Mitja Mitrovic

List of Publications by Year in Descending Order

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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.

15 4,402 10 18 g-index

18 5,188 17 3.57 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
15	Birth characteristics and risk of febrile seizures. <i>Acta Neurologica Scandinavica</i> , 2021 , 144, 51-57	3.8	3
14	Genome-wide association studies of multiple sclerosis. <i>Clinical and Translational Immunology</i> , 2018 , 7, e1018	6.8	36
13	Multiple sclerosis. Handbook of Clinical Neurology / Edited By P J Vinken and G W Bruyn, 2018 , 148, 723-	-73 ₅ 0	13
12	Low-Frequency and Rare-Coding Variation Contributes to Multiple Sclerosis Risk. <i>Cell</i> , 2018 , 175, 1679	-1 68 Ze	2 7 72
11	NR1H3 p.Arg415Gln Is Not Associated to Multiple Sclerosis Risk. <i>Neuron</i> , 2016 , 92, 333-335	13.9	19
10	Regulatory polymorphisms modulate the expression of HLA class II molecules and promote autoimmunity. <i>ELife</i> , 2016 , 5,	8.9	75
9	AKT isoforms modulate Th1-like Treg generation and function in human autoimmune disease. <i>EMBO Reports</i> , 2016 , 17, 1169-83	6.5	48
8	Genetic variants associated with autoimmunity drive NFB signaling and responses to inflammatory stimuli. <i>Science Translational Medicine</i> , 2015 , 7, 291ra93	17.5	63
7	Dense genotyping of immune-related disease regions identifies nine new risk loci for primary sclerosing cholangitis. <i>Nature Genetics</i> , 2013 , 45, 670-5	36.3	267
6	Host-microbe interactions have shaped the genetic architecture of inflammatory bowel disease. <i>Nature</i> , 2012 , 491, 119-24	50.4	3239
5	Limited evidence for parent-of-origin effects in inflammatory bowel disease associated loci. <i>PLoS ONE</i> , 2012 , 7, e45287	3.7	9
4	High-Resolution Melting Curve Analysis for High-Throughput Genotyping of NOD2/CARD15 Mutations and Distribution of These Mutations in Slovenian Inflammatory Bowel Diseases Patients. <i>Disease Markers</i> , 2011 , 30, 265-274	3.2	7
3	The quest for genetic risk factors for Crohnd disease in the post-GWAS era. <i>Genome Medicine</i> , 2011 , 3, 13	14.4	10
2	Dense genotyping identifies and localizes multiple common and rare variant association signals in celiac disease. <i>Nature Genetics</i> , 2011 , 43, 1193-201	36.3	535
1	High-resolution melting curve analysis for high-throughput genotyping of NOD2/CARD15 mutations and distribution of these mutations in Slovenian inflammatory bowel diseases patients. <i>Disease Markers</i> , 2011 , 30, 265-74	3.2	5