

Rowmika Ravi

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1434562/publications.pdf>

Version: 2024-02-01

10
papers

105
citations

1937685

4
h-index

1372567

10
g-index

10
all docs

10
docs citations

10
times ranked

201
citing authors

#	ARTICLE	IF	CITATIONS
1	Saliva microbiome in primary Sjögren's syndrome reveals distinct set of disease-associated microbes. Oral Diseases, 2020, 26, 295-301.	3.0	39
2	Pharmacogenetic landscape of <i>DPYD</i> variants in south Asian populations by integration of genome-scale data. Pharmacogenomics, 2018, 19, 227-241.	1.3	25
3	SAGE: a comprehensive resource of genetic variants integrating South Asian whole genomes and exomes. Database: the Journal of Biological Databases and Curation, 2018, 2018, 1-10.	3.0	20
4	Egyptian tale from India: application of whole-exome sequencing in diagnosis of atypical familial Mediterranean fever. International Journal of Rheumatic Diseases, 2017, 20, 1770-1775.	1.9	7
5	Understanding the complexity of epimorphic regeneration in zebrafish caudal fin tissue: A transcriptomic and proteomic approach. Genomics, 2022, 114, 110300.	2.9	5
6	Case Report: Whole exome sequencing reveals a novel frameshift deletion mutation p.G2254fs in COL7A1 associated with autosomal recessive dystrophic epidermolysis bullosa. F1000Research, 2016, 5, 900.	1.6	3
7	Case Report: Whole exome sequencing reveals a novel frameshift deletion mutation p.G2254fs in COL7A1 associated with autosomal recessive dystrophic epidermolysis bullosa. F1000Research, 2016, 5, 900.	1.6	2
8	Case Report: Whole exome sequencing identifies variation c.2308G>A p.E770K in RAG1 associated with B- T- NK+ severe combined immunodeficiency. F1000Research, 2016, 5, 2532.	1.6	2
9	Case Report: Whole exome sequencing identifies variation c.2308G>A p.E770K in RAG1 associated with B- T- NK+ severe combined immunodeficiency. F1000Research, 2016, 5, 2532.	1.6	1
10	Metabolite Signature in the Carriers of Pathogenic Genetic Variants for Cardiomyopathy: A Population-Based METSIM Study. Metabolites, 2022, 12, 437.	2.9	1