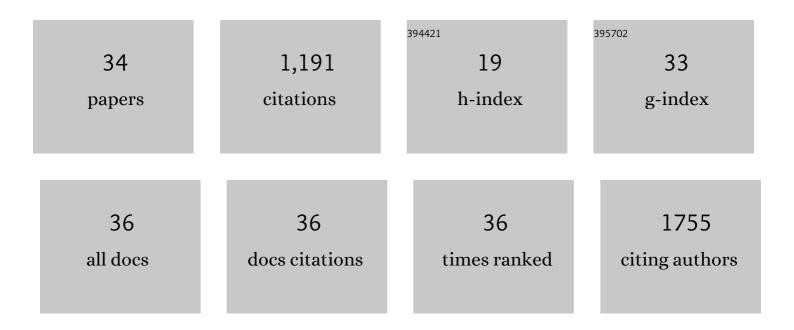
Emmanuel Gonzalez

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Altered microbiome composition in individuals with fibromyalgia. Pain, 2019, 160, 2589-2602.	4.2	130
2	GenPipes: an open-source framework for distributed and scalable genomic analyses. GigaScience, 2019, 8, .	6.4	121
3	Copper Corroles Are Inherently Saddled. Inorganic Chemistry, 2009, 48, 7794-7799.	4.0	103
4	Trees, fungi and bacteria: tripartite metatranscriptomics of a root microbiome responding to soil contamination. Microbiome, 2018, 6, 53.	11.1	88
5	Molecular structures and electron distributions of higher-valent iron and manganese porphyrins: Density functional theory calculations and some preliminary open-shell coupled-cluster results. Journal of Porphyrins and Phthalocyanines, 2001, 05, 345-356.	0.8	80
6	The Response of a 16S Ribosomal RNA Gene Fragment Amplified Community to Lead, Zinc, and Copper Pollution in a Shanghai Field Trial. Frontiers in Microbiology, 2018, 9, 366.	3.5	73
7	Theoretical Studies of Low-Spin Six-Coordinate Iron(III) Porphyrins Relevant to Cytochromes b: Variable Electronic Configurations, Ligand Noninnocence, and Macrocycle Ruffling. Journal of Physical Chemistry B, 1999, 103, 1363-1367.	2.6	60
8	Absorption spectroscopy of octahedral nickel(II) complexes: A case study of interactions between multiple electronic excited states. Coordination Chemistry Reviews, 2007, 251, 351-363.	18.8	60
9	Theoretical studies on high-valent manganese porphyrins: Toward a deeper understanding of the energetics, electron distributions, and structural features of the reactive intermediates of enzymatic and synthetic manganese-catalyzed oxidative processes. Israel Journal of Chemistry, 2000, 40, 1-8.	2.3	38
10	Co-cropping with three phytoremediation crops influences rhizosphere microbiome community in contaminated soil. Science of the Total Environment, 2020, 711, 135067.	8.0	38
11	Transcriptomic Response of Purple Willow (Salix purpurea) to Arsenic Stress. Frontiers in Plant Science, 2017, 8, 1115.	3.6	32
12	ANCHOR: a 16S rRNA gene amplicon pipeline for microbial analysis of multiple environmental samples. Environmental Microbiology, 2019, 21, 2440-2468.	3.8	30
13	Valence tautomerism and macrocycle ruffling in nickel(III) porphyrins. Journal of Inorganic Biochemistry, 2000, 78, 79-82.	3.5	28
14	Reductive coupling and protonation leading to diboron corroles with a B–H–B bridge. Dalton Transactions, 2010, 39, 4032.	3.3	26
15	Intranasal Application of Lactococcus lactis W136 Is Safe in Chronic Rhinosinusitis Patients With Previous Sinus Surgery. Frontiers in Cellular and Infection Microbiology, 2020, 10, 440.	3.9	26
16	Distinct Changes Occur in the Human Breast Milk Microbiome Between Early and Established Lactation in Breastfeeding Guatemalan Mothers. Frontiers in Microbiology, 2021, 12, 557180.	3.5	26
17	Upregulated IL-32 Expression And Reduced Gut Short Chain Fatty Acid Caproic Acid in People Living With HIV With Subclinical Atherosclerosis. Frontiers in Immunology, 2021, 12, 664371.	4.8	25
18	Meta-transcriptomics indicates biotic cross-tolerance in willow trees cultivated on petroleum hydrocarbon contaminated soil. BMC Plant Biology, 2015, 15, 246.	3.6	21

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#	Article	IF	CITATIONS
19	DFT at Its Best: Metal- versus Ligand-Centered Reduction in Nickel Hydroporphyrins. Journal of Physical Chemistry B, 2008, 112, 15158-15173.	2.6	20
20	Impact of RNA-seq attributes on false positive rates in differential expression analysis of de novo assembled transcriptomes. BMC Research Notes, 2013, 6, 503.	1.4	20
21	Comparative Transcriptomic Approaches Exploring Contamination Stress Tolerance in <i>Salix</i> sp. Reveal the Importance for a Metaorganismal de Novo Assembly Approach for Nonmodel Plants. Plant Physiology, 2016, 171, 3-24.	4.8	20
22	Mapping the dâ^'d Excited-State Manifolds of Transition Metal β-Diiminatoâ^'Imido Complexes. Comparison of Density Functional Theory and CASPT2 Energetics. Journal of Physical Chemistry A, 2008, 112, 12792-12798.	2.5	19
23	Models of High-Valent Intermediates of Non-Heme Diiron Alkane Monooxygenases: Electronic Structure of a Bis(1¼-oxo)diron(IV) Complex with Locally Low-Spin Metal Centers. Angewandte Chemie - International Edition, 2004, 43, 834-838.	13.8	17
24	Oral iron supplementation after antibiotic exposure induces a deleterious recovery of the gut microbiota. BMC Microbiology, 2021, 21, 259.	3.3	15
25	Reanalysis of the Mars500 experiment reveals common gut microbiome alterations in astronauts induced by long-duration confinement. Computational and Structural Biotechnology Journal, 2021, 19, 2223-2235.	4.1	12
26	Density Functional Theory Calculations on Ruthenium(IV) Bis(amido) Porphyrins: Search for a Broader Perspective of Heme Protein Compound II Intermediates. Journal of Physical Chemistry B, 2010, 114, 15380-15388.	2.6	10
27	Lowâ€dose and longâ€ŧerm azithromycin significantly decreases <i>Staphylococcus aureus</i> in the microbiome of refractory CRS patients. International Forum of Allergy and Rhinology, 2021, 11, 93-105.	2.8	10
28	Phytochelatin and coumarin enrichment in root exudates of arsenicâ€ŧreated white lupin. Plant, Cell and Environment, 2022, 45, 936-954.	5.7	10
29	Snow crab (Chionoecetes opilio) hepatopancreas transcriptome: Identification and testing of candidate molecular biomarkers of seismic survey impact. Fisheries Research, 2021, 234, 105794.	1.7	7
30	Human Milk Microbiota in an Indigenous Population Is Associated with Maternal Factors, Stage of Lactation, and Breastfeeding Practices. Current Developments in Nutrition, 2021, 5, nzab013.	0.3	5
31	Spectroscopic effects resulting from interacting singlet and triplet excited states: vibronic structure involving the O–H stretching mode in d–d absorption bands of Ni(H ₂ O) ₆ ²⁺ . Dalton Transactions, 2014, 43, 17864-17870.	3.3	4
32	Dietary Intake Is Unlikely to Explain Symptom Severity and Syndrome-Specific Microbiome Alterations in a Cohort of Women with Fibromyalgia. International Journal of Environmental Research and Public Health, 2022, 19, 3254.	2.6	4
33	Microbial community origin and fate through a rural wastewater treatment plant. Environmental Microbiology, 2022, 24, 2516-2542.	3.8	4
34	Models of the ox1 State of Methylcoenzymeâ€M Reductase: Where are the Electrons?. Chemistry - A European Journal, 2008, 14, 9981-9989.	3.3	3