

Emanuele Panza

List of Publications by Year in descending order

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Version: 2024-02-01

26
papers

1,008
citations

567281

15
h-index

580821

25
g-index

26
all docs

26
docs citations

26
times ranked

1365
citing authors

#	ARTICLE	IF	CITATIONS
1	The clear cell sarcoma functional genomic landscape. <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	15
2	Benign albeit glycolytic: MCT4 expression and lactate release in giant cell tumour of bone. <i>Bone</i> , 2020, 134, 115302.	2.9	4
3	γ-Pyrroline-5-carboxylate synthetase deficiency: An emergent multifaceted urea cycle-related disorder. <i>Journal of Inherited Metabolic Disease</i> , 2020, 43, 657-670.	3.6	20
4	Seizures and Cardiomyopathy in a Patient with Pallister-Killian Syndrome due to Hexasomy 12p Mosaicism. <i>Molecular Syndromology</i> , 2020, 11, 125-129.	0.8	0
5	P5CS expression study in a new family with <i>ALDH18A1</i> -associated hereditary spastic paraplegia SPG9. <i>Annals of Clinical and Translational Neurology</i> , 2019, 6, 1533-1540.	3.7	14
6	Hereditary Spastic Paraplegia Is a Common Phenotypic Finding in <i>ARG1</i> Deficiency, P5CS Deficiency and HHH Syndrome: Three Inborn Errors of Metabolism Caused by Alteration of an Interconnected Pathway of Glutamate and Urea Cycle Metabolism. <i>Frontiers in Neurology</i> , 2019, 10, 131.	2.4	24
7	Delineation of the 9q31 deletion syndrome: Genomic microarray characterization of two patients with overlapping deletions. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2901-2906.	1.2	4
8	<i>ALDH18A1</i> gene mutations cause dominant spastic paraplegia SPG9: loss of function effect and plausibility of a dominant negative mechanism. <i>Brain</i> , 2016, 139, e3-e3.	7.6	42
9	New perspectives in the diagnosis and management of enteric neuropathies. <i>Nature Reviews Gastroenterology and Hepatology</i> , 2013, 10, 206-218.	17.8	97
10	Genetics of human enteric neuropathies. <i>Progress in Neurobiology</i> , 2012, 96, 176-189.	5.7	36
11	Mutations responsible for MYH9-related thrombocytopenia impair SDF-1-driven migration of megakaryoblastic cells. <i>Thrombosis and Haemostasis</i> , 2011, 106, 693-704.	3.4	24
12	Heavy chain myosin 9-related disease (MYH9-RD): Neutrophil inclusions of myosin-9 as a pathognomonic sign of the disorder. <i>Thrombosis and Haemostasis</i> , 2010, 103, 826-832.	3.4	81
13	A heritable cause of cleft lip and palate—Van der Woude syndrome caused by a novel <i>IRF6</i> mutation. Review of the literature and of the differential diagnosis. <i>European Journal of Pediatrics</i> , 2010, 169, 223-228.	2.7	7
14	<i>MYH9</i> related disease: four novel mutations of the tail domain of myosin-9 correlating with a mild clinical phenotype. <i>European Journal of Haematology</i> , 2010, 84, 291-297.	2.2	32
15	A family with autosomal dominant leukodystrophy linked to 5q23.2-q23.3 without lamin B1 mutations. <i>European Journal of Neurology</i> , 2010, 17, 541-549.	3.3	36
16	Autosomal recessive hereditary spastic paraplegia with thin corpus callosum: a novel mutation in the <i>SPG11</i> gene and further evidence for genetic heterogeneity. <i>European Journal of Neurology</i> , 2009, 16, 121-126.	3.3	13
17	Association of hereditary thrombocythemia and distal limb defects with a thrombopoietin gene mutation. <i>Blood</i> , 2009, 114, 1655-1657.	1.4	21
18	Position of nonmuscle myosin heavy chain IIA (NMMHC-IIA) mutations predicts the natural history of MYH9-related disease. <i>Human Mutation</i> , 2008, 29, 409-417.	2.5	172

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19	Refinement of the SPG9 locus on chromosome 10q23.3â€”24.2 and exclusion of candidate genes. <i>European Journal of Neurology</i> , 2008, 15, 520-524.	3.3	8
20	Transfection of the mutant MYH9 cDNA reproduces the most typical cellular phenotype of MYH9-related disease in different cell lines. <i>PathoGenetics</i> , 2008, 1, 5.	5.7	5
21	Genetic Predisposition to Familial Neuroblastoma: Identification of Two Novel Genomic Regions at 2p and 12p. <i>Human Heredity</i> , 2007, 63, 205-211.	0.8	34
22	The breakpoint identified in a balanced de novo translocation t(7;9)(p14.1;q31.3) disrupts the A-kinase (PRKA) anchor protein 2 gene (AKAP2) on chromosome 9 in a patient with Kallmann syndrome and bone anomalies. <i>International Journal of Molecular Medicine</i> , 2007, 19, 429.	4.0	4
23	A novel locus for syndromic chronic idiopathic intestinal pseudo-obstruction maps to chromosome 8q23â€”q24. <i>European Journal of Human Genetics</i> , 2007, 15, 889-897.	2.8	29
24	Application of a Fluorescent PCR Method for Molecular Diagnosis of Posttransplant Lymphoproliferative Disorders on Routine Tissue Sections. <i>Diagnostic Molecular Pathology</i> , 2005, 14, 170-176.	2.1	1
25	Title is missing!. <i>Medicine (United States)</i> , 2003, 82, 203-215.	1.0	30
26	MYH9-Related Disease. <i>Medicine (United States)</i> , 2003, 82, 203-215.	1.0	255