

Wayne A Cabral

List of Publications by Year in descending order

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Version: 2024-02-01

21
papers

2,292
citations

516710

16
h-index

713466

21
g-index

21
all docs

21
docs citations

21
times ranked

2230
citing authors

#	ARTICLE	IF	CITATIONS
1	Consortium for osteogenesis imperfecta mutations in the helical domain of type I collagen: regions rich in lethal mutations align with collagen binding sites for integrins and proteoglycans. <i>Human Mutation</i> , 2007, 28, 209-221.	2.5	620
2	Prolyl 3-hydroxylase 1 deficiency causes a recessive metabolic bone disorder resembling lethal/severe osteogenesis imperfecta. <i>Nature Genetics</i> , 2007, 39, 359-365.	21.4	429
3	In vivo base editing rescues Hutchinsonâ€™Gilford progeria syndrome in mice. <i>Nature</i> , 2021, 589, 608-614.	27.8	275
4	Lack of Cyclophilin B in Osteogenesis Imperfecta with Normal Collagen Folding. <i>New England Journal of Medicine</i> , 2010, 362, 521-528.	27.0	158
5	Mutations Near Amino End of Î±1(I) Collagen Cause Combined Osteogenesis Imperfecta/Ehlers-Danlos Syndrome by Interference with N-propeptide Processing. <i>Journal of Biological Chemistry</i> , 2005, 280, 19259-19269.	3.4	118
6	Null mutations in LEPRE1 and CRTAP cause severe recessive osteogenesis imperfecta. <i>Cell and Tissue Research</i> , 2010, 339, 59-70.	2.9	108
7	Abnormal Type I Collagen Post-translational Modification and Crosslinking in a Cyclophilin B KO Mouse Model of Recessive Osteogenesis Imperfecta. <i>PLoS Genetics</i> , 2014, 10, e1004465.	3.5	98
8	Structural Heterogeneity of Type I Collagen Triple Helix and Its Role in Osteogenesis Imperfecta. <i>Journal of Biological Chemistry</i> , 2008, 283, 4787-4798.	3.4	81
9	Prolyl 3-hydroxylase 1 and CRTAP are mutually stabilizing in the endoplasmic reticulum collagen prolyl 3-hydroxylation complex. <i>Human Molecular Genetics</i> , 2010, 19, 223-234.	2.9	73
10	Somatic activating mutations in MAP2K1 cause melorheostosis. <i>Nature Communications</i> , 2018, 9, 1390.	12.8	56
11	A targeted antisense therapeutic approach for Hutchinsonâ€™Gilford progeria syndrome. <i>Nature Medicine</i> , 2021, 27, 536-545.	30.7	55
12	Cyclophilin-B Modulates Collagen Cross-linking by Differentially Affecting Lysine Hydroxylation in the Helical and Telo-peptidyl Domains of Tendon Type I Collagen. <i>Journal of Biological Chemistry</i> , 2016, 291, 9501-9512.	3.4	46
13	Type I Collagen Triplet Duplication Mutation in Lethal Osteogenesis Imperfecta Shifts Register of Î± Chains throughout the Helix and Disrupts Incorporation of Mutant Helices into Fibrils and Extracellular Matrix. <i>Journal of Biological Chemistry</i> , 2003, 278, 10006-10012.	3.4	29
14	Genetic reduction of mTOR extends lifespan in a mouse model of Hutchinsonâ€™Gilford Progeria syndrome. <i>Aging Cell</i> , 2021, 20, e13457.	6.7	27
15	Somatic <i>SMAD3</i> -activating mutations cause melorheostosis by up-regulating the TGF-Î²/SMAD pathway. <i>Journal of Experimental Medicine</i> , 2020, 217, .	8.5	24
16	Evaluation of musculoskeletal phenotype of the G608G progeria mouse model with lonafarnib, pravastatin, and zoledronic acid as treatment groups. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 12029-12040.	7.1	20
17	Cyclophilin B control of lysine post-translational modifications of skin type I collagen. <i>PLoS Genetics</i> , 2019, 15, e1008196.	3.5	18
18	COL1A1 C-propeptide mutations cause ER mislocalization of procollagen and impair C-terminal procollagen processing. <i>Biochimica Et Biophysica Acta - Molecular Basis of Disease</i> , 2019, 1865, 2210-2223.	3.8	18

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19	Melorheostotic Bone Lesions Caused by Somatic Mutations in <i>MAP2K1</i> Have Deteriorated Microarchitecture and Periosteal Reaction. <i>Journal of Bone and Mineral Research</i> , 2019, 34, 883-895.	2.8	16
20	Cyclophilin B Deficiency Causes Abnormal Dentin Collagen Matrix. <i>Journal of Proteome Research</i> , 2017, 16, 2914-2923.	3.7	12
21	Substitution of murine type I collagen A1 3-hydroxylation site alters matrix structure but does not recapitulate osteogenesis imperfecta bone dysplasia. <i>Matrix Biology</i> , 2020, 90, 20-39.	3.6	11