

# Hilde Van Esch

## List of Publications by Year in descending order

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Version: 2024-02-01

75  
papers

3,048  
citations

136950

32  
h-index

189892

50  
g-index

76  
all docs

76  
docs citations

76  
times ranked

7169  
citing authors

#	ARTICLE	IF	CITATIONS
1	RNF170 mutation causes autosomal dominant sensory ataxia with variable pyramidal involvement. <i>European Journal of Neurology</i> , 2022, 29, 345-349.	3.3	2
2	Human Brain Models of Intellectual Disability: Experimental Advances and Novelties. <i>International Journal of Molecular Sciences</i> , 2022, 23, 6476.	4.1	0
3	Comprehensive analysis of neuronal guidance cue expression regulation during monocyte-to-macrophage differentiation reveals post-transcriptional regulation of semaphorin7A by the RNA-binding protein quaking. <i>Innate Immunity</i> , 2021, 27, 118-132.	2.4	2
4	<sc><i>IQSEC2</i></sc> disorder: A new disease entity or a Rett spectrum continuum?. <i>Clinical Genetics</i> , 2021, 99, 462-474.	2.0	11
5	<sc><i>MED27</i></sc> Variants Cause Developmental Delay, Dystonia, and Cerebellar Hypoplasia. <i>Annals of Neurology</i> , 2021, 89, 828-833.	5.3	14
6	ERBB4 exonic deletions on chromosome 2q34 in patients with intellectual disability or epilepsy. <i>European Journal of Human Genetics</i> , 2021, 29, 1377-1383.	2.8	9
7	MAPRE2 mutations result in altered human cranial neural crest migration, underlying craniofacial malformations in CSC-KT syndrome. <i>Scientific Reports</i> , 2021, 11, 4976.	3.3	10
8	Phenotypes and genotypes in non- consanguineous and consanguineous primary microcephaly: High incidence of epilepsy. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2021, 9, e1768.	1.2	6
9	Loss-of-function variants in exon 4 of TAB2 cause a recognizable multisystem disorder with cardiovascular, facial, cutaneous, and musculoskeletal involvement. <i>Genetics in Medicine</i> , 2021, , .	2.4	1
10	cGAS-mediated induction of type I interferon due to inborn errors of histone pre-mRNA processing. <i>Nature Genetics</i> , 2020, 52, 1364-1372.	21.4	105
11	De Novo Variants in LMNB1 Cause Pronounced Syndromic Microcephaly and Disruption of Nuclear Envelope Integrity. <i>American Journal of Human Genetics</i> , 2020, 107, 753-762.	6.2	30
12	Use of Multimodal Imaging and Clinical Biomarkers in Presymptomatic Carriers of <i>C9orf72</i> Repeat Expansion. <i>JAMA Neurology</i> , 2020, 77, 1008.	9.0	45
13	Challenges in molecular diagnosis of X-linked Intellectual disability. <i>British Medical Bulletin</i> , 2020, 133, 36-48.	6.9	10
14	Biological concepts in human sodium channel epilepsies and their relevance in clinical practice. <i>Epilepsia</i> , 2020, 61, 387-399.	5.1	65
15	The clinical relevance of intragenic NRXN1 deletions. <i>Journal of Medical Genetics</i> , 2020, 57, 347-355.	3.2	11
16	De Novo SOX6 Variants Cause a Neurodevelopmental Syndrome Associated with ADHD, Craniosynostosis, and Osteochondromas. <i>American Journal of Human Genetics</i> , 2020, 106, 830-845.	6.2	17
17	CTCF variants in 39 individuals with a variable neurodevelopmental disorder broaden the mutational and clinical spectrum. <i>Genetics in Medicine</i> , 2019, 21, 2723-2733.	2.4	48
18	Paralog Studies Augment Gene Discovery: DDX and DHX Genes. <i>American Journal of Human Genetics</i> , 2019, 105, 302-316.	6.2	56

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19	Enabling Global Clinical Collaborations on Identifiable Patient Data: The Minerva Initiative. <i>Frontiers in Genetics</i> , 2019, 10, 611.	2.3	14
20	Maternal copy-number variations in the DMD gene as secondary findings in noninvasive prenatal screening. <i>Genetics in Medicine</i> , 2019, 21, 2774-2780.	2.4	16
21	Mutation update for the <i>SATB2</i> gene. <i>Human Mutation</i> , 2019, 40, 1013-1029.	2.5	38
22	Defective DNA Polymerase $\delta$ -Primase Leads to X-Linked Intellectual Disability Associated with Severe Growth Retardation, Microcephaly, and Hypogonadism. <i>American Journal of Human Genetics</i> , 2019, 104, 957-967.	6.2	32
23	Diagnostic implications of genetic copy number variation in epilepsy plus. <i>Epilepsia</i> , 2019, 60, 689-706.	5.1	61
24	The Role of AKT3 Copy Number Changes in Brain Abnormalities and Neurodevelopmental Disorders: Four New Cases and Literature Review. <i>Frontiers in Genetics</i> , 2019, 10, 58.	2.3	7
25	Clinical exome sequencing reveals locus heterogeneity and phenotypic variability of cohesinopathies. <i>Genetics in Medicine</i> , 2019, 21, 663-675.	2.4	52
26	Further delineation of the <i>MECP2</i> duplication syndrome phenotype in 59 French male patients, with a particular focus on morphological and neurological features. <i>Journal of Medical Genetics</i> , 2018, 55, 359-371.	3.2	45
27	Predicting fetoplacental chromosomal mosaicism during noninvasive prenatal testing. <i>Prenatal Diagnosis</i> , 2018, 38, 258-266.	2.3	58
28	ZNF462 and KLF12 are disrupted by a de novo translocation in a patient with syndromic intellectual disability and autism spectrum disorder. <i>European Journal of Medical Genetics</i> , 2018, 61, 376-383.	1.3	13
29	FRMPD4 mutations cause X-linked intellectual disability and disrupt dendritic spine morphogenesis. <i>Human Molecular Genetics</i> , 2018, 27, 589-600.	2.9	20
30	Novel CASK mutations in cases with syndromic microcephaly. <i>Human Mutation</i> , 2018, 39, 993-1001.	2.5	16
31	Dual Molecular Effects of Dominant RORA Mutations Cause Two Variants of Syndromic Intellectual Disability with Either Autism or Cerebellar Ataxia. <i>American Journal of Human Genetics</i> , 2018, 102, 744-759.	6.2	51
32	A genotype-first approach identifies an intellectual disability-overweight syndrome caused by PHIP haploinsufficiency. <i>European Journal of Human Genetics</i> , 2018, 26, 54-63.	2.8	32
33	Interstitial microdeletion of 17q11.2 is associated with hypotonia, fatigue, intellectual disability, and a subtle facial phenotype in three unrelated patients. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 209-213.	1.2	2
34	Mapping the landscape of tandem repeat variability by targeted long read single molecule sequencing in familial X-linked intellectual disability. <i>BMC Medical Genomics</i> , 2018, 11, 123.	1.5	5
35	Detecting AGG Interruptions in Females With a FMR1 Premutation by Long-Read Single-Molecule Sequencing: A 1 Year Clinical Experience. <i>Frontiers in Genetics</i> , 2018, 9, 150.	2.3	26
36	A novel SYN1 missense mutation in non-syndromic X-linked intellectual disability affects synaptic vesicle life cycle, clustering and mobility. <i>Human Molecular Genetics</i> , 2017, 26, 4699-4714.	2.9	37

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37	Novel STIL Compound Heterozygous Mutations Cause Severe Fetal Microcephaly and Centriolar Lengthening. <i>Molecular Syndromology</i> , 2017, 8, 282-293.	0.8	15
38	Towards a 21st-century roadmap for biomedical research and drug discovery: consensus report and recommendations. <i>Drug Discovery Today</i> , 2017, 22, 327-339.	6.4	64
39	Accuracy and clinical value of maternal incidental findings during noninvasive prenatal testing for fetal aneuploidies. <i>Genetics in Medicine</i> , 2017, 19, 306-313.	2.4	47
40	Detecting AGG Interruptions in Male and Female FMR1 Premutation Carriers by Single-Molecule Sequencing. <i>Human Mutation</i> , 2017, 38, 324-331.	2.5	37
41	Optic Nerve Hypoplasia Is a Pervasive Subcortical Pathology of Visual System in Neonates. , 2017, 58, 5485.		11
42	A Distinct Class of Chromoanagenesis Events Characterized by Focal Copy Number Gains. <i>Human Mutation</i> , 2016, 37, 661-668.	2.5	30
43	Gain-of-function <i>FHF1</i> mutation causes early-onset epileptic encephalopathy with cerebellar atrophy. <i>Neurology</i> , 2016, 86, 2162-2170.	1.1	57
44	Mutations in <i>SNORD118</i> cause the cerebral microangiopathy leukoencephalopathy with calcifications and cysts. <i>Nature Genetics</i> , 2016, 48, 1185-1192.	21.4	114
45	Quaking promotes monocyte differentiation into pro-atherogenic macrophages by controlling pre-mRNA splicing and gene expression. <i>Nature Communications</i> , 2016, 7, 10846.	12.8	87
46	Mutations in <i>CEP120</i> cause Joubert syndrome as well as complex ciliopathy phenotypes. <i>Journal of Medical Genetics</i> , 2016, 53, 608-615.	3.2	55
47	Identification of Intellectual Disability Genes in Female Patients with a Skewed X-Inactivation Pattern. <i>Human Mutation</i> , 2016, 37, 804-811.	2.5	92
48	PIK3CA-associated developmental disorders exhibit distinct classes of mutations with variable expression and tissue distribution. <i>JCI Insight</i> , 2016, 1, .	5.0	134
49	A novel fragile X syndrome mutation reveals a conserved role for the carboxy-terminus in <i>FMRP</i> localization and function. <i>EMBO Molecular Medicine</i> , 2015, 7, 423-437.	6.9	41
50	A homozygous deletion of exon 1 in <i>WISP3</i> causes progressive pseudorheumatoid dysplasia in two siblings. <i>Human Genome Variation</i> , 2015, 2, 15049.	0.7	8
51	No evidence of locus heterogeneity in familial microcephaly with or without chorioretinopathy, lymphedema, or mental retardation syndrome. <i>Orphanet Journal of Rare Diseases</i> , 2015, 10, 52.	2.7	16
52	Mutations in Either <i>TUBB</i> or <i>MAPRE2</i> Cause Circumferential Skin Creases Kunze Type. <i>American Journal of Human Genetics</i> , 2015, 97, 790-800.	6.2	63
53	Increased <i>STAG2</i> dosage defines a novel cohesinopathy with intellectual disability and behavioral problems. <i>Human Molecular Genetics</i> , 2015, 24, 7171-7181.	2.9	28
54	Focus group discussions on secondary variants and next-generation sequencing technologies. <i>European Journal of Medical Genetics</i> , 2015, 58, 249-257.	1.3	14

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55	Noninvasive prenatal testing using a novel analysis pipeline to screen for all autosomal fetal aneuploidies improves pregnancy management. <i>European Journal of Human Genetics</i> , 2015, 23, 1286-1293.	2.8	108
56	Mutations in DDX3X Are a Common Cause of Unexplained Intellectual Disability with Gender-Specific Effects on Wnt Signaling. <i>American Journal of Human Genetics</i> , 2015, 97, 343-352.	6.2	230
57	THOC2 Mutations Implicate mRNA-Export Pathway in X-Linked Intellectual Disability. <i>American Journal of Human Genetics</i> , 2015, 97, 302-310.	6.2	82
58	Microdeletions on 6p22.3 are associated with mesomelic dysplasia Savarirayan type. <i>Journal of Medical Genetics</i> , 2015, 52, 476-483.	3.2	27
59	3 generation pedigree with paternal transmission of the 22q11.2 deletion syndrome: Intrafamilial phenotypic variability. <i>European Journal of Medical Genetics</i> , 2015, 58, 244-248.	1.3	15
60	Microdeletion of the escape genes KDM5C and IQSEC2 in a girl with severe intellectual disability and autistic features. <i>European Journal of Medical Genetics</i> , 2015, 58, 324-327.	1.3	36
61	Expanding the phenotypic spectrum of PORCN variants in two males with syndromic microphthalmia. <i>European Journal of Human Genetics</i> , 2015, 23, 551-554.	2.8	24
62	Homozygous missense mutation in STYXL1 associated with moderate intellectual disability, epilepsy and behavioural complexities. <i>European Journal of Medical Genetics</i> , 2015, 58, 205-210.	1.3	11
63	Genes that Affect Brain Structure and Function Identified by Rare Variant Analyses of Mendelian Neurologic Disease. <i>Neuron</i> , 2015, 88, 499-513.	8.1	258
64	Positron Emission Tomography (PET) Quantification of GABAA Receptors in the Brain of Fragile X Patients. <i>PLoS ONE</i> , 2015, 10, e0131486.	2.5	52
65	Pseudoautosomal Region 1 Length Polymorphism in the Human Population. <i>PLoS Genetics</i> , 2014, 10, e1004578.	3.5	24
66	A prospective study of the clinical utility of prenatal chromosomal microarray analysis in fetuses with ultrasound abnormalities and an exploration of a framework for reporting unclassified variants and risk factors. <i>Genetics in Medicine</i> , 2014, 16, 469-476.	2.4	66
67	Presenting symptoms in adults with the 22q11 deletion syndrome. <i>European Journal of Medical Genetics</i> , 2014, 57, 157-162.	1.3	24
68	Developmental delay and connective tissue disorder in four patients sharing a common microdeletion at 6q13-14. <i>Journal of Medical Genetics</i> , 2010, 47, 717-720.	3.2	22
69	Early myoclonic encephalopathy caused by a disruption of the neuregulin-1 receptor ErbB4. <i>European Journal of Human Genetics</i> , 2009, 17, 378-382.	2.8	40
70	Very early premature ovarian failure in two sisters compound heterozygous for the FMR1 premutation. <i>European Journal of Medical Genetics</i> , 2009, 52, 37-40.	1.3	15
71	Congenital diaphragmatic hernia is part of the new 15q24 microdeletion syndrome. <i>European Journal of Medical Genetics</i> , 2009, 52, 153-156.	1.3	33
72	Screening for FMR-1 premutations in 122 older Flemish males presenting with ataxia. <i>European Journal of Human Genetics</i> , 2005, 13, 121-123.	2.8	58

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73	X-linked mental retardation, short stature, microcephaly and hypogonadism maps to Xp22.1-p21.3 in a Belgian family. <i>European Journal of Medical Genetics</i> , 2005, 48, 145-152.	1.3	7
74	A boy with an unusual association of ventral midline anomalies including a trunk-like umbilicus. <i>Clinical Dysmorphology</i> , 2004, 13, 261-263.	0.3	2
75	Partial DiGeorge syndrome in two patients with a 10p rearrangement. <i>Clinical Genetics</i> , 1999, 55, 269-276.	2.0	34