

# Sree Rohit Raj Kolora

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/1360191/publications.pdf>

Version: 2024-02-01

15  
papers

116  
citations

1937685

4  
h-index

1588992

8  
g-index

16  
all docs

16  
docs citations

16  
times ranked

220  
citing authors

#	ARTICLE	IF	CITATIONS
1	Origins and evolution of extreme life span in Pacific Ocean rockfishes. <i>Science</i> , 2021, 374, 842-847.	12.6	71
2	Divergent evolution in the genomes of closely-related lacertids, <i>Lacerta viridis</i> and <i>L. bilineata</i> and implications for speciation. <i>GigaScience</i> , 2018, 8, .	6.4	10
3	Identification of potential therapeutic targets in <i>Neisseria gonorrhoeae</i> by an in-silico approach. <i>Journal of Theoretical Biology</i> , 2020, 490, 110172.	1.7	8
4	Elephant seal muscle cells adapt to sustained glucocorticoid exposure by shifting their metabolic phenotype. <i>American Journal of Physiology - Regulatory Integrative and Comparative Physiology</i> , 2021, 321, R413-R428.	1.8	7
5	A crowdsourced set of curated structural variants for the human genome. <i>PLoS Computational Biology</i> , 2020, 16, e1007933.	3.2	6
6	The complete mitochondrial genome of <i>Lacerta bilineata</i> and comparison with its closely related congener <i>L. Viridis</i> . <i>Mitochondrial DNA Part A: DNA Mapping, Sequencing, and Analysis</i> , 2017, 28, 116-118.	0.7	5
7	Introducing evolutionary biologists to the analysis of big data: guidelines to organize extended bioinformatics training courses. <i>Evolution: Education and Outreach</i> , 2018, 11, .	0.8	3
8	Prioritizing long range interactions in noncoding regions using GWAS and deletions perturbed TADs. <i>Computational and Structural Biotechnology Journal</i> , 2020, 18, 2945-2952.	4.1	2
9	Mitogenome evolution in the <i>Lacerta viridis</i> complex (Lacertidae, Squamata) reveals phylogeny of diverging clades. <i>Systematics and Biodiversity</i> , 0, , 1-12.	1.2	2
10	Accelerated Evolution of Tissue-Specific Genes Mediates Divergence Amidst Gene Flow in European Green Lizards. <i>Genome Biology and Evolution</i> , 2021, 13, .	2.5	1
11	The third international hackathon for applying insights into large-scale genomic composition to use cases in a wide range of organisms. <i>F1000Research</i> , 0, 11, 530.	1.6	1
12	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
13	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
14	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0
15	A crowdsourced set of curated structural variants for the human genome. , 2020, 16, e1007933.		0