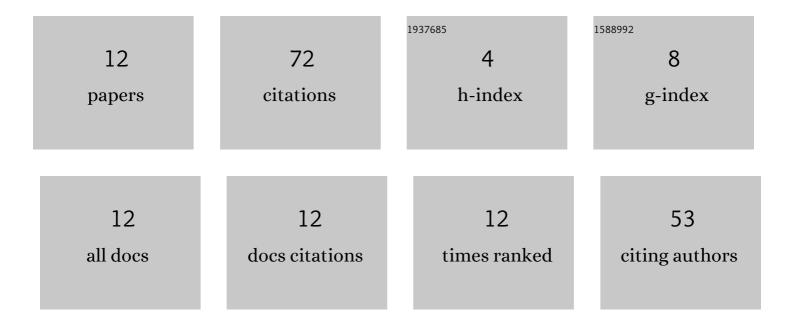
## Chia Ling Chang

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4101104/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Enthesitis-related arthritis is the most common category of juvenile idiopathic arthritis in Taiwan and presents persistent active disease. Pediatric Rheumatology, 2019, 17, 58.	2.1	29
2	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.	0.6	6
3	<i><scp>HLAâ€DQB1</scp>*05:02:12</i> , an <i><scp>HLAâ€DQB1</scp>*05:02:01:01</i> variant, identified in Taiwanese individual. Hla, 2020, 96, 551-552.	<sup>a</sup> 0.6	6
4	<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.	0.6	5
5	Identification of a novel <i>HLAâ€DQB1</i> allele, <i>DQB1*06:111</i> , by sequenceâ€based typing in a Taiwanese individual. Hla, 2018, 92, 257-258.	0.6	4
6	Different clinical features of patients with pulmonary disease caused by various Mycobacterium avium–intracellulare complex subspecies and antimicrobial susceptibility. International Journal of Infectious Diseases, 2020, 98, 33-40.	3.3	4
7	A novel allele, <i>HLAâ€B*15:259</i> , was identified in a Taiwanese individual by sequenceâ€based typing. Hla, 2016, 87, 103-104.	0.6	3
8	<i>HLAâ€A*11:134</i> , a novel <i>HLAâ€A*11</i> variant, identified by sequenceâ€based typing in a Taiwanese individual. Hla, 2016, 88, 195-196.	0.6	3
9	<i><scp>HLA</scp>â€B*40:247</i> , a novel <i><scp>HLA</scp>â€B*40</i> variant, identified by sequenceâ€based typing in a Taiwanese individual. Hla, 2017, 90, 121-122.	0.6	3
10	Detection of a novel <i>HLAâ€B*46:01</i> variant, <i>HLAâ€B*46:01:19</i> , in a Taiwanese individual. Hla, 2018, 92, 414-415.	0.6	3
11	Characterization of the novel <i><scp>HLA </scp>*03:294</i> allele by sequencingâ€based typing in a Taiwanese individual. Hla, 2022, 99, 215-216.	0.6	3
12	Identification of the novel <i><scp>HLA </scp>*07:446</i> allele in a volunteer bone marrow donor. Hla, 2022, 99, 397-399.	0.6	3