

Sergio Guerrero-Castillo

List of Publications by Year in descending order

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Version: 2024-02-01

38
papers

1,882
citations

304743

22
h-index

315739

38
g-index

42
all docs

42
docs citations

42
times ranked

2947
citing authors

#	ARTICLE	IF	CITATIONS
1	The Assembly Pathway of Mitochondrial Respiratory Chain Complex I. <i>Cell Metabolism</i> , 2017, 25, 128-139.	16.2	325
2	The m-AAA Protease Associated with Neurodegeneration Limits MCU Activity in Mitochondria. <i>Molecular Cell</i> , 2016, 64, 148-162.	9.7	153
3	Evolution and structural organization of the mitochondrial contact site (MICOS) complex and the mitochondrial intermembrane space bridging (MIB) complex. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2016, 1863, 91-101.	4.1	150
4	Unraveling the complexity of mitochondrial complex I assembly: A dynamic process. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2016, 1857, 980-990.	1.0	110
5	MR-1S Interacts with PET100 and PET117 in Module-Based Assembly of Human Cytochrome c Oxidase. <i>Cell Reports</i> , 2017, 18, 1727-1738.	6.4	86
6	Mutations in ATP6V1E1 or ATP6V1A Cause Autosomal-Recessive Cutis Laxa. <i>American Journal of Human Genetics</i> , 2017, 100, 216-227.	6.2	82
7	Locking loop movement in the ubiquinone pocket of complex I disengages the proton pumps. <i>Nature Communications</i> , 2018, 9, 4500.	12.8	80
8	A salvage pathway maintains highly functional respiratory complex I. <i>Nature Communications</i> , 2020, 11, 1643.	12.8	80
9	Accessory NUMM (NDUFS6) subunit harbors a Zn-binding site and is essential for biogenesis of mitochondrial complex I. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 5685-5690.	7.1	64
10	A homozygous missense mutation in ERAL1, encoding a mitochondrial rRNA chaperone, causes Perrault syndrome. <i>Human Molecular Genetics</i> , 2017, 26, 2541-2550.	2.9	61
11	Bi-allelic Mutations in the Mitochondrial Ribosomal Protein MRPS2 Cause Sensorineural Hearing Loss, Hypoglycemia, and Multiple OXPHOS Complex Deficiencies. <i>American Journal of Human Genetics</i> , 2018, 102, 685-695.	6.2	61
12	Barth syndrome cells display widespread remodeling of mitochondrial complexes without affecting metabolic flux distribution. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2018, 1864, 3650-3658.	3.8	53
13	Mutations in Complex I Assembly Factor TMEM126B Result in Muscle Weakness and Isolated Complex I Deficiency. <i>American Journal of Human Genetics</i> , 2016, 99, 208-216.	6.2	51
14	In <i>Yarrowia lipolytica</i> mitochondria, the alternative NADH dehydrogenase interacts specifically with the cytochrome complexes of the classic respiratory pathway. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2009, 1787, 75-85.	1.0	42
15	Physiological uncoupling of mitochondrial oxidative phosphorylation. Studies in different yeast species. <i>Journal of Bioenergetics and Biomembranes</i> , 2011, 43, 323-331.	2.3	38
16	COMplexome Profiling ALIGNment (COPAL) reveals remodeling of mitochondrial protein complexes in Barth syndrome. <i>Bioinformatics</i> , 2019, 35, 3083-3091.	4.1	37
17	Mitochondrial Unselective Channels throughout the eukaryotic domain. <i>Mitochondrion</i> , 2011, 11, 382-390.	3.4	35
18	<i>Staphylococcus epidermidis</i> : metabolic adaptation and biofilm formation in response to different oxygen concentrations. <i>Pathogens and Disease</i> , 2016, 74, ftv111.	2.0	35

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19	Identification and evolutionary analysis of tissue-specific isoforms of mitochondrial complex I subunit NDUFV3. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2017, 1858, 208-217.	1.0	33
20	TMEM70 functions in the assembly of complexes I and V. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2020, 1861, 148202.	1.0	31
21	CEDAR, an online resource for the reporting and exploration of complexome profiling data. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2021, 1862, 148411.	1.0	27
22	Mitochondria from the salt-tolerant yeast <i>Debaryomyces hansenii</i> (halophilic organelles?). <i>Journal of Bioenergetics and Biomembranes</i> , 2010, 42, 11-19.	2.3	25
23	During the stationary growth phase, <i>Yarrowia lipolytica</i> prevents the overproduction of reactive oxygen species by activating an uncoupled mitochondrial respiratory pathway. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2012, 1817, 353-362.	1.0	25
24	Deactivation of mitochondrial complex I after hypoxiaâ€“ischemia in the immature brain. <i>Journal of Cerebral Blood Flow and Metabolism</i> , 2019, 39, 1790-1802.	4.3	24
25	Neutropenia and intellectual disability are hallmarks of biallelic and de novo CLPB deficiency. <i>Genetics in Medicine</i> , 2021, 23, 1705-1714.	2.4	22
26	Complexome Profilingâ€”Exploring Mitochondrial Protein Complexes in Health and Disease. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 796128.	3.7	20
27	The branched mitochondrial respiratory chain from <i>Debaryomyces hansenii</i> : Components and supramolecular organization. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2014, 1837, 73-84.	1.0	19
28	Mitochondrial Dysfunction during the Early Stages of Excitotoxic Spinal Motor Neuron Degeneration in Vivo. <i>ACS Chemical Neuroscience</i> , 2016, 7, 886-896.	3.5	18
29	Complexome analysis of the nitrite-dependent methanotroph <i>Methylomirabilis lanthanidiphila</i> . <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2019, 1860, 734-744.	1.0	18
30	Ablation of mitochondrial DNA results in widespread remodeling of the mitochondrial complexome. <i>EMBO Journal</i> , 2021, 40, e108648.	7.8	18
31	Effect of glycolysis inhibition on mitochondrial function in rat brain. <i>Journal of Biochemical and Molecular Toxicology</i> , 2012, 26, 206-211.	3.0	12
32	The <i>Saccharomyces cerevisiae</i> mitochondrial unselective channel behaves as a physiological uncoupling system regulated by Ca ²⁺ , Mg ²⁺ , phosphate and ATP. <i>Journal of Bioenergetics and Biomembranes</i> , 2015, 47, 477-491.	2.3	10
33	Novel defect in phosphatidylinositol 4â€“kinase type 2â€“alpha (<i>PI4K2A</i>) at the membraneâ€“enzyme interface is associated with metabolic cutis laxa. <i>Journal of Inherited Metabolic Disease</i> , 2020, 43, 1382-1391.	3.6	7
34	Multiplexed complexome profiling using tandem mass tags. <i>Biochimica Et Biophysica Acta - Bioenergetics</i> , 2021, 1862, 148448.	1.0	6
35	Oxidative phosphorylation in <i>Debaryomyces hansenii</i> : Physiological uncoupling at different growth phases. <i>Biochimie</i> , 2014, 102, 124-136.	2.6	5
36	The plastid proteome of the nonphotosynthetic chlorophycean alga <i>Polytomella parva</i> . <i>Microbiological Research</i> , 2021, 243, 126649.	5.3	5

#	ARTICLE	IF	CITATIONS
37	ZBTB11 dysfunction: spectrum of brain abnormalities, biochemical signature and cellular consequences. <i>Brain</i> , 2022, 145, 2602-2616.	7.6	5
38	A critical tyrosine residue determines the uncoupling protein-like activity of the yeast mitochondrial oxaloacetate carrier. <i>Biochemical Journal</i> , 2012, 443, 317-325.	3.7	4