

Michael D Briggs

List of Publications by Year in descending order

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35
papers

2,148
citations

218677

26
h-index

377865

34
g-index

35
all docs

35
docs citations

35
times ranked

1805
citing authors

#	ARTICLE	IF	CITATIONS
1	Pseudoachondroplasia and multiple epiphyseal dysplasia: Mutation review, molecular interactions, and genotype to phenotype correlations. <i>Human Mutation</i> , 2002, 19, 465-478.	2.5	261
2	Cartilage Oligomeric Matrix Protein Interacts with Type IX Collagen, and Disruptions to These Interactions Identify a Pathogenetic Mechanism in a Bone Dysplasia Family. <i>Journal of Biological Chemistry</i> , 2001, 276, 6046-6055.	3.4	188
3	Mutations in the region encoding the von Willebrand factor A domain of matrilin-3 are associated with multiple epiphyseal dysplasia. <i>Nature Genetics</i> , 2001, 28, 393-396.	21.4	183
4	Targeted Induction of Endoplasmic Reticulum Stress Induces Cartilage Pathology. <i>PLoS Genetics</i> , 2009, 5, e1000691.	3.5	127
5	The unfolded protein response and its relevance to connective tissue diseases. <i>Cell and Tissue Research</i> , 2010, 339, 197-211.	2.9	124
6	Pseudoachondroplasia and multiple epiphyseal dysplasia: A 7-year comprehensive analysis of the known disease genes identify novel and recurrent mutations and provides an accurate assessment of their relative contribution. <i>Human Mutation</i> , 2012, 33, 144-157.	2.5	104
7	Reduced cell proliferation and increased apoptosis are significant pathological mechanisms in a murine model of mild pseudoachondroplasia resulting from a mutation in the C-terminal domain of COMP. <i>Human Molecular Genetics</i> , 2007, 16, 2072-2088.	2.9	84
8	Decreased chondrocyte proliferation and dysregulated apoptosis in the cartilage growth plate are key features of a murine model of epiphyseal dysplasia caused by a <i>matn3</i> mutation. <i>Human Molecular Genetics</i> , 2007, 16, 1728-1741.	2.9	67
9	Identification of Novel pro- α 2(I)X Collagen Gene Mutations in Two Families with Distinctive Oligo-Epiphyseal Forms of Multiple Epiphyseal Dysplasia. <i>American Journal of Human Genetics</i> , 1999, 65, 31-38.	6.2	64
10	Multiple epiphyseal dysplasia mutations in <i>MATN3</i> cause misfolding of the A-domain and prevent secretion of mutant matrilin-3. <i>Human Mutation</i> , 2005, 26, 557-565.	2.5	63
11	The aggrecanopathies; an evolving phenotypic spectrum of human genetic skeletal diseases. <i>Orphanet Journal of Rare Diseases</i> , 2016, 11, 86.	2.7	63
12	Armet/Manf and Creld2 are components of a specialized ER stress response provoked by inappropriate formation of disulphide bonds: implications for genetic skeletal diseases. <i>Human Molecular Genetics</i> , 2013, 22, 5262-5275.	2.9	62
13	Multiple epiphyseal dysplasia: radiographic abnormalities correlated with genotype. <i>Pediatric Radiology</i> , 2001, 31, 10-18.	2.0	61
14	An unfolded protein response is the initial cellular response to the expression of mutant matrilin-3 in a mouse model of multiple epiphyseal dysplasia. <i>Cell Stress and Chaperones</i> , 2010, 15, 835-849.	2.9	59
15	Clinical and radiographic findings in multiple epiphyseal dysplasia caused by <i>MATN3</i> mutations: Description of 12 patients. , 2004, 125A, 278-284.		55
16	Increased intracellular proteolysis reduces disease severity in an ER stress-associated dwarfism. <i>Journal of Clinical Investigation</i> , 2017, 127, 3861-3865.	8.2	50
17	Novel and recurrent mutations in the C-terminal domain of COMP cluster in two distinct regions and result in a spectrum of phenotypes within the pseudoachondroplasia - multiple epiphyseal dysplasia disease group. <i>Human Mutation</i> , 2005, 25, 593-594.	2.5	44
18	Type IX collagen gene mutations can result in multiple epiphyseal dysplasia that is associated with osteochondritis dissecans and a mild myopathy. <i>American Journal of Medical Genetics, Part A</i> , 2010, 152A, 863-869.	1.2	44

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19	A novel form of chondrocyte stress is triggered by a COMP mutation causing pseudoachondroplasia. <i>Human Mutation</i> , 2012, 33, 218-231.	2.5	42
20	Genotype to phenotype correlations in cartilage oligomeric matrix protein associated chondrodysplasias. <i>European Journal of Human Genetics</i> , 2014, 22, 1278-1282.	2.8	40
21	Structural and Functional Characterization of Recombinant Matrilin-3 A-domain and Implications for Human Genetic Bone Diseases. <i>Journal of Biological Chemistry</i> , 2007, 282, 34634-34643.	3.4	39
22	New therapeutic targets in rare genetic skeletal diseases. <i>Expert Opinion on Orphan Drugs</i> , 2015, 3, 1137-1154.	0.8	34
23	Mutations in the known genes are not the major cause of MED; distinctive phenotypic entities among patients with no identified mutations. <i>European Journal of Human Genetics</i> , 2005, 13, 292-301.	2.8	32
24	Increased Classical Endoplasmic Reticulum Stress Is Sufficient to Reduce Chondrocyte Proliferation Rate in the Growth Plate and Decrease Bone Growth. <i>PLoS ONE</i> , 2015, 10, e0117016.	2.5	32
25	Exon skipping mutation in the COL9A2 gene in a family with multiple epiphyseal dysplasia. <i>Matrix Biology</i> , 2000, 19, 121-128.	3.6	30
26	Preselection of cases through expert clinical and radiological review significantly increases mutation detection rate in multiple epiphyseal dysplasia. <i>European Journal of Human Genetics</i> , 2007, 15, 150-154.	2.8	28
27	Abnormal Chondrocyte Apoptosis in the Cartilage Growth Plate is Influenced by Genetic Background and Deletion of CHOP in a Targeted Mouse Model of Pseudoachondroplasia. <i>PLoS ONE</i> , 2014, 9, e85145.	2.5	27
28	Clinical and radiographic features of multiple epiphyseal dysplasia not linked to the COMP or type IX collagen genes. <i>European Journal of Human Genetics</i> , 2001, 9, 606-612.	2.8	26
29	The utility of mouse models to provide information regarding the pathomolecular mechanisms in human genetic skeletal diseases: The emerging role of endoplasmic reticulum stress (Review). <i>International Journal of Molecular Medicine</i> , 2015, 35, 1483-1492.	4.0	23
30	Novel mutations in exon 2 of <i>MATN3</i> affect residues within the \pm -helices of the A-domain and can result in the intracellular retention of mutant matrilin-3. <i>Human Mutation</i> , 2008, 29, 330-330.	2.5	18
31	New developments in chondrocyte ER-stress and related diseases. <i>F1000Research</i> , 2020, 9, 290.	1.6	17
32	XBP1 signalling is essential for alleviating mutant protein aggregation in ER-stress related skeletal disease. <i>PLoS Genetics</i> , 2019, 15, e1008215.	3.5	16
33	Multiple epiphyseal dysplasia and related disorders: Molecular genetics, disease mechanisms, and therapeutic avenues. <i>Developmental Dynamics</i> , 2021, 250, 345-359.	1.8	15
34	Cartilage endoplasmic reticulum stress may influence the onset but not the progression of experimental osteoarthritis. <i>Arthritis Research and Therapy</i> , 2019, 21, 206.	3.5	14
35	<i>CRELD2</i> Is a Novel <i>LRP1</i> Chaperone That Regulates Noncanonical <i>WNT</i> Signaling in Skeletal Development. <i>Journal of Bone and Mineral Research</i> , 2020, 35, 1452-1469.	2.8	12