

# Eveline Boudin

## List of Publications by Year in descending order

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Version: 2024-02-01

45  
papers

1,319  
citations

471061

17  
h-index

360668

35  
g-index

48  
all docs

48  
docs citations

48  
times ranked

2047  
citing authors

#	ARTICLE	IF	CITATIONS
1	Bone Overgrowth-associated Mutations in the LRP4 Gene Impair Sclerostin Facilitator Function. <i>Journal of Biological Chemistry</i> , 2011, 286, 19489-19500.	1.6	255
2	A look behind the scenes: the risk and pathogenesis of primary osteoporosis. <i>Nature Reviews Rheumatology</i> , 2015, 11, 462-474.	3.5	204
3	Wnt signaling: A win for bone. <i>Archives of Biochemistry and Biophysics</i> , 2008, 473, 112-116.	1.4	110
4	A Novel Domain-Specific Mutation in a Sclerosteosis Patient Suggests a Role of LRP4 as an Anchor for Sclerostin in Human Bone. <i>Journal of Bone and Mineral Research</i> , 2016, 31, 874-881.	3.1	65
5	The role of extracellular modulators of canonical Wnt signaling in bone metabolism and diseases. <i>Seminars in Arthritis and Rheumatism</i> , 2013, 43, 220-240.	1.6	62
6	WNT Signaling and Bone: Lessons From Skeletal Dysplasias and Disorders. <i>Frontiers in Endocrinology</i> , 2020, 11, 165.	1.5	61
7	Genetic control of bone mass. <i>Molecular and Cellular Endocrinology</i> , 2016, 432, 3-13.	1.6	59
8	Mutations in the latent TGF-beta binding protein 3 (LTBP3) gene cause brachyolmia with amelogenesis imperfecta. <i>Human Molecular Genetics</i> , 2015, 24, 3038-3049.	1.4	40
9	Variation in the Kozak sequence of WNT16 results in an increased translation and is associated with osteoporosis related parameters. <i>Bone</i> , 2014, 59, 57-65.	1.4	39
10	Buried in the Middle but Guilty: Intronic Mutations in the <i>TCIRG1</i> Gene Cause Human Autosomal Recessive Osteopetrosis. <i>Journal of Bone and Mineral Research</i> , 2015, 30, 1814-1821.	3.1	39
11	Germline and Mosaic Variants in PRKACA and PRKACB Cause a Multiple Congenital Malformation Syndrome. <i>American Journal of Human Genetics</i> , 2020, 107, 977-988.	2.6	33
12	MECHANISMS IN ENDOCRINOLOGY: Genetics of human bone formation. <i>European Journal of Endocrinology</i> , 2017, 177, R69-R83.	1.9	29
13	Identification of the first deletion in the LRP5 gene in a patient with Autosomal Dominant Osteopetrosis type I. <i>Bone</i> , 2011, 49, 568-571.	1.4	27
14	The <i>Lrp4</i> <sup>R1170Q</sup> Homozygous Knock-In Mouse Recapitulates the Bone Phenotype of Sclerosteosis in Humans. <i>Journal of Bone and Mineral Research</i> , 2017, 32, 1739-1749.	3.1	27
15	Bi-allelic Loss-of-Function Mutations in the NPR-C Receptor Result in Enhanced Growth and Connective Tissue Abnormalities. <i>American Journal of Human Genetics</i> , 2018, 103, 288-295.	2.6	25
16	Camurati's "Engelmann Disease. <i>Calcified Tissue International</i> , 2019, 104, 554-560.	1.5	25
17	Novel SOST gene mutation in a sclerosteosis patient and her parents. <i>Bone</i> , 2013, 52, 707-710.	1.4	21
18	A common LRP4 haplotype is associated with bone mineral density and hip geometry in men "Data from the Odense Androgen Study (OAS). <i>Bone</i> , 2013, 53, 414-420.	1.4	17

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19	Common Genetic Variation in the DKK1 Gene is Associated with Hip Axis Length but not with Bone Mineral Density and Bone Turnover Markers in Young Adult Men: Results from the Odense Androgen Study. <i>Calcified Tissue International</i> , 2010, 86, 271-281.	1.5	16
20	Sclerosing bone dysplasias. <i>Best Practice and Research in Clinical Endocrinology and Metabolism</i> , 2018, 32, 707-723.	2.2	15
21	Human Genetics of Sclerosing Bone Disorders. <i>Current Osteoporosis Reports</i> , 2018, 16, 256-268.	1.5	13
22	Conditional mouse models support the role of SLC39A14 (ZIP14) in Hyperostosis Cranialis Interna and in bone homeostasis. <i>PLoS Genetics</i> , 2018, 14, e1007321.	1.5	13
23	Sclerosing Bone Dysplasias: Leads Toward Novel Osteoporosis Treatments. <i>Current Osteoporosis Reports</i> , 2014, 12, 243-251.	1.5	12
24	DNA Methylation Profiling and Genomic Analysis in 20 Children with Short Stature Who Were Born Small for Gestational Age. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, e4730-e4741.	1.8	12
25	Genetic association study of WNT10B polymorphisms with BMD and adiposity parameters in Danish and Belgian males. <i>Endocrine</i> , 2013, 44, 247-254.	1.1	11
26	Resistin polymorphisms show associations with obesity, but not with bone parameters in men: results from the Odense Androgen Study. <i>Molecular Biology Reports</i> , 2013, 40, 2467-2472.	1.0	10
27	Single nucleotide polymorphisms in sFRP4 are associated with bone and body composition related parameters in Danish but not in Belgian men. <i>Molecular Genetics and Metabolism</i> , 2012, 106, 366-374.	0.5	9
28	WNT16 Requires G $\beta$ Subunits as Intracellular Partners for Both Its Canonical and Non-Canonical WNT Signalling Activity in Osteoblasts. <i>Calcified Tissue International</i> , 2020, 106, 294-302.	1.5	9
29	No mutations in the serotonin related TPH1 and HTR1B genes in patients with monogenic sclerosing bone disorders. <i>Bone</i> , 2013, 55, 52-56.	1.4	7
30	Genetic Screening of WNT4 and WNT5B in Two Populations with Deviating Bone Mineral Densities. <i>Calcified Tissue International</i> , 2017, 100, 244-249.	1.5	6
31	A multi-omics approach expands the mutational spectrum of MAP2K1-related melorheostosis. <i>Bone</i> , 2020, 137, 115406.	1.4	6
32	Association study of common variants in the sFRP1 gene region and parameters of bone strength and body composition in two independent healthy Caucasian male cohorts. <i>Molecular Genetics and Metabolism</i> , 2012, 105, 508-515.	0.5	5
33	Genetic Variation in RIN3 in the Belgian Population Supports Its Involvement in the Pathogenesis of Paget's Disease of Bone and Modifies the Age of Onset. <i>Calcified Tissue International</i> , 2019, 104, 613-621.	1.5	5
34	Functional Assessment of Coding and Regulatory Variants From the <i>DKK1</i> Locus. <i>JBMR Plus</i> , 2020, 4, e10423.	1.3	5
35	An Activating Deletion Variant in the Submembrane Region of Natriuretic Peptide Receptor-B Causes Tall Stature. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, 2354-2366.	1.8	5
36	Mutations in sFRP1 or sFRP4 are not a common cause of craniotubular hyperostosis. <i>Bone</i> , 2013, 52, 292-295.	1.4	4

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37	Familial Paget's disease of bone: Long-term follow-up of unaffected relatives with and without Sequestosome 1 mutations. <i>Bone</i> , 2019, 128, 115044.	1.4	4
38	Delineation of a new fibrillino-2-pathway with evidence for a role of FBN2 in the pathogenesis of carpal tunnel syndrome. <i>Journal of Medical Genetics</i> , 2020, 58, jmedgenet-2020-107085.	1.5	4
39	Identification of Compound Heterozygous Variants in LRP4 Demonstrates That a Pathogenic Variant outside the Third Î²-Propeller Domain Can Cause Sclerosteosis. <i>Genes</i> , 2022, 13, 80.	1.0	3
40	Broadening the spectrum of loss-of-function variants in NPR-C-related extreme tall stature. <i>Journal of the Endocrine Society</i> , 2022, 6, bvac019.	0.1	2
41	Sclerosing Bone Disorders. , 2018, , 507-521.		1
42	Expression analysis of mesenchymal KS483 cells during differentiation towards osteoblasts. <i>Bone Abstracts</i> , 0, , .	0.0	0
43	No mutations in the serotonin related TPH1 and HTR1B genes in patients with monogenic sclerosing bone disorders. <i>Bone Abstracts</i> , 0, , .	0.0	0
44	Rs55710688 in the Kozak sequence of WNT16 increases translation efficiency and is associated with osteoporosis related parameters. <i>Bone Abstracts</i> , 0, , .	0.0	0
45	Common variants in Rspo 1,2 and 3 do not associate with BMD in stratified subpopulations of the Odense Androgen Study and mutations in these genes are not a common cause of craniotubular hyperostosis. <i>Bone Abstracts</i> , 0, , .	0.0	0