

# Brendan H Lee

## List of Publications by Year in descending order

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Version: 2024-02-01

226  
papers

14,451  
citations

19657

61  
h-index

23533

111  
g-index

265  
all docs

265  
docs citations

265  
times ranked

16812  
citing authors

#	ARTICLE	IF	CITATIONS
1	Molecular alterations due to <i>Col5a1</i> haploinsufficiency in a mouse model of classic Ehlers-Danlos syndrome. <i>Human Molecular Genetics</i> , 2022, 31, 1325-1335.	2.9	1
2	What Has the Undiagnosed Diseases Network Taught Us About the Clinical Applications of Genomic Testing?. <i>Annual Review of Medicine</i> , 2022, 73, 575-585.	12.2	11
3	Targeting TGF- $\beta$ 2 for treatment of osteogenesis imperfecta. <i>Journal of Clinical Investigation</i> , 2022, 132, .	8.2	26
4	<i>PRUNE1</i> c.933G>A synonymous variant induces exon 7 skipping, disrupts the <i>DHHA2</i> domain, and leads to an atypical <i>NMIHBA</i> syndrome presentation: Case report and review of the literature. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 1868-1874.	1.2	2
5	Genome sequencing reveals novel noncoding variants in <i>PLA2G6</i> and <i>LMNB1</i> causing progressive neurologic disease. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2022, 10, e1892.	1.2	4
6	DDRGK1 is required for the proper development and maintenance of the growth plate cartilage. <i>Human Molecular Genetics</i> , 2022, 31, 2820-2830.	2.9	5
7	A novel, de novo intronic variant in <i>POGZ</i> causes <i>White-Sutton</i> syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 2198-2203.	1.2	4
8	<i>miRNA-34c</i> Suppresses Osteosarcoma Progression In Vivo by Targeting Notch and <i>E2F</i> . <i>JBM Plus</i> , 2022, 6, e10623.	2.7	8
9	The microRNA processor <i>DROSHA</i> is a candidate gene for a severe progressive neurological disorder. <i>Human Molecular Genetics</i> , 2022, 31, 2934-2950.	2.9	6
10	Yap and Taz promote osteogenesis in neural crest cells by preventing chondrogenesis. <i>FASEB Journal</i> , 2022, 36, .	0.5	1
11	Widespread disturbance in extracellular matrix collagen biomarker responses to teriparatide therapy in osteogenesis imperfecta. <i>Bone</i> , 2021, 142, 115703.	2.9	4
12	The transcriptional cofactor <i>Jab1/Cops5</i> is crucial for BMP-mediated mouse chondrocyte differentiation by repressing p53 activity. <i>Journal of Cellular Physiology</i> , 2021, 236, 5686-5697.	4.1	4
13	Transcriptome-directed analysis for Mendelian disease diagnosis overcomes limitations of conventional genomic testing. <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	87
14	Health-related quality of life in adults with osteogenesis imperfecta. <i>Clinical Genetics</i> , 2021, 99, 772-779.	2.0	4
15	Nitric oxide modulates bone anabolism through regulation of osteoblast glycolysis and differentiation. <i>Journal of Clinical Investigation</i> , 2021, 131, .	8.2	39
16	Evaluation of extracorporeal cardiopulmonary resuscitation eligibility criteria for out-of-hospital cardiac arrest patients. <i>BMC Research Notes</i> , 2021, 14, 139.	1.4	1
17	Heterozygous variants in <i>SPTBN1</i> cause intellectual disability and autism. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 2037-2045.	1.2	9
18	A novel de novo intronic variant in <i>ITPR1</i> causes Gillespie syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 2315-2324.	1.2	2

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19	Tendon and motor phenotypes in the Crtap <sup>-/-</sup> mouse model of recessive osteogenesis imperfecta. <i>ELife</i> , 2021, 10, .	6.0	11
20	A Novel Mouse Model for SNP in Steroid Receptor Co-Activator-1 Reveals Role in Bone Density and Breast Cancer Metastasis. <i>Endocrinology</i> , 2021, 162, .	2.8	5
21	Yap and Taz function as the osteochondrogenic determinant in neural crest cells. <i>FASEB Journal</i> , 2021, 35, .	0.5	0
22	Localized chondro-ossification underlies joint dysfunction and motor deficits in the <i>Fkbp10</i> mouse model of osteogenesis imperfecta. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, .	7.1	3
23	Pregnancy in women with osteogenesis imperfecta: pregnancy characteristics, maternal, and neonatal outcomes. <i>American Journal of Obstetrics &amp; Gynecology MFM</i> , 2021, 3, 100362.	2.6	11
24	COPB2 loss of function causes a coatopathy with osteoporosis and developmental delay. <i>American Journal of Human Genetics</i> , 2021, 108, 1710-1724.	6.2	18
25	Novel assessment of leukocyte-rich platelet-rich plasma on functional and patient-reported outcomes in knee osteoarthritis: a pilot study. <i>Regenerative Medicine</i> , 2021, 16, 823-832.	1.7	8
26	Double-Spin Leukocyte-Rich Platelet-Rich Plasma Is Predominantly Lymphocyte Rich With Notable Concentrations of Other White Blood Cell Subtypes. <i>Arthroscopy, Sports Medicine, and Rehabilitation</i> , 2021, 4, e335-e341.	1.7	7
27	Using the Delphi method to identify clinicians' perceived importance of pediatric exome sequencing results. <i>Genetics in Medicine</i> , 2020, 22, 69-76.	2.4	11
28	Genetic Burden Contributing to Extremely Low or High Bone Mineral Density in a Senior Male Population From the Osteoporotic Fractures in Men Study (MrOS). <i>JBMR Plus</i> , 2020, 4, e10335.	2.7	1
29	Hearing loss in individuals with osteogenesis imperfecta in North America: Results from a multicenter study. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 697-704.	1.2	17
30	Pediatric Outcomes Data Collection Instrument is a Useful Patient-Reported Outcome Measure for Physical Function in Children with Osteogenesis Imperfecta. <i>Genetics in Medicine</i> , 2020, 22, 581-589.	2.4	14
31	Assessment of longitudinal bone growth in osteogenesis imperfecta using metacarpophalangeal pattern profiles. <i>Bone</i> , 2020, 140, 115547.	2.9	8
32	Further delineation of the clinical spectrum of KAT6B disorders and allelic series of pathogenic variants. <i>Genetics in Medicine</i> , 2020, 22, 1338-1347.	2.4	25
33	<i>ATRAID</i> regulates the action of nitrogen-containing bisphosphonates on bone. <i>Science Translational Medicine</i> , 2020, 12, .	12.4	15
34	The crucial p53-dependent oncogenic role of JAB1 in osteosarcoma in vivo. <i>Oncogene</i> , 2020, 39, 4581-4591.	5.9	22
35	A global Slc7a7 knockout mouse model demonstrates characteristic phenotypes of human lysinuric protein intolerance. <i>Human Molecular Genetics</i> , 2020, 29, 2171-2184.	2.9	15
36	De novo EIF2AK1 and EIF2AK2 Variants Are Associated with Developmental Delay, Leukoencephalopathy, and Neurologic Decompensation. <i>American Journal of Human Genetics</i> , 2020, 106, 570-583.	6.2	37

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37	Loss- or Gain-of-Function Mutations in ACOX1 Cause Axonal Loss via Different Mechanisms. <i>Neuron</i> , 2020, 106, 589-606.e6.	8.1	71
38	Leukocyte-dependent effects of platelet-rich plasma on cartilage loss and thermal hyperalgesia in a mouse model of post-traumatic osteoarthritis. <i>Osteoarthritis and Cartilage</i> , 2020, 28, 1385-1393.	1.3	17
39	Notch Signaling in Skeletal Development, Homeostasis and Pathogenesis. <i>Biomolecules</i> , 2020, 10, 332.	4.0	43
40	Fracture Healing in Collagen-Related Preclinical Models of Osteogenesis Imperfecta. <i>Journal of Bone and Mineral Research</i> , 2020, 35, 1132-1148.	2.8	13
41	Exome sequencing compared with standard genetic tests for critically ill infants with suspected genetic conditions. <i>Genetics in Medicine</i> , 2020, 22, 1303-1310.	2.4	21
42	Sites of Cre-recombinase activity in mouse lines targeting skeletal cells. <i>Journal of Bone and Mineral Research</i> , 2020, 36, 1661-1679.	2.8	24
43	Chronic liver disease and impaired hepatic glycogen metabolism in argininosuccinate lyase deficiency. <i>JCI Insight</i> , 2020, 5, .	5.0	10
44	Targeted and sustained Sox9 expression in mouse hypertrophic chondrocytes causes severe and spontaneous osteoarthritis by perturbing cartilage homeostasis. <i>American Journal of Translational Research (discontinued)</i> , 2020, 12, 1056-1069.	0.0	4
45	4-PBA Treatment Improves Bone Phenotypes in the <i>Aga2</i> Mouse Model of Osteogenesis Imperfecta. <i>Journal of Bone and Mineral Research</i> , 2020, 37, 675-686.	2.8	14
46	Clinical Application of Genome and Exome Sequencing as a Diagnostic Tool for Pediatric Patients: a Scoping Review of the Literature. <i>Genetics in Medicine</i> , 2019, 21, 3-16.	2.4	96
47	Growth characteristics in individuals with osteogenesis imperfecta in North America: results from a multicenter study. <i>Genetics in Medicine</i> , 2019, 21, 275-283.	2.4	34
48	Mutations in ANAPC1, Encoding a Scaffold Subunit of the Anaphase-Promoting Complex, Cause Rothmund-Thomson Syndrome Type 1. <i>American Journal of Human Genetics</i> , 2019, 105, 625-630.	6.2	42
49	<i>Bmi1</i> Suppresses Adipogenesis in the Hematopoietic Stem Cell Niche. <i>Stem Cell Reports</i> , 2019, 13, 545-558.	4.8	28
50	Untargeted metabolomic profiling reveals multiple pathway perturbations and new clinical biomarkers in urea cycle disorders. <i>Genetics in Medicine</i> , 2019, 21, 1977-1986.	2.4	47
51	A Multicenter Observational Cohort Study to Evaluate the Effects of Bisphosphonate Exposure on Bone Mineral Density and Other Health Outcomes in Osteogenesis Imperfecta. <i>JBMR Plus</i> , 2019, 3, e10118.	2.7	22
52	<i>Trps1</i> Regulates Development of Craniofacial Skeleton and Is Required for the Initiation of Palatal Shelves Fusion. <i>Frontiers in Physiology</i> , 2019, 10, 513.	2.8	5
53	Impact of Diagnosis and Therapy on Cognitive Function in Urea Cycle Disorders. <i>Annals of Neurology</i> , 2019, 86, 116-128.	5.3	42
54	Effects of Aspirin on Growth Factor Release From Freshly Isolated Leukocyte-Rich Platelet-Rich Plasma in Healthy Men: A Prospective Fixed-Sequence Controlled Laboratory Study. <i>American Journal of Sports Medicine</i> , 2019, 47, 1223-1229.	4.2	33

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55	Mobility in osteogenesis imperfecta: a multicenter North American study. <i>Genetics in Medicine</i> , 2019, 21, 2311-2318.	2.4	15
56	Caries prevalence and experience in individuals with osteogenesis imperfecta: A cross-sectional multicenter study. <i>Special Care in Dentistry</i> , 2019, 39, 214-219.	0.8	11
57	Bi-allelic Variants in TONSL Cause SPONASTRIME Dysplasia and a Spectrum of Skeletal Dysplasia Phenotypes. <i>American Journal of Human Genetics</i> , 2019, 104, 422-438.	6.2	27
58	Identification of Functionally Distinct Mx1+ $\alpha$ SMA+ Periosteal Skeletal Stem Cells. <i>Cell Stem Cell</i> , 2019, 25, 784-796.e5.	11.1	128
59	Osteogenesis imperfecta: advancements in genetics and treatment. <i>Current Opinion in Pediatrics</i> , 2019, 31, 708-715.	2.0	84
60	Microdeletions excluding YWHAE and PFAH1B1 cause a unique leukoencephalopathy: further delineation of the 17p13.3 microdeletion spectrum. <i>Genetics in Medicine</i> , 2019, 21, 1652-1656.	2.4	8
61	Expanding the Spectrum of BAF-Related Disorders: De Novo Variants in SMARCC2 Cause a Syndrome with Intellectual Disability and Developmental Delay. <i>American Journal of Human Genetics</i> , 2019, 104, 164-178.	6.2	59
62	Recurrent mosaic MTOR c.5930C > T (p.Thr1977Ile) variant causing megalencephaly, asymmetric polymicrogyria, and cutaneous pigmented mosaicism: Case report and review of the literature. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 475-479.	1.2	11
63	Oro-dental and cranio-facial characteristics of osteogenesis imperfecta type V. <i>European Journal of Medical Genetics</i> , 2019, 62, 103606.	1.3	11
64	Dental and craniofacial characteristics caused by the p.Ser40Leu mutation in <i>IFITM5</i> . <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 65-70.	1.2	7
65	Alterations in non-type I collagen biomarkers in osteogenesis imperfecta. <i>Bone</i> , 2019, 120, 70-74.	2.9	11
66	SAT-LB088 Assessing Metacarpal Cortical Thickness as a Tool to Evaluate Bone Density Compared to DXA in Osteogenesis Imperfecta. <i>Journal of the Endocrine Society</i> , 2019, 3, .	0.2	0
67	Autosomal recessive Noonan syndrome associated with biallelic LZTR1 variants. <i>Genetics in Medicine</i> , 2018, 20, 1175-1185.	2.4	133
68	Whole-Exome Sequencing Identifies an Intronic Cryptic Splice Site in <i>SERPINF1</i> Causing Osteogenesis Imperfecta Type VI. <i>JBMR Plus</i> , 2018, 2, 235-239.	2.7	6
69	Gene therapy for repair and regeneration of bone and cartilage. <i>Current Opinion in Pharmacology</i> , 2018, 40, 59-66.	3.5	56
70	TGF- $\beta$ 2 Family Signaling in Mesenchymal Differentiation. <i>Cold Spring Harbor Perspectives in Biology</i> , 2018, 10, a022202.	5.5	175
71	Serum Sclerostin Levels in Adults With Osteogenesis Imperfecta: Comparison With Normal Individuals and Response to Teriparatide Therapy. <i>Journal of Bone and Mineral Research</i> , 2018, 33, 307-315.	2.8	12
72	Pathogenic Variants in Fucokinase Cause a Congenital Disorder of Glycosylation. <i>American Journal of Human Genetics</i> , 2018, 103, 1030-1037.	6.2	18

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73	Heterozygous <i>WNT1</i> variant causing a variable bone phenotype. American Journal of Medical Genetics, Part A, 2018, 176, 2419-2424.	1.2	11
74	Effect of Genetic Diagnosis on Patients with Previously Undiagnosed Disease. New England Journal of Medicine, 2018, 379, 2131-2139.	27.0	261
75	Oral health-related quality of life in children and adolescents with osteogenesis imperfecta: cross-sectional study. Orphanet Journal of Rare Diseases, 2018, 13, 187.	2.7	19
76	A multicenter study to evaluate pulmonary function in osteogenesis imperfecta. Clinical Genetics, 2018, 94, 502-511.	2.0	33
77	Mechanical properties of infant bone. Bone, 2018, 113, 151-160.	2.9	18
78	2017 Victor A. McKusick Leadership Award Introduction: Arthur L. Beaudet. American Journal of Human Genetics, 2018, 102, 359-360.	6.2	0
79	Heterozygous Truncating Variants in POMP Escape Nonsense-Mediated Decay and Cause a Unique Immune Dysregulatory Syndrome. American Journal of Human Genetics, 2018, 102, 1126-1142.	6.2	128
80	Further evidence for the involvement of <i>EFL1</i> in a Shwachmanâ€Diamond-like syndrome and expansion of the phenotypic features. Journal of Physical Education and Sports Management, 2018, 4, a003046.	1.2	29
81	De Novo Missense Variants in TRAF7 Cause Developmental Delay, Congenital Anomalies, and Dysmorphic Features. American Journal of Human Genetics, 2018, 103, 154-162.	6.2	56
82	Prospects of Gene Therapy for Skeletal Diseases. , 2018, , 119-137.		1
83	Arginase overexpression in neurons and its effect on traumatic brain injury. Molecular Genetics and Metabolism, 2018, 125, 112-117.	1.1	22
84	Argininosuccinate Lyase Deficiency Causes an Endothelial-Dependent Form of Hypertension. American Journal of Human Genetics, 2018, 103, 276-287.	6.2	39
85	Atypical Alexander disease with dystonia, retinopathy, and a brain mass mimicking astrocytoma. Neurology: Genetics, 2018, 4, e248.	1.9	7
86	IRF2BPL Is Associated with Neurological Phenotypes. American Journal of Human Genetics, 2018, 103, 245-260.	6.2	69
87	The Undiagnosed Diseases Network: Accelerating Discovery about Health and Disease. American Journal of Human Genetics, 2017, 100, 185-192.	6.2	142
88	Kaufman oculo-cerebro-facial syndrome in a child with small and absent terminal phalanges and absent nails. Journal of Human Genetics, 2017, 62, 465-471.	2.3	14
89	A Recurrent De Novo Variant in NACC1 Causes a Syndrome Characterized by Infantile Epilepsy, Cataracts, and Profound Developmental Delay. American Journal of Human Genetics, 2017, 100, 343-351.	6.2	35
90	P3h3-null and Sc65-null Mice Phenocopy the Collagen Lysine Under-hydroxylation and Cross-linking Abnormality of Ehlers-Danlos Syndrome Type VIA. Journal of Biological Chemistry, 2017, 292, 3877-3887.	3.4	32

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91	<i>Fkbp10</i> Deletion in Osteoblasts Leads to Qualitative Defects in Bone. <i>Journal of Bone and Mineral Research</i> , 2017, 32, 1354-1367.	2.8	16
92	Genetic causes and mechanisms of Osteogenesis Imperfecta. <i>Bone</i> , 2017, 102, 40-49.	2.9	82
93	2016 Curt Stern Award Address: From Rare to Common Diseases: Translating Genetic Discovery to Therapy 1. <i>American Journal of Human Genetics</i> , 2017, 100, 397-400.	6.2	0
94	A Chaperone Complex Formed by HSP47, FKBP65, and BiP Modulates Telopeptide Lysyl Hydroxylation of Type I Procollagen. <i>Journal of Bone and Mineral Research</i> , 2017, 32, 1309-1319.	2.8	50
95	Correlations Between Bone Mechanical Properties and Bone Composition Parameters in Mouse Models of Dominant and Recessive Osteogenesis Imperfecta and the Response to Anti-TGF- $\beta$ 2 Treatment. <i>Journal of Bone and Mineral Research</i> , 2017, 32, 347-359.	2.8	24
96	MicroRNA miR-23a cluster promotes osteocyte differentiation by regulating TGF- $\beta$ 2 signalling in osteoblasts. <i>Nature Communications</i> , 2017, 8, 15000.	12.8	91
97	Generalized metabolic bone disease and fracture risk in Rothmund-Thomson syndrome. <i>Human Molecular Genetics</i> , 2017, 26, 3046-3055.	2.9	13
98	Sc11 deficiency causes cystic kidney by activating ERK and STAT3 signaling. <i>Human Molecular Genetics</i> , 2017, 26, 2949-2960.	2.9	28
99	MARRVEL: Integration of Human and Model Organism Genetic Resources to Facilitate Functional Annotation of the Human Genome. <i>American Journal of Human Genetics</i> , 2017, 100, 843-853.	6.2	181
100	An exome sequencing study of Moebius syndrome including atypical cases reveals an individual with CFEOM3A and a <i>TUBB3</i> mutation. <i>Journal of Physical Education and Sports Management</i> , 2017, 3, a000984.	1.2	18
101	A Syndromic Neurodevelopmental Disorder Caused by De Novo Variants in EBF3. <i>American Journal of Human Genetics</i> , 2017, 100, 128-137.	6.2	96
102	Use of Exome Sequencing for Infants in Intensive Care Units. <i>JAMA Pediatrics</i> , 2017, 171, e173438.	6.2	348
103	Neonatal fractures as a presenting feature of <i>LMOD3</i> -associated congenital myopathy. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2789-2794.	1.2	17
104	Milder clinical and biochemical phenotypes associated with the c.482G > A (p.Arg161Gln) pathogenic variant in cobalamin C disease: Implications for management and screening. <i>Molecular Genetics and Metabolism</i> , 2017, 122, 60-66.	1.1	20
105	A non-mosaic PORCN mutation in a male with severe congenital anomalies overlapping focal dermal hypoplasia. <i>Molecular Genetics and Metabolism Reports</i> , 2017, 12, 57-61.	1.1	13
106	Therapeutic Antibody Targeting Tumor- and Osteoblastic Niche-Derived Jagged1 Sensitizes Bone Metastasis to Chemotherapy. <i>Cancer Cell</i> , 2017, 32, 731-747.e6.	16.8	133
107	mTORC1 Signaling is a Critical Regulator of Postnatal Tendon Development. <i>Scientific Reports</i> , 2017, 7, 17175.	3.3	19
108	Long-term use of angiotensin-converting enzyme inhibitors protects against bone loss in African-American elderly men. <i>Archives of Osteoporosis</i> , 2017, 12, 94.	2.4	13

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109	Mutations in Fibronectin Cause a Subtype of Spondylometaphyseal Dysplasia with “Corner Fractures”. American Journal of Human Genetics, 2017, 101, 815-823.	6.2	37
110	Heterozygous variants in <i>ACTL6A</i> , encoding a component of the BAF complex, are associated with intellectual disability. Human Mutation, 2017, 38, 1365-1371.	2.5	27
111	Corner fracture type spondylometaphyseal dysplasia: Overlap with type II collagenopathies. American Journal of Medical Genetics, Part A, 2017, 173, 733-739.	1.2	8
112	Phenotypic and molecular characterisation of CDK13-related congenital heart defects, dysmorphic facial features and intellectual developmental disorders. Genome Medicine, 2017, 9, 73.	8.2	39
113	Osteocyte-specific WNT1 regulates osteoblast function during bone homeostasis. Journal of Clinical Investigation, 2017, 127, 2678-2688.	8.2	143
114	Clinically severe CACNA1A alleles affect synaptic function and neurodegeneration differentially. PLoS Genetics, 2017, 13, e1006905.	3.5	80
115	A recurrent p.Arg92Trp variant in steroidogenic factor-1 (NR5A1) can act as a molecular switch in human sex development. Human Molecular Genetics, 2016, 25, 3446-3453.	2.9	90
116	Sclerostin Antibody Treatment Improves the Bone Phenotype of <i>Crtp</i> Mice, a Model of Recessive Osteogenesis Imperfecta. Journal of Bone and Mineral Research, 2016, 31, 1030-1040.	2.8	70
117	Barriers to drug adherence in the treatment of urea cycle disorders: Assessment of patient, caregiver and provider perspectives. Molecular Genetics and Metabolism Reports, 2016, 8, 43-47.	1.1	27
118	Pharmacological and biological therapeutic strategies for osteogenesis imperfecta. American Journal of Medical Genetics, Part C: Seminars in Medical Genetics, 2016, 172, 367-383.	1.6	53
119	Protein and calorie intakes in adult and pediatric subjects with urea cycle disorders participating in clinical trials of glycerol phenylbutyrate. Molecular Genetics and Metabolism Reports, 2016, 6, 34-40.	1.1	14
120	Dissatisfaction with maintenance of certification in academic pediatrics. Pediatric Research, 2016, 79, 240-242.	2.3	12
121	Restoration of the serum level of SERPINF1 does not correct the bone phenotype in <i>Serpinf1</i> null mice. Molecular Genetics and Metabolism, 2016, 117, 378-382.	1.1	12
122	Glutamine and hyperammonemic crises in patients with urea cycle disorders. Molecular Genetics and Metabolism, 2016, 117, 27-32.	1.1	16
123	A Transgenic Mouse Model of OI Type V Supports a Neomorphic Mechanism of the <i>IFITM5</i> Mutation. Journal of Bone and Mineral Research, 2015, 30, 489-498.	2.8	30
124	Adult presentation of X-linked Conradi-Hünermann-Happle syndrome. American Journal of Medical Genetics, Part A, 2015, 167, 1309-1314.	1.2	6
125	The Osteogenic Niche Promotes Early-Stage Bone Colonization of Disseminated Breast Cancer Cells. Cancer Cell, 2015, 27, 193-210.	16.8	308
126	Losartan increases bone mass and accelerates chondrocyte hypertrophy in developing skeleton. Molecular Genetics and Metabolism, 2015, 115, 53-60.	1.1	19



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127	Post-translationally Abnormal Collagens of Prolyl 3-Hydroxylase-2 Null Mice Offer a Pathobiological Mechanism for the High Myopia Linked to Human LEPREL1 Mutations. <i>Journal of Biological Chemistry</i> , 2015, 290, 8613-8622.	3.4	44
128	Autosomal-Dominant Multiple Pterygium Syndrome Is Caused by Mutations in MYH3. <i>American Journal of Human Genetics</i> , 2015, 96, 841-849.	6.2	55
129	RECQL4 Regulates p53 Function In Vivo During Skeletogenesis. <i>Journal of Bone and Mineral Research</i> , 2015, 30, 1077-1089.	2.8	30
130	Self-reported treatment-associated symptoms among patients with urea cycle disorders participating in glycerol phenylbutyrate clinical trials. <i>Molecular Genetics and Metabolism</i> , 2015, 116, 29-34.	1.1	12
131	Human recombinant arginase enzyme reduces plasma arginine in mouse models of arginase deficiency. <i>Human Molecular Genetics</i> , 2015, 24, 6417-6427.	2.9	40
132	Blood ammonia and glutamine as predictors of hyperammonemic crises in patients with urea cycle disorder. <i>Genetics in Medicine</i> , 2015, 17, 561-568.	2.4	30
133	Catelâ€“Manzke syndrome: Further delineation of the phenotype associated with pathogenic variants in TGDS. <i>Molecular Genetics and Metabolism Reports</i> , 2015, 4, 89-91.	1.1	7
134	Abstract 050: Loss of Argininosuccinate Lyase Leads to Nitric Oxide Deficiency, Endothelial Dysfunction, Impaired Angiogenesis, and Hypertension. <i>Hypertension</i> , 2015, 66, .	2.7	0
135	SERPINF1 as a Cause of Osteogenesis Imperfecta Type VI. , 2014, , 167-172.		1
136	The swaying mouse as a model of osteogenesis imperfecta caused by WNT1 mutations. <i>Human Molecular Genetics</i> , 2014, 23, 4035-4042.	2.9	66
137	Dual Role of the Trps1 Transcription Factor in Dentin Mineralization. <i>Journal of Biological Chemistry</i> , 2014, 289, 27481-27493.	3.4	27
138	Differential Effects of Collagen Prolyl 3-Hydroxylation on Skeletal Tissues. <i>PLoS Genetics</i> , 2014, 10, e1004121.	3.5	31
139	Recessive Osteogenesis Imperfecta Due to Mutations in CRTAP, LEPRE1 and PPIB. , 2014, , 141-150.		0
140	The genetic basis of DOORS syndrome: an exome-sequencing study. <i>Lancet Neurology</i> , The, 2014, 13, 44-58.	10.2	108
141	Notch Signaling in Skeletal Stem Cells. <i>Calcified Tissue International</i> , 2014, 94, 68-77.	3.1	42
142	Branched-chain amino acid metabolism: from rare Mendelian diseases to more common disorders. <i>Human Molecular Genetics</i> , 2014, 23, R1-R8.	2.9	234
143	Connective tissue alterations in Fkbp10âˆ“/âˆ“ mice. <i>Human Molecular Genetics</i> , 2014, 23, 4822-4831.	2.9	54
144	Excessive transforming growth factor-Î² signaling is a common mechanism in osteogenesis imperfecta. <i>Nature Medicine</i> , 2014, 20, 670-675.	30.7	237

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145	Notch Activation as a Driver of Osteogenic Sarcoma. <i>Cancer Cell</i> , 2014, 26, 390-401.	16.8	115
146	Genotypeâ€œPhenotype Correlation â€œ Promiscuity in the Era of Next-Generation Sequencing. <i>New England Journal of Medicine</i> , 2014, 371, 593-596.	27.0	86
147	Sodium phenylbutyrate decreases plasma branched-chain amino acids in patients with urea cycle disorders. <i>Molecular Genetics and Metabolism</i> , 2014, 113, 131-135.	1.1	58
148	ADAMTS-7 forms a positive feedback loop with TNF-Î± in the pathogenesis of osteoarthritis. <i>Annals of the Rheumatic Diseases</i> , 2014, 73, 1575-1584.	0.9	64
149	Mice Expressing Mutant <i>Trpv4</i> Recapitulate the Human <i>TRPV4</i> Disorders. <i>Journal of Bone and Mineral Research</i> , 2014, 29, 1815-1822.	2.8	18
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