## Alaguraj Veluchamy

List of Publications by Year in descending order

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257450 3,057 32 24 citations h-index papers

32 g-index 37 37 37 5014 docs citations times ranked citing authors all docs

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#	Article	IF	CITATIONS
1	Exploring the biological behavior of Heat shock protein (HSPs) for understanding the Anti-ischemic stroke in humans. Journal of Infection and Public Health, 2022, 15, 379-388.	4.1	O
2	Evaluating the structural and immune mechanism of Interleukin-6 for the investigation of goat milk peptides as potential treatments for COVID-19. Journal of King Saud University - Science, 2022, 34, 101924.	3 <b>.</b> 5	6
3	Genome wide natural variation of H3K27me3 selectively marks genes predicted to be important for cell differentiation in <i>Phaeodactylum tricornutum</i> . New Phytologist, 2021, 229, 3208-3220.	7.3	19
4	Information theoretic perspective on genome clustering. Saudi Journal of Biological Sciences, 2021, 28, 1867-1889.	3.8	2
5	A genomics approach reveals the global genetic polymorphism, structure, and functional diversity of ten accessions of the marine model diatom <i>Phaeodactylum tricornutum</i> . ISME Journal, 2020, 14, 347-363.	9.8	50
6	Genome-enabled phylogenetic and functional reconstruction of an araphid pennate diatom Plagiostriata sp. CCMP470, previously assigned as a radial centric diatom, and its bacterial commensal. Scientific Reports, 2020, 10, 9449.	3.3	25
7	GCN5 modulates salicylic acid homeostasis by regulating H3K14ac levels at the $5\hat{a} \in \mathbb{Z}^2$ and $3\hat{a} \in \mathbb{Z}^2$ ends of its target genes. Nucleic Acids Research, 2020, 48, 5953-5966.	14.5	44
8	Probing the Diversity of Polycomb and Trithorax Proteins in Cultured and Environmentally Sampled Microalgae. Frontiers in Marine Science, 2020, 7, .	2.5	26
9	Alternative splicing and allosteric regulation modulate the chromatin binding of UHRF1. Nucleic Acids Research, 2020, 48, 7728-7747.	14.5	16
10	R-Loop Mediated trans Action of the APOLO Long Noncoding RNA. Molecular Cell, 2020, 77, 1055-1065.e4.	9.7	164
11	Wheat chromatin architecture is organized in genome territories and transcription factories. Genome Biology, 2020, 21, 104.	8.8	99
12	INDETERMINATE-DOMAIN 4 (IDD4) coordinates immune responses with plant-growth in Arabidopsis thaliana. PLoS Pathogens, 2019, 15, e1007499.	4.7	17
13	Community‣evel Responses to Iron Availability in Open Ocean Plankton Ecosystems. Global Biogeochemical Cycles, 2019, 33, 391-419.	4.9	76
14	Thermopriming triggers splicing memory in Arabidopsis. Journal of Experimental Botany, 2018, 69, 2659-2675.	4.8	119
15	Evolutionary genomics of the cold-adapted diatom Fragilariopsis cylindrus. Nature, 2017, 541, 536-540.	27.8	332
16	Ploidy-dependent changes in the epigenome of symbiotic cells correlate with specific patterns of gene expression. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, 4543-4548.	7.1	50
17	Arabidopsis ATRX Modulates H3.3 Occupancy and Fine-Tunes Gene Expression. Plant Cell, 2017, 29, 1773-1793.	6.6	35
18	The Arabidopsis SWI/SNF protein BAF60 mediates seedling growth control by modulating DNA accessibility. Genome Biology, 2017, 18, 114.	8.8	53

#	Article	IF	Citations
19	The quest for epigenetic regulation underlying unisexual flower development in Cucumis melo. Epigenetics and Chromatin, 2017, 10, 22.	3.9	27
20	MAPK-triggered chromatin reprogramming by histone deacetylase in plant innate immunity. Genome Biology, 2017, 18, 131.	8.8	73
21	Insights into global diatom distribution and diversity in the world's ocean. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E1516-25.	7.1	561
22	Put your 3D glasses on: plant chromatin is on show. Journal of Experimental Botany, 2016, 67, 3205-3221.	4.8	59
23	Reverse transcriptase genes are highly abundant and transcriptionally active in marine plankton assemblages. ISME Journal, 2016, 10, 1134-1146.	9.8	35
24	LHP1 Regulates H3K27me3 Spreading and Shapes the Three-Dimensional Conformation of the Arabidopsis Genome. PLoS ONE, 2016, 11, e0158936.	2.5	97
25	An integrative analysis of post-translational histone modifications in the marine diatom Phaeodactylum tricornutum. Genome Biology, 2015, 16, 102.	8.8	107
26	Chromerid genomes reveal the evolutionary path from photosynthetic algae to obligate intracellular parasites. ELife, 2015, 4, e06974.	6.0	198
27	A Novel Protein, Ubiquitous in Marine Phytoplankton, Concentrates Iron at the Cell Surface and Facilitates Uptake. Current Biology, 2015, 25, 364-371.	3.9	90
28	Oil Accumulation by the Oleaginous Diatom <i>Fistulifera solaris</i> as Revealed by the Genome and Transcriptome. Plant Cell, 2015, 27, 162-176.	6.6	149
29	Insights into the role of DNA methylation in diatoms by genome-wide profiling in Phaeodactylum tricornutum. Nature Communications, 2013, 4, 2091.	12.8	113
30	A Holistic Approach to Marine Eco-Systems Biology. PLoS Biology, 2011, 9, e1001177.	5.6	353
31	HNHDb: A database on pattern based classification of HNH domains reveals functional relevance of sequence patterns and domain associations Bioinformation, 2009, 4, 80-83.	0.5	12
32	Identification of Prophages in Bacterial Genomes by Dinucleotide Relative Abundance Difference. PLoS ONE, 2007, 2, e1193.	2.5	44