

Mary D Fortune

List of Publications by Year in descending order

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11
papers

1,136
citations

932766

10
h-index

1199166

12
g-index

14
all docs

14
docs citations

14
times ranked

3195
citing authors

#	ARTICLE	IF	CITATIONS
1	Personalized Model to Predict Keratoconus Progression From Demographic, Topographic, and Genetic Data. <i>American Journal of Ophthalmology</i> , 2022, 240, 321-329.	1.7	7
2	Specificity and positive predictive value of SARS-CoV-2 nucleic acid amplification testing in a low-prevalence setting. <i>Clinical Microbiology and Infection</i> , 2021, 27, 469.e9-469.e15.	2.8	52
3	Diagnostic tool or screening programme? Asymptomatic testing for SARS-CoV-2 needs clear goals and protocols. <i>Lancet Regional Health - Europe</i> , The, 2021, 1, 100002.	3.0	19
4	Systematic review on the outcomes of poller screw augmentation in intramedullary nailing of long bone fracture. <i>EFORT Open Reviews</i> , 2020, 5, 189-203.	1.8	17
5	Stochastic search and joint fine-mapping increases accuracy and identifies previously unreported associations in immune-mediated diseases. <i>Nature Communications</i> , 2019, 10, 3216.	5.8	24
6	simGWAS: a fast method for simulation of large scale case-control GWAS summary statistics. <i>Bioinformatics</i> , 2019, 35, 1901-1906.	1.8	23
7	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.	1.5	80
8	Fine mapping of type 1 diabetes susceptibility loci and evidence for colocalization of causal variants with lymphoid gene enhancers. <i>Nature Genetics</i> , 2015, 47, 381-386.	9.4	589
9	Statistical colocalization of genetic risk variants for related autoimmune diseases in the context of common controls. <i>Nature Genetics</i> , 2015, 47, 839-846.	9.4	128
10	Integration of disease association and eQTL data using a Bayesian colocalisation approach highlights six candidate causal genes in immune-mediated diseases. <i>Human Molecular Genetics</i> , 2015, 24, 3305-3313.	1.4	134
11	A Method for Gene-Based Pathway Analysis Using Genomewide Association Study Summary Statistics Reveals Nine New Type 1 Diabetes Associations. <i>Genetic Epidemiology</i> , 2014, 38, 661-670.	0.6	54