

# Chia Ling Chang

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4101104/publications.pdf>

Version: 2024-02-01

12  
papers

72  
citations

1937685  
4  
h-index

1588992  
8  
g-index

12  
all docs

12  
docs citations

12  
times ranked

53  
citing authors

#	ARTICLE	IF	CITATIONS
1	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
2	Identification of a novel <i>HLA-DQB1</i> allele, <i>HLA-DQB1*03:168</i> , by sequence-based typing in a Taiwanese individual. <i>Hla</i> , 2020, 96, 546-547.	0.6	6
3	<i>HLA-DQB1*05:02:12</i> , an <i>HLA-DQB1*05:02:01:01</i> variant, identified in a Taiwanese individual. <i>Hla</i> , 2020, 96, 551-552.	0.6	6
4	<i>HLA-A*33:74N</i> , a novel <i>HLA-A*33</i> variant, identified by sequence-based typing in a Taiwanese individual. <i>Hla</i> , 2017, 90, 365-366.	0.6	5
5	Identification of a novel <i>HLA-DQB1</i> allele, <i>DQB1*06:111</i> , by sequence-based typing in a Taiwanese individual. <i>Hla</i> , 2018, 92, 257-258.	0.6	4
6	Different clinical features of patients with pulmonary disease caused by various <i>Mycobacterium avium</i> "intracellulare complex" subspecies and antimicrobial susceptibility. <i>International Journal of Infectious Diseases</i> , 2020, 98, 33-40.	3.3	4
7	A novel allele, <i>HLA-B*15:259</i> , was identified in a Taiwanese individual by sequence-based typing. <i>Hla</i> , 2016, 87, 103-104.	0.6	3
8	<i>HLA-A*11:134</i> , a novel <i>HLA-A*11</i> variant, identified by sequence-based typing in a Taiwanese individual. <i>Hla</i> , 2016, 88, 195-196.	0.6	3
9	<i>HLA-B*40:247</i> , a novel <i>HLA-B*40</i> variant, identified by sequence-based typing in a Taiwanese individual. <i>Hla</i> , 2017, 90, 121-122.	0.6	3
10	Detection of a novel <i>HLA-B*46:01</i> variant, <i>HLA-B*46:01:19</i> , in a Taiwanese individual. <i>Hla</i> , 2018, 92, 414-415.	0.6	3
11	Characterization of the novel <i>HLA-C*03:294</i> allele by sequencing-based typing in a Taiwanese individual. <i>Hla</i> , 2022, 99, 215-216.	0.6	3
12	Identification of the novel <i>HLA-C*07:446</i> allele in a volunteer bone marrow donor. <i>Hla</i> , 2022, 99, 397-399.	0.6	3