

Chun Ying Lin

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/8282836/publications.pdf>

Version: 2024-02-01

11
papers

68
citations

1937685
4
h-index

1588992
8
g-index

11
all docs

11
docs citations

11
times ranked

53
citing authors

#	ARTICLE	IF	CITATIONS
1	Characterization of the novel <i><i><sc>HLAâ€C</sc>*03:294</i></i> allele by sequencingâ€based typing in a Taiwanese individual. <i>Hla</i> , 2022, 99, 215-216.	0.6	3
2	Identification of the novel <i><i><sc>HLAâ€C</sc>*07:446</i></i> allele in a volunteer bone marrow donor. <i>Hla</i> , 2022, 99, 397-399.	0.6	3
3	Identification of a novel <i><sc>HLAâ€DQB1</sc></i> allele, <i><sc><i>HLAâ€DQB1*03:168</i></sc></i> , by sequenceâ€based typing in a Taiwanese individual. <i>Hla</i> , 2020, 96, 546-547.	0.6	6
4	<i><i><sc>HLAâ€DQB1</sc>*05:02:12</i></i> , an <i><i><sc>HLAâ€DQB1</sc>*05:02:01:01</i></i> variant, identified in a Taiwanese individual. <i>Hla</i> , 2020, 96, 551-552.	0.6	6
5	Enthesitis-related arthritis is the most common category of juvenile idiopathic arthritis in Taiwan and presents persistent active disease. <i>Pediatric Rheumatology</i> , 2019, 17, 58.	2.1	29
6	Detection of a novel <i><i>HLAâ€B*46:01</i></i> variant, <i><i>HLAâ€B*46:01:19</i></i> , in a Taiwanese individual. <i>Hla</i> , 2018, 92, 414-415.	0.6	3
7	Identification of a novel <i><i>HLAâ€DQB1</i></i> allele, <i><i>DQB1*06:111</i></i> , by sequenceâ€based typing in a Taiwanese individual. <i>Hla</i> , 2018, 92, 257-258.	0.6	4
8	<i><i><sc>HLA</sc>â€B*40:247</i></i> , a novel <i><i><sc>HLA</sc>â€B*40</i></i> variant, identified by sequenceâ€based typing in a Taiwanese individual. <i>Hla</i> , 2017, 90, 121-122.	0.6	3
9	<i><i><sc>HLA</sc>â€A*33:<sc>74N</sc></i></i> , a novel <i><i><sc>HLA</sc>â€A*33</i></i> variant, identified by sequenceâ€based typing in a Taiwanese individual. <i>Hla</i> , 2017, 90, 365-366.	0.6	5
10	A novel allele, <i><i>HLAâ€B*15:259</i></i> , was identified in a Taiwanese individual by sequenceâ€based typing. <i>Hla</i> , 2016, 87, 103-104.	0.6	3
11	<i><i>HLAâ€A*11:134</i></i> , a novel <i><i>HLAâ€A*11</i></i> variant, identified by sequenceâ€based typing in a Taiwanese individual. <i>Hla</i> , 2016, 88, 195-196.	0.6	3