

# Bernt Popp

## List of Publications by Year in descending order

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Version: 2024-02-01

50  
papers

2,015  
citations

361413

20  
h-index

265206

42  
g-index

61  
all docs

61  
docs citations

61  
times ranked

4634  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mutations in GRIN2A and GRIN2B encoding regulatory subunits of NMDA receptors cause variable neurodevelopmental phenotypes. <i>Nature Genetics</i> , 2010, 42, 1021-1026.	21.4	431
2	Haploinsufficiency of ARID1B, a Member of the SWI/SNF-A Chromatin-Remodeling Complex, Is a Frequent Cause of Intellectual Disability. <i>American Journal of Human Genetics</i> , 2012, 90, 565-572.	6.2	225
3	Clinical relevance of systematic phenotyping and exome sequencing in patients with short stature. <i>Genetics in Medicine</i> , 2018, 20, 630-638.	2.4	101
4	Exome Pool-Seq in neurodevelopmental disorders. <i>European Journal of Human Genetics</i> , 2017, 25, 1364-1376.	2.8	77
5	De novo missense mutations in the NAA10 gene cause severe non-syndromic developmental delay in males and females. <i>European Journal of Human Genetics</i> , 2015, 23, 602-609.	2.8	72
6	Expanding the Phenotype Associated with NAA10-Related N-Terminal Acetylation Deficiency. <i>Human Mutation</i> , 2016, 37, 755-764.	2.5	70
7	Need for high-resolution Genetic Analysis in iPSC: Results and Lessons from the ForIPS Consortium. <i>Scientific Reports</i> , 2018, 8, 17201.	3.3	70
8	Mutations in MBOAT7, Encoding Lysophosphatidylinositol Acyltransferase I, Lead to Intellectual Disability Accompanied by Epilepsy and Autistic Features. <i>American Journal of Human Genetics</i> , 2016, 99, 912-916.	6.2	69
9	Mutations in the BAF-Complex Subunit DPF2 Are Associated with Coffin-Siris Syndrome. <i>American Journal of Human Genetics</i> , 2018, 102, 468-479.	6.2	63
10	Genetic and phenotypic spectrum associated with IFIH1 gain-of-function. <i>Human Mutation</i> , 2020, 41, 837-849.	2.5	63
11	Haploinsufficiency of the Chromatin Remodeler BPTF Causes Syndromic Developmental and Speech Delay, Postnatal Microcephaly, and Dysmorphic Features. <i>American Journal of Human Genetics</i> , 2017, 101, 503-515.	6.2	61
12	The mutational and phenotypic spectrum of TUBA1A-associated tubulinopathy. <i>Orphanet Journal of Rare Diseases</i> , 2019, 14, 38.	2.7	48
13	Behavioral phenotype in five individuals with de novo mutations within the GRIN2B gene. <i>Behavioral and Brain Functions</i> , 2013, 9, 20.	3.3	47
14	<i>TRIM28</i> haploinsufficiency predisposes to Wilms tumor. <i>International Journal of Cancer</i> , 2019, 145, 941-951.	5.1	45
15	CUSHAW3: Sensitive and Accurate Base-Space and Color-Space Short-Read Alignment with Hybrid Seeding. <i>PLoS ONE</i> , 2014, 9, e86869.	2.5	45
16	Single molecule real time sequencing in ADTKD-MUC1 allows complete assembly of the VNTR and exact positioning of causative mutations. <i>Scientific Reports</i> , 2018, 8, 4170.	3.3	40
17	New insights into the clinical and molecular spectrum of the novel CYFIP2-related neurodevelopmental disorder and impairment of the WRC-mediated actin dynamics. <i>Genetics in Medicine</i> , 2021, 23, 543-554.	2.4	32
18	The genetic landscape of intellectual disability and epilepsy in adults and the elderly: a systematic genetic work-up of 150 individuals. <i>Genetics in Medicine</i> , 2021, 23, 1492-1497.	2.4	31

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19	Matching clinical and genetic diagnoses in autosomal dominant polycystic kidney disease reveals novel phenocopies and potential candidate genes. <i>Genetics in Medicine</i> , 2020, 22, 1374-1383.	2.4	30
20	DYNC2LI1 mutations broaden the clinical spectrum of dynein-2 defects. <i>Scientific Reports</i> , 2015, 5, 11649.	3.3	28
21	Biallelic Expression of Mucin-1 in Autosomal Dominant Tubulointerstitial Kidney Disease: Implications for Nongenetic Disease Recognition. <i>Journal of the American Society of Nephrology: JASN</i> , 2018, 29, 2298-2309.	6.1	25
22	De novo and inherited variants in ZNF292 underlie a neurodevelopmental disorder with features of autism spectrum disorder. <i>Genetics in Medicine</i> , 2020, 22, 538-546.	2.4	24
23	Exome first approach to reduce diagnostic costs and time – retrospective analysis of 111 individuals with rare neurodevelopmental disorders. <i>European Journal of Human Genetics</i> , 2022, 30, 117-125.	2.8	22
24	Is MED13L-related intellectual disability a recognizable syndrome?. <i>European Journal of Medical Genetics</i> , 2019, 62, 129-136.	1.3	21
25	De novo variants in SNAP25 cause an early-onset developmental and epileptic encephalopathy. <i>Genetics in Medicine</i> , 2021, 23, 653-660.	2.4	20
26	Targeted sequencing of FH-deficient uterine leiomyomas reveals biallelic inactivating somatic fumarase variants and allows characterization of missense variants. <i>Modern Pathology</i> , 2020, 33, 2341-2353.	5.5	19
27	SWI/SNF protein expression status in fumarate hydratase-deficient renal cell carcinoma: immunohistochemical analysis of 32 tumors from 28 patients. <i>Human Pathology</i> , 2018, 77, 139-146.	2.0	18
28	Clinical, neuroimaging, and molecular spectrum of <i>TECP2</i> – associated hereditary sensory and autonomic neuropathy with intellectual disability. <i>Human Mutation</i> , 2021, 42, 762-776.	2.5	18
29	Prenatal diagnosis of <i>HNF1B</i> – associated renal cysts: Is there a need to differentiate intragenic variants from 17q12 microdeletion syndrome?. <i>Prenatal Diagnosis</i> , 2019, 39, 1136-1147.	2.3	16
30	DLG4-related synaptopathy: a new rare brain disorder. <i>Genetics in Medicine</i> , 2021, 23, 888-899.	2.4	16
31	Genotype-phenotype correlation at codon 1740 of <i>SETD2</i> . <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 2037-2048.	1.2	14
32	Do the exome: A case of Williams-Beuren syndrome with severe epilepsy due to a truncating de novo variant in GABRA1. <i>European Journal of Medical Genetics</i> , 2016, 59, 549-553.	1.3	11
33	A biallelic truncating <i>AEBP1</i> variant causes connective tissue disorder in two siblings. <i>American Journal of Medical Genetics, Part A</i> , 2019, 179, 50-56.	1.2	11
34	Evolutionary conserved networks of human height identify multiple Mendelian causes of short stature. <i>European Journal of Human Genetics</i> , 2019, 27, 1061-1071.	2.8	11
35	High-throughput imaging of ATG9A distribution as a diagnostic functional assay for adaptor protein complex 4-associated hereditary spastic paraplegia. <i>Brain Communications</i> , 2021, 3, fcab221.	3.3	11
36	De novo variants in ATP2B1 lead to neurodevelopmental delay. <i>American Journal of Human Genetics</i> , 2022, 109, 944-952.	6.2	11

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37	Whole exome sequencing reveals a novel de novo FOXC1 mutation in a patient with unrecognized Axenfeld-Rieger syndrome and glaucoma. <i>Gene</i> , 2015, 568, 76-80.	2.2	10
38	Differential Coassembly of $\gamma$ -GABA <sub>A</sub> Rs Associated with Epileptic Encephalopathy. <i>Journal of Neuroscience</i> , 2020, 40, 5518-5530.	3.6	10
39	Diverse molecular causes of unsolved autosomal dominant tubulointerstitial kidney diseases. <i>Kidney International</i> , 2022, 102, 405-420.	5.2	10
40	BDV Syndrome: an Emerging Syndrome With Profound Obesity and Neurodevelopmental Delay Resembling Prader-Willi Syndrome. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2021, 106, 3413-3427.	3.6	9
41	Complementing the phenotypical spectrum of TUBA1A tubulinopathy and its role in early-onset epilepsies. <i>European Journal of Human Genetics</i> , 2022, 30, 298-306.	2.8	9
42	Breast MRI texture analysis for prediction of BRCA-associated genetic risk. <i>BMC Medical Imaging</i> , 2020, 20, 86.	2.7	8
43	Biallelic pathogenic variants in roundabout guidance receptor 1 associate with syndromic congenital anomalies of the kidney and urinary tract. <i>Kidney International</i> , 2022, 101, 1039-1053.	5.2	8
44	Discovering a new part of the phenotypic spectrum of Coffin-Siris syndrome in a fetal cohort. <i>Genetics in Medicine</i> , 2022, 24, 1753-1760.	2.4	6
45	QRICH1 variants in Ververi-Brady syndrome delineation of the genotypic and phenotypic spectrum. <i>Clinical Genetics</i> , 2021, 99, 199-207.	2.0	5
46	ZMYND11 variants are a novel cause of centrotemporal and generalised epilepsies with neurodevelopmental disorder. <i>Clinical Genetics</i> , 2021, 100, 412-429.	2.0	5
47	Biallelic ANKS6 mutations cause late-onset ciliopathy with chronic kidney disease through YAP dysregulation. <i>Human Molecular Genetics</i> , 2022, 31, 1357-1369.	2.9	5
48	Prenatal phenotype of PNKP-related primary microcephaly associated with variants affecting both the FHA and phosphatase domain. <i>European Journal of Human Genetics</i> , 2022, 30, 101-110.	2.8	3
49	The Dilemma of Regularly Missed Diagnoses: ADTKD. <i>Archives of Clinical and Medical Case Reports</i> , 2019, 03, .	0.1	2
50	Dissecting TSC2-mutated renal and hepatic angiomyolipomas in an individual with ARID1B-associated intellectual disability. <i>BMC Cancer</i> , 2019, 19, 435.	2.6	1