## Pierre Cardol

## List of Publications by Year in descending order

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83 6,403 33
papers citations h-index

74163 75 g-index

90 all docs

90 docs citations 90 times ranked 6454 citing authors

#	Article	IF	CITATIONS
1	Beyond being an energy supplier, ATP synthase is a sculptor of mitochondrial cristae. Biochimica Et Biophysica Acta - Bioenergetics, 2022, 1863, 148569.	1.0	4
2	Automated Open-Hardware Multiwell Imaging Station for Microorganisms Observation. Micromachines, 2022, 13, 833.	2.9	3
3	Supramolecular associations between atypical oxidative phosphorylation complexes of Euglena gracilis. Journal of Bioenergetics and Biomembranes, 2021, 53, 351-363.	2.3	2
4	Open-hardware wireless controller and 3D-printed pumps for efficient liquid manipulation. HardwareX, 2021, 9, e00199.	2.2	13
5	De Novo Transcriptome Meta-Assembly of the Mixotrophic Freshwater Microalga Euglena gracilis. Genes, 2021, 12, 842.	2.4	9
6	Long-term acclimation to cadmium exposure reveals extensive phenotypic plasticity in Chlamydomonas. Plant Physiology, 2021, 187, 1653-1678.	4.8	7
7	Response of dimethylsulfoniopropionate (DMSP) and dimethylsulfoxide (DMSO) cell quotas to oxidative stress in three phytoplankton species. Journal of Plankton Research, 2021, 43, 673-690.	1.8	2
8	Trophic state alters the mechanism whereby energetic coupling between photosynthesis and respiration occurs in <i>Euglena gracilis</i> New Phytologist, 2021, 232, 1603-1617.	7.3	11
9	In vivo assessment of mitochondrial respiratory alternative oxidase activity and cyclic electron flow around photosystem I on small coral fragments. Scientific Reports, 2020, 10, 17514.	3.3	9
10	Different levels of energetic coupling between photosynthesis and respiration do not determine the occurrence of adaptive responses of Symbiodiniaceae to global warming. New Phytologist, 2020, 228, 855-868.	7.3	12
11	Photosynthetic capacity of the endosymbiotic dinoflagellate Cladocopium sp. is preserved during digestion of its jellyfish host Mastigias papua by the anemone Entacmaea medusivora. FEMS Microbiology Ecology, 2019, 95, .	2.7	8
12	Alternative Photosynthetic Electron Transfers and Bleaching Phenotypes Upon Acute Heat Stress in Symbiodinium and Breviolum spp. (Symbiodiniaceae) in Culture. Frontiers in Marine Science, 2019, 6, .	2.5	17
13	The mechanism of cyclic electron flow. Biochimica Et Biophysica Acta - Bioenergetics, 2019, 1860, 433-438.	1.0	90
14	Maximal cyclic electron flow rate is independent of PGRL1 in Chlamydomonas. Biochimica Et Biophysica Acta - Bioenergetics, 2019, 1860, 425-432.	1.0	50
15	The peculiar NPQ regulation in the stramenopile Phaeomonas sp. challenges the xanthophyll cycle dogma. Biochimica Et Biophysica Acta - Bioenergetics, 2018, 1859, 491-500.	1.0	16
16	Oxidative phosphorylation supercomplexes and respirasome reconstitution of the colorless alga Polytomella sp Biochimica Et Biophysica Acta - Bioenergetics, 2018, 1859, 434-444.	1.0	13
17	Host-pathogen biotic interactions shaped vitamin K metabolism in Archaeplastida. Scientific Reports, 2018, 8, 15243.	3.3	14
18	The atypical subunit composition of respiratory complexes I and IV is associated with original extra structural domains in Euglena gracilis. Scientific Reports, 2018, 8, 9698.	3.3	12

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19	Atypical composition and structure of the mitochondrial dimeric ATP synthase from Euglena gracilis. Biochimica Et Biophysica Acta - Bioenergetics, 2017, 1858, 267-275.	1.0	21
20	Near-neighbor interactions of the membrane-embedded subunits of the mitochondrial ATP synthase of a chlorophycean alga. Biochimica Et Biophysica Acta - Bioenergetics, 2017, 1858, 497-509.	1.0	5
21	From light capture to metabolic needs, oxygenic photosynthesis is an everâ€expanding field of study in plants, algae and cyanobacteria. Physiologia Plantarum, 2017, 161, 2-5.	5.2	2
22	Near-Neighbor Relationships of the Atypical Subunits that Form the Peripheral Stalk of the Mitochondrial ATP Synthase in Chlorophycean Algae. Biophysical Journal, 2017, 112, 2a-3a.	0.5	0
23	<i>In vivo</i> chlorophyll fluorescence screening allows the isolation of a <i>Chlamydomonas</i> mutant defective for <scp>NDUFAF</scp> 3, an assembly factor involved in mitochondrial complex I assembly. Plant Journal, 2017, 92, 584-595.	5.7	11
24	Isolation and characterization of mutants corresponding to the <scp>MENA</scp> , <scp> MENB</scp> , <scp> MENC</scp> and <scp>MENE</scp> enzymatic steps of 5′â€monohydroxyphylloquinone biosynthesis in <i>Chlamydomonas reinhardtii</i> i>. Plant Journal, 2017, 89, 141-154.	5.7	19
25	Mitochondrial Bioenergetics Pathways in Chlamydomonas. Microbiology Monographs, 2017, , 59-95.	0.6	2
26	Carbon Supply and Photoacclimation Cross Talk in the Green Alga <i>Chlamydomonas reinhardtii</i> Plant Physiology, 2016, 172, 1494-1505.	4.8	65
27	Subunit Asa1 spans all the peripheral stalk of the mitochondrial ATP synthase of the chlorophycean alga Polytomella sp Biochimica Et Biophysica Acta - Bioenergetics, 2016, 1857, 359-369.	1.0	10
28	Dissecting the peripheral stalk of the mitochondrial ATP synthase of chlorophycean algae. Biochimica Et Biophysica Acta - Bioenergetics, 2016, 1857, 1183-1190.	1.0	18
29	Isolation of Chlamydomonas reinhardtii mutants with altered mitochondrial respiration by chlorophyll fluorescence measurement. Journal of Biotechnology, 2015, 215, 27-34.	3.8	17
30	Imbalance between oxygen photoreduction and antioxidant capacities in Symbiodinium cells exposed to combined heat and high light stress. Coral Reefs, 2015, 34, 1063-1073.	2.2	57
31	Energetic coupling between plastids and mitochondria drives CO2 assimilation in diatoms. Nature, 2015, 524, 366-369.	27.8	311
32	Induction of Photosynthetic Carbon Fixation in Anoxia Relies on Hydrogenase Activity and Proton-Gradient Regulation-Like1-Mediated Cyclic Electron Flow in <i>Chlamydomonas reinhardtii</i> Plant Physiology, 2015, 168, 648-658.	4.8	59
33	The Involvement of Hydrogen-producing and ATP-dependent NADPH-consuming Pathways in Setting the Redox Poise in the Chloroplast of Chlamydomonas reinhardtii in Anoxia. Journal of Biological Chemistry, 2015, 290, 8666-8676.	3.4	27
34	Regulation of Electron Transport in Photosynthesis. , 2014, , 437-464.		7
35	Respiratory-deficient mutants of the unicellular green alga Chlamydomonas: A review. Biochimie, 2014, 100, 207-218.	2.6	31
36	Inactivation of genes coding for mitochondrial Nd7 and Nd9 complex I subunits in Chlamydomonas reinhardtii. Impact of complex I loss on respiration and energetic metabolism. Mitochondrion, 2014, 19, 365-374.	3.4	10

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37	<scp>PSI</scp> Mehler reaction is the main alternative photosynthetic electron pathway in <i>Symbiodinium</i> sp., symbiotic dinoflagellates of cnidarians. New Phytologist, 2014, 204, 81-91.	7.3	131
38	Lack of isocitrate lyase in <i><scp>C</scp>hlamydomonas</i> leads to changes in carbon metabolism and in the response to oxidative stress under mixotrophic growth. Plant Journal, 2014, 77, 404-417.	5.7	73
39	Interactions of subunits Asa2, Asa4 and Asa7 in the peripheral stalk of the mitochondrial ATP synthase of the chlorophycean alga Polytomella sp Biochimica Et Biophysica Acta - Bioenergetics, 2014, 1837, 1-13.	1.0	14
40	The mitochondrial respiratory chain of the secondary green alga Euglena gracilis shares many additional subunits with parasitic Trypanosomatidae. Mitochondrion, 2014, 19, 338-349.	3.4	59
41	A Dual Strategy to Cope with High Light in <i>Chlamydomonas reinhardtii</i> Â. Plant Cell, 2013, 25, 545-557.	6.6	193
42	A novel screening method for hydrogenase-deficient mutants in Chlamydomonas reinhardtii based on inÂvivo chlorophyll fluorescence and photosystem II quantum yield. International Journal of Hydrogen Energy, 2013, 38, 1826-1836.	7.1	16
43	Function of the Chloroplast Hydrogenase in the Microalga Chlamydomonas: The Role of Hydrogenase and State Transitions during Photosynthetic Activation in Anaerobiosis. PLoS ONE, 2013, 8, e64161.	2.5	47
44	Complexes I in the Green Lineage. , 2012, , 219-244.		0
45	Characterization of an internal type-II NADH dehydrogenase from Chlamydomonas reinhardtii mitochondria. Current Genetics, 2012, 58, 205-216.	1.7	19
46	Functional analysis of hydrogen photoproduction in respiratory-deficient mutants of Chlamydomonas reinhardtii. International Journal of Hydrogen Energy, 2011, 36, 9562-9570.	7.1	12
47	Regulation of electron transport in microalgae. Biochimica Et Biophysica Acta - Bioenergetics, 2011, 1807, 912-918.	1.0	129
48	Mitochondrial NADH:ubiquinone oxidoreductase (complex I) in eukaryotes: A highly conserved subunit composition highlighted by mining of protein databases. Biochimica Et Biophysica Acta - Bioenergetics, 2011, 1807, 1390-1397.	1.0	78
49	Knock-down of the COX3 and COX17 gene expression of cytochrome c oxidase in the unicellular green alga Chlamydomonas reinhardtii. Plant Molecular Biology, 2010, 74, 223-233.	3.9	29
50	Electrochromism: a useful probe to study algal photosynthesis. Photosynthesis Research, 2010, 106, 179-189.	2.9	184
51	Eukaryotic algae: where lies the diversity of oxygenic photosynthesis. Photosynthesis Research, 2010, 106, 1-2.	2.9	2
52	The onset of NPQ and $\hat{l}^{"}\hat{l}^{1}/4H+$ upon illumination of tobacco plants studied through the influence of mitochondrial electron transport. Biochimica Et Biophysica Acta - Bioenergetics, 2010, 1797, 177-188.	1.0	34
53	Subunit–subunit interactions and overall topology of the dimeric mitochondrial ATP synthase of Polytomella sp Biochimica Et Biophysica Acta - Bioenergetics, 2010, 1797, 1439-1448.	1.0	26
54	Loss of mitochondrial ATP synthase subunit beta (Atp2) alters mitochondrial and chloroplastic function and morphology in Chlamydomonas. Biochimica Et Biophysica Acta - Bioenergetics, 2010, 1797, 1533-1539.	1.0	37

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55	An atypical member of the light-harvesting complex stress-related protein family modulates diatom responses to light. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 18214-18219.	7.1	258
56	Atypical Subunit Composition of the Chlorophycean Mitochondrial F1FO-ATP Synthase and Role of Asa7 Protein in Stability and Oligomycin Resistance of the Enzyme. Molecular Biology and Evolution, 2010, 27, 1630-1644.	8.9	47
57	Proteomic and Functional Characterization of a <i>Chlamydomonas reinhardtii</i> Mutant Lacking the Mitochondrial Alternative Oxidase 1. Journal of Proteome Research, 2010, 9, 2825-2838.	3.7	29
58	Oxidative Phosphorylation., 2009,, 469-502.		12
59	The Mitochondrial Genome. , 2009, , 445-467.		3
60	Impaired respiration discloses the physiological significance of state transitions in <i>Chlamydomonas</i> . Proceedings of the National Academy of Sciences of the United States of America, 2009, 106, 15979-15984.	7.1	115
61	The fully-active and structurally-stable form of the mitochondrial ATP synthase of Polytomella sp. is dimeric. Journal of Bioenergetics and Biomembranes, 2009, 41, 1-13.	2.3	23
62	Eukaryotic complex I: functional diversity and experimental systems to unravel the assembly process. Molecular Genetics and Genomics, 2008, 280, 93-110.	2.1	51
63	In Chlamydomonas, the loss of ND5 subunit prevents the assembly of whole mitochondrial complex I and leads to the formation of a low abundant 700ÂkDa subcomplex. Biochimica Et Biophysica Acta - Bioenergetics, 2008, 1777, 388-396.	1.0	33
64	Alternative photosynthetic electron flow to oxygen in marine Synechococcus. Biochimica Et Biophysica Acta - Bioenergetics, 2008, 1777, 269-276.	1.0	155
65	S13.45 Chlamydomonas reinhardtii mitoproteome adaptation in response to inactivation of the energy-dissipating alternative oxidase 1 by RNA interference. Biochimica Et Biophysica Acta - Bioenergetics, 2008, 1777, S99.	1.0	0
66	A type II NAD(P)H dehydrogenase mediates light-independent plastoquinone reduction in the chloroplast of <i>Chlamydomonas</i> . Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 20546-20551.	7.1	187
67	An original adaptation of photosynthesis in the marine green alga <i>Ostreococcus</i> ). Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 7881-7886.	7.1	154
68	The <i>Chlamydomonas</i> Genome Reveals the Evolution of Key Animal and Plant Functions. Science, 2007, 318, 245-250.	12.6	2,354
69	The mitochondrial ATP synthase of chlorophycean algae contains eight subunits of unknown origin involved in the formation of an atypical stator-stalk and in the dimerization of the complex. Journal of Bioenergetics and Biomembranes, 2006, 38, 271-282.	2.3	62
70	High-efficiency biolistic transformation of Chlamydomonas mitochondria can be used to insert mutations in complex I genes. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 4771-4776.	7.1	159
71	ND3 and ND4L Subunits of Mitochondrial Complex I, Both Nucleus Encoded in Chlamydomonas reinhardtii , Are Required for Activity and Assembly of the Enzyme. Eukaryotic Cell, 2006, 5, 1460-1467.	3.4	44
72	The Mitochondrial Oxidative Phosphorylation Proteome of Chlamydomonas reinhardtii Deduced from the Genome Sequencing Project: Table I Plant Physiology, 2005, 137, 447-459.	4.8	78

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73	Impact of a mutation in the mitochondrial LSU rRNA gene from Chlamydomonas reinhardtii on the activity and the assembly of respiratory-chain complexes. Current Genetics, 2004, 45, 323-330.	1.7	12
74	Higher plant-like subunit composition of mitochondrial complex I from Chlamydomonas reinhardtii: 31 conserved components among eukaryotes. Biochimica Et Biophysica Acta - Bioenergetics, 2004, 1658, 212-224.	1.0	107
75	Structural organization of mitochondrial human complex I: role of the ND4 and ND5 mitochondria-encoded subunits and interaction with prohibitin. Biochemical Journal, 2004, 383, 491-499.	3.7	117
76	Photosynthesis and State Transitions in Mitochondrial Mutants of Chlamydomonas reinhardtii Affected in Respiration. Plant Physiology, 2003, 133, 2010-2020.	4.8	119
77	Impact of Mutations Affecting ND Mitochondria-encoded Subunits on the Activity and Assembly of Complex I in Chlamydomonas. Implication for the Structural Organization of the Enzyme. Journal of Molecular Biology, 2002, 319, 1211-1221.	4.2	119
78	Mutations inactivating mitochondrial genes in Chlamydomonas reinhardtii. Biochemical Society Transactions, 2001, 29, 442-446.	3.4	41
79	Abnormal gene expression and assembly of the multimeric complex I in the <i>dum24</i> deletion mitochondrial mutant of <i>Chlamydomonas reinhardtii</i> Biochemical Society Transactions, 2001, 29, A52-A52.	3.4	0
80	Abnormal gene expression and assembly of the multimeric complex I in the <i>dum24</i> deletion mitochondrial mutant of <i>Chlamydomonas reinhardtii</i> Biochemical Society Transactions, 2001, 29, A65-A65.	3.4	0
81	<i>Chlamydomonas reinhardtii</i> mitochondrial mutants lacking complex I activity: characterization of the mutations and assembly of the multimeric complex I. Biochemical Society Transactions, 2001, 29, A66-A66.	3.4	0
82	Structure of the telomeric ends of mtÂDNA, transcriptional analysis and complexÂl assembly in the dum24 mitochondrial mutant of Chlamydomonas reinhardtii. Molecular Genetics and Genomics, 2001, 266, 109-114.	2.1	23
83	Mutants of <i>Chlamydomonas reinhardtii</i> Deficient in Mitochondrial Complex I: Characterization of Two Mutations Affecting the <i>nd1</i> Coding Sequence. Genetics, 2001, 158, 1051-1060.	2.9	47