Anne De Paepe

List of Publications by Year in descending order

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Version: 2024-02-01

60 papers 5,295 citations

147801 31 h-index 58 g-index

63 all docs 63 docs citations

63 times ranked

7490 citing authors

#	Article	IF	CITATIONS
1	Phenotypic and Molecular Heterogeneity in Mandibulofacial Dysostoses: A Case Series From India. Cleft Palate-Craniofacial Journal, 2022, 59, 1346-1351.	0.9	3
2	SOX11 variants cause a neurodevelopmental disorder with infrequent ocular malformations and hypogonadotropic hypogonadism and with distinct DNA methylation profile. Genetics in Medicine, 2022, 24, 1261-1273.	2.4	14
3	A clinical scoring system for congenital contractural arachnodactyly. Genetics in Medicine, 2020, 22, 124-131.	2.4	17
4	<scp><i>VEGFA</i></scp> variants as prognostic markers for the retinopathy in pseudoxanthoma elasticum. Clinical Genetics, 2020, 98, 74-79.	2.0	8
5	A homozygous pathogenic missense variant broadens the phenotypic and mutational spectrum of CREB3L1-related osteogenesis imperfecta. Human Molecular Genetics, 2019, 28, 1801-1809.	2.9	21
6	Bi-allelic AEBP1 mutations in two patients with Ehlers–Danlos syndrome. Human Molecular Genetics, 2019, 28, 1853-1864.	2.9	29
7	Arterial tortuosity syndrome: 40 new families and literature review. Genetics in Medicine, 2018, 20, 1236-1245.	2.4	66
8	Type III collagen affects dermal and vascular collagen fibrillogenesis and tissue integrity in a mutant Col3a1 transgenic mouse model. Matrix Biology, 2018, 70, 72-83.	3.6	48
9	Genetic analysis of osteogenesis imperfecta in the <scp>P</scp> alestinian population: molecular screening of 49 affected families. Molecular Genetics & Enough Commitment (1988)	1.2	26
10	Pathogenic variants in the <i>ABCC6</i> gene are associated with an increased risk for ischemic stroke. Brain Pathology, 2018, 28, 822-831.	4.1	28
11	CRISPR/Cas9-mediated homology-directed repair by ssODNs in zebrafish induces complex mutational patterns resulting from genomic integration of repair-template fragments. DMM Disease Models and Mechanisms, 2018, 11, .	2.4	77
12	Zebrafish type I collagen mutants faithfully recapitulate human type I collagenopathies. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E8037-E8046.	7.1	77
13	Mutations in ATP6V1E1 or ATP6V1A Cause Autosomal-Recessive Cutis Laxa. American Journal of Human Genetics, 2017, 100, 216-227.	6.2	82
14	Tissueâ€specific mosaicism for a lethal osteogenesis imperfecta ⟨i⟩COL1A1⟨/i⟩ mutation causes mild Ol/EDS overlap syndrome. American Journal of Medical Genetics, Part A, 2017, 173, 1047-1050.	1.2	13
15	Osteogenesis imperfecta. Nature Reviews Disease Primers, 2017, 3, 17052.	30.5	481
16	BATCH-GE: Batch analysis of Next-Generation Sequencing data for genome editing assessment. Scientific Reports, 2016, 6, 30330.	3.3	82
17	Orthostatic intolerance and fatigue in the hypermobility type of Ehlers-Danlos Syndrome. Rheumatology, 2016, 55, 1412-1420.	1.9	35
18	Loss of Type I Collagen Telopeptide Lysyl Hydroxylation Causes Musculoskeletal Abnormalities in a Zebrafish Model of Bruck Syndrome. Journal of Bone and Mineral Research, 2016, 31, 1930-1942.	2.8	65

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19	7p22.1 microdeletions involving ACTB associated with developmental delay, short stature, and microcephaly. European Journal of Medical Genetics, 2016, 59, 502-506.	1.3	10
20	Expanding the clinical and mutational spectrum of the Ehlers–Danlos syndrome, dermatosparaxis type. Genetics in Medicine, 2016, 18, 882-891.	2.4	37
21	Clinical utility gene card for: Hereditary thoracic aortic aneurysm and dissection including next-generation sequencing-based approaches. European Journal of Human Genetics, 2016, 24, 146-150.	2.8	28
22	An Exploratory Caseâ€Control Study on the Impact of <scp><i>IL</i></scp> <i>i≥8€1</i> Gene Polymorphisms on Early Implant Failure. Clinical Implant Dentistry and Related Research, 2016, 18, 234-240.	3.7	19
23	Attitudes of cystic fibrosis patients and their parents towards direct-to-consumer genetic testing for carrier status. Personalized Medicine, 2015, 12, 99-107.	1.5	6
24	Ehlers-Danlos Syndrome, Hypermobility Type, Is Linked to Chromosome 8p22-8p21.1 in an Extended Belgian Family. Disease Markers, 2015, 2015, 1-9.	1.3	28
25	Defective Proteolytic Processing of Fibrillar Procollagens and Prodecorin Due to Biallelic <i>BMP1</i> Mutations Results in a Severe, Progressive Form of Osteogenesis Imperfecta. Journal of Bone and Mineral Research, 2015, 30, 1445-1456.	2.8	42
26	Gene panel sequencing in heritable thoracic aortic disorders and related entities $\hat{a} \in ``results of comprehensive testing in a cohort of 264 patients. Orphanet Journal of Rare Diseases, 2015, 10, 9.$	2.7	62
27	Genetics of the Ehlers–Danlos syndrome: more than collagen disorders. Expert Opinion on Orphan Drugs, 2015, 3, 379-392.	0.8	3
28	Efficiency of Exome Sequencing for the Molecular Diagnosis of Pseudoxanthoma Elasticum. Journal of Investigative Dermatology, 2015, 135, 992-998.	0.7	25
29	<i>RNF216</i> mutations as a novel cause of autosomal recessive Huntington-like disorder. Neurology, 2015, 84, 1760-1766.	1.1	59
30	Association of Type and Location of <i> BRCA1 < /i > and <i> BRCA2 < /i > Mutations With Risk of Breast and Ovarian Cancer. JAMA - Journal of the American Medical Association, 2015, 313, 1347.</i></i>	7.4	390
31	Genetic Defects in TAPT1 Disrupt Ciliogenesis and Cause a Complex Lethal Osteochondrodysplasia. American Journal of Human Genetics, 2015, 97, 521-534.	6.2	39
32	Congenital contractural arachnodactyly due to a novel splice site mutation in the FBN2 gene. Journal of Pediatric Genetics, 2015, 03, 163-166.	0.7	3
33	Intrinsic cardiomyopathy in Marfan syndrome: results from in-vivo and ex-vivo studies of the Fbn1C1039G/+ model and longitudinal findings in humans. Pediatric Research, 2015, 78, 256-263.	2.3	45
34	Absence of Cardiovascular Manifestations in a Haploinsufficient Tgfbr1 Mouse Model. PLoS ONE, 2014, 9, e89749.	2.5	9
35	Expressed Repeat Elements Improve RT-qPCR Normalization across a Wide Range of Zebrafish Gene Expression Studies. PLoS ONE, 2014, 9, e109091.	2.5	38
36	Novel pathogenic COL11A1/COL11A2 variants in Stickler syndrome detected by targeted NGS and exome sequencing. Molecular Genetics and Metabolism, 2014, 113, 230-235.	1.1	48

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37	Reference Values for Echocardiographic Assessment of the Diameter of the Aortic Root and Ascending Aorta Spanning All Age Categories. American Journal of Cardiology, 2014, 114, 914-920.	1.6	151
38	Occipital horn syndrome and classical Menkes Syndrome caused by deep intronic mutations, leading to the activation of ATP7A pseudo-exon. European Journal of Human Genetics, 2014, 22, 517-521.	2.8	22
39	Dysautonomia and its underlying mechanisms in the hypermobility type of Ehlers–Danlos syndrome. Seminars in Arthritis and Rheumatism, 2014, 44, 93-100.	3.4	116
40	Pharmacologic activation of wild-type p53 by nutlin therapy in childhood cancer. Cancer Letters, 2014, 344, 157-165.	7.2	39
41	Autonomic symptom burden in the hypermobility type of Ehlers–Danlos syndrome: A comparative study with two other EDS types, fibromyalgia, and healthy controls. Seminars in Arthritis and Rheumatism, 2014, 44, 353-361.	3.4	81
42	Molecular Docking Simulations Provide Insights in the Substrate Binding Sites and Possible Substrates of the ABCC6 Transporter. PLoS ONE, 2014, 9, e102779.	2.5	25
43	Reply to the letter to the editor by Marc Williams. Genetics in Medicine, 2011, 13, 77-78.	2.4	O
44	Downregulation of MiR-449a Is Essential for the Survival of EVI1 Positive Leukemic Cells through Modulation of NOTCH1 and BCL2 Blood, 2009, 114, 361-361.	1.4	0
45	MicroRNA Profiling of EVI1 Deregulated Myeloid Leukemia Blood, 2007, 110, 4146-4146.	1.4	O
46	Bleeding and bruising in patients with Ehlers–Danlos syndrome and other collagen vascular disorders. British Journal of Haematology, 2004, 127, 491-500.	2.5	190
47	Novel Types of Mutation Responsible for the Dermatosparactic Type of Ehlers–Danlos Syndrome (Type) Tj ETQq 123, 656-663.	1 1 0.784 0.7	314 rgBT /C 116
48	The natural history, including orofacial features of three patients with Ehlers-Danlos syndrome, dermatosparaxis type (EDS type VIIC). American Journal of Medical Genetics Part A, 2004, 131A, 18-28.	2.4	75
49	Frequent allelic loss at 10q23 but low incidence of PTEN mutations in merkel cell carcinoma. International Journal of Cancer, 2001, 92, 409-413.	5.1	63
50	Combined M-FISH and CGH analysis allows comprehensive description of genetic alterations in neuroblastoma cell lines. Genes Chromosomes and Cancer, 2001, 32, 126-135.	2.8	46
51	Molecular cytogenetic and clinical findings in ETV6/ABL1-positive leukemia. Genes Chromosomes and Cancer, 2001, 30, 274-282.	2.8	103
52	Molecular cytogenetic and clinical findings in ETV6ABL1â€positive leukemia. Genes Chromosomes and Cancer, 2001, 30, 274-282.	2.8	3
53	Chromosome 2 short arm translocations revealed by M-FISH analysis of neuroblastoma cell lines. Medical and Pediatric Oncology, 2000, 35, 538-540.	1.0	17
54	Exhaustive mutation analysis of the NF1 gene allows identification of 95% of mutations and reveals a high frequency of unusual splicing defects. Human Mutation, 2000, 15, 541-555.	2.5	477

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55	Exhaustive mutation analysis of the NF1 gene allows identification of 95% of mutations and reveals a high frequency of unusual splicing defects., 2000, 15, 541.		4
56	Exhaustive mutation analysis of the NF1 gene allows identification of 95% of mutations and reveals a high frequency of unusual splicing defects. Human Mutation, 2000, 15, 541.	2.5	6
57	Ehlers-Danlos syndromes: Revised nosology, Villefranche, 1997. American Journal of Medical Genetics Part A, 1998, 77, 31-37.	2.4	1,624
58	Ehlersâ€Danlos syndromes: Revised nosology, Villefranche, 1997. American Journal of Medical Genetics Part A, 1998, 77, 31-37.	2.4	40
59	Substitution of valine for glycine 793 in type III procollagen in Ehlers-Danlos syndrome type IV. Human Mutation, 1995, 5, 179-181.	2.5	6
60	Pseudoxanthoma elasticum: Similar autosomal recessive subtype in Belgian and Afrikaner families. American Journal of Medical Genetics Part A, 1991, 38, 16-20.	2.4	24