

Samuel A Lambert

List of Publications by Year in descending order

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12
papers

5,241
citations

759233

12
h-index

1125743

13
g-index

22
all docs

22
docs citations

22
times ranked

9842
citing authors

#	ARTICLE	IF	CITATIONS
1	The Human Transcription Factors. <i>Cell</i> , 2018, 172, 650-665.	28.9	2,048
2	Determination and Inference of Eukaryotic Transcription Factor Sequence Specificity. <i>Cell</i> , 2014, 158, 1431-1443.	28.9	1,515
3	Towards clinical utility of polygenic risk scores. <i>Human Molecular Genetics</i> , 2019, 28, R133-R142.	2.9	381
4	The Polygenic Score Catalog as an open database for reproducibility and systematic evaluation. <i>Nature Genetics</i> , 2021, 53, 420-425.	21.4	293
5	Improving reporting standards for polygenic scores in risk prediction studies. <i>Nature</i> , 2021, 591, 211-219.	27.8	265
6	Similarity regression predicts evolution of transcription factor sequence specificity. <i>Nature Genetics</i> , 2019, 51, 981-989.	21.4	105
7	Mapping and analysis of <i>Caenorhabditis elegans</i> transcription factor sequence specificities. <i>ELife</i> , 2015, 4, .	6.0	85
8	Individual Saturated and Monounsaturated Fatty Acids Trigger Distinct Transcriptional Networks in Differentiated 3T3-L1 Preadipocytes. <i>Journal of Nutrigenetics and Nutrigenomics</i> , 2013, 6, 1-15.	1.3	43
9	Integrative analysis of the plasma proteome and polygenic risk of cardiometabolic diseases. <i>Nature Metabolism</i> , 2021, 3, 1476-1483.	11.9	43
10	Comparison of ChIP-Seq Data and a Reference Motif Set for Human KRAB C2H2 Zinc Finger Proteins. <i>G3: Genes, Genomes, Genetics</i> , 2018, 8, 219-229.	1.8	22
11	Motif comparison based on similarity of binding affinity profiles. <i>Bioinformatics</i> , 2016, 32, 3504-3506.	4.1	16
12	Transcription factors read epigenetics. <i>Science</i> , 2017, 356, 489-490.	12.6	16