Angela Pyle

List of Publications by Year in descending order

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Version: 2024-02-01

186265 189892 2,850 50 28 50 h-index citations g-index papers 52 52 52 5374 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Pathogenic SLC25A26 variants impair SAH transport activity causing mitochondrial disease. Human Molecular Genetics, 2022, 31, 2049-2062.	2.9	3
2	Interactions between nuclear and mitochondrial SNPs and Parkinson's disease risk. Mitochondrion, 2022, 63, 85-88.	3.4	6
3	A subcellular cookie cutter for spatial genomics in human tissue. Analytical and Bioanalytical Chemistry, 2022, 414, 5483-5492.	3.7	6
4	Heteroplasmic mitochondrial DNA variants in cardiovascular diseases. PLoS Genetics, 2022, 18, e1010068.	3.5	19
5	Recent advances in understanding the molecular genetic basis of mitochondrial disease. Journal of Inherited Metabolic Disease, 2020, 43, 36-50.	3.6	113
6	Clinical presentation and proteomic signature of patients with <i>TANGO2</i> mutations. Journal of Inherited Metabolic Disease, 2020, 43, 297-308.	3.6	43
7	Effects of thyroid hormone on mitochondria and metabolism of human preimplantation embryos. Stem Cells, 2020, 38, 369-381.	3.2	20
8	The Human Coronavirus Receptor <scp>ANPEP</scp> (<scp>CD13</scp>) Is Overexpressed in Parkinson's Disease. Movement Disorders, 2020, 35, 2134-2136.	3.9	4
9	Metabolic effects of bezafibrate in mitochondrial disease. EMBO Molecular Medicine, 2020, 12, e11589.	6.9	45
10	Cell-free mitochondrial DNA in progressive multiple sclerosis. Mitochondrion, 2019, 46, 307-312.	3.4	32
11	Germline selection shapes human mitochondrial DNA diversity. Science, 2019, 364, .	12.6	178
12	NAD(P)HX dehydratase (NAXD) deficiency: a novel neurodegenerative disorder exacerbated by febrile illnesses. Brain, 2019, 142, 50-58.	7.6	51
13	Defective mitochondrial protease LonP1 can cause classical mitochondrial disease. Human Molecular Genetics, 2018, 27, 1743-1753.	2.9	46
14	Mitochondrial oxodicarboxylate carrier deficiency is associated with mitochondrial DNA depletion and spinal muscular atrophyâ \in "like disease. Genetics in Medicine, 2018, 20, 1224-1235.	2.4	31
15	A novel mechanism causing imbalance of mitochondrial fusion and fission in human myopathies. Human Molecular Genetics, 2018, 27, 1186-1195.	2.9	52
16	Segregation of mitochondrial DNA heteroplasmy through a developmental genetic bottleneck in human embryos. Nature Cell Biology, 2018, 20, 144-151.	10.3	182
17	Exposure of Monocytic Cells to Lipopolysaccharide Induces Coordinated Endotoxin Tolerance, Mitochondrial Biogenesis, Mitophagy, and Antioxidant Defenses. Frontiers in Immunology, 2018, 9, 2217.	4.8	45
18	Clonal expansion of mtDNA deletions: different disease models assessed by digital droplet PCR in single muscle cells. Scientific Reports, 2018, 8, 11682.	3.3	21

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19	Genetic heterogeneity of motor neuropathies. Neurology, 2017, 88, 1226-1234.	1.1	81
20	Hypomorphic mutations in POLR3A are a frequent cause of sporadic and recessive spastic ataxia. Brain, 2017, 140, 1561-1578.	7.6	85
21	Mitochondrial DNA depletion induces innate immune dysfunction rescued by IFN-γ. Journal of Allergy and Clinical Immunology, 2017, 140, 1461-1464.e8.	2.9	5
22	De novo <i>CTBP1</i> variant is associated with decreased mitochondrial respiratory chain activities. Neurology: Genetics, 2017, 3, e187.	1.9	11
23	Opening One's Eyes to Mosaicism in Progressive External Ophthalmoplegia. Neurology: Genetics, 2017, 3, e202.	1.9	1
24	Homozygous deletion in <i>MICU1</i> presenting with fatigue and lethargy in childhood. Neurology: Genetics, 2016, 2, e59.	1.9	86
25	Phenotypic convergence of Menkes and Wilson disease. Neurology: Genetics, 2016, 2, e119.	1.9	18
26	A Woman With Intellectual Disability, Amenorrhoea, Seizures, and Balance Problems. JAMA Neurology, 2016, 73, 1494.	9.0	1
27	Recessive Mutations in TRMT10C Cause Defects in Mitochondrial RNA Processing and Multiple Respiratory Chain Deficiencies. American Journal of Human Genetics, 2016, 98, 993-1000.	6.2	89
28	Riboflavin-Responsive and -Non-responsive Mutations in FAD Synthase Cause Multiple Acyl-CoA Dehydrogenase and Combined Respiratory-Chain Deficiency. American Journal of Human Genetics, 2016, 98, 1130-1145.	6.2	118
29	Adult Onset Leigh Syndrome in the Intensive Care Setting: A Novel Presentation of a C12orf65 Related Mitochondrial Disease. Journal of Neuromuscular Diseases, 2015, 2, 409-419.	2.6	22
30	Clonal Expansion of Secondary Mitochondrial DNA Deletions Associated With Spinocerebellar Ataxia Type 28. JAMA Neurology, 2015, 72, 106.	9.0	41
31	Respiratory chain deficiency in nonmitochondrial disease. Neurology: Genetics, 2015, 1, e6.	1.9	23
32	Reply: Evaluation of exome sequencing variation in undiagnosed ataxias. Brain, 2015, 138, e384-e384.	7.6	2
33	TRMT5 Mutations Cause a Defect in Post-transcriptional Modification of Mitochondrial tRNA Associated with Multiple Respiratory-Chain Deficiencies. American Journal of Human Genetics, 2015, 97, 319-328.	6.2	83
34	Extreme-Depth Re-sequencing of Mitochondrial DNA Finds No Evidence of Paternal Transmission in Humans. PLoS Genetics, 2015, 11, e1005040.	3.5	62
35	<i>SPG7</i> mutations are a common cause of undiagnosed ataxia. Neurology, 2015, 84, 1174-1176.	1.1	87
36	Phenotypic variability of TRPV4 related neuropathies. Neuromuscular Disorders, 2015, 25, 516-521.	0.6	27

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37	Clinical heterogeneity of primary familial brain calcification due to a novel mutation in <i>PDGFB</i> Neurology, 2015, 84, 1818-1820.	1.1	14
38	The p.Ser107Leu in <i>BICD2</i> is a mutation †hot spot†causing distal spinal muscular atrophy. Brain, 2015, 138, e391-e391.	7.6	13
39	<i>SCP2</i> mutations and neurodegeneration with brain iron accumulation. Neurology, 2015, 85, 1909-1911.	1.1	39
40	Exome sequencing in undiagnosed inherited and sporadic ataxias. Brain, 2015, 138, 276-283.	7.6	120
41	1353The Induction of Endotoxin Tolerance is Associated with the Activation of Mitochondrial Biogenesis in THP-1 Cells. Open Forum Infectious Diseases, 2014, 1, S354-S354.	0.9	0
42	Mutations in GTPBP3 Cause a Mitochondrial Translation Defect Associated with Hypertrophic Cardiomyopathy, Lactic Acidosis, and Encephalopathy. American Journal of Human Genetics, 2014, 95, 708-720.	6.2	123
43	Accurate mitochondrial DNA sequencing using off-target reads provides a single test to identify pathogenic point mutations. Genetics in Medicine, 2014, 16, 962-971.	2.4	64
44	Defective i6A37 Modification of Mitochondrial and Cytosolic tRNAs Results from Pathogenic Mutations in TRIT1 and Its Substrate tRNA. PLoS Genetics, 2014, 10, e1004424.	3.5	112
45	EXOSC8 mutations alter mRNA metabolism and cause hypomyelination with spinal muscular atrophy and cerebellar hypoplasia. Nature Communications, 2014, 5, 4287.	12.8	120
46	Use of Whole-Exome Sequencing to Determine the Genetic Basis of Multiple Mitochondrial Respiratory Chain Complex Deficiencies. JAMA - Journal of the American Medical Association, 2014, 312, 68.	7.4	304
47	Synaptotagmin 2 Mutations Cause an Autosomal-Dominant Form of Lambert-Eaton Myasthenic Syndrome and Nonprogressive Motor Neuropathy. American Journal of Human Genetics, 2014, 95, 332-339.	6.2	96
48	Behr's Syndrome is Typically Associated with Disturbed Mitochondrial Translation and Mutations in the C12orf65 Gene. Journal of Neuromuscular Diseases, 2014, 1, 55-63.	2.6	20
49	Prominent Sensorimotor Neuropathy Due to SACS Mutations Revealed by Whole-Exome Sequencing. Archives of Neurology, 2012, 69, 1351-4.	4.5	21
50	Fall in circulating mononuclear cell mitochondrial DNA content in human sepsis. Intensive Care Medicine, 2010, 36, 956-962.	8.2	62