

W Ford Doolittle

List of Publications by Year in descending order

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Version: 2024-02-01

103
papers

10,441
citations

53794

45
h-index

33894

99
g-index

108
all docs

108
docs citations

108
times ranked

7986
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 1 | All about levels: transposable elements as selfish DNAs and drivers of evolution. <i>Biology and Philosophy</i> , 2022, 37, . | 1.4 | 2 |
| 2 | “Species”™ without species. <i>Studies in History and Philosophy of Science Part A</i> , 2021, 87, 72-80. | 1.2 | 15 |
| 3 | The role of purifying selection in the origin and maintenance of complex function. <i>Studies in History and Philosophy of Science Part A</i> , 2021, 87, 125-135. | 1.2 | 12 |
| 4 | Life and life only: a radical alternative to life definitionism. <i>Synthese</i> , 2020, 197, 2975-2989. | 1.1 | 28 |
| 5 | Horizontal persistence and the complexity hypothesis. <i>Biology and Philosophy</i> , 2020, 35, 1. | 1.4 | 17 |
| 6 | Getting clear about the F-word in genomics. <i>PLoS Genetics</i> , 2020, 16, e1008702. | 3.5 | 22 |
| 7 | Could this pandemic usher in evolution’s next major transition?. <i>Current Biology</i> , 2020, 30, R846-R848. | 3.9 | 0 |
| 8 | Evolution: Two Domains of Life or Three?. <i>Current Biology</i> , 2020, 30, R177-R179. | 3.9 | 17 |
| 9 | Neutral evolution of cellular phenotypes. <i>Current Opinion in Genetics and Development</i> , 2019, 58-59, 87-94. | 3.3 | 17 |
| 10 | How microbes “jeopardize” the modern synthesis. <i>PLoS Genetics</i> , 2019, 15, e1008166. | 3.5 | 6 |
| 11 | Making Evolutionary Sense of Gaia. <i>Trends in Ecology and Evolution</i> , 2019, 34, 889-894. | 8.7 | 23 |
| 12 | Mutationism, not Lamarckism, captures the novelty of CRISPR-Cas. <i>Biology and Philosophy</i> , 2019, 34, 1. | 1.4 | 5 |
| 13 | Processes and patterns of interaction as units of selection: An introduction to ITSNTS thinking. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, 4006-4014. | 7.1 | 95 |
| 14 | The generality of Constructive Neutral Evolution. <i>Biology and Philosophy</i> , 2018, 33, 1. | 1.4 | 20 |
| 15 | We simply cannot go on being so vague about “function”™. <i>Genome Biology</i> , 2018, 19, 223. | 8.8 | 22 |
| 16 | Darwinizing Gaia. <i>Journal of Theoretical Biology</i> , 2017, 434, 11-19. | 1.7 | 55 |
| 17 | Making the Most of Clade Selection. <i>Philosophy of Science</i> , 2017, 84, 275-295. | 1.0 | 32 |
| 18 | The coupling of taxonomy and function in microbiomes. <i>Biology and Philosophy</i> , 2017, 32, 1225-1243. | 1.4 | 36 |

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|----|------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|-----|-----------|
| 19 | It's the song, not the singer: an exploration of holobiosis and evolutionary theory. <i>Biology and Philosophy</i> , 2017, 32, 5-24. | 1.4 | 155 |
| 20 | On causal roles and selected effects: our genome is mostly junk. <i>BMC Biology</i> , 2017, 15, 116. | 3.8 | 29 |
| 21 | Microbial Evolution: Xenology (Apparently) Trumps Paralogy. <i>Current Biology</i> , 2016, 26, R1181-R1183. | 3.9 | 7 |
| 22 | Molecular Phylogenetics and the Perennial Problem of Homology. <i>Journal of Molecular Evolution</i> , 2016, 83, 184-192. | 1.8 | 29 |
| 23 | The Modern Synthesis in the Light of Microbial Genomics. <i>Annual Review of Microbiology</i> , 2016, 70, 279-297. | 7.3 | 58 |
| 24 | What Is the Tree of Life?. <i>PLoS Genetics</i> , 2016, 12, e1005912. | 3.5 | 35 |
| 25 | Microbial Diversity: A Bonanza of Phyla. <i>Current Biology</i> , 2015, 25, R227-R230. | 3.9 | 18 |
| 26 | Eukaryogenesis, how special really?. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 10278-10285. | 7.1 | 83 |
| 27 | Archaea. <i>Current Biology</i> , 2015, 25, R851-R855. | 3.9 | 45 |
| 28 | Multilevel Selection Theory and the Evolutionary Functions of Transposable Elements: Fig. 1. <i>Genome Biology and Evolution</i> , 2015, 7, 2445-2457. | 2.5 | 36 |
| 29 | Reply to Lane and Martin: Being and becoming eukaryotes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, E4824-E4824. | 7.1 | 10 |
| 30 | Eukaryotes first: how could that be?. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2015, 370, 20140322. | 4.0 | 20 |
| 31 | Getting "function" right. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, E3365. | 7.1 | 22 |
| 32 | Natural selection through survival alone, and the possibility of Gaia. <i>Biology and Philosophy</i> , 2014, 29, 415-423. | 1.4 | 61 |
| 33 | The trouble with (group II) introns. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 6536-6537. | 7.1 | 19 |
| 34 | Distinguishing between "Function" and "Effect" in Genome Biology. <i>Genome Biology and Evolution</i> , 2014, 6, 1234-1237. | 2.5 | 79 |
| 35 | How Natural a Kind Is "Eukaryote?". <i>Cold Spring Harbor Perspectives in Biology</i> , 2014, 6, a015974-a015974. | 5.5 | 3 |
| 36 | <i>Rhodoluna laticola</i> gen. nov., sp. nov., a planktonic freshwater bacterium with stream-lined genome. <i>International Journal of Systematic and Evolutionary Microbiology</i> , 2014, 64, 3254-3263. | 1.7 | 66 |

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|----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 37 | Carl R. Woese (1928–2012). <i>Current Biology</i> , 2013, 23, R183-R185. | 3.9 | 4 |
| 38 | Sixty years of genome biology. <i>Genome Biology</i> , 2013, 14, 113. | 9.6 | 6 |
| 39 | Microbial neopleomorphism. <i>Biology and Philosophy</i> , 2013, 28, 351-378. | 1.4 | 21 |
| 40 | Is junk DNA bunk? A critique of ENCODE. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 5294-5300. | 7.1 | 333 |
| 41 | A ratchet for protein complexity. <i>Nature</i> , 2012, 481, 270-271. | 27.8 | 34 |
| 42 | Population Genomics: How Bacterial Species Form and Why They Don't Exist. <i>Current Biology</i> , 2012, 22, R451-R453. | 3.9 | 36 |
| 43 | Coral-mucus-associated <i>Vibrio</i> integrons in the Great Barrier Reef: genomic hotspots for environmental adaptation. <i>ISME Journal</i> , 2011, 5, 962-972. | 9.8 | 37 |
| 44 | How a neutral evolutionary ratchet can build cellular complexity. <i>IUBMB Life</i> , 2011, 63, 528-537. | 3.4 | 160 |
| 45 | Comment on "Does constructive neutral evolution play an important role in the origin of cellular complexity?" DOI 10.1002/bies.201100010. <i>BioEssays</i> , 2011, 33, 427-429. | 2.5 | 10 |
| 46 | The attempt on the life of the Tree of Life: science, philosophy and politics. <i>Biology and Philosophy</i> , 2010, 25, 455-473. | 1.4 | 22 |
| 47 | Irremediable Complexity?. <i>Science</i> , 2010, 330, 920-921. | 12.6 | 204 |
| 48 | Intertwined Evolutionary Histories of Marine Synechococcus and Prochlorococcus marinus. <i>Genome Biology and Evolution</i> , 2009, 1, 325-339. | 2.5 | 80 |
| 49 | The practice of classification and the theory of evolution, and what the demise of Charles Darwin's tree of life hypothesis means for both of them. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2009, 364, 2221-2228. | 4.0 | 63 |
| 50 | On the chimeric nature, thermophilic origin, and phylogenetic placement of the Thermotogales. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 5865-5870. | 7.1 | 221 |
| 51 | On the origin of prokaryotic species. <i>Genome Research</i> , 2009, 19, 744-756. | 5.5 | 207 |
| 52 | Microbial Evolution: Stalking the Wild Bacterial Species. <i>Current Biology</i> , 2008, 18, R565-R567. | 3.9 | 15 |
| 53 | The Impact of Reticulate Evolution on Genome Phylogeny. <i>Systematic Biology</i> , 2008, 57, 844-856. | 5.6 | 47 |
| 54 | Searching for species in haloarchaea. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 14092-14097. | 7.1 | 128 |

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|----|-----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 55 | Pattern pluralism and the Tree of Life hypothesis. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 2043-2049. | 7.1 | 440 |
| 56 | Systematic overestimation of gene gain through false diagnosis of gene absence. Genome Biology, 2007, 8, 402. | 9.6 | 23 |
| 57 | Evolution: Reducible Complexity – The Case for Bacterial Flagella. Current Biology, 2007, 17, R510-R512. | 3.9 | 15 |
| 58 | Genomics and the bacterial species problem. Genome Biology, 2006, 7, 116. | 9.6 | 200 |
| 59 | Microbial rhodopsins: functional versatility and genetic mobility. Trends in Microbiology, 2006, 14, 463-469. | 7.7 | 193 |
| 60 | Environmental genomics of "Haloquadratum walsbyi" in a saltern crystallizer indicates a large pool of accessory genes in an otherwise coherent species. BMC Genomics, 2006, 7, 171. | 2.8 | 128 |
| 61 | Recombination in Thermotoga: Implications for Species Concepts and Biogeography. Genetics, 2006, 172, 759-769. | 2.9 | 87 |
| 62 | Phylogenetic analyses of cyanobacterial genomes: Quantification of horizontal gene transfer events. Genome Research, 2006, 16, 1099-1108. | 5.5 | 278 |
| 63 | Computing prokaryotic gene ubiquity: Rescuing the core from extinction. Genome Research, 2004, 14, 2469-2477. | 5.5 | 170 |
| 64 | Phylogenetic reconstruction and lateral gene transfer. Trends in Microbiology, 2004, 12, 406-411. | 7.7 | 124 |
| 65 | Diversity of bacteriorhodopsins in different hypersaline waters from a single Spanish saltern. Environmental Microbiology, 2003, 5, 1039-1045. | 3.8 | 29 |
| 66 | Lateral Gene Transfer and the Origins of Prokaryotic Groups. Annual Review of Genetics, 2003, 37, 283-328. | 7.6 | 357 |
| 67 | Some thoughts on the tree of life. Harvey Lectures, 2003, 99, 111-28. | 0.2 | 0 |
| 68 | Prokaryotic Evolution in Light of Gene Transfer. Molecular Biology and Evolution, 2002, 19, 2226-2238. | 8.9 | 858 |
| 69 | Novel syntaxin gene sequences from <i>Giardia</i> , <i>Trypanosoma</i> and algae: implications for the ancient evolution of the eukaryotic endomembrane system. Journal of Cell Science, 2002, 115, 1635-1642. | 2.0 | 64 |
| 70 | Defining the Core of Nontransferable Prokaryotic Genes: The Euryarchaeal Core. Journal of Molecular Evolution, 2001, 53, 340-350. | 1.8 | 74 |
| 71 | Tubulins in <i>Trichomonas vaginalis</i> : Molecular Characterization of alpha-Tubulin Genes, Posttranslational Modifications, and Homology Modeling of the Tubulin Dimer. Journal of Eukaryotic Microbiology, 2001, 48, 647-654. | 1.7 | 6 |
| 72 | GENOMICS: Enhanced: Are There Bugs in Our Genome?. Science, 2001, 292, 1848-1850. | 12.6 | 79 |

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|----|-------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 73 | The role of lateral gene transfer in the evolution of isoprenoid biosynthesis pathways. <i>Molecular Microbiology</i> , 2000, 37, 703-716. | 2.5 | 247 |
| 74 | The nature of the universal ancestor and the evolution of the proteome. <i>Current Opinion in Structural Biology</i> , 2000, 10, 355-358. | 5.7 | 123 |
| 75 | An Updated and Comprehensive rRNA Phylogeny of (Crown) Eukaryotes Based on Rate-Calibrated Evolutionary Distances. <i>Journal of Molecular Evolution</i> , 2000, 51, 565-576. | 1.8 | 136 |
| 76 | Gene Descent, Duplication, and Horizontal Transfer in the Evolution of Glutamyl- and Glutaminyl-tRNA Synthetases. <i>Journal of Molecular Evolution</i> , 1999, 49, 485-495. | 1.8 | 99 |
| 77 | You are what you eat: a gene transfer ratchet could account for bacterial genes in eukaryotic nuclear genomes. <i>Trends in Genetics</i> , 1998, 14, 307-311. | 6.7 | 571 |
| 78 | <i>Sulfolobus islandicus</i> plasmids pRN1 and pRN2 share distant but common evolutionary ancestry. <i>Extremophiles</i> , 1998, 2, 391-393. | 2.3 | 44 |
| 79 | Evidence for the early divergence of tryptophanyl- and tyrosyl-tRNA synthetases. <i>Journal of Molecular Evolution</i> , 1997, 45, 9-16. | 1.8 | 67 |
| 80 | Archaeobacterial genomics: The complete genome sequence of <i>Methanococcus jannaschii</i> . <i>BioEssays</i> , 1997, 19, 1-4. | 2.5 | 17 |
| 81 | The <i>Sulfolobus solfataricus</i> P2 genome project. <i>FEBS Letters</i> , 1996, 389, 88-91. | 2.8 | 21 |
| 82 | Linked Genes for Calmodulin and E2 Ubiquitin-Conjugating Enzyme in <i>Trichomonas vaginalis</i> . <i>Journal of Eukaryotic Microbiology</i> , 1996, 43, 468-474. | 1.7 | 8 |
| 83 | Evidence for the Heterolobosea from Phylogenetic Analysis of Genes Encoding Glycerinaldehyde-3-Phosphate Dehydrogenase. <i>Journal of Eukaryotic Microbiology</i> , 1996, 43, 475-485. | 1.7 | 53 |
| 84 | Why introns-in-pieces?. <i>Nature</i> , 1993, 364, 289-290. | 27.8 | 40 |
| 85 | Genes-in-pieces revisited. <i>Nature</i> , 1993, 361, 403-403. | 27.8 | 26 |
| 86 | Sol's world, the RNA world, our world.. <i>FASEB Journal</i> , 1993, 7, 1-2. | 0.5 | 7 |
| 87 | TRANSFORMATION OF A BOP-HOP-SOP-I-SOP-II-Halobacterium halobium MUTANT TO BOP+: EFFECTS OF BACTERIORHODOPSIN PHOTOACTIVATION ON CELLULAR PROTON FLUXES AND SWIMMING BEHAVIOR. <i>Photochemistry and Photobiology</i> , 1992, 56, 553-561. | 2.5 | 16 |
| 88 | Biochemical Evolution and the History of Life. <i>Short Courses in Paleontology</i> , 1988, 1, 138-145. | 0.2 | 0 |
| 89 | The Evolutionary Significance of the Archaeobacteria. <i>Annals of the New York Academy of Sciences</i> , 1987, 503, 72-77. | 3.8 | 3 |
| 90 | Systematics of basidiomycetes based on 5S rRNA sequences and other data (reply). <i>Nature</i> , 1983, 303, 732-732. | 27.8 | 3 |

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|-----|----------------------------------------------------------------------------------------------------------------------------------------------------------------------------------|------|-----------|
| 91 | Unusual physical organization of the Halobacterium genome. Nature, 1982, 295, 384-389. | 27.8 | 128 |
| 92 | High-frequency genomic rearrangements involving archaeobacterial repeat sequence elements. Nature, 1982, 299, 182-185. | 27.8 | 103 |
| 93 | Redividing the basidiomycetes on the basis of 5S rRNA sequences. Nature, 1982, 299, 723-724. | 27.8 | 101 |
| 94 | Selfish genes, the phenotype paradigm and genome evolution. Nature, 1980, 284, 601-603. | 27.8 | 1,738 |
| 95 | Nucleotide Sequences of Wheatâ€™Embryo Cytosol 5â€™ and 5.8â€™ Ribosomal Ribonucleic Acids. FEBS Journal, 1980, 112, 561-576. | 0.2 | 83 |
| 96 | Genes in pieces: were they ever together?. Nature, 1978, 272, 581-582. | 27.8 | 453 |
| 97 | Transmission of Creutzfeldt-Jakob disease from man to the guinea pig. Science, 1975, 190, 571-572. | 12.6 | 193 |
| 98 | Novel Ribonucleic Acid Species Accumulated in the Dark in the Blue-Green Alga Anacystis nidulans. Journal of Bacteriology, 1974, 118, 351-357. | 2.2 | 12 |
| 99 | Mutational Analysis of Dark Endogenous Metabolism in the Blue-Green Bacterium <i>Anacystis nidulans</i> . Journal of Bacteriology, 1974, 119, 677-683. | 2.2 | 50 |
| 100 | Postmaturational Cleavage of 23 s Ribosomal Ribonucleic Acid and Its Metabolic Control in the Blue-Green Alga Anacystis nidulans. Journal of Bacteriology, 1973, 113, 1256-1263. | 2.2 | 63 |
| 101 | Ribosomal Ribonucleic Acid Synthesis and Maturation in the Blue-Green Alga Anacystis nidulans. Journal of Bacteriology, 1972, 111, 316-324. | 2.2 | 41 |
| 102 | The Root of the Tree: Lateral Gene Transfer and the Nature of the Domains. , 0, , 29-37. | | 1 |
| 103 | A Chemostat Model for Evolution by Persistence: Clade Selection and its Explanatory Autonomy. Philosophy of Science, 0, , 1-47. | 1.0 | 0 |