

# Yoshihiro Onouchi

## List of Publications by Year in Descending Order

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**Version:** 2024-04-27

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

13  
papers

979  
citations

9  
h-index

15  
g-index

15  
ext. papers

1,237  
ext. citations

14.9  
avg, IF

4.07  
L-index

#	Paper	IF	Citations
13	Markers of Memory CD8 T Cells Depicting the Effect of the BNT162b2 mRNA COVID-19 Vaccine in Japan.. <i>Frontiers in Immunology</i> , <b>2022</b> , 13, 836923	8.4	
12	Association of an IGHV3-66 gene variant with Kawasaki disease. <i>Journal of Human Genetics</i> , <b>2021</b> , 66, 475-489	4.3	9
11	Global Comparison of Changes in the Number of Test-Positive Cases and Deaths by Coronavirus Infection (COVID-19) in the World. <i>Journal of Clinical Medicine</i> , <b>2020</b> , 9,	5.1	9
10	Population-specific and trans-ancestry genome-wide analyses identify distinct and shared genetic risk loci for coronary artery disease. <i>Nature Genetics</i> , <b>2020</b> , 52, 1169-1177	36.3	51
9	Efficacy of primary treatment with immunoglobulin plus ciclosporin for prevention of coronary artery abnormalities in patients with Kawasaki disease predicted to be at increased risk of non-response to intravenous immunoglobulin (KAICA): a randomised controlled, open-label, blinded-endpoints, phase 3 trial. <i>Lancet. The</i> , <b>2019</b> , 393, 1128-1137	4.0	70
8	Investigation of novel variations of ORAI1 gene and their association with Kawasaki disease. <i>Journal of Human Genetics</i> , <b>2019</b> , 64, 511-519	4.3	6
7	The genetics of Kawasaki disease. <i>International Journal of Rheumatic Diseases</i> , <b>2018</b> , 21, 26-30	2.3	71
6	A genome-wide association analysis identifies NMNAT2 and HCP5 as susceptibility loci for Kawasaki disease. <i>Journal of Human Genetics</i> , <b>2017</b> , 62, 1023-1029	4.3	29
5	Genetic Variation in the SLC8A1 Calcium Signaling Pathway Is Associated With Susceptibility to Kawasaki Disease and Coronary Artery Abnormalities. <i>Circulation: Cardiovascular Genetics</i> , <b>2016</b> , 9, 559-568		33
4	Variations in ORAI1 Gene Associated with Kawasaki Disease. <i>PLoS ONE</i> , <b>2016</b> , 11, e0145486	3.7	30
3	A genome-wide association study identifies three new risk loci for Kawasaki disease. <i>Nature Genetics</i> , <b>2012</b> , 44, 517-21	36.3	217
2	Common variants in CASP3 confer susceptibility to Kawasaki disease. <i>Human Molecular Genetics</i> , <b>2010</b> , 19, 2898-906	5.6	114
1	ITPKC functional polymorphism associated with Kawasaki disease susceptibility and formation of coronary artery aneurysms. <i>Nature Genetics</i> , <b>2008</b> , 40, 35-42	36.3	339