

# Mark Daly

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/11835373/publications.pdf>

Version: 2024-02-01

14  
papers

33,461  
citations

687363  
13  
h-index

1058476  
14  
g-index

18  
all docs

18  
docs citations

18  
times ranked

65909  
citing authors

#	ARTICLE	IF	CITATIONS
1	The Genome Analysis Toolkit: A MapReduce framework for analyzing next-generation DNA sequencing data. <i>Genome Research</i> , 2010, 20, 1297-1303.	5.5	21,358
2	Initial sequencing and comparative analysis of the mouse genome. <i>Nature</i> , 2002, 420, 520-562.	27.8	6,319
3	The common PPAR $\gamma$ Pro12Ala polymorphism is associated with decreased risk of type 2 diabetes. <i>Nature Genetics</i> , 2000, 26, 76-80.	21.4	1,672
4	Replicating genotype-phenotype associations. <i>Nature</i> , 2007, 447, 655-660.	27.8	1,509
5	Modeling Linkage Disequilibrium Increases Accuracy of Polygenic Risk Scores. <i>American Journal of Human Genetics</i> , 2015, 97, 576-592.	6.2	1,098
6	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
7	Polygenic and clinical risk scores and their impact on age at onset and prediction of cardiometabolic diseases and common cancers. <i>Nature Medicine</i> , 2020, 26, 549-557.	30.7	281
8	A recurrent germline PAX5 mutation confers susceptibility to pre-B cell acute lymphoblastic leukemia. <i>Nature Genetics</i> , 2013, 45, 1226-1231.	21.4	270
9	The role of polygenic risk and susceptibility genes in breast cancer over the course of life. <i>Nature Communications</i> , 2020, 11, 6383.	12.8	101
10	Exome Sequencing Analysis Reveals Variants in Primary Immunodeficiency Genes in Patients With Very Early Onset Inflammatory Bowel Disease. <i>Gastroenterology</i> , 2015, 149, 1415-1424.	1.3	99
11	The effect of LRRK2 loss-of-function variants in humans. <i>Nature Medicine</i> , 2020, 26, 869-877.	30.7	79
12	Protective Low-Frequency Variants for Preeclampsia in the Fms Related Tyrosine Kinase 1 Gene in the Finnish Population. <i>Hypertension</i> , 2017, 70, 365-371.	2.7	37
13	Principles and methods of in-silico prioritization of non-coding regulatory variants. <i>Human Genetics</i> , 2018, 137, 15-30.	3.8	37
14	Polygenic burden has broader impact on health, cognition, and socioeconomic outcomes than most rare and high-risk copy number variants. <i>Molecular Psychiatry</i> , 2021, 26, 4884-4895.	7.9	8