## Johan Ã-ckinger

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Critical role for calcium mobilization in activation of the NLRP3 inflammasome. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 11282-11287.	7.1	709
2	Combined sequence-based and genetic mapping analysis of complex traits in outbred rats. Nature Genetics, 2013, 45, 767-775.	21.4	176
3	The lung microbiota in early rheumatoid arthritis and autoimmunity. Microbiome, 2016, 4, 60.	11.1	158
4	A resource for the simultaneous high-resolution mapping of multiple quantitative trait loci in rats: The NIH heterogeneous stock. Genome Research, 2009, 19, 150-158.	5.5	72
5	<i>IL-22RA2</i> Associates with Multiple Sclerosis and Macrophage Effector Mechanisms in Experimental Neuroinflammation. Journal of Immunology, 2010, 185, 6883-6890.	0.8	68
6	Tobacco smoking induces changes in true DNA methylation, hydroxymethylation and gene expression in bronchoalveolar lavage cells. EBioMedicine, 2019, 46, 290-304.	6.1	48
7	Expression of Ccl11 Associates with Immune Response Modulation and Protection against Neuroinflammation in Rats. PLoS ONE, 2012, 7, e39794.	2.5	46
8	Expression of MATE1, P-gp, OCTN1 and OCTN2, in epithelial and immune cells in the lung of COPD and healthy individuals. Respiratory Research, 2018, 19, 68.	3.6	27
9	Genetic variants of CC chemokine genes in experimental autoimmune encephalomyelitis, multiple sclerosis and rheumatoid arthritis. Genes and Immunity, 2010, 11, 142-154.	4.1	23
10	Fine-Mapping Resolves Eae23 into Two QTLs and Implicates ZEB1 as a Candidate Gene Regulating Experimental Neuroinflammation in Rat. PLoS ONE, 2010, 5, e12716.	2.5	23
11	Definition of a 1.06-Mb Region Linked to Neuroinflammation in Humans, Rats and Mice. Genetics, 2006, 173, 1539-1545.	2.9	20
12	Multiple loci comprising immune-related genes regulate experimental neuroinflammation. Genes and Immunity, 2010, 11, 21-36.	4.1	20
13	<i>Vra4</i> Congenic Rats with Allelic Differences in the Class II Transactivator Gene Display Altered Susceptibility to Experimental Autoimmune Encephalomyelitis. Journal of Immunology, 2008, 180, 3289-3296.	0.8	18
14	Advanced Intercross Line Mapping Suggests That Ncf1 (Ean6) Regulates Severity in an Animal Model of Guillain-Barré Syndrome. Journal of Immunology, 2009, 182, 4432-4438.	0.8	18
15	T-cell activation and HLA-regulated response to smoking in the deep airways of patients with multiple sclerosis. Clinical Immunology, 2016, 169, 114-120.	3.2	17
16	Parent-of-Origin Effects Implicate Epigenetic Regulation of Experimental Autoimmune Encephalomyelitis and Identify Imprinted Dlk1 as a Novel Risk Gene. PLoS Genetics, 2014, 10, e1004265.	3.5	16
17	Methylome and transcriptome signature of bronchoalveolar cells from multiple sclerosis patients in relation to smoking. Multiple Sclerosis Journal, 2021, 27, 1014-1026.	3.0	12
18	Combining genetic mapping with genome-wide expression in experimental autoimmune encephalomyelitis highlights a gene network enriched for T cell functions and candidate genes regulating autoimmunity. Human Molecular Genetics, 2013, 22, 4952-4966.	2.9	11

#	Article	IF	CITATIONS
19	Hospital diagnosed pneumonia before age 20 years and multiple sclerosis risk. BMJ Neurology Open, 2020, 2, e000044.	1.6	4
20	Distinctive Regulatory T Cells and Altered Cytokine Profile Locally in the Airways of Young Smokers with Normal Lung Function. PLoS ONE, 2016, 11, e0164751.	2.5	2