## James R Garey

## List of Publications by Year in descending order

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		218677	182427
54	6,605	26	51
papers	citations	h-index	g-index
55	55	55	5518
all docs	docs citations	times ranked	citing authors

#	Article	IF	CITATIONS
1	A molecular evolutionary framework for the phylum Nematoda. Nature, 1998, 392, 71-75.	27.8	1,697
2	Evidence for a clade of nematodes, arthropods and other moulting animals. Nature, 1997, 387, 489-493.	27.8	1,502
3	Evolution of the chordate body plan: New insights from phylogenetic analyses of deuterostome phyla. Proceedings of the National Academy of Sciences of the United States of America, 2000, 97, 4469-4474.	7.1	380
4	Ecdysozoan phylogeny and Bayesian inference: first use of nearly complete 28S and 18S rRNA gene sequences to classify the arthropods and their kin. Molecular Phylogenetics and Evolution, 2004, 31, 178-191.	2.7	315
5	Platyhelminth mitochondrial DNA: Evidence for early evolutionary origin of a tRNAserAGN that contains a dihydrouridine arm replacement loop, and of serine-specifying AGA and AGG codons. Journal of Molecular Evolution, 1989, 28, 374-387.	1.8	246
6	Urochordates Are Monophyletic Within the Deuterostomes. Systematic Biology, 2000, 49, 52-64.	5.6	218
7	The position of the Arthropoda in the phylogenetic system. Journal of Morphology, 1998, 238, 263-285.	1.2	186
8	Globalâ€scale patterns of assemblage structure of soil nematodes in relation to climate and ecosystem properties. Global Ecology and Biogeography, 2014, 23, 968-978.	5.8	171
9	Molecular study of worldwide distribution and diversity of soil animals. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 17720-17725.	7.1	165
10	Molecular evidence for Acanthocephala as a subtaxon of Rotifera. Journal of Molecular Evolution, 1996, 43, 287-292.	1.8	163
11	The evolution of bacterial LuxI and LuxR quorum sensing regulators. Microbiology (United Kingdom), 2001, 147, 2379-2387.	1.8	161
12	Large-Scale Taxonomic Profiling of Eukaryotic Model Organisms: A Comparison of Orthologous Proteins Encoded by the Human, Fly, Nematode, and Yeast Genomes. Genome Research, 1998, 8, 590-598.	5.5	156
13	Phylogenetic Relationships of the Acanthocephala Inferred from 18S Ribosomal DNA Sequences. Molecular Phylogenetics and Evolution, 1998, 10, 287-298.	2.7	137
14	Comparative Evolutionary Genomics Unveils the Molecular Mechanism of Reassignment of the CTG Codon in Candida spp Genome Research, 2003, 13, 544-557.	5.5	111
15	Molecular Analysis Supports a Tardigrade-Arthropod Association. Invertebrate Biology, 1996, 115, 79.	0.9	110
16	The evolutionary relationships of rotifers and acanthocephalans. Hydrobiologia, 1998, 387/387, 83-91.	2.0	97
17	Ecdysozoa: The Relationship between Cycloneuralia and Panarthropoda. Zoologischer Anzeiger, 2001, 240, 321-330.	0.9	70
18	The Essential Role of "Minor―Phyla in Molecular Studies of Animal Evolution. American Zoologist, 1998, 38, 907-917.	0.7	59

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19	Systematic relationships of Nematomorpha based on molecular and morphological data. Invertebrate Biology, 2002, 121, 357-364.	0.9	52
20	Analysis of uroporphyrinogen decarboxylase complementary DNAs in sporadic porphyria cutanea tarda. Gastroenterology, 1993, 105, 165-169.	1.3	49
21	Molecular profiling of soil animal diversity in natural ecosystems: Incongruence of molecular and morphological results. Soil Biology and Biochemistry, 2009, 41, 849-857.	8.8	48
22	A Family Level Analysis of Tardigrade Phylogeny. Hydrobiologia, 2006, 558, 53-60.	2.0	45
23	Structure and function of hemoglobin from Urechis caupo. Archives of Biochemistry and Biophysics, 1984, 228, 320-331.	3.0	38
24	Exon-Intron Organization in Genes of Earthworm and Vertebrate Globins. Science, 1988, 240, 334-336.	12.6	38
25	Phylogeny, Biogeography, and Infrageneric Classification of <i>Harrisia</i> (Cactaceae). Systematic Botany, 2013, 38, 210-223.	0.5	36
26	A Comparative Genomics Analysis of Codon Reassignments Reveals a Link with Mitochondrial Proteome Size and a Mechanism of Genetic Code Change Via Suppressor tRNAs. Journal of Molecular Evolution, 2007, 64, 399-410.	1.8	32
27	Uroporphyrinogen decarboxylase in Saccharomyces cerevisiae. HEM12 gene sequence and evidence for two conserved glycines essential for enzymatic activity. FEBS Journal, 1992, 205, 1011-1016.	0.2	27
28	Microbial Function and Hydrochemistry within a Stratified Anchialine Sinkhole: A Window into Coastal Aquifer Interactions. Water (Switzerland), 2018, 10, 972.	2.7	24
29	Comparison of an inactive submarine spring with an active nearshore anchialine spring in Florida. Hydrobiologia, 2011, 677, 65-87.	2.0	21
30	Karst estuaries are governed by interactions between inland hydrological conditions and sea level. Journal of Hydrology, 2015, 527, 718-733.	5.4	21
31	A natural laboratory to study arsenic geobiocomplexity. Eos, 2006, 87, 221.	0.1	20
32	Sequencing and analysis of the gastrula transcriptome of the brittle star Ophiocoma wendtii. EvoDevo, 2012, 3, 19.	3.2	19
33	The position of the Arthropoda in the phylogenetic system. Journal of Morphology, 1998, 238, 263-285.	1.2	19
34	The Florida amphioxus (Cephalochordata) hosts larvae of the tapeworm <i>Acanthobothrium brevissime</i> : natural history, anatomy and taxonomic identification of the parasite. Acta Zoologica, 2009, 90, 75-86.	0.8	18
35	Changes in Benthic Macrofauna Associated with a Shallow-Water Hydrothermal Vent Gradient in Papua New Guinea. Pacific Science, 2010, 64, 391-404.	0.6	18
36	Global diversity of tardigrades (Tardigrada) in freshwater. Hydrobiologia, 2008, 595, 101-106.	2.0	16

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37	The transition of a freshwater karst aquifer to an anoxic marine system. Estuaries and Coasts, 2005, 28, 686-693.	1.7	14
38	Lowâ€copy nuclear primers and <i>ycf1</i> primers in Cactaceae. American Journal of Botany, 2012, 99, e405-7.	1.7	14
39	Megraft: a software package to graft ribosomal small subunit (16S/18S) fragments onto full-length sequences for accurate species richness and sequencing depth analysis in pyrosequencing-length metagenomes and similar environmental datasets. Research in Microbiology, 2012, 163, 407-412.	2.1	12
40	Aquifer Discharge Drives Microbial Community Change in Karst Estuaries. Estuaries and Coasts, 2018, 41, 430-443.	2.2	12
41	The fate of urban springs: Pumping-induced seawater intrusion in a phreatic cave. Journal of Hydrology, 2018, 564, 230-245.	5.4	10
42	Surface runoff alters cave microbial community structure and function. PLoS ONE, 2020, 15, e0232742.	2.5	9
43	Relationships and dispersal of the Caribbean species of <i>Harrisia</i> (sect. <i>Harrisia</i> ; Cactaceae) using AFLPs and seven DNA regions. Taxon, 2013, 62, 486-497.	0.7	8
44	Molecular Evidence for Acanthocephala as a Subtaxon of Rotifera. Journal of Molecular Evolution, 1996, 43, 287-292.	1.8	8
45	The Lesser-Known Protostome Taxa: An Introduction and a Tribute to Robert P. Higgins. Integrative and Comparative Biology, 2002, 42, 611-618.	2.0	6
46	Molecular analyses of microbial abundance and diversity in the water column of anchialine caves in Mallorca, Spain. International Journal of Speleology, 2014, 43, 217-226.	1.0	6
47	Source of saline groundwater on tidally influenced blue holes on San Salvador Island, Bahamas. Hydrogeology Journal, 2021, 29, 429-441.	2.1	6
48	Relationship between aquifer biofilms and unattached microbial indicators of urban groundwater contamination. Molecular Ecology, 2021, 30, 324-342.	3.9	6
49	The Biology of Tardigrades: An Introduction to the 9th International Symposium on Tardigrada*. Hydrobiologia, 2006, 558, 1-3.	2.0	4
50	Expansion of the genus Imleria in North America to include Imleria floridana, sp. nov., and Imleria pallida, comb. nov Mycologia, 2020, 112, 423-437.	1.9	2
51	Changes in Eukaryotic and Bacterial Communities along a 120 m Transect Associated with a Shallow Marine Hydrothermal Vent. Frontiers in Marine Science, 2017, 4, .	2.5	1
52	Global diversity of tardigrades (Tardigrada) in freshwater., 2007,, 101-106.		1
53	MICROBIAL TEMPORAL DYNAMICS OF A NOVEL GYPSUM KARST SULFIDIC SPRING. , 2017, , .		1
54	Review paper: The evolutionary relationships of rotifers and acanthocephalans., 1998,, 83-91.		0