

# Jonathan Baets

## List of Publications by Year in descending order

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Version: 2024-02-01

41  
papers

1,378  
citations

361413

20  
h-index

361022

35  
g-index

41  
all docs

41  
docs citations

41  
times ranked

2983  
citing authors

#	ARTICLE	IF	CITATIONS
1	A Recurrent <i>KPNA3</i> Missense Variant Causing Infantile Pure Spastic Paraplegia. <i>Annals of Neurology</i> , 2022, 91, 298-299.	5.3	0
2	Clinical practice guideline for the management of paediatric Charcot-Marie-Tooth disease. <i>Journal of Neurology, Neurosurgery and Psychiatry</i> , 2022, 93, 530-538.	1.9	10
3	De Novo and Dominantly Inherited <i>SPTAN1</i> Mutations Cause Spastic Paraplegia and Cerebellar Ataxia. <i>Movement Disorders</i> , 2022, 37, 1175-1186.	3.9	9
4	<i>RFC1</i> repeat expansions: A recurrent cause of sensory and autonomic neuropathy with cough and ataxia. <i>European Journal of Neurology</i> , 2022, 29, 2156-2161.	3.3	14
5	<i>PCYT2</i> mutations disrupting etherlipid biosynthesis: phenotypes converging on the CDP-ethanolamine pathway. <i>Brain</i> , 2021, 144, e17-e17.	7.6	6
6	Biallelic variants in <i>HPDL</i> cause pure and complicated hereditary spastic paraplegia. <i>Brain</i> , 2021, 144, 1422-1434.	7.6	22
7	High prevalence of sporadic late-onset nemaline myopathy in a cohort of whole-exome sequencing negative myopathy patients. <i>Neuromuscular Disorders</i> , 2021, 31, 1154-1160.	0.6	4
8	Unrestrained poly-ADP-ribosylation provides insights into chromatin regulation and human disease. <i>Molecular Cell</i> , 2021, 81, 2640-2655.e8.	9.7	52
9	Characterization of <i>HNRNPA1</i> mutations defines diversity in pathogenic mechanisms and clinical presentation. <i>JCI Insight</i> , 2021, 6, .	5.0	38
10	Biallelic <i>ADPRHL2</i> mutations in complex neuropathy affect ADP ribosylation and DNA damage response. <i>Life Science Alliance</i> , 2021, 4, e202101057.	2.8	11
11	Assessment of Sacsin Turnover in Patients With ARSACS. <i>Neurology</i> , 2021, 97, e2315-e2327.	1.1	11
12	Multisystem proteinopathy due to a homozygous p.Arg159His <i>VCP</i> mutation. <i>Neurology</i> , 2020, 94, e785-e796.	1.1	15
13	Pathogenic Variants in the Myosin Chaperone <i>UNC-45B</i> Cause Progressive Myopathy with Eccentric Cores. <i>American Journal of Human Genetics</i> , 2020, 107, 1078-1095.	6.2	24
14	The expanding genetic landscape of hereditary motor neuropathies. <i>Brain</i> , 2020, 143, 3540-3563.	7.6	12
15	De Novo and Inherited Variants in <i>GBF1</i> are Associated with Axonal Neuropathy Caused by Golgi Fragmentation. <i>American Journal of Human Genetics</i> , 2020, 107, 763-777.	6.2	14
16	Reply: A homozygous <i>GDAP2</i> loss-of-function variant in a patient with adult-onset cerebellar ataxia; and Novel <i>GDAP2</i> pathogenic variants cause autosomal recessive spinocerebellar ataxia-27 (SCAR27) in a Chinese family. <i>Brain</i> , 2020, 143, e51-e51.	7.6	1
17	Clinico-Genetic, Imaging and Molecular Delineation of <i>COQ8A</i> -Ataxia: A Multicenter Study of 59 Patients. , 2020, 88, 251.		1
18	Reply: <i>De novo SPTAN1</i> mutation in axonal sensorimotor neuropathy and developmental disorder. <i>Brain</i> , 2020, 143, e105-e105.	7.6	1

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19	Safety and efficacy of intravenous bimagrumab in inclusion body myositis (RESILIENT): a randomised, double-blind, placebo-controlled phase 2b trial. <i>Lancet Neurology</i> , The, 2019, 18, 834-844.	10.2	91
20	Nonsense mutations in alpha-II spectrin in three families with juvenile onset hereditary motor neuropathy. <i>Brain</i> , 2019, 142, 2605-2616.	7.6	29
21	Functional characterization of GYG1 variants in two patients with myopathy and glycogenin-1 deficiency. <i>Neuromuscular Disorders</i> , 2019, 29, 951-960.	0.6	8
22	Defects in Axonal Transport in Inherited Neuropathies. <i>Journal of Neuromuscular Diseases</i> , 2019, 6, 401-419.	2.6	23
23	Muscular dystrophy with arrhythmia caused by loss-of-function mutations in <i>BVES</i> . <i>Neurology: Genetics</i> , 2019, 5, e321.	1.9	26
24	Truncating SLC5A7 mutations underlie a spectrum of dominant hereditary motor neuropathies. <i>Neurology: Genetics</i> , 2018, 4, e222.	1.9	19
25	Beyond ALS and FTD: the phenotypic spectrum of TBK1 mutations includes PSP-like and cerebellar phenotypes. <i>Neurobiology of Aging</i> , 2018, 62, 244.e9-244.e13.	3.1	30
26	Detection of variants in dystroglycanopathy-associated genes through the application of targeted whole-exome sequencing analysis to a large cohort of patients with unexplained limb-girdle muscle weakness. <i>Skeletal Muscle</i> , 2018, 8, 23.	4.2	40
27	No supportive evidence for TIA1 gene mutations in a European cohort of ALS-FTD spectrum patients. <i>Neurobiology of Aging</i> , 2018, 69, 293.e9-293.e11.	3.1	15
28	Investigating the role of ALS genes CHCHD10 and TUBA4A in Belgian FTD-ALS spectrum patients. <i>Neurobiology of Aging</i> , 2017, 51, 177.e9-177.e16.	3.1	60
29	Clinical Evidence of Disease Anticipation in Families Segregating a <i>C9orf72</i> Repeat Expansion. <i>JAMA Neurology</i> , 2017, 74, 445.	9.0	56
30	Multisystemic <i>SYNE1</i> ataxia: confirming the high frequency and extending the mutational and phenotypic spectrum. <i>Brain</i> , 2016, 139, e46-e46.	7.6	40
31	<i>SYNE1</i> ataxia is a common recessive ataxia with major non-cerebellar features: a large multi-centre study. <i>Brain</i> , 2016, 139, 1378-1393.	7.6	87
32	Clinical features of <i>TBK1</i> carriers compared with <i>C9orf72</i> , <i>GRN</i> and non-mutation carriers in a Belgian cohort. <i>Brain</i> , 2016, 139, 452-467.	7.6	86
33	Megaconial muscular dystrophy caused by mitochondrial membrane homeostasis defect, new insights from skeletal and heart muscle analyses. <i>Mitochondrion</i> , 2016, 27, 32-38.	3.4	9
34	Transcriptional regulator PRDM12 is essential for human pain perception. <i>Nature Genetics</i> , 2015, 47, 803-808.	21.4	137
35	Loss of function mutations in <i>HARS</i> cause a spectrum of inherited peripheral neuropathies. <i>Brain</i> , 2015, 138, 2161-2172.	7.6	71
36	A novel <i>AARS</i> mutation in a family with dominant myeloneuropathy. <i>Neurology</i> , 2015, 84, 2040-2047.	1.1	33

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37	Defects of mutant DNMT1 are linked to a spectrum of neurological disorders. <i>Brain</i> , 2015, 138, 845-861.	7.6	94
38	Truncating and Missense Mutations in IGHMBP2 Cause Charcot-Marie Tooth Disease Type 2. <i>American Journal of Human Genetics</i> , 2014, 95, 590-601.	6.2	75
39	Partial deletion of <i>AFG3L2</i> causing spinocerebellar ataxia type 28. <i>Neurology</i> , 2014, 82, 2092-2100.	1.1	24
40	Recent advances in Charcot-Marie-Tooth disease. <i>Current Opinion in Neurology</i> , 2014, 27, 532-540.	3.6	60
41	Autosomal Recessive Axonal Neuropathy With Neuromyotonia: A Rare Entity. <i>Pediatric Neurology</i> , 2014, 50, 104-107.	2.1	20