

Daniel H Cohn

List of Publications by Year in Descending Order

Source: <https://exaly.com/author-pdf/7366588/daniel-h-cohn-publications-by-year.pdf>

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

67
papers

4,207
citations

31
h-index

64
g-index

67
ext. papers

4,870
ext. citations

11.4
avg, IF

4.44
L-index

#	Paper	IF	Citations
67	Mutations in GRK2 cause Jeune syndrome by impairing Hedgehog and canonical Wnt signaling. <i>EMBO Molecular Medicine</i> , 2020 , 12, e11739	12	7
66	Biallelic mutations in LAMA5 disrupts a skeletal noncanonical focal adhesion pathway and produces a distinct bent bone dysplasia. <i>EBioMedicine</i> , 2020 , 62, 103075	8.8	3
65	Nosology and classification of genetic skeletal disorders: 2019 revision. <i>American Journal of Medical Genetics, Part A</i> , 2019 , 179, 2393-2419	2.5	216
64	The α chain of type IX collagen is essential for type IX collagen biosynthesis. <i>American Journal of Medical Genetics, Part A</i> , 2019 , 179, 1672-1677	2.5	1
63	Dominant-negative SOX9 mutations in campomelic dysplasia. <i>Human Mutation</i> , 2019 , 40, 2344-2352	4.7	8
62	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	11	10
61	Expanding the genetic architecture and phenotypic spectrum in the skeletal ciliopathies. <i>Human Mutation</i> , 2018 , 39, 152-166	4.7	51
60	Autosomal recessive Stickler syndrome resulting from a COL9A3 mutation. <i>American Journal of Medical Genetics, Part A</i> , 2018 , 176, 2887-2891	2.5	14
59	The PTH/PTHrP-SIK3 pathway affects skeletogenesis through altered mTOR signaling. <i>Science Translational Medicine</i> , 2018 , 10,	17.5	21
58	A postnatal role for embryonic myosin revealed by MYH3 mutations that alter TGF β signaling and cause autosomal dominant spondylacropotarsal synostosis. <i>Scientific Reports</i> , 2017 , 7, 41803	4.9	17
57	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	6.3	41
56	Mutations in IFT-A satellite core component genes and produce short rib polydactyly syndrome with distinctive campomelia. <i>Cilia</i> , 2017 , 6, 7	5.5	16
55	Genes uniquely expressed in human growth plate chondrocytes uncover a distinct regulatory network. <i>BMC Genomics</i> , 2017 , 18, 983	4.5	7
54	MED resulting from recessively inherited mutations in the gene encoding calcium-activated nucleotidase CANT1. <i>American Journal of Medical Genetics, Part A</i> , 2017 , 173, 2415-2421	2.5	14
53	Somatic mosaicism for a lethal TRPV4 mutation results in non-lethal metatropic dysplasia. <i>American Journal of Medical Genetics, Part A</i> , 2016 , 170, 3298-3302	2.5	6
52	Destabilization of the IFT-B cilia core complex due to mutations in IFT81 causes a Spectrum of Short-Rib Polydactyly Syndrome. <i>Scientific Reports</i> , 2016 , 6, 34232	4.9	30
51	Altered mRNA Splicing, Chondrocyte Gene Expression and Abnormal Skeletal Development due to SF3B4 Mutations in Rodriguez Acrofacial Dysostosis. <i>PLoS Genetics</i> , 2016 , 12, e1006307	6	22

50	Clinical and radiographic delineation of Bent Bone Dysplasia-FGFR2 type or Bent Bone Dysplasia with Distinctive Clavicles and Angel-shaped Phalanges. <i>American Journal of Medical Genetics, Part A</i> , 2016 , 170, 2652-61	2.5	6
49	Variable brain phenotype primarily affects the brainstem and cerebellum in patients with osteogenesis imperfecta caused by recessive WNT1 mutations. <i>Journal of Medical Genetics</i> , 2016 , 53, 427-30	5.8	30
48	The ciliopathy-associated CPLANE proteins direct basal body recruitment of intraflagellar transport machinery. <i>Nature Genetics</i> , 2016 , 48, 648-56	36.3	78
47	An inactivating mutation in intestinal cell kinase, ICK, impairs hedgehog signalling and causes short rib-polydactyly syndrome. <i>Human Molecular Genetics</i> , 2016 , 25, 3998-4011	5.6	31
46	IFT52 mutations destabilize anterograde complex assembly, disrupt ciliogenesis and result in short rib polydactyly syndrome. <i>Human Molecular Genetics</i> , 2016 , 25, 4012-4020	5.6	25
45	Mutations in DYNC2LI1 disrupt cilia function and cause short rib polydactyly syndrome. <i>Nature Communications</i> , 2015 , 6, 7092	17.4	55
44	HSP47 and FKBP65 cooperate in the synthesis of type I procollagen. <i>Human Molecular Genetics</i> , 2015 , 24, 1918-28	5.6	39
43	A second locus for Schneckenbecken dysplasia identified by a mutation in the gene encoding inositol polyphosphate phosphatase-like 1 (INPPL1). <i>American Journal of Medical Genetics, Part A</i> , 2015 , 167A, 2470-3	2.5	8
42	Mice expressing mutant Trpv4 recapitulate the human TRPV4 disorders. <i>Journal of Bone and Mineral Research</i> , 2014 , 29, 1815-1822	6.3	15
41	Diagnosis of ALG12-CDG by exome sequencing in a case of severe skeletal dysplasia. <i>Molecular Genetics and Metabolism Reports</i> , 2014 , 1, 213-219	1.8	14
40	Follistatin in chondrocytes: the link between TRPV4 channelopathies and skeletal malformations. <i>FASEB Journal</i> , 2014 , 28, 2525-37	0.9	24
39	Opsismodysplasia resulting from an insertion mutation in the SH2 domain, which destabilizes INPPL1. <i>American Journal of Medical Genetics, Part A</i> , 2014 , 164A, 2407-11	2.5	10
38	WDR34 mutations that cause short-rib polydactyly syndrome type III/severe asphyxiating thoracic dysplasia reveal a role for the NF- κ B pathway in cilia. <i>American Journal of Human Genetics</i> , 2013 , 93, 926-31 ¹¹		66
37	Whole-genome analysis reveals that mutations in inositol polyphosphate phosphatase-like 1 cause opsismodysplasia. <i>American Journal of Human Genetics</i> , 2013 , 92, 137-43	11	46
36	WNT1 mutations in early-onset osteoporosis and osteogenesis imperfecta. <i>New England Journal of Medicine</i> , 2013 , 368, 1809-16	59.2	253
35	Dominant and recessive forms of fibrochondrogenesis resulting from mutations at a second locus, COL11A2. <i>American Journal of Medical Genetics, Part A</i> , 2012 , 158A, 309-14	2.5	14
34	Exome sequencing identifies PDE4D mutations in acrodysostosis. <i>American Journal of Human Genetics</i> , 2012 , 90, 746-51	11	114
33	Fibrochondrogenesis results from mutations in the COL11A1 type XI collagen gene. <i>American Journal of Human Genetics</i> , 2010 , 87, 708-12	11	56

32	Dominant TRPV4 mutations in nonlethal and lethal metatropic dysplasia. <i>American Journal of Medical Genetics, Part A</i> , 2010 , 152A, 1169-77	2.5	71
31	A recessive skeletal dysplasia, SEMD aggrecan type, results from a missense mutation affecting the C-type lectin domain of aggrecan. <i>American Journal of Human Genetics</i> , 2009 , 84, 72-9	11	94
30	Mutations in the gene encoding the calcium-permeable ion channel TRPV4 produce spondylometaphyseal dysplasia, Kozlowski type and metatropic dysplasia. <i>American Journal of Human Genetics</i> , 2009 , 84, 307-15	11	148
29	Ciliary abnormalities due to defects in the retrograde transport protein DYNC2H1 in short-rib polydactyly syndrome. <i>American Journal of Human Genetics</i> , 2009 , 84, 542-9	11	131
28	Gain-of-function mutations in TRPV4 cause autosomal dominant brachyolmia. <i>Nature Genetics</i> , 2008 , 40, 999-1003	36.3	295
27	Nucleotide-sugar transporter SLC35D1 is critical to chondroitin sulfate synthesis in cartilage and skeletal development in mouse and human. <i>Nature Medicine</i> , 2007 , 13, 1363-7	50.5	91
26	Cartilage-selective genes identified in genome-scale analysis of non-cartilage and cartilage gene expression. <i>BMC Genomics</i> , 2007 , 8, 165	4.5	17
25	MED, COMP, multilayered and NEIN: an overview of multiple epiphyseal dysplasia. <i>Pediatric Radiology</i> , 2005 , 35, 116-23	2.8	43
24	A transcriptional profile of human fetal cartilage. <i>Matrix Biology</i> , 2004 , 23, 299-307	11.4	17
23	Mental retardation and abnormal skeletal development (Dyggve-Melchior-Clausen dysplasia) due to mutations in a novel, evolutionarily conserved gene. <i>American Journal of Human Genetics</i> , 2003 , 72, 419-28	11	43
22	Analysis of clones from a human cartilage cDNA library provides insight into chondrocyte gene expression and identifies novel candidate genes for the osteochondrodysplasias. <i>Molecular Genetics and Metabolism</i> , 2003 , 79, 34-42	3.7	10
21	Defects in extracellular matrix structural proteins in the osteochondrodysplasias. <i>Novartis Foundation Symposium</i> , 2001 , 232, 195-210; discussion 210-2		7
20	Double heterozygosity for pseudoachondroplasia and spondyloepiphyseal dysplasia congenita. <i>American Journal of Medical Genetics Part A</i> , 2001 , 104, 140-6		37
19	Widely distributed mutations in the COL2A1 gene produce achondrogenesis type II/hypochondrogenesis. <i>American Journal of Medical Genetics Part A</i> , 2000 , 92, 95-100		52
18	Rapid determination of COL2A1 mutations in individuals with Stickler syndrome: analysis of potential premature termination codons. <i>American Journal of Medical Genetics Part A</i> , 2000 , 94, 141-8		24
17	Heterozygous mutations in the gene encoding noggin affect human joint morphogenesis. <i>Nature Genetics</i> , 1999 , 21, 302-4	36.3	285
16	Small deletions in the type II collagen triple helix produce Kniest dysplasia 1999 , 85, 105-112		47
15	Small deletions in the type II collagen triple helix produce Kniest dysplasia 1999 , 85, 105		1

14	Identification of five novel mutations in cartilage oligomeric matrix protein gene in pseudoachondroplasia and multiple epiphyseal dysplasia. <i>Human Mutation</i> , 1998 , Suppl 1, S10-7	4.7	29
13	Distinct, autosomal recessive form of spondyloepimetaphyseal dysplasia segregating in an inbred Pakistani kindred. <i>American Journal of Medical Genetics Part A</i> , 1998 , 78, 468-473		44
12	Correlation of linkage data with phenotype in eight families with Stickler syndrome. <i>American Journal of Medical Genetics Part A</i> , 1998 , 80, 121-127		40
11	Radiographic and morphologic findings in a previously undescribed type of mesomelic dysplasia resembling atelosteogenesis type II. <i>American Journal of Medical Genetics Part A</i> , 1998 , 80, 247-251		11
10	Diverse mutations in the gene for cartilage oligomeric matrix protein in the pseudoachondroplasia-multiple epiphyseal dysplasia disease spectrum. <i>American Journal of Human Genetics</i> , 1998 , 62, 311-9	11	148
9	Clinical, morphological, and biochemical phenotype of a new case of Ehlers-Danlos syndrome type VIIC. <i>American Journal of Medical Genetics Part A</i> , 1997 , 68, 25-8		24
8	Sulfate transport in chondrodysplasia. <i>Annals of the New York Academy of Sciences</i> , 1996 , 785, 131-6	6.5	14
7	Achondrogenesis type IB is caused by mutations in the diastrophic dysplasia sulphate transporter gene. <i>Nature Genetics</i> , 1996 , 12, 100-2	36.3	189
6	Thanatophoric dysplasia (types I and II) caused by distinct mutations in fibroblast growth factor receptor 3. <i>Nature Genetics</i> , 1995 , 9, 321-8	36.3	524
5	Dominant mutations in the type II collagen gene, COL2A1, produce spondyloepimetaphyseal dysplasia, Strudwick type. <i>Nature Genetics</i> , 1995 , 11, 87-9	36.3	76
4	Homology-mediated recombination between type I collagen gene exons results in an internal tandem duplication and lethal osteogenesis imperfecta. <i>Human Mutation</i> , 1993 , 2, 21-7	4.7	10
3	First-trimester prenatal diagnosis of osteogenesis imperfecta type II by DNA analysis and sonography. <i>Prenatal Diagnosis</i> , 1993 , 13, 589-96	3.2	30
2	A type X collagen mutation causes Schmid metaphyseal chondrodysplasia. <i>Nature Genetics</i> , 1993 , 5, 79-83	36.3	219
1	Recurrence of lethal osteogenesis imperfecta due to parental mosaicism for a mutation in the COL1A2 gene of type I collagen. The mosaic parent exhibits phenotypic features of a mild form of the disease. <i>Human Mutation</i> , 1992 , 1, 47-54	4.7	68