James E Peters

List of Publications by Citations

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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.

26 1,208 13 34 g-index

38 2,118 13.9 avg, IF L-index

#	Paper	IF	Citations
26	Genomic atlas of the human plasma proteome. <i>Nature</i> , 2018 , 558, 73-79	50.4	529
25	Association of LPA Variants With Risk of Coronary Disease and the Implications for Lipoprotein(a)-Lowering Therapies: A Mendelian Randomization Analysis. <i>JAMA Cardiology</i> , 2018 , 3, 61	9 -627	235
24	Genome-wide association study of eosinophilic granulomatosis with polyangiitis reveals genomic loci stratified by ANCA status. <i>Nature Communications</i> , 2019 , 10, 5120	17.4	71
23	Genomic and drug target evaluation of 90 cardiovascular proteins in 30,931 individuals. <i>Nature Metabolism</i> , 2020 , 2, 1135-1148	14.6	61
22	Insight into Genotype-Phenotype Associations through eQTL Mapping in Multiple Cell Types in Health and Immune-Mediated Disease. <i>PLoS Genetics</i> , 2016 , 12, e1005908	6	58
21	Genetic Determinants of Lipids and Cardiovascular Disease Outcomes: A Wide-Angled Mendelian Randomization Investigation. <i>Circulation Genomic and Precision Medicine</i> , 2019 , 12, e002711	5.2	41
20	Discovery of rare variants associated with blood pressure regulation through meta-analysis of 1.3 million individuals. <i>Nature Genetics</i> , 2020 , 52, 1314-1332	36.3	26
19	Interleukin-6 Receptor Signaling and Abdominal Aortic Aneurysm Growth Rates. <i>Circulation Genomic and Precision Medicine</i> , 2019 , 12, e002413	5.2	25
18	MT-HESS: an efficient Bayesian approach for simultaneous association detection in OMICS datasets, with application to eQTL mapping in multiple tissues. <i>Bioinformatics</i> , 2016 , 32, 523-32	7.2	20
17	Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19. <i>Nature Medicine</i> , 2021 , 27, 668-676	50.5	19
16	Reduced monocyte and macrophage TNFSF15/TL1A expression is associated with susceptibility to inflammatory bowel disease. <i>PLoS Genetics</i> , 2018 , 14, e1007458	6	18
15	Longitudinal proteomic profiling of dialysis patients with COVID-19 reveals markers of severity and predictors of death. <i>ELife</i> , 2021 , 10,	8.9	17
14	Three cases of rheumatoid arthritis with laryngeal stridor. <i>Clinical Rheumatology</i> , 2011 , 30, 723-7	3.9	14
13	Temporal changes in complement activation in haemodialysis patients with COVID-19 as a predictor of disease progression. <i>CKJ: Clinical Kidney Journal</i> , 2020 , 13, 889-896	4.5	13
12	Severe localised granulomatosis with polyangiitis (Wegener\fout) granulomatosis) manifesting with extensive cranial nerve palsies and cranial diabetes insipidus: a case report and literature review. <i>BMC Neurology</i> , 2018 , 18, 59	3.1	11
11	A genome-wide meta-analysis yields 46 new loci associating with biomarkers of iron homeostasis. <i>Communications Biology</i> , 2021 , 4, 156	6.7	11
10	Targeted genomic analysis reveals widespread autoimmune disease association with regulatory variants in the TNF superfamily cytokine signalling network. <i>Genome Medicine</i> , 2016 , 8, 76	14.4	10

LIST OF PUBLICATIONS

9	Immuno-proteomic profiling reveals aberrant immune cell regulation in the airways of individuals with ongoing post-COVID-19 respiratory disease <i>Immunity</i> , 2022 ,	2.3	9
8	Novel Approach to Imaging Active Takayasu Arteritis Using Somatostatin Receptor Positron Emission Tomography/Magnetic Resonance Imaging. <i>Circulation: Cardiovascular Imaging</i> , 2020 , 13, e01038	89	8
7	Genetic Analyses of Blood Cell Structure for Biological and Pharmacological Inference		3
6	Genetically distinct clinical subsets, and associations with asthma and eosinophil abundance, within Eosinophilic Granulomatosis with Polyangiitis		3
5	Elucidating mechanisms of genetic cross-disease associations: an integrative approach implicates protein C as a causal pathway in arterial and venous diseases		2
4	Actionable druggable genome-wide Mendelian randomization identifies repurposing opportunities for COVID-19		1
3	A critical beat in eosinophilic granulomatosis with polyangiitis. <i>Rheumatology</i> , 2021 , 60, iii28-iii31	.9	1
2	Elucidating mechanisms of genetic cross-disease associations at the PROCR vascular disease locus Nature Communications, 2022, 13, 1222	7.4	O
1	Autoimmunity Due to Unicentric Castleman Disease Cured by Resection of a Hepatic Mass. <i>Journal of Clinical Rheumatology</i> , 2021 , 27, e32-e33	.1	