

# Hilde Peeters

## List of Publications by Year in descending order

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Version: 2024-02-01

70  
papers

3,962  
citations

218381

26  
h-index

138251

58  
g-index

71  
all docs

71  
docs citations

71  
times ranked

7988  
citing authors

#	ARTICLE	IF	CITATIONS
1	Orthognathic surgery in patients with systemic diseases. <i>Oral and Maxillofacial Surgery</i> , 2023, 27, 235-243.	0.6	3
2	Multi-Scale Part-Based Syndrome Classification of 3D Facial Images. <i>IEEE Access</i> , 2022, 10, 23450-23462.	2.6	3
3	Knowledge, attitudes and preferences regarding reproductive genetic carrier screening among reproductive-aged men and women in Flanders (Belgium). <i>European Journal of Human Genetics</i> , 2022, , .	1.4	5
4	Reasons affecting the uptake of reproductive genetic carrier screening among nonpregnant reproductive-aged women in Flanders (Belgium). <i>Journal of Genetic Counseling</i> , 2022, 31, 1043-1053.	0.9	5
5	Insights into the genetic architecture of the human face. <i>Nature Genetics</i> , 2021, 53, 45-53.	9.4	94
6	The Intersection of the Genetic Architectures of Orofacial Clefts and Normal Facial Variation. <i>Frontiers in Genetics</i> , 2021, 12, 626403.	1.1	10
7	Maxillofacial and oral surgery in patients with thrombophilia: safe territory for the oral surgeon? A single-center retrospective study. <i>Oral Surgery, Oral Medicine, Oral Pathology and Oral Radiology</i> , 2021, 132, 514-522.	0.2	1
8	8p21.3 deletions are rare causes of non-syndromic autism spectrum disorder. <i>Neurogenetics</i> , 2021, 22, 207-213.	0.7	1
9	Shared heritability of human face and brain shape. <i>Nature Genetics</i> , 2021, 53, 830-839.	9.4	57
10	Rare deleterious mutations of HNRNP genes result in shared neurodevelopmental disorders. <i>Genome Medicine</i> , 2021, 13, 63.	3.6	50
11	3D facial phenotyping by biometric sibling matching used in contemporary genomic methodologies. <i>PLoS Genetics</i> , 2021, 17, e1009528.	1.5	13
12	Large-scale open-source three-dimensional growth curves for clinical facial assessment and objective description of facial dysmorphism. <i>Scientific Reports</i> , 2021, 11, 12175.	1.6	17
13	3D analysis of facial morphology in Dutch children with cancer. <i>Computer Methods and Programs in Biomedicine</i> , 2021, 205, 106093.	2.6	1
14	Expanded carrier screening in Flanders (Belgium): an online survey on the perspectives of nonpregnant reproductive-aged women. <i>Personalized Medicine</i> , 2021, 18, 361-373.	0.8	6
15	NRXN1 <sup>±</sup> is associated with increased excitability in ASD iPSC-derived neurons. <i>BMC Neuroscience</i> , 2021, 22, 56.	0.8	14
16	Biallelic MADD variants cause a phenotypic spectrum ranging from developmental delay to a multisystem disorder. <i>Brain</i> , 2020, 143, 2437-2453.	3.7	21
17	A de-novo 15q24.2 deletion involving SIN3A is associated with emotional, behavioural, motor problems and hypersensitivity in a girl with above average intelligence and typical facial features.. <i>Clinical Dysmorphology</i> , 2020, 29, 210-213.	0.1	0
18	NCKAP1 Disruptive Variants Lead to a Neurodevelopmental Disorder with Core Features of Autism. <i>American Journal of Human Genetics</i> , 2020, 107, 963-976.	2.6	18

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19	Frequency and Management of Craniofacial Syndromes. <i>Journal of Craniofacial Surgery</i> , 2020, 31, 1091-1097.	0.3	2
20	Interest in expanded carrier screening among individuals and couples in the general population: systematic review of the literature. <i>Human Reproduction Update</i> , 2020, 26, 335-355.	5.2	36
21	The clinical relevance of intragenic NRXN1 deletions. <i>Journal of Medical Genetics</i> , 2020, 57, 347-355.	1.5	11
22	Large-scale targeted sequencing identifies risk genes for neurodevelopmental disorders. <i>Nature Communications</i> , 2020, 11, 4932.	5.8	105
23	Abstract P6-08-03: Germline mutational landscape in 5422 individuals at risk for hereditary breast and ovarian cancer who underwent multi-gene panel testing. , 2020, , .		0
24	Usefulness of fragile X checklist and CGG distribution in specialized institutions in Kinshasa, DR Congo. <i>Journal of Community Genetics</i> , 2019, 10, 153-159.	0.5	2
25	The East Flanders Prospective Twin Survey (EFPTS): 55 Years Later. <i>Twin Research and Human Genetics</i> , 2019, 22, 454-459.	0.3	23
26	Maternal copy-number variations in the DMD gene as secondary findings in noninvasive prenatal screening. <i>Genetics in Medicine</i> , 2019, 21, 2774-2780.	1.1	16
27	Predicting fetoplacental chromosomal mosaicism during noninvasive prenatal testing. <i>Prenatal Diagnosis</i> , 2018, 38, 258-266.	1.1	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.	0.7	13
29	Noise-robust assessment of SNP array based CNV calls through local noise estimation of log R ratios. <i>Statistical Applications in Genetics and Molecular Biology</i> , 2018, 17, .	0.2	0
30	The ethics of patenting autism genes. <i>Nature Reviews Genetics</i> , 2018, 19, 247-248.	7.7	3
31	Six NSCL/P Loci Show Associations With Normal-Range Craniofacial Variation. <i>Frontiers in Genetics</i> , 2018, 9, 502.	1.1	20
32	Spatially Dense 3D Facial Heritability and Modules of Co-heritability in a Father-Offspring Design. <i>Frontiers in Genetics</i> , 2018, 9, 554.	1.1	12
33	Olfactory function in patients with nonsyndromic orofacial clefts and their unaffected relatives. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 2375-2381.	0.7	1
34	NFIB Haploinsufficiency Is Associated with Intellectual Disability and Macrocephaly. <i>American Journal of Human Genetics</i> , 2018, 103, 752-768.	2.6	40
35	A comprehensive clinical and genetic study in 127 patients with ID in Kinshasa, DR Congo. <i>American Journal of Medical Genetics, Part A</i> , 2018, 176, 1897-1909.	0.7	7
36	Compound heterozygous loss-of-function mutations in KIF20A are associated with a novel lethal congenital cardiomyopathy in two siblings. <i>PLoS Genetics</i> , 2018, 14, e1007138.	1.5	18

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37	Targeted sequencing identifies 91 neurodevelopmental-disorder risk genes with autism and developmental-disability biases. <i>Nature Genetics</i> , 2017, 49, 515-526.	9.4	443
38	Variants in TTC25 affect autistic trait in patients with autism spectrum disorder and general population. <i>European Journal of Human Genetics</i> , 2017, 25, 982-987.	1.4	5
39	Hotspots of missense mutation identify neurodevelopmental disorder genes and functional domains. <i>Nature Neuroscience</i> , 2017, 20, 1043-1051.	7.1	152
40	Testing the face shape hypothesis in twins discordant for nonsyndromic orofacial clefting. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2886-2892.	0.7	7
41	Accuracy and Clinical Value of Maternal Incidental Findings During Noninvasive Prenatal Testing for Fetal Aneuploidies. <i>Obstetrical and Gynecological Survey</i> , 2017, 72, 469-470.	0.2	1
42	Executive functioning and local/global visual processing: candidate endophenotypes for autism spectrum disorder?. <i>Journal of Child Psychology and Psychiatry and Allied Disciplines</i> , 2017, 58, 258-269.	3.1	21
43	Accuracy and clinical value of maternal incidental findings during noninvasive prenatal testing for fetal aneuploidies. <i>Genetics in Medicine</i> , 2017, 19, 306-313.	1.1	47
44	The ethics of complexity. Genetics and autism, a literature review. <i>American Journal of Medical Genetics Part B: Neuropsychiatric Genetics</i> , 2016, 171, 305-316.	1.1	23
45	<i>CREBBP</i> mutations in individuals without Rubinstein-Taybi syndrome phenotype. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 2681-2693.	0.7	43
46	Genetic testing and counseling in the case of an autism diagnosis: A caregivers perspective. <i>European Journal of Medical Genetics</i> , 2016, 59, 452-458.	0.7	64
47	Shooting a moving target. Researching autism genes: An interview study with professionals. <i>European Journal of Medical Genetics</i> , 2016, 59, 32-38.	0.7	53
48	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.	1.4	108
49	A Potential Contributory Role for Ciliary Dysfunction in the 16p11.2 600 kb BP4-BP5 Pathology. <i>American Journal of Human Genetics</i> , 2015, 96, 784-796.	2.6	53
50	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.	0.7	11
51	The Influence of Microdeletions and Microduplications of 16p11.2 on Global Transcription Profiles. <i>Journal of Child Neurology</i> , 2015, 30, 1947-1953.	0.7	13
52	Platelet studies in autism spectrum disorder patients and first-degree relatives. <i>Molecular Autism</i> , 2015, 6, 57.	2.6	28
53	Refinement of the critical 2p25.3 deletion region: the role of MYT1L in intellectual disability and obesity. <i>Genetics in Medicine</i> , 2015, 17, 460-466.	1.1	45
54	Refining analyses of copy number variation identifies specific genes associated with developmental delay. <i>Nature Genetics</i> , 2014, 46, 1063-1071.	9.4	583

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55	Disruptive CHD8 Mutations Define a Subtype of Autism Early in Development. <i>Cell</i> , 2014, 158, 263-276.	13.5	637
56	Homozygous loss-of-function mutation in ALMS1 causes the lethal disorder mitogenic cardiomyopathy in two siblings. <i>European Journal of Medical Genetics</i> , 2014, 57, 532-535.	0.7	31
57	Homozygous and heterozygous disruptions of ANK3: at the crossroads of neurodevelopmental and psychiatric disorders. <i>Human Molecular Genetics</i> , 2013, 22, 1960-1970.	1.4	137
58	Multiple pilomatricomas with somatic <i>CTNNB1</i> mutations in children with constitutive mismatch repair deficiency. <i>Genes Chromosomes and Cancer</i> , 2013, 52, 656-664.	1.5	40
59	The East Flanders Prospective Twin Survey (EFPTS): An Actual Perception. <i>Twin Research and Human Genetics</i> , 2013, 16, 58-63.	0.3	34
60	Haploinsufficiency of ANKRD11 causes mild cognitive impairment, short stature and minor dysmorphisms. <i>European Journal of Human Genetics</i> , 2012, 20, 131-133.	1.4	45
61	Mutations in lectin complement pathway genes COLEC11 and MASP1 cause 3MC syndrome. <i>Nature Genetics</i> , 2011, 43, 197-203.	9.4	229
62	Narrowing the critical deletion region for autism spectrum disorders on 16p11.2. , 2011, 156, 243-245.		44
63	Time trends in the natural dizygotic twinning rate. <i>Human Reproduction</i> , 2011, 26, 2247-2252.	0.4	15
64	Aarskog syndrome: from prenatal features towards postnatal diagnosis. <i>Ultrasound</i> , 2011, 19, 107-109.	0.3	0
65	Focal preauricular dermal dysplasia: distinctive congenital lesions with a bilateral and symmetric distribution. <i>European Journal of Medical Genetics</i> , 2006, 49, 135-139.	0.7	11
66	Human laterality disorders. <i>European Journal of Medical Genetics</i> , 2006, 49, 349-362.	0.7	134
67	Two female siblings with congenital heart disease, postaxial polydactyly, ectopic neuropituitary gland, hair anomalies and characteristic facial features: a new syndrome?. <i>Clinical Dysmorphology</i> , 2006, 15, 71-74.	0.1	1
68	Sesn1 is a novel gene for left-right asymmetry and mediating nodal signaling. <i>Human Molecular Genetics</i> , 2006, 15, 3369-3377.	1.4	16
69	Recurrent involvement of chromosomal region 6q21 in heterotaxy. <i>American Journal of Medical Genetics Part A</i> , 2001, 103, 44-47.	2.4	11
70	Validation of a telephone zygosity questionnaire in twins of known zygosity. <i>Behavior Genetics</i> , 1998, 28, 159-163.	1.4	185