## Deepak R Unni

List of Publications by Year in descending order

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840776 1058476 3,718 14 11 14 citations h-index g-index papers 16 16 16 6274 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	The Gene Ontology resource: enriching a GOld mine. Nucleic Acids Research, 2021, 49, D325-D334.	14.5	2,416
2	The National COVID Cohort Collaborative (N3C): Rationale, design, infrastructure, and deployment. Journal of the American Medical Informatics Association: JAMIA, 2021, 28, 427-443.	4.4	342
3	How many rare diseases are there?. Nature Reviews Drug Discovery, 2020, 19, 77-78.	46.4	204
4	Apollo: Democratizing genome annotation. PLoS Computational Biology, 2019, 15, e1006790.	3.2	179
5	The Monarch Initiative in 2019: an integrative data and analytic platform connecting phenotypes to genotypes across species. Nucleic Acids Research, 2020, 48, D704-D715.	14.5	178
6	Hymenoptera Genome Database: integrating genome annotations in HymenopteraMine. Nucleic Acids Research, 2016, 44, D793-D800.	14.5	105
7	Bovine Genome Database: new tools for gleaning function from the <i>Bos taurus </i> Acids Research, 2016, 44, D834-D839.	14.5	87
8	KG-COVID-19: A Framework to Produce Customized Knowledge Graphs for COVID-19 Response. Patterns, 2021, 2, 100155.	5.9	62
9	Novel dual cyclooxygenase and lipoxygenase inhibitors targeting hyaluronan–CD44v6 pathway and inducing cytotoxicity in colon cancer cells. Bioorganic and Medicinal Chemistry, 2013, 21, 2551-2559.	3.0	41
10	Biolink Model: A universal schema for knowledge graphs in clinical, biomedical, and translational science. Clinical and Translational Science, 2022, 15, 1848-1855.	3.1	38
11	Bovine Genome Database: new annotation tools for a new reference genome. Nucleic Acids Research, 2019, 48, D676-D681.	14.5	18
12	MaizeMine: A Data Mining Warehouse for the Maize Genetics and Genomics Database. Frontiers in Plant Science, 2020, 11, 592730.	3.6	13
13	Hymenoptera Genome Database: Using HymenopteraMine to Enhance Genomic Studies of Hymenopteran Insects. Methods in Molecular Biology, 2018, 1757, 513-556.	0.9	12
14	Bovine Genome Database: Tools for Mining the Bos taurus Genome. Methods in Molecular Biology, 2018, 1757, 211-249.	0.9	5