## David R Nelson

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Molecular Mechanisms behind Safranal's Toxicity to HepG2 Cells from Dual Omics. Antioxidants, 2022, 11, 1125.	5.1	31
2	Large-scale genome sequencing reveals the driving forces of viruses in microalgal evolution. Cell Host and Microbe, 2021, 29, 250-266.e8.	11.0	48
3	Alternative glycosylation controls endoplasmic reticulum dynamics and tubular extension in mammalian cells. Science Advances, 2021, 7, .	10.3	8
4	Protocol to generate and characterize biofouling transformants of a model marine diatom. STAR Protocols, 2021, 2, 100716.	1.2	0
5	A high-quality genome assembly and annotation of the gray mangrove, <i>Avicennia marina</i> . G3: Genes, Genomes, Genetics, 2021, 11, .	1.8	16
6	High-Throughput Metabolic Profiling for Model Refinements of Microalgae. Journal of Visualized Experiments, 2021, , .	0.3	0
7	GPCR Genes as Activators of Surface Colonization Pathways in a Model Marine Diatom. IScience, 2020, 23, 101424.	4.1	7
8	Prospects for the Study and Improvement of Abiotic Stress Tolerance in Date Palms in the Post-genomics Era. Frontiers in Plant Science, 2020, 11, 293.	3.6	34
9	The genome of pest Rhynchophorus ferrugineus reveals gene families important at the plant-beetle interface. Communications Biology, 2020, 3, 323.	4.4	44
10	Positive correlation between wood δ <sup>15</sup> N and stream nitrate concentrations in two temperate deciduous forests. Environmental Research Communications, 2020, 2, 025003.	2.3	8
11	Advances in microalgal research and engineering development. Current Opinion in Biotechnology, 2019, 59, 157-164.	6.6	73
12	Potential for Heightened Sulfur-Metabolic Capacity in Coastal Subtropical Microalgae. IScience, 2019, 11, 450-465.	4.1	23
13	Combined artificial high-silicate medium and LED illumination promote carotenoid accumulation in the marine diatom Phaeodactylum tricornutum. Microbial Cell Factories, 2019, 18, 209.	4.0	27
14	Safranal induces DNA double-strand breakage and ER-stress-mediated cell death in hepatocellular carcinoma cells. Scientific Reports, 2018, 8, 16951.	3.3	82
15	Mapping of HKT1;5 Gene in Barley Using GWAS Approach and Its Implication in Salt Tolerance Mechanism. Frontiers in Plant Science, 2018, 9, 156.	3.6	95
16	Intracellular spectral recompositioning of light enhances algal photosynthetic efficiency. Science Advances, 2017, 3, e1603096.	10.3	42
17	The genome and phenome of the green alga Chloroidium sp. UTEX 3007 reveal adaptive traits for desert acclimatization. ELife, 2017, 6, .	6.0	16
18	Algal Cell Factories: Approaches, Applications, and Potentials. Marine Drugs, 2016, 14, 225.	4.6	65

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19	Systems level analysis of the Chlamydomonas reinhardtii metabolic network reveals variability in evolutionary co-conservation. Molecular BioSystems, 2016, 12, 2394-2407.	2.9	12
20	Single-Cell Characterization of Microalgal Lipid Contents with Confocal Raman Microscopy. Series in Bioengineering, 2016, , 363-382.	0.6	3
21	An integrative Raman microscopy-based workflow for rapid in situ analysis of microalgal lipid bodies. Biotechnology for Biofuels, 2015, 8, 164.	6.2	58
22	Toward Applications of Genomics and Metabolic Modeling to Improve Algal Biomass Productivity. Biofuel and Biorefinery Technologies, 2015, , 173-189.	0.3	5
23	Whole-Genome Resequencing Reveals Extensive Natural Variation in the Model Green Alga <i>Chlamydomonas reinhardtii</i> . Plant Cell, 2015, 27, 2353-2369.	6.6	92
24	Microalgal Metabolic Network Model Refinement through High-Throughput Functional Metabolic Profiling. Frontiers in Bioengineering and Biotechnology, 2014, 2, 68.	4.1	29
25	Heterococcus sp. DN1 draft genome: focus on cold tolerance and lipid production. Microbiome Science and Medicine, 2013, 1, .	0.3	3
26	The <i>Chlamydomonas</i> Genome Reveals the Evolution of Key Animal and Plant Functions. Science, 2007, 318, 245-250.	12.6	2,354