

# Michael Steffens

## List of Publications by Year in descending order

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Version: 2024-02-01

41  
papers

5,807  
citations

304743

22  
h-index

330143

37  
g-index

42  
all docs

42  
docs citations

42  
times ranked

11649  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study identifies 30 loci associated with bipolar disorder. <i>Nature Genetics</i> , 2019, 51, 793-803.	21.4	1,191
2	A mega-analysis of genome-wide association studies for major depressive disorder. <i>Molecular Psychiatry</i> , 2013, 18, 497-511.	7.9	1,002
3	Key susceptibility locus for nonsyndromic cleft lip with or without cleft palate on chromosome 8q24. <i>Nature Genetics</i> , 2009, 41, 473-477.	21.4	415
4	Recurrent microdeletions at 15q11.2 and 16p13.11 predispose to idiopathic generalized epilepsies. <i>Brain</i> , 2010, 133, 23-32.	7.6	406
5	Genome-wide association study identifies two susceptibility loci for nonsyndromic cleft lip with or without cleft palate. <i>Nature Genetics</i> , 2010, 42, 24-26.	21.4	379
6	Genome-wide Association Study of Alcohol Dependence. <i>Archives of General Psychiatry</i> , 2009, 66, 773.	12.3	354
7	Genome-wide Association Study Identifies Genetic Variation in Neurocan as a Susceptibility Factor for Bipolar Disorder. <i>American Journal of Human Genetics</i> , 2011, 88, 372-381.	6.2	257
8	Occupational Noise, Smoking, and a High Body Mass Index are Risk Factors for Age-related Hearing Impairment and Moderate Alcohol Consumption is Protective: A European Population-based Multicenter Study. <i>JARO - Journal of the Association for Research in Otolaryngology</i> , 2008, 9, 264-276.	1.8	214
9	Genome-Wide Association-, Replication-, and Neuroimaging Study Implicates HOMER1 in the Etiology of Major Depression. <i>Biological Psychiatry</i> , 2010, 68, 578-585.	1.3	156
10	Genome-wide significant association between alcohol dependence and a variant in the <i>ADH</i> gene cluster. <i>Addiction Biology</i> , 2012, 17, 171-180.	2.6	154
11	Genome-wide association analysis of genetic generalized epilepsies implicates susceptibility loci at 1q43, 2p16.1, 2q22.3 and 17q21.32. <i>Human Molecular Genetics</i> , 2012, 21, 5359-5372.	2.9	134
12	INTERSNP: genome-wide interaction analysis guided by a priori information. <i>Bioinformatics</i> , 2009, 25, 3275-3281.	4.1	129
13	SNP-Based Analysis of Genetic Substructure in the German Population. <i>Human Heredity</i> , 2006, 62, 20-29.	0.8	121
14	The grainyhead like 2 gene (GRHL2), alias TFCP2L3, is associated with age-related hearing impairment. <i>Human Molecular Genetics</i> , 2008, 17, 159-169.	2.9	121
15	Susceptibility variants for male-pattern baldness on chromosome 20p11. <i>Nature Genetics</i> , 2008, 40, 1279-1281.	21.4	119
16	Association of the OPRM1 Variant rs1799971 (A118G) with Non-Specific Liability to Substance Dependence in a Collaborative de novo Meta-Analysis of European-Ancestry Cohorts. <i>Behavior Genetics</i> , 2016, 46, 151-169.	2.1	98
17	Polymorphisms in the chemokine (C-X-C motif) ligand 10 are associated with invasive aspergillosis after allogeneic stem-cell transplantation and influence CXCL10 expression in monocyte-derived dendritic cells. <i>Blood</i> , 2008, 111, 534-536.	1.4	97
18	CaMKII Autophosphorylation Controls the Establishment of Alcohol Drinking Behavior. <i>Neuropsychopharmacology</i> , 2013, 38, 1636-1647.	5.4	63

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19	Polymorphisms in the Genes Encoding Chemokine Receptor 5, Interleukin-10, and Monocyte Chemoattractant Protein 1 Contribute to Cytomegalovirus Reactivation and Disease after Allogeneic Stem Cell Transplantation. <i>Journal of Clinical Microbiology</i> , 2006, 44, 1847-1850.	3.9	58
20	Cytokine regulation by epidermal growth factor receptor inhibitors and epidermal growth factor receptor inhibitor associated skin toxicity in cancer patients. <i>European Journal of Cancer</i> , 2014, 50, 1855-1863.	2.8	46
21	The contribution of GJB2 (Connexin 26) 35delG to age-related hearing impairment and noise-induced hearing loss. <i>Otology and Neurotology</i> , 2007, 28, 970-5.	1.3	37
22	Genome-wide association data provide further support for an association between 5-HTTLPR and major depressive disorder. <i>Journal of Affective Disorders</i> , 2013, 146, 438-440.	4.1	24
23	Feasible and Successful: Genome-Wide Interaction Analysis Involving All 1.9 Å— 10<sup>6</sup>; Pair-Wise Interaction Tests. <i>Human Heredity</i> , 2010, 69, 268-284.	0.8	22
24	CHL1,ITGB3andSLC6A4gene expression and antidepressant drug response: results from the Munich Antidepressant Response Signature (MARS) study. <i>Pharmacogenomics</i> , 2015, 16, 689-701.	1.3	22
25	Dosing to rash? â€œ The role of erlotinib metabolic ratio from patient serum in the search of predictive biomarkers for EGFR inhibitor-mediated skin rash. <i>European Journal of Cancer</i> , 2016, 55, 131-139.	2.8	19
26	Familial Aggregation of Pure Tone Hearing Thresholds in an Aging European Population. <i>Otology and Neurotology</i> , 2013, 34, 838-844.	1.3	15
27	Personalising drug safetyâ€”results from the multi-centre prospective observational study on Adverse Drug Reactions in Emergency Departments (ADRED). <i>European Journal of Clinical Pharmacology</i> , 2020, 76, 439-448.	1.9	15
28	Role of Glycogen Synthase Kinase 3 (GSK-3) in innate immune response of human immature dendritic cells to <i>Aspergillus fumigatus</i> . <i>Medical Mycology</i> , 2010, 48, 589-597.	0.7	14
29	Cluster-Localized Sparse Logistic Regression for SNP Data. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2012, 11, .	0.6	14
30	The phenotype of adverse drug effects: Do emergency visits due to adverse drug reactions look different in older people? Results from the ADRED study. <i>British Journal of Clinical Pharmacology</i> , 2020, 86, 2144-2154.	2.4	14
31	XRCC5 as a Risk Gene for Alcohol Dependence: Evidence from a Genome-Wide Gene-Set-Based Analysis and Follow-up Studies in <i>Drosophila</i> and Humans. <i>Neuropsychopharmacology</i> , 2015, 40, 361-371.	5.4	12
32	Predictive blood plasma biomarkers for EGFR inhibitor-induced skin rash. <i>Oncotarget</i> , 2017, 8, 35193-35204.	1.8	10
33	Adverse Drug Reactions in the Emergency Department: Is There a Role for Pharmacogenomic Profiles at Risk?â€”Results from the ADRED Study. <i>Journal of Clinical Medicine</i> , 2020, 9, 1801.	2.4	8
34	High-throughput screening identified inherited genetic variations in the EGFR pathway contributing to skin toxicity of EGFR inhibitors. <i>Pharmacogenomics</i> , 2015, 16, 1605-1619.	1.3	7
35	Citalopram-induced pathways regulation and tentative treatment-outcome-predicting biomarkers in lymphoblastoid cell lines from depression patients. <i>Translational Psychiatry</i> , 2020, 10, 210.	4.8	7
36	CYP2D6 in the Brain: Potential Impact on Adverse Drug Reactions in the Central Nervous Systemâ€”Results From the ADRED Study. <i>Frontiers in Pharmacology</i> , 2021, 12, 624104.	3.5	6

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37	Molecular Genetic Screening in Patients With ACE Inhibitor/Angiotensin Receptor Blocker-Induced Angioedema to Explore the Role of Hereditary Angioedema Genes. <i>Frontiers in Genetics</i> , 0, 13, .	2.3	4
38	Influence of metabolic profiles on the safety of drug therapy in routine care in Germany: protocol of the cohort study EMPAR. <i>BMJ Open</i> , 2020, 10, e032624.	1.9	1
39	Association between miRNA signatures in serum samples from epidermal growth factor inhibitor treated patients and skin toxicity. <i>Oncotarget</i> , 2021, 12, 982-995.	1.8	0
40	Pharmacogenetics of Oral Anticoagulation - VKORC1-Haplotypes Determine the Inter-Individual and Inter-Ethnic Variability.. <i>Blood</i> , 2006, 108, 719-719.	1.4	0
41	Evaluation of the EMPAR study population on the basis of metabolic phenotypes of selected pharmacogenes. <i>Pharmacogenomics Journal</i> , 2022, , .	2.0	0