

# Christian Hammer

## List of Publications by Year in descending order

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Version: 2024-02-01

50  
papers

18,161  
citations

172457

29  
h-index

182427

51  
g-index

63  
all docs

63  
docs citations

63  
times ranked

28594  
citing authors

#	ARTICLE	IF	CITATIONS
1	Biological insights from 108 schizophrenia-associated genetic loci. <i>Nature</i> , 2014, 511, 421-427.	27.8	6,934
2	LD Score regression distinguishes confounding from polygenicity in genome-wide association studies. <i>Nature Genetics</i> , 2015, 47, 291-295.	21.4	3,905
3	Schizophrenia risk from complex variation of complement component 4. <i>Nature</i> , 2016, 530, 177-183.	27.8	1,915
4	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. <i>American Journal of Human Genetics</i> , 2015, 97, 576-592.	6.2	1,098
5	Partitioning Heritability of Regulatory and Cell-Type-Specific Variants across 11 Common Diseases. <i>American Journal of Human Genetics</i> , 2014, 95, 535-552.	6.2	569
6	Antigen presentation in cancer: insights into tumour immunogenicity and immune evasion. <i>Nature Reviews Cancer</i> , 2021, 21, 298-312.	28.4	553
7	Contrasting genetic architectures of schizophrenia and other complex diseases using fast variance-components analysis. <i>Nature Genetics</i> , 2015, 47, 1385-1392.	21.4	431
8	Memory B Cells Activate Brain-Homing, Autoreactive CD4+ T Cells in Multiple Sclerosis. <i>Cell</i> , 2018, 175, 85-100.e23.	28.9	350
9	Seroprevalence of autoantibodies against brain antigens in health and disease. <i>Annals of Neurology</i> , 2014, 76, 82-94.	5.3	301
10	Neuropsychiatric disease relevance of circulating anti-NMDA receptor autoantibodies depends on blood-brain barrier integrity. <i>Molecular Psychiatry</i> , 2014, 19, 1143-1149.	7.9	293
11	Natural variation in the parameters of innate immune cells is preferentially driven by genetic factors. <i>Nature Immunology</i> , 2018, 19, 302-314.	14.5	205
12	Estimation of Genetic Correlation via Linkage Disequilibrium Score Regression and Genomic Restricted Maximum Likelihood. <i>American Journal of Human Genetics</i> , 2018, 102, 1185-1194.	6.2	119
13	Performance of Serum microRNAs -122, -192 and -21 as Biomarkers in Patients with Non-Alcoholic Steatohepatitis. <i>PLoS ONE</i> , 2015, 10, e0142661.	2.5	116
14	Human genetic variants and age are the strongest predictors of humoral immune responses to common pathogens and vaccines. <i>Genome Medicine</i> , 2018, 10, 59.	8.2	113
15	Severe viral respiratory infections in children with <i>IFIH1</i> loss-of-function mutations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 8342-8347.	7.1	111
16	Accumulated environmental risk determining age at schizophrenia onset: a deep phenotyping-based study. <i>Lancet Psychiatry</i> , 2014, 1, 444-453.	7.4	84
17	Serotonin type 3 receptor genes: <i>HTR3A, B, C, D, E</i> . <i>Pharmacogenomics</i> , 2008, 9, 501-504.	1.3	80
18	The brain as immunoprecipitator of serum autoantibodies against N-methyl-D-aspartate receptor subunit NR1. <i>Annals of Neurology</i> , 2016, 79, 144-151.	5.3	75

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19	The HTR3A Polymorphism c. -42C>T Is Associated With Amygdala Responsiveness in Patients With Irritable Bowel Syndrome. <i>Gastroenterology</i> , 2011, 140, 1943-1951.	1.3	73
20	Polygenic risk for skin autoimmunity impacts immune checkpoint blockade in bladder cancer. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 12288-12294.	7.1	65
21	Amino Acid Variation in HLA Class II Proteins Is a Major Determinant of Humoral Response to Common Viruses. <i>American Journal of Human Genetics</i> , 2015, 97, 738-743.	6.2	63
22	Mechanisms of immune-related adverse events associated with immune checkpoint blockade: using germline genetics to develop a personalized approach. <i>Genome Medicine</i> , 2019, 11, 39.	8.2	62
23	HIV-1 Vpu is a potent transcriptional suppressor of NF- $\kappa$ B-elicited antiviral immune responses. <i>ELife</i> , 2019, 8, .	6.0	53
24	A Coding Variant of ANO10, Affecting Volume Regulation of Macrophages, Is Associated with Borrelia Seropositivity. <i>Molecular Medicine</i> , 2015, 21, 26-37.	4.4	49
25	Replication of functional serotonin receptor type 3A and B variants in bipolar affective disorder: a European multicenter study. <i>Translational Psychiatry</i> , 2012, 2, e103-e103.	4.8	42
26	A single gene defect causing claustrophobia. <i>Translational Psychiatry</i> , 2013, 3, e254-e254.	4.8	41
27	Genetic variation associated with thyroid autoimmunity shapes the systemic immune response to PD-1 checkpoint blockade. <i>Nature Communications</i> , 2021, 12, 3355.	12.8	40
28	Hallmarks of Resistance to Immune-Checkpoint Inhibitors. <i>Cancer Immunology Research</i> , 2022, 10, 372-383.	3.4	36
29	Functional variants of the serotonin receptor type 3A and B gene are associated with eating disorders. <i>Pharmacogenetics and Genomics</i> , 2009, 19, 790-799.	1.5	35
30	Polygenic determinants of white matter volume derived from GWAS lack reproducibility in a replicate sample. <i>Translational Psychiatry</i> , 2014, 4, e362-e362.	4.8	35
31	Naturally occurring variants in the HTR3B gene significantly alter properties of human heteromeric 5-hydroxytryptamine-3A/B receptors. <i>Pharmacogenetics and Genomics</i> , 2008, 18, 793-802.	1.5	34
32	RIC-3 Exclusively Enhances the Surface Expression of Human Homomeric 5-Hydroxytryptamine Type 3A (5-HT3A) Receptors Despite Direct Interactions with 5-HT3A, -C, -D, and -E Subunits. <i>Journal of Biological Chemistry</i> , 2010, 285, 26956-26965.	3.4	31
33	Polymorphism in <i>HTR3D</i> shows different risks for acute chemotherapy-induced vomiting after anthracycline chemotherapy. <i>Pharmacogenomics</i> , 2010, 11, 943-950.	1.3	29
34	Apolipoprotein E4 carrier status plus circulating anti-NMDAR1 autoantibodies: association with schizoaffective disorder. <i>Molecular Psychiatry</i> , 2014, 19, 1054-1056.	7.9	27
35	Two naturally occurring variants of the serotonin receptor gene <i>HTR3C</i> are associated with nausea in pregnancy. <i>Acta Obstetrica Et Gynecologica Scandinavica</i> , 2010, 89, 7-14.	2.8	24
36	Mild expression differences of MECP2 influencing aggressive social behavior. <i>EMBO Molecular Medicine</i> , 2014, 6, 662-684.	6.9	23

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37	<i>In silico</i> tools for accurate HLA and KIR inference from clinical sequencing data empower immunogenetics on individual-patient and population scales. <i>Briefings in Bioinformatics</i> , 2021, 22, .	6.5	19
38	Catecholaminergic Gene Polymorphisms Are Associated with GI Symptoms and Morphological Brain Changes in Irritable Bowel Syndrome. <i>PLoS ONE</i> , 2015, 10, e0135910.	2.5	18
39	Correcting for Population Stratification Reduces False Positive and False Negative Results in Joint Analyses of Host and Pathogen Genomes. <i>Frontiers in Genetics</i> , 2018, 9, 266.	2.3	14
40	MiDASâ€”Meaningful Immunogenetic Data at Scale. <i>PLoS Computational Biology</i> , 2021, 17, e1009131.	3.2	12
41	Human genomics of the humoral immune response against polyomaviruses. <i>Virus Evolution</i> , 2021, 7, veab058.	4.9	9
42	The influence of human genetic variation on Epsteinâ€”Barr virus sequence diversity. <i>Scientific Reports</i> , 2021, 11, 4586.	3.3	8
43	Disparities in Tumor Mutational Burden, Immunotherapy Use, and Outcomes Based on Genomic Ancestry in Nonâ€”Small-Cell Lung Cancer. <i>JCO Global Oncology</i> , 2021, 7, 1537-1546.	1.8	8
44	Impact of Genetic and Nongenetic Factors on Body Mass Index and Waist-Hip Ratio Change in HIV-Infected Individuals Initiating Antiretroviral Therapy. <i>Open Forum Infectious Diseases</i> , 2020, 7, ofz464.	0.9	7
45	Allelic variation in <i>HLAâ€”DRB1</i> is associated with development of a <i>drug</i> antibodies in cancer patients treated with atezolizumab that are neutralizing in vitro. <i>Clinical and Translational Science</i> , 2022, 15, 1393-1399.	3.1	6
46	The alternative serotonin transporter promoter P2 impacts gene function in females with irritable bowel syndrome. <i>Journal of Cellular and Molecular Medicine</i> , 2021, 25, 8047-8061.	3.6	5
47	Coming of Age: Human Genomics and the Cancerâ€”Immune Set Point. <i>Cancer Immunology Research</i> , 2022, 10, 674-679.	3.4	5
48	Human genomics of acute liver failure due to hepatitis B virus infection: An exome sequencing study in liver transplant recipients. <i>Journal of Viral Hepatitis</i> , 2019, 26, 271-277.	2.0	4
49	Genetic variation near CXCL12 is associated with susceptibility to HIV-related non-Hodgkin lymphoma. <i>Haematologica</i> , 2021, 106, 2233-2241.	3.5	4
50	A common microdeletion affecting a hippocampusâ€”and amygdalaâ€”specific isoform of tryptophan hydroxylase 2 is not associated with affective disorders. <i>Bipolar Disorders</i> , 2014, 16, 764-768.	1.9	2