Benjamin P Fairfax

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/1155397/publications.pdf

Version: 2024-02-01

44 papers

6,737 citations

218677 26 h-index 243625 44 g-index

52 all docs 52 docs citations

times ranked

52

16711 citing authors

#	Article	IF	CITATIONS
1	Systematic identification of trans eQTLs as putative drivers of known disease associations. Nature Genetics, 2013, 45, 1238-1243.	21.4	1,544
2	Genetic association analyses implicate aberrant regulation of innate and adaptive immunity genes in the pathogenesis of systemic lupus erythematosus. Nature Genetics, 2015, 47, 1457-1464.	21.4	730
3	Innate Immune Activity Conditions the Effect of Regulatory Variants upon Monocyte Gene Expression. Science, 2014, 343, 1246949.	12.6	706
4	Large-scale cis- and trans-eQTL analyses identify thousands of genetic loci and polygenic scores that regulate blood gene expression. Nature Genetics, 2021, 53, 1300-1310.	21.4	590
5	Genetics of gene expression in primary immune cells identifies cell type–specific master regulators and roles of HLA alleles. Nature Genetics, 2012, 44, 502-510.	21.4	445
6	A common haplotype lowers PU.1 expression in myeloid cells and delays onset of Alzheimer's disease. Nature Neuroscience, 2017, 20, 1052-1061.	14.8	330
7	Meta-analysis of genome-wide association studies identifies ten loci influencing allergic sensitization. Nature Genetics, 2013, 45, 902-906.	21.4	221
8	Peripheral CD8+ T cell characteristics associated with durable responses to immune checkpoint blockade in patients with metastatic melanoma. Nature Medicine, 2020, 26, 193-199.	30.7	211
9	Phospho-Dependent Functional Modulation of GABAB Receptors by the Metabolic Sensor AMP-Dependent Protein Kinase. Neuron, 2007, 53, 233-247.	8.1	167
10	A blood atlas of COVID-19 defines hallmarks of disease severity and specificity. Cell, 2022, 185, 916-938.e58.	28.9	164
11	A genetics-led approach defines the drug target landscape of 30 immune-related traits. Nature Genetics, 2019, 51, 1082-1091.	21.4	157
12	Leprosy and the Adaptation of Human Toll-Like Receptor 1. PLoS Pathogens, 2010, 6, e1000979.	4.7	139
13	Genomic modulators of gene expression in human neutrophils. Nature Communications, 2015, 6, 7545.	12.8	120
14	Cell Specific eQTL Analysis without Sorting Cells. PLoS Genetics, 2015, 11, e1005223.	3.5	115
15	Pathogenic implications for autoimmune mechanisms derived by comparative eQTL analysis of CD4+ versus CD8+ T cells. PLoS Genetics, 2017, 13, e1006643.	3.5	110
16	An immunodominant NP105–113-B*07:02 cytotoxic T cell response controls viral replication and is associated with less severe COVID-19 disease. Nature Immunology, 2022, 23, 50-61.	14.5	110
17	Phosphorylation and Chronic Agonist Treatment Atypically Modulate GABAB Receptor Cell Surface Stability. Journal of Biological Chemistry, 2004, 279, 12565-12573.	3.4	99
18	Interferon-Gamma–Producing CD8+ Tissue Resident Memory T Cells Are a Targetable Hallmark of Immune Checkpoint Inhibitor–Colitis. Gastroenterology, 2021, 161, 1229-1244.e9.	1.3	87

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19	Pervasive haplotypic variation in the spliceo-transcriptome of the human major histocompatibility complex. Genome Research, 2011, 21, 1042-1054.	5. 5	63
20	Distinct Transcriptional and Anti-Mycobacterial Profiles of Peripheral Blood Monocytes Dependent on the Ratio of Monocytes: Lymphocytes. EBioMedicine, 2015, 2, 1619-1626.	6.1	61
21	Genetics of gene expression in immunity to infection. Current Opinion in Immunology, 2014, 30, 63-71.	5.5	54
22	HLA-C Level Is Regulated by a Polymorphic Oct1 Binding Site in the HLA-C Promoter Region. American Journal of Human Genetics, 2016, 99, 1353-1358.	6.2	49
23	Unravelling the unusual signalling properties of the GABAB receptor. Biochemical Pharmacology, 2004, 68, 1527-1536.	4.4	46
24	Increased prevalence of sex chromosome aneuploidies in specific language impairment and dyslexia. Developmental Medicine and Child Neurology, 2014, 56, 346-353.	2.1	42
25	The GABAB2 subunit is critical for the trafficking and function of native GABAB receptors. Biochemical Pharmacology, 2004, 68, 1655-1666.	4.4	41
26	Immune checkpoint blockade sensitivity and progression-free survival associates with baseline CD8 ⁺ T cell clone size and cytotoxicity. Science Immunology, 2021, 6, eabj8825.	11.9	41
27	Context-specific regulation of surface and soluble IL7R expression by an autoimmune risk allele. Nature Communications, 2019, 10, 4575.	12.8	37
28	Genomic mapping of the MHC transactivator CIITA using an integrated ChIP-seq and genetical genomics approach. Genome Biology, 2014, 15, 494.	8.8	32
29	A functional SNP associated with atopic dermatitis controls cell type-specific methylation of the VSTM1 gene locus. Genome Medicine, 2017, 9, 18.	8.2	30
30	Risk of nontyphoidal Salmonella bacteraemia in African children is modified by STAT4. Nature Communications, 2018, 9, 1014.	12.8	29
31	An integrated expression phenotype mapping approach defines common variants in LEP, ALOX15 and CAPNS1 associated with induction of IL-6. Human Molecular Genetics, 2010, 19, 720-730.	2.9	23
32	Toward a better understanding of TÂcells in cancer. Cancer Cell, 2021, 39, 1549-1552.	16.8	21
33	Checkpoint-blocker-induced autoimmunity is associated with favourable outcome in metastatic melanoma and distinct T-cell expression profiles. British Journal of Cancer, 2021, 124, 1661-1669.	6.4	20
34	Fine mapping genetic determinants of the highly variably expressed MHC gene ZFP57. European Journal of Human Genetics, 2014, 22, 568-571.	2.8	16
35	A Common Haplotype of the TNF Receptor 2 Gene Modulates Endotoxin Tolerance. Journal of Immunology, 2011, 186, 3058-3065.	0.8	12
36	Natural Killer cells demonstrate distinct eQTL and transcriptome-wide disease associations, highlighting their role in autoimmunity. Nature Communications, 2022, 13, .	12.8	10

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37	Whole-genome sequencing identifies homozygous <i>BRCA2</i> deletion guiding treatment in dedifferentiated prostate cancer. Journal of Physical Education and Sports Management, 2017, 3, a001362.	1.2	9
38	A global-local approach for detecting hotspots in multiple-response regression. Annals of Applied Statistics, 2020, 14, 905-928.	1.1	8
39	EPISPOT: An epigenome-driven approach for detecting and interpreting hotspots in molecular QTL studies. American Journal of Human Genetics, 2021, 108, 983-1000.	6.2	6
40	Genetic variants associated with non-typhoidal Salmonella bacteraemia in African children. Lancet, The, 2015, 385, S13.	13.7	5
41	Abiraterone acetate: a potential source of interference in testosterone assays. Clinical Chemistry and Laboratory Medicine, 2018, 56, e138-e140.	2.3	5
42	Changes in epigenetic profiles throughout early childhood and their relationship to the response to pneumococcal vaccination. Clinical Epigenetics, 2021, 13, 29.	4.1	4
43	Dissecting genetic determinants of variation in human immune responses. Current Opinion in Immunology, 2020, 65, 74-78.	5.5	3
44	Studying the Localization, Surface Stability and Endocytosis of Neurotransmitter Receptors by Antibody Labeling and Biotinylation Approaches. Frontiers in Neuroscience, 2006, , 91-118.	0.0	1