

# John M Archibald

## List of Publications by Year in descending order

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107  
papers

7,434  
citations

66343

42  
h-index

60623

81  
g-index

112  
all docs

112  
docs citations

112  
times ranked

6758  
citing authors

| #  | ARTICLE   | IF   | CITATIONS |
|----|---|------|-----------|
| 1  | The Earth BioGenome Project 2020: Starting the clock. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .   | 7.1  | 124       |
| 2  | Standards recommendations for the Earth BioGenome Project. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .  | 7.1  | 33        |
| 3  | Why sequence all eukaryotes?. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .   | 7.1  | 51        |
| 4  | TreeTuner: A pipeline for minimizing redundancy and complexity in large phylogenetic datasets. <i>STAR Protocols</i> , 2022, 3, 101175.   | 1.2  | 0         |
| 5  | Submergence of the filamentous Zygnematophyceae <i>Mougeotia</i> induces differential gene expression patterns associated with core metabolism and photosynthesis. <i>Protoplasma</i> , 2022, 259, 1157-1174. | 2.1  | 12        |
| 6  | Evolutionary Dynamics and Lateral Gene Transfer in Raphidophyceae Plastid Genomes. <i>Frontiers in Plant Science</i> , 2022, 13, .  | 3.6  | 3         |
| 7  | Mitochondrial Genome Evolution in Pelagophyte Algae. <i>Genome Biology and Evolution</i> , 2021, 13, .  | 2.5  | 10        |
| 8  | The past, present and future of the tree of life. <i>Current Biology</i> , 2021, 31, R314-R321.   | 3.9  | 18        |
| 9  | Re-examination of two diatom reference genomes using long-read sequencing. <i>BMC Genomics</i> , 2021, 22, 379.   | 2.8  | 22        |
| 10 | RNA-Seq analysis reveals potential regulators of programmed cell death and leaf remodelling in lace plant ( <i>Aponogeton madagascariensis</i> ). <i>BMC Plant Biology</i> , 2021, 21, 375.                   | 3.6  | 5         |
| 11 | Genomic analysis finds no evidence of canonical eukaryotic DNA processing complexes in a free-living protist. <i>Nature Communications</i> , 2021, 12, 6003.  | 12.8 | 17        |
| 12 | Cryptomonads. <i>Current Biology</i> , 2020, 30, R1114-R1116.   | 3.9  | 4         |
| 13 | Comparative Plastid Genomics of Non-Photosynthetic Chrysophytes: Genome Reduction and Compaction. <i>Frontiers in Plant Science</i> , 2020, 11, 572703.   | 3.6  | 8         |
| 14 | Comparative analyses of saprotrophy in <i>Salisapilia sapeloensis</i> and diverse plant pathogenic oomycetes reveal lifestyle-specific gene expression. <i>FEMS Microbiology Ecology</i> , 2020, 96, .        | 2.7  | 4         |
| 15 | Lateral Gene Transfer Mechanisms and Pan-genomes in Eukaryotes. <i>Trends in Parasitology</i> , 2020, 36, 927-941.  | 3.3  | 41        |
| 16 | Genomic Insights into Plastid Evolution. <i>Genome Biology and Evolution</i> , 2020, 12, 978-990.   | 2.5  | 79        |
| 17 | Phagocytosis in a Shape-shifting Bacterium. <i>Trends in Microbiology</i> , 2020, 28, 428-430.  | 7.7  | 0         |
| 18 | Comparative Plastid Genomics of <i>Cryptomonas</i> Species Reveals Fine-Scale Genomic Responses to Loss of Photosynthesis. <i>Genome Biology and Evolution</i> , 2020, 12, 3926-3937.                         | 2.5  | 27        |

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|----|--|-----|-----------|
| 19 | Heat stress response in the closest algal relatives of land plants reveals conserved stress signaling circuits. <i>Plant Journal</i> , 2020, 103, 1025-1048.                                   | 5.7 | 65        |
| 20 | Evolutionary Biology: Viral Rhodopsins Illuminate Algal Evolution. <i>Current Biology</i> , 2020, 30, R1469-R1471.   | 3.9 | 4         |
| 21 | Genomics reveals alga-associated cyanobacteria hiding in plain sight. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 15757-15759.         | 7.1 | 2         |
| 22 | Evolution: New Protist Predators under the Sun. <i>Current Biology</i> , 2019, 29, R936-R938.  | 3.9 | 2         |
| 23 | Ubiquitin fusion proteins in algae: implications for cell biology and the spread of photosynthesis. <i>BMC Genomics</i> , 2019, 20, 38.  | 2.8 | 9         |
| 24 | Comparative plastid genomics of Synurophyceae: inverted repeat dynamics and gene content variation. <i>BMC Evolutionary Biology</i> , 2019, 19, 20.  | 3.2 | 27        |
| 25 | Relative Mutation Rates in Nucleomorph-Bearing Algae. <i>Genome Biology and Evolution</i> , 2019, 11, 1045-1053.   | 2.5 | 8         |
| 26 | Nucleomorph Small RNAs in Cryptophyte and Chlorarachniophyte Algae. <i>Genome Biology and Evolution</i> , 2019, 11, 1117-1134.   | 2.5 | 1         |
| 27 | Symbiosis in the microbial world: from ecology to genome evolution. <i>Biology Open</i> , 2018, 7, .   | 1.2 | 34        |
| 28 | 10KP: A phylodiverse genome sequencing plan. <i>GigaScience</i> , 2018, 7, 1-9.  | 6.4 | 169       |
| 29 | Opportunistic but Lethal: The Mystery of Paramoebae. <i>Trends in Parasitology</i> , 2018, 34, 404-419.  | 3.3 | 41        |
| 30 | Plant evolution: landmarks on the path to terrestrial life. <i>New Phytologist</i> , 2018, 217, 1428-1434.   | 7.3 | 236       |
| 31 | Plastid genomes. <i>Current Biology</i> , 2018, 28, R336-R337.   | 3.9 | 22        |
| 32 | Embryophyte stress signaling evolved in the algal progenitors of land plants. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E3471-E3480. | 7.1 | 164       |
| 33 | Nuclear genome sequence of the plastid-lacking cryptomonad <i>Goniomonas avonlea</i> provides insights into the evolution of secondary plastids. <i>BMC Biology</i> , 2018, 16, 137.           | 3.8 | 42        |
| 34 | Massive mitochondrial DNA content in diplomonid and kinetoplastid protists. <i>IUBMB Life</i> , 2018, 70, 1267-1274.   | 3.4 | 39        |
| 35 | On plant defense signaling networks and early land plant evolution. <i>Communicative and Integrative Biology</i> , 2018, 11, 1-14.   | 1.4 | 54        |
| 36 | Comparative mitochondrial genomics of cryptophyte algae: gene shuffling and dynamic mobile genetic elements. <i>BMC Genomics</i> , 2018, 19, 275.  | 2.8 | 23        |

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|----|---|------|-----------|
| 37 | Lateral Gene Transfer in the Adaptation of the Anaerobic Parasite <i>Blastocystis</i> to the Gut. <i>Current Biology</i> , 2017, 27, 807-820.                             | 3.9  | 94        |
| 38 | Diversity and Evolution of <i>Paramoeba</i> spp. and their Kinetoplastid Endosymbionts. <i>Journal of Eukaryotic Microbiology</i> , 2017, 64, 598-607.                    | 1.7  | 14        |
| 39 | Endosymbiosis: Did Plastids Evolve from a Freshwater Cyanobacterium?. <i>Current Biology</i> , 2017, 27, R103-R105.   | 3.9  | 56        |
| 40 | More protist genomes needed. <i>Nature Ecology and Evolution</i> , 2017, 1, 145.  | 7.8  | 78        |
| 41 | The New Red Algal Subphylum Proteorhodophytina Comprises the Largest and Most Divergent Plastid Genomes Known. <i>Current Biology</i> , 2017, 27, 1677-1684.e4.           | 3.9  | 89        |
| 42 | A Non-photosynthetic Diatom Reveals Early Steps of Reductive Evolution in Plastids. <i>Molecular Biology and Evolution</i> , 2017, 34, 2355-2366.                         | 8.9  | 52        |
| 43 | How Embryophytic is the Biosynthesis of Phenylpropanoids and their Derivatives in Streptophyte Algae?. <i>Plant and Cell Physiology</i> , 2017, 58, 934-945.              | 3.1  | 102       |
| 44 | Evolution: Protein Import in a Nascent Photosynthetic Organelle. <i>Current Biology</i> , 2017, 27, R1004-R1006.  | 3.9  | 2         |
| 45 | Genome sequencing reveals metabolic and cellular interdependence in an amoeba-kinetoplastid symbiosis. <i>Scientific Reports</i> , 2017, 7, 11688.                        | 3.3  | 44        |
| 46 | Evolutionary Dynamics of Cryptophyte Plastid Genomes. <i>Genome Biology and Evolution</i> , 2017, 9, 1859-1872.   | 2.5  | 51        |
| 47 | Probing the evolution, ecology and physiology of marine protists using transcriptomics. <i>Nature Reviews Microbiology</i> , 2017, 15, 6-20.                              | 28.6 | 176       |
| 48 | The Carboxy Terminus of YCF1 Contains a Motif Conserved throughout >500 Myr of Streptophyte Evolution. <i>Genome Biology and Evolution</i> , 2017, 9, 473-479.            | 2.5  | 14        |
| 49 | Extreme genome diversity in the hyper-prevalent parasitic eukaryote <i>Blastocystis</i> . <i>PLoS Biology</i> , 2017, 15, e2003769.                                       | 5.6  | 99        |
| 50 | Heme pathway evolution in kinetoplastid protists. <i>BMC Evolutionary Biology</i> , 2016, 16, 109.  | 3.2  | 19        |
| 51 | Comparative genomics of mitochondria in chlorarachniophyte algae: endosymbiotic gene transfer and organellar genome dynamics. <i>Scientific Reports</i> , 2016, 6, 21016. | 3.3  | 23        |
| 52 | Evolution: Plumbing the Depths of Diplonemid Diversity. <i>Current Biology</i> , 2016, 26, R1290-R1292.   | 3.9  | 11        |
| 53 | Streptophyte Terrestrialization in Light of Plastid Evolution. <i>Trends in Plant Science</i> , 2016, 21, 467-476.  | 8.8  | 136       |
| 54 | Gene Loss and Error-Prone RNA Editing in the Mitochondrion of <i>Perkinsela</i> , an Endosymbiotic Kinetoplastid. <i>MBio</i> , 2015, 6, e01498-15.                       | 4.1  | 28        |

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|----|--|------|-----------|
| 55 | Genomic perspectives on the birth and spread of plastids. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, 10147-10153.   | 7.1  | 121       |
| 56 | Localization and Evolution of Putative Triose Phosphate Translocators in the Diatom <i>Phaeodactylum tricornutum</i> . Genome Biology and Evolution, 2015, 7, 2955-2969.   | 2.5  | 53        |
| 57 | Endosymbiosis and Eukaryotic Cell Evolution. Current Biology, 2015, 25, R911-R921.   | 3.9  | 426       |
| 58 | Gene transfer in complex cells. Nature, 2015, 524, 423-424.  | 27.8 | 9         |
| 59 | Dual Organellar Targeting of Aminoacyl-tRNA Synthetases in Diatoms and Cryptophytes. Genome Biology and Evolution, 2015, 7, 1728-1742.   | 2.5  | 46        |
| 60 | Reduced Nuclear Genomes Maintain High Gene Transcription Levels. Molecular Biology and Evolution, 2014, 31, 625-635.   | 8.9  | 20        |
| 61 | Overexpression of Molecular Chaperone Genes in Nucleomorph Genomes. Molecular Biology and Evolution, 2014, 31, 1437-1443.  | 8.9  | 12        |
| 62 | Alternatives to vitamin B1 uptake revealed with discovery of riboswitches in multiple marine eukaryotic lineages. ISME Journal, 2014, 8, 2517-2529.  | 9.8  | 69        |
| 63 | The Marine Microbial Eukaryote Transcriptome Sequencing Project (MMETSP): Illuminating the Functional Diversity of Eukaryotic Life in the Oceans through Transcriptome Sequencing. PLoS Biology, 2014, 12, e1001889.                 | 5.6  | 885       |
| 64 | Complete genome of a nonphotosynthetic cyanobacterium in a diatom reveals recent adaptations to an intracellular lifestyle. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, 11407-11412. | 7.1  | 121       |
| 65 | Nucleomorph and plastid genome sequences of the chlorarachniophyte <i>Lotharella oceanica</i> : convergent reductive evolution and frequent recombination in nucleomorph-bearing algae. BMC Genomics, 2014, 15, 374.                 | 2.8  | 32        |
| 66 | Nucleomorph Comparative Genomics. , 2014, , 197-213.   |      | 8         |
| 67 | Ultrastructure and Molecular Phylogeny of the Cryptomonad <i>Goniomonas avonlea</i> sp. nov.. Protist, 2013, 164, 160-182.   | 1.5  | 33        |
| 68 | Treertrimmer: a method for phylogenetic dataset size reduction. BMC Research Notes, 2013, 6, 145.  | 1.4  | 25        |
| 69 | Algal genomes reveal evolutionary mosaicism and the fate of nucleomorphs. Nature, 2012, 492, 59-65.  | 27.8 | 377       |
| 70 | Nucleomorph Genome Sequence of the Cryptophyte Alga <i>Chroomonas mesostigmatica</i> CCMP1168 Reveals Lineage-Specific Gene Loss and Genome Complexity. Genome Biology and Evolution, 2012, 4, 1162-1175.                            | 2.5  | 50        |
| 71 | Complete Nucleomorph Genome Sequence of the Nonphotosynthetic Alga <i>Cryptomonas paramecium</i> Reveals a Core Nucleomorph Gene Set. Genome Biology and Evolution, 2011, 3, 44-54.  | 2.5  | 62        |
| 72 | Origin of eukaryotic cells: 40 years on. Symbiosis, 2011, 54, 69-86.   | 2.3  | 32        |

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|----|---|------|-----------|
| 73 | Eukaryote-to-eukaryote gene transfer gives rise to genome mosaicism in euglenids. <i>BMC Evolutionary Biology</i> , 2011, 11, 105.  | 3.2  | 53        |
| 74 | Genomic Characterization of <i>Neoparamoeba pemaquidensis</i> (Amoebozoa) and Its Kinetoplastid Endosymbiont. <i>Eukaryotic Cell</i> , 2011, 10, 1143-1146.   | 3.4  | 20        |
| 75 | Gene transfer: anything goes in plant mitochondria. <i>BMC Biology</i> , 2010, 8, 147.  | 3.8  | 32        |
| 76 | Large-Scale Phylogenomic Analyses Reveal That Two Enigmatic Protist Lineages, <i>Telonemia</i> and <i>Centroheliozoa</i> , Are Related to Photosynthetic Chromalveolates. <i>Genome Biology and Evolution</i> , 2009, 1, 231-238. | 2.5  | 143       |
| 77 | The Complete Plastid Genome Sequence of the Secondarily Nonphotosynthetic Alga <i>Cryptomonas paramecium</i> : Reduction, Compaction, and Accelerated Evolutionary Rate. <i>Genome Biology and Evolution</i> , 2009, 1, 439-448.  | 2.5  | 70        |
| 78 | Going, Going, Not Quite Gone: Nucleomorphs as a Case Study in Nuclear Genome Reduction. <i>Journal of Heredity</i> , 2009, 100, 582-590.  | 2.4  | 38        |
| 79 | The Puzzle of Plastid Evolution. <i>Current Biology</i> , 2009, 19, R81-R88.  | 3.9  | 413       |
| 80 | Green Evolution, Green Revolution. <i>Science</i> , 2009, 324, 191-192.   | 12.6 | 11        |
| 81 | Nucleomorph Genomes. <i>Annual Review of Genetics</i> , 2009, 43, 251-264.  | 7.6  | 80        |
| 82 | <i>Lotharella oceanica</i> sp. nov. – a new planktonic chlorarachniophyte studied by light and electron microscopy. <i>Phycologia</i> , 2009, 48, 315-323.  | 1.4  | 19        |
| 83 | The origin and spread of eukaryotic photosynthesis: evolving views in light of genomics. <i>Botanica Marina</i> , 2009, 52, 95-103.   | 1.2  | 8         |
| 84 | NUCLEOMORPH KARYOTYPE DIVERSITY IN THE FRESHWATER CRYPTOPHYTE GENUS <i>CRYPTOMONAS</i> <sup>1</sup> . <i>Journal of Phycology</i> , 2008, 44, 11-14.  | 2.3  | 15        |
| 85 | NEW MARINE MEMBERS OF THE GENUS <i>HEMISELMIS</i> (CRYPTOMONADALES.) Tj ETQq1 1 0.784314 rgBT /Overlock 10 Tf 50 2 2.3 37   |      |           |
| 86 | Complete Sequence and Analysis of the Mitochondrial Genome of <i>Hemiselms andersenii</i> CCMP644 (Cryptophyceae). <i>BMC Genomics</i> , 2008, 9, 215.  | 2.8  | 49        |
| 87 | Plastid Evolution: Remnant Algal Genes in Ciliates. <i>Current Biology</i> , 2008, 18, R663-R665.   | 3.9  | 18        |
| 88 | The eukaryotic tree of life: endosymbiosis takes its TOL. <i>Trends in Ecology and Evolution</i> , 2008, 23, 268-275.   | 8.7  | 267       |
| 89 | The eocyte hypothesis and the origin of eukaryotic cells. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 20049-20050.  | 7.1  | 21        |
| 90 | Lateral transfer of introns in the cryptophyte plastid genome. <i>Nucleic Acids Research</i> , 2008, 36, 3043-3053.   | 14.5 | 34        |

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|-----|--|-----|-----------|
| 91  | Nucleomorph genome of <i>Hemiselmis andersenii</i> reveals complete intron loss and compaction as a driver of protein structure and function. Proceedings of the National Academy of Sciences of the United States of America, 2007, 104, 19908-19913. | 7.1 | 139       |
| 92  | Nucleomorph genomes: structure, function, origin and evolution. BioEssays, 2007, 29, 392-402.  | 2.5 | 103       |
| 93  | Plastid Genome Sequence of the Cryptophyte Alga <i>Rhodomonas salina</i> CCMP1319: Lateral Transfer of Putative DNA Replication Machinery and a Test of Chromist Plastid Phylogeny. Molecular Biology and Evolution, 2007, 24, 1832-1842.              | 8.9 | 100       |
| 94  | Endosymbiosis: Double-Take on Plastid Origins. Current Biology, 2006, 16, R690-R692.   | 3.9 | 24        |
| 95  | Algal Genomics: Exploring the Imprint of Endosymbiosis. Current Biology, 2006, 16, R1033-R1035.  | 3.9 | 14        |
| 96  | Insight into the Diversity and Evolution of the Cryptomonad Nucleomorph Genome. Molecular Biology and Evolution, 2006, 23, 856-865.  | 8.9 | 42        |
| 97  | Genome complexity in a lean, mean photosynthetic machine. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 11433-11434.   | 7.1 | 7         |
| 98  | Jumping Genes and Shrinking Genomes – Probing the Evolution of Eukaryotic Photosynthesis with Genomics. IUBMB Life, 2005, 57, 539-547.   | 3.4 | 45        |
| 99  | Phagotrophy in chlorarachniophyte algae: implications for eukaryotic genome evolution. Journal of Eukaryotic Microbiology, 2005, 52, 7S-27S.   | 1.7 | 0         |
| 100 | Actin and Ubiquitin Protein Sequences Support a Cercozoan/Foraminiferan Ancestry for the Plasmodiophorid Plant Pathogens. Journal of Eukaryotic Microbiology, 2004, 51, 113-118.   | 1.7 | 62        |
| 101 | Novel Ubiquitin Fusion Proteins: Ribosomal Protein P1 and Actin. Journal of Molecular Biology, 2003, 328, 771-778.   | 4.2 | 28        |
| 102 | A Novel Polyubiquitin Structure in Cercozoa and Foraminifera: Evidence for a New Eukaryotic Supergroup. Molecular Biology and Evolution, 2003, 20, 62-66.  | 8.9 | 87        |
| 103 | Lateral gene transfer and the evolution of plastid-targeted proteins in the secondary plastid-containing alga <i>Bigeloviella natans</i> . Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 7678-7683.      | 7.1 | 241       |
| 104 | The Chaperonin Genes of Jakobid and Jakobid-Like Flagellates: Implications for Eukaryotic Evolution. Molecular Biology and Evolution, 2002, 19, 422-431.   | 8.9 | 59        |
| 105 | Recycled plastids: a “green movement” in eukaryotic evolution. Trends in Genetics, 2002, 18, 577-584.  | 6.7 | 212       |
| 106 | Gene Conversion and the Evolution of Euryarchaeal Chaperonins: A Maximum Likelihood-Based Method for Detecting Conflicting Phylogenetic Signals. Journal of Molecular Evolution, 2002, 55, 232-245.  | 1.8 | 30        |
| 107 | Molecular Chaperones Encoded by a Reduced Nucleus: The Cryptomonad Nucleomorph. Journal of Molecular Evolution, 2001, 52, 490-501.   | 1.8 | 27        |