

# Peter D Turnpenny

## List of Publications by Year in descending order

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Version: 2024-02-01

53  
papers

3,114  
citations

218592

26  
h-index

182361

51  
g-index

58  
all docs

58  
docs citations

58  
times ranked

6242  
citing authors

#	ARTICLE	IF	CITATIONS
1	Mutations in the human Delta homologue, DLL3, cause axial skeletal defects in spondylocostal dysostosis. <i>Nature Genetics</i> , 2000, 24, 438-441.	9.4	362
2	Evidence for 28 genetic disorders discovered by combining healthcare and research data. <i>Nature</i> , 2020, 586, 757-762.	13.7	343
3	Alagille syndrome: pathogenesis, diagnosis and management. <i>European Journal of Human Genetics</i> , 2012, 20, 251-257.	1.4	319
4	Mutations in the histone methyltransferase gene KMT2B cause complex early-onset dystonia. <i>Nature Genetics</i> , 2017, 49, 223-237.	9.4	186
5	Quantifying the contribution of recessive coding variation to developmental disorders. <i>Science</i> , 2018, 362, 1161-1164.	6.0	158
6	Abnormal vertebral segmentation and the notch signaling pathway in man. <i>Developmental Dynamics</i> , 2007, 236, 1456-1474.	0.8	143
7	Discovery of four recessive developmental disorders using probabilistic genotype and phenotype matching among 4,125 families. <i>Nature Genetics</i> , 2015, 47, 1363-1369.	9.4	133
8	Delineation of two distinct 6p deletion syndromes. <i>Human Genetics</i> , 1999, 104, 64-72.	1.8	108
9	Polycystic Kidney Disease with Hyperinsulinemic Hypoglycemia Caused by a Promoter Mutation in Phosphomannomutase 2. <i>Journal of the American Society of Nephrology: JASN</i> , 2017, 28, 2529-2539.	3.0	99
10	Clinical and genetic aspects of KBG syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 2835-2846.	0.7	87
11	The Tatton-Brown-Rahman Syndrome: A clinical study of 55 individuals with de novo constitutive DNMT3A variants. <i>Wellcome Open Research</i> , 2018, 3, 46.	0.9	75
12	Whole exome sequencing in family trios reveals <i>de novo</i> mutations in <i>PURA</i> as a cause of severe neurodevelopmental delay and learning disability. <i>Journal of Medical Genetics</i> , 2014, 51, 806-813.	1.5	73
13	Recontacting patients in clinical genetics services: recommendations of the European Society of Human Genetics. <i>European Journal of Human Genetics</i> , 2019, 27, 169-182.	1.4	65
14	Chromosome 17q12 microdeletions but not intragenic HNF1B mutations link developmental kidney disease and psychiatric disorder. <i>Kidney International</i> , 2016, 90, 203-211.	2.6	64
15	Diagnosis of lethal or prenatal-onset autosomal recessive disorders by parental exome sequencing. <i>Prenatal Diagnosis</i> , 2018, 38, 33-43.	1.1	64
16	MYT1L mutations cause intellectual disability and variable obesity by dysregulating gene expression and development of the neuroendocrine hypothalamus. <i>PLoS Genetics</i> , 2017, 13, e1006957.	1.5	60
17	Fetal anticonvulsant syndrome and mutation in the maternal MTHFR gene. <i>Clinical Genetics</i> , 1999, 56, 216-220.	1.0	58
18	Opposite Modulation of RAC1 by Mutations in TRIO Is Associated with Distinct, Domain-Specific Neurodevelopmental Disorders. <i>American Journal of Human Genetics</i> , 2020, 106, 338-355.	2.6	58

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19	The CHD4-related syndrome: a comprehensive investigation of the clinical spectrum, genotype-phenotype correlations, and molecular basis. <i>Genetics in Medicine</i> , 2020, 22, 389-397.	1.1	53
20	An exome sequencing strategy to diagnose lethal autosomal recessive disorders. <i>European Journal of Human Genetics</i> , 2015, 23, 401-404.	1.4	51
21	De novo gain-of-function variants in <i>KCNT2</i> as a novel cause of developmental and epileptic encephalopathy. <i>Annals of Neurology</i> , 2018, 83, 1198-1204.	2.8	41
22	Recontact in clinical practice: a survey of clinical genetics services in the United Kingdom. <i>Genetics in Medicine</i> , 2016, 18, 876-881.	1.1	40
23	A 'joint venture' model of recontacting in clinical genomics: challenges for responsible implementation. <i>European Journal of Medical Genetics</i> , 2017, 60, 403-409.	0.7	36
24	Recontacting in clinical practice: an investigation of the views of healthcare professionals and clinical scientists in the United Kingdom. <i>European Journal of Human Genetics</i> , 2017, 25, 275-279.	1.4	35
25	Delineation of dominant and recessive forms of <i>LZTR1</i> -associated Noonan syndrome. <i>Clinical Genetics</i> , 2019, 95, 693-703.	1.0	35
26	Disruption of RFX family transcription factors causes autism, attention-deficit/hyperactivity disorder, intellectual disability, and dysregulated behavior. <i>Genetics in Medicine</i> , 2021, 23, 1028-1040.	1.1	34
27	Recontacting or not recontacting? A survey of current practices in clinical genetics centres in Europe. <i>European Journal of Human Genetics</i> , 2018, 26, 946-954.	1.4	33
28	Diagnosis and management of individuals with Fetal Valproate Spectrum Disorder; a consensus statement from the European Reference Network for Congenital Malformations and Intellectual Disability. <i>Orphanet Journal of Rare Diseases</i> , 2019, 14, 180.	1.2	33
29	Factors determining penetrance in familial atypical haemolytic uraemic syndrome. <i>Journal of Medical Genetics</i> , 2014, 51, 756-764.	1.5	28
30	Recontacting in clinical practice: the views and expectations of patients in the United Kingdom. <i>European Journal of Human Genetics</i> , 2017, 25, 1106-1112.	1.4	23
31	Bilateral microphthalmia, esophageal atresia, and cryptorchidism: The anophthalmia-esophageal-genital syndrome. , 1997, 70, 171-173.		22
32	Clinical findings of 21 previously unreported probands with <i>HNRNPU</i> -related syndrome and comprehensive literature review. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 1637-1654.	0.7	19
33	Primrose syndrome: Characterization of the phenotype in 42 patients. <i>Clinical Genetics</i> , 2020, 97, 890-901.	1.0	18
34	Missense Mutations of the Pro65 Residue of PCGF2 Cause a Recognizable Syndrome Associated with Craniofacial, Neurological, Cardiovascular, and Skeletal Features. <i>American Journal of Human Genetics</i> , 2018, 103, 786-793.	2.6	17
35	Recontacting in clinical genetics and genomic medicine? We need to talk about it. <i>European Journal of Human Genetics</i> , 2017, 25, 520-521.	1.4	16
36	Delineating the molecular and phenotypic spectrum of the SETD1B-related syndrome. <i>Genetics in Medicine</i> , 2021, 23, 2122-2137.	1.1	16

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37	Extending the clinical and genetic spectrum of ARID2 related intellectual disability. A case series of 7 patients. <i>European Journal of Medical Genetics</i> , 2019, 62, 27-34.	0.7	13
38	Bi-allelic variants in SPATA5L1 lead to intellectual disability, spastic-dystonic cerebral palsy, epilepsy, and hearing loss. <i>American Journal of Human Genetics</i> , 2021, 108, 2006-2016.	2.6	11
39	27 years of prenatal diagnosis for Huntington disease in the United Kingdom. <i>Genetics in Medicine</i> , 2019, 21, 1639-1643.	1.1	9
40	Of eponyms, acronyms and ... orthonyms. <i>Nature Reviews Genetics</i> , 2003, 4, 152-156.	7.7	8
41	Defective Somitogenesis and Abnormal Vertebral Segmentation in Man. <i>Advances in Experimental Medicine and Biology</i> , 2008, 638, 164-189.	0.8	8
42	Final Exon Frameshift Biallelic PTPN23 Variants Are Associated with Microcephalic Complex Hereditary Spastic Paraplegia. <i>Brain Sciences</i> , 2021, 11, 614.	1.1	5
43	Amniotic band sequence in paternal half-siblings with vascular Ehlers-Danlos syndrome. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 553-556.	0.7	5
44	Natural history of NF1 c.2970_2972del p.(Met992del): confirmation of a low risk of complications in a longitudinal study. <i>European Journal of Human Genetics</i> , 2022, 30, 291-297.	1.4	5
45	Dimensions of responsibility in medical genetics: exploring the complexity of the "duty to recontact". <i>New Genetics and Society</i> , 2018, 37, 187-206.	0.7	4
46	Delineating the Smith-Kingsmore syndrome phenotype: Investigation of 16 patients with the <i>MTOR</i> c.5395G>A p.(Glu1799Lys) missense variant. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 2445-2454.	0.7	4
47	Consolidating biallelic SDHD variants as a cause of mitochondrial complex II deficiency. <i>European Journal of Human Genetics</i> , 2021, 29, 1570-1576.	1.4	3
48	Further delineation of phenotypic spectrum of <i>SCN2A</i> -related disorder. <i>American Journal of Medical Genetics, Part A</i> , 2022, 188, 867-877.	0.7	3
49	Understanding the new <i>BRD4</i> -related syndrome: Clinical and genomic delineation with an international cohort study. <i>Clinical Genetics</i> , 2022, 102, 117-122.	1.0	3
50	Ectrodactyly-ectodermal dysplasia-clefting syndrome presenting with bilateral choanal atresia and rectal stenosis. <i>American Journal of Medical Genetics, Part A</i> , 2020, 182, 1939-1943.	0.7	2
51	Human Genetics and Genetic Enhancement. , 0, , 241-264.		1
52	From the Ottomans to the present day: 150 years of Scottish medical charity in the Holy Land. <i>BMJ, The</i> , 2013, 347, f6994-f6994.	3.0	0
53	Digesting GWAS. <i>Cellular and Molecular Gastroenterology and Hepatology</i> , 2016, 2, 542-543.	2.3	0