novel KIR2DL3 Alleles in the Chinese Han Population. Scandinavian Journal of Immunology, 2015, 81, 265-271.	1.3	1
227	c.426G>C mutation in <i>ABO*A1.02</i> allele was associated with Aw phenotype. Transfusion, 2019, 59, E4-E5.	0.8	1
228	Two novel <i>A</i> alleles with c.322C>T or c.410C>T mutations on the <i>ABO*A1.02</i> allele were identified in the Chinese individuals. Transfusion, 2020, 60, E38-E39.	0.8	1
229	Characterization of the novel KIR3DL2 allele, <i>KIR3DL2*113</i> . Hla, 2020, 95, 594-596.	0.4	1
230	Identification of a novel <i>A</i> allele with a c.731T>C mutation on the <i>ABO*A1.02</i> allele. Transfusion, 2020, 60, E30-E31.	0.8	1
231	Identification of the novel <i>KIR3DL2*114</i> allele in a Chinese individual by polymerase chain reaction sequencing-based typing. Hla, 2020, 95, 596-598.	0.4	1
232	Identification of a novel <i>B</i> allele with c.<math>10\text{â€}14\text{dupGTGT}</math> and c.<math>17\text{G}>\text{A}</math> variants in a Chinese individual with a weak B phenotype. Transfusion, 2021, 61, E67-E68.	0.8	1
233	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
234	A novel allele, <i>HLA*A*33:97</i> was identified in a Chinese bone marrow donor. Hla, 2016, 88, 305-306.	0.4	0

#	ARTICLE	IF	CITATIONS
235	Identification of the novel <i><sc>KIR2DL2</sc>*00103</i> allele in a Chinese individual by sequence-based typing. Hla, 2016, 87, 476-477.	0.4	0
236	c.518T>>C missense mutation in the B glycosyltransferase gene responsible for a weak B variant. Transfusion, 2018, 58, 269-270.	0.8	0
237	c.125_126insT mutation in <i>ABO*B.01</i> allele responsible for Bel phenotype. Transfusion, 2018, 58, 2467-2468.	0.8	0
238	Identification of a novel B allele with a nucleotide deletion (c.3_4 del G) in the ABO gene associated with a Bx phenotype individual. Transfusion, 2019, 59, 793-794.	0.8	0
239	Identification of a novel B allele with a c.586T>>C mutation on the <i>ABO*B.01</i> allele. Transfusion, 2020, 60, E1-E2.	0.8	0
240	Identification of the novel <i><sc>KIR3DL2</sc>*00711</i> allele by sequencing-based typing in a Chinese individual. Hla, 2021, 98, 416-418.	0.4	0
241	Identification of a novel B allele with a c.<sc>256G</sc>>A mutation on the <i><sc>ABO</sc>*B.01</i> allele. Transfusion, 2022, 62, .	0.8	0