

Hong-Yan Zou

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Full length sequence of a novel HLA-B*3818 allele differs from HLA-B*380201 allele in exon 4 and intron 5. Tissue Antigens, 2009, 74, 439-440.	1.0	18
2	Characterization of a new HLA-C allele in a Chinese family by sequence-based typing: HLA-Cw*0348. Tissue Antigens, 2009, 73, 616-618.	1.0	7
3	Genomic full-length sequence of the <i><sc>HLA-A</sc>*11:334</i> allele, identified by cloning and sequencing. Hla, 2020, 96, 495-496.	0.6	6
4	Characterization of the novel variant allele, <i><sc>HLA-B</sc>*13:82</i>, identified in a Chinese Han individual. Hla, 2020, 96, 510-511.	0.6	6
5	Characterization of a novel variant allele, HLA-A*11:155, identified in a Chinese Han individual. Hla, 2020, 96, 208-209.	0.6	6
6	Identification of a novel HLA-DPB1 allele, <i>HLA-DPB1*612:01</i>, in a Chinese individual. Hla, 2018, 92, 428-429.	0.6	5
7	Characterization of a novel variant allele, HLA-C*08:125, identified in a Chinese Han individual. Hla, 2019, 94, 78-80.	0.6	5
8	Identification of a novel <i><sc>HLA</sc>-B*51</i> allele, <i><sc>HLA</sc>-B*51:144</i>, in a Chinese individual. Tissue Antigens, 2014, 84, 416-417.	1.0	3
9	Identification of a novel <i><sc>HLA</sc>-A*33</i> allele, <i><sc>HLA</sc>-A*33:03:13</i>, in a Chinese family. Tissue Antigens, 2015, 85, 136-138.	1.0	3
10	Identification of a novel <i>HLA-A*02</i> allele</i>, <i>HLA-A*02:432</i>, in a Chinese individual. Hla, 2016, 87, 37-38.	0.6	3
11	Full-length sequences of 3 <sc>HLA</sc>-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.	0.6	3
12	Full-length sequences of 3 <sc>HLA</sc>-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.	0.6	3
13	Full-length sequences of <i><sc>HLA</sc>-B*39:05:01</i> and <i>B*39:<sc>38Q</sc></i>, confirmed by cloning and sequencing. Hla, 2017, 89, 159-162.	0.6	3
14	Full-length sequences of 3 <i><sc>HLA</sc>-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.	0.6	3
15	Full-length sequences of 4 <i><sc>HLA</sc>-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.	0.6	3
16	Full-length sequences of 4 <i><sc>HLA</sc>-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.	0.6	3
17	Full-length sequences of <i><sc>HLA</sc>-B*67:01:01</i> and <i>B*67:01:02</i>, confirmed by cloning and sequencing. Hla, 2017, 89, 324-327.	0.6	3
18	Full-length sequence of <i><sc>HLA</sc>-B*55:02:01:01</i>, confirmed by cloning and sequencing. Hla, 2017, 90, 45-48.	0.6	3

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19	Full-length sequences of 4 <i>HLA-B*15</i> alleles, <i>B*15:07:01:01</i> , <i>B*15:27:01</i> , <i>B*15:32:01</i> and <i>B*15:58</i> , confirmed by cloning and sequencing. <i>Hla</i> , 2017, 89, 313-317.	0.6	3
20	Full-length sequences of 3 <i>HLA-B</i> alleles, <i>HLA-B*27:04:01</i> , <i>B*27:07:01</i> and <i>B*27:25</i> , confirmed by cloning and sequencing. <i>Hla</i> , 2017, 90, 40-43.	0.6	3
21	The full-length sequence of the <i>HLA-C</i> allele, <i>HLA-C*15:13:01:01</i> . <i>Hla</i> , 2018, 91, 216-220.	0.6	3
22	The full-length sequence of the <i>HLA-C</i> allele, <i>HLA-C*03:02:01</i> . <i>Hla</i> , 2018, 91, 204-207.	0.6	3
23	The full-length sequence of the <i>HLA-C</i> allele, <i>HLA-C*03:40:01</i> . <i>Hla</i> , 2018, 91, 208-211.	0.6	3
24	The <i>HLA-DRB1*11:23:02</i> allele confirmed in a Chinese individual by next-generation sequencing. <i>Hla</i> , 2021, 97, 85-86.	0.6	3
25	The full-length sequence of the novel <i>HLA-C*08:190</i> allele, identified by cloning and sequencing. <i>Hla</i> , 2021, 97, 80-81.	0.6	3
26	Characterization of the novel <i>HLA-DQB1*03:362</i> allele in a Chinese family. <i>Hla</i> , 2021, 98, 410-412.	0.6	3
27	Discovery of the novel <i>HLA-C*01:179</i> allele in a southern Chinese patient. <i>Hla</i> , 2021, 98, 165-166.	0.6	3
28	A substitution in exon 2 resulted in the novel <i>HLA-A*30:140</i> variant identified in a Chinese individual. <i>Hla</i> , 2021, 98, 226-228.	0.6	3
29	A novel <i>HLA-C*15</i> allele, <i>HLA-C*15:192</i> , identified by next generation sequencing in a Chinese individual. <i>Hla</i> , 2021, 98, 485-486.	0.6	3
30	Discovery of the <i>HLA-C*03:561</i> allele, a variant of <i>HLA-C*03</i> , in a Chinese individual. <i>Hla</i> , 2022, 99, 127-129.	0.6	3
31	Characterization of the novel <i>HLA-C*01:213</i> allele by next-generation sequencing in a Chinese family. <i>Hla</i> , 2022, 99, 125-127.	0.6	3
32	Characterization of a novel variant allele, <i>HLA-C*03:587</i> , identified in a Chinese Han individual. <i>Hla</i> , 2022, 100, 82-83.	0.6	3
33	Full-length sequence of 2 <i>HLA-B</i> alleles, <i>B*52:01:01:01</i> and <i>B*52:01:02:01</i> , identified by cloning and sequencing. <i>Hla</i> , 2017, 89, 163-165.	0.6	2
34	The full-length sequence of <i>HLA-B*59:01:01:01</i> confirmed by cloning and sequencing. <i>Hla</i> , 2017, 90, 255-258.	0.6	2
35	Confirmed the full-length sequence of <i>HLA-B*44:03:02</i> by cloning and sequencing. <i>Hla</i> , 2017, 90, 125-127.	0.6	2
36	Full-length sequences of 2 <i>HLA-B</i> alleles, <i>B*48:03:01</i> and <i>B*48:04:01</i> , confirmed by cloning and sequencing. <i>Hla</i> , 2017, 90, 128-130.	0.6	2

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37	Characterization of the novel <i>HLA*QB1*06:01:22</i> allele by next-generation sequencing. Hla, 2019, 94, 543-545.	0.6	2
38	Characterization of the novel HLA*15:435 allele by next-generation sequencing in a Chinese family. Hla, 2019, 93, 108-109.	0.6	2
39	<i>HLA*C*08:01:25</i>, a novel HLA allele, which has arisen by a silent mutation in codon 271. Hla, 2019, 94, 162-163.	0.6	2
40	Identification of <i>HLA*31:150</i> by next-generation sequencing in a Chinese Han individual. Hla, 2019, 94, 373-375.	0.6	2
41	Identification of the novel <i>HLA*B*46:78</i> allele by next generation sequencing in a Chinese individual. Hla, 2020, 95, 137-138.	0.6	2