

Hong-Yan Zou

List of Publications by Year in descending order

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#	ARTICLE	IF	CITATIONS
1	Discovery of the <sc>HLAâ€C</sc>*03:561 allele, a variant of <sc>HLAâ€C</sc>*03, in a Chinese individual. Hla, 2022, 99, 127-129.	0.4	3
2	Characterization of the novel <sc><i>HLAâ€C*01:213</i></sc> allele by nextâ€generation sequencing in a Chinese family. Hla, 2022, 99, 125-127.	0.4	3
3	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
4	The <i><sc>HLAâ€DRB1</sc>*11:23:02</i> allele confirmed in a Chinese individual by nextâ€generation sequencing. Hla, 2021, 97, 85-86.	0.4	3
5	The fullâ€length sequence of the novel <i><sc>HLAâ€C</sc>*08:190</i> allele, identified by cloning and sequencing. Hla, 2021, 97, 80-81.	0.4	3
6	Characterization of the novel <i>HLAâ€DQB1*03:362</i> allele in a Chinese family. Hla, 2021, 98, 410-412.	0.4	3
7	Discovery of the novel <i><sc>HLAâ€C</sc>*01:179</i> allele in a southern Chinese patient. Hla, 2021, 98, 165-166.	0.4	3
8	A substitution in exon 2 resulted in the novel <i><sc>HLAâ€A</sc>*30:140</i> variant identified in a Chinese individual. Hla, 2021, 98, 226-228.	0.4	3
9	A novel <i><sc>HLAâ€C</sc>*15</i> allele, <i><sc>HLAâ€C</sc>*15:192</i>, identified by next generation sequencing in a Chinese individual. Hla, 2021, 98, 485-486.	0.4	3
10	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
11	Genomic fullâ€length sequence of the <i><sc>HLAâ€A</sc>*11:334</i> allele, identified by cloning and sequencing. Hla, 2020, 96, 495-496.	0.4	6
12	Characterization of the novel variant allele, <i><sc>HLAâ€B</sc>*13:82</i>, identified in a Chinese Han individual. Hla, 2020, 96, 510-511.	0.4	6
13	Characterization of a novel variant allele, HLAâ€A*11:155 , identified in a Chinese Han individual. Hla, 2020, 96, 208-209.	0.4	6
14	Characterization of the novel <i>HLAâ€DQB1*06:01:22</i> allele by nextâ€generation sequencing. Hla, 2019, 94, 543-545.	0.4	2
15	Characterization of the novel HLAâ€B*15:435 allele by nextâ€generation sequencing in a Chinese family. Hla, 2019, 93, 108-109.	0.4	2
16	<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
17	Identification of <i>HLAâ€A*31:150</i> by nextâ€generation sequencing in a Chinese Han individual. Hla, 2019, 94, 373-375.	0.4	2
18	Characterization of a novel variant allele, HLAâ€C*08:125 , identified in a Chinese Han individual. Hla, 2019, 94, 78-80.	0.4	5

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19	The full-length sequence of the HLA-B allele, <i>HLA-B*15:13:01:01</i>. Hla, 2018, 91, 216-220.	0.4	3
20	The full-length sequence of the HLA-B allele, <i>HLA-B*03:02:01</i>. Hla, 2018, 91, 204-207.	0.4	3
21	The full-length sequence of the HLA-B allele, <i>HLA-B*03:40:01</i>. Hla, 2018, 91, 208-211.	0.4	3
22	Identification of a novel HLA-DPB1 allele, <i>HLA-DPB1*612:01</i>, in a Chinese individual. Hla, 2018, 92, 428-429.	0.4	5
23	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.	0.4	3
24	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.	0.4	3
25	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.	0.4	3
26	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.	0.4	3
27	Full-length sequence of 2 <sc>HLA</sc>-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
28	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.	0.4	3
29	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.	0.4	3
30	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.	0.4	3
31	Full-length sequence of <i><sc>HLA</sc>-B*55:02:01:01</i>, confirmed by cloning and sequencing. Hla, 2017, 90, 45-48.	0.4	3
32	The full-length sequence of <i><sc>HLA</sc>-B*59:01:01:01</i> confirmed by cloning and sequencing. Hla, 2017, 90, 255-258.	0.4	2
33	Confirmed the full-length sequence of <i><sc>HLA</sc>-B*44:03:02</i> by cloning and sequencing. Hla, 2017, 90, 125-127.	0.4	2
34	Full-length sequences of 4 <i><sc>HLA</sc>-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. Hla, 2017, 89, 313-317.	0.4	3
35	Full-length sequences of 3 <sc>HLA</sc>-B alleles, <i><sc>HLA</sc>-B*27:04:01</i>, <i><sc>B</sc>*27:07:01</i> and <i><sc>B</sc>*27:25</i>, confirmed by cloning and sequencing. Hla, 2017, 90, 40-43.	0.4	3
36	Full-length sequences of 2 <sc>HLA</sc>-B alleles, <i>B*48:03:01</i> and <i>B*48:04:01</i>, confirmed by cloning and sequencing. Hla, 2017, 90, 128-130.	0.4	2

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37	Identification of a novel <i>HLA*02</i> allele, <i>HLA*02:432</i> , in a Chinese individual. <i>Hla</i> , 2016, 87, 37-38.	0.4	3
38	Identification of a novel <i>HLA*33</i> allele, <i>HLA*33:03:13</i> , in a Chinese family. <i>Tissue Antigens</i> , 2015, 85, 136-138.	1.0	3
39	Identification of a novel <i>HLA*51</i> allele, <i>HLA*51:144</i> , in a Chinese individual. <i>Tissue Antigens</i> , 2014, 84, 416-417.	1.0	3
40	Characterization of a new HLA-C allele in a Chinese family by sequence-based typing: HLA-Cw*0348. <i>Tissue Antigens</i> , 2009, 73, 616-618.	1.0	7
41	Full length sequence of a novel HLA*3818 allele differs from HLA*380201 allele in exon 4 and intron 5. <i>Tissue Antigens</i> , 2009, 74, 439-440.	1.0	18