## Richard Caswell

List of Publications by Year in descending order

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471061 433756 1,767 34 17 31 citations h-index g-index papers 36 36 36 3778 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Activating germline mutations in STAT3 cause early-onset multi-organ autoimmune disease. Nature Genetics, 2014, 46, 812-814.	9.4	411
2	Recessive mutations in a distal PTF1A enhancer cause isolated pancreatic agenesis. Nature Genetics, 2014, 46, 61-64.	9.4	255
3	An in-frame deletion at the polymerase active site of POLD1 causes a multisystem disorder with lipodystrophy. Nature Genetics, 2013, 45, 947-950.	9.4	151
4	Germline or somatic GPR101 duplication leads to X-linked acrogigantism: a clinico-pathological and genetic study. Acta Neuropathologica Communications, 2016, 4, 56.	2.4	110
5	Polycystic Kidney Disease with Hyperinsulinemic Hypoglycemia Caused by a Promoter Mutation in Phosphomannomutase 2. Journal of the American Society of Nephrology: JASN, 2017, 28, 2529-2539.	3.0	99
6	<i>MAFA</i> missense mutation causes familial insulinomatosis and diabetes mellitus. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, 1027-1032.	3.3	88
7	Dominant ER Stress–Inducing <i>WFS1</i> Mutations Underlie a Genetic Syndrome of Neonatal/Infancy-Onset Diabetes, Congenital Sensorineural Deafness, and Congenital Cataracts. Diabetes, 2017, 66, 2044-2053.	0.3	77
8	A <i>CACNA1D</i> mutation in a patient with persistent hyperinsulinaemic hypoglycaemia, heart defects, and severe hypotonia. Pediatric Diabetes, 2017, 18, 320-323.	1.2	67
9	Diagnosis of lethal or prenatalâ€onset autosomal recessive disorders by parental exome sequencing. Prenatal Diagnosis, 2018, 38, 33-43.	1.1	64
10	An exome sequencing strategy to diagnose lethal autosomal recessive disorders. European Journal of Human Genetics, 2015, 23, 401-404.	1.4	51
11	Exome sequencing reveals a de novo POLD1 mutation causing phenotypic variability in mandibular hypoplasia, deafness, progeroid features, and lipodystrophy syndrome (MDPL). Metabolism: Clinical and Experimental, 2017, 71, 213-225.	1.5	43
12	A Specific CNOT1 Mutation Results in a Novel Syndrome of Pancreatic Agenesis and Holoprosencephaly through Impaired Pancreatic and Neurological Development. American Journal of Human Genetics, 2019, 104, 985-989.	2.6	43
13	Biallelic RFX6 mutations can cause childhood as well as neonatal onset diabetes mellitus. European Journal of Human Genetics, 2015, 23, 1744-1748.	1.4	34
14	De Novo Mutations in <i>EIF2B1</i> Affecting eIF2 Signaling Cause Neonatal/Early-Onset Diabetes and Transient Hepatic Dysfunction. Diabetes, 2020, 69, 477-483.	0.3	29
15	Analysis of cellâ€free fetal <scp>DNA</scp> for nonâ€invasive prenatal diagnosis in a family with neonatal diabetes. Diabetic Medicine, 2017, 34, 582-585.	1.2	27
16	Mitochondrial Retinopathy. Ophthalmology Retina, 2022, 6, 65-79.	1.2	26
17	Isolated Pancreatic Aplasia Due to a Hypomorphic <i>PTF1A</i> Mutation. Diabetes, 2016, 65, 2810-2815.	0.3	22
18	Classification and correlation of RYR2 missense variants in individuals with catecholaminergic polymorphic ventricular tachycardia reveals phenotypic relationships. Journal of Human Genetics, 2020, 65, 531-539.	1.1	20

#	Article	IF	CITATIONS
19	A comparison of mitochondrial DNA isolation methods in frozen post-mortem human brain tissueâ€"applications for studies of mitochondrial genetics in brain disorders. BioTechniques, 2015, 59, 241-246.	0.8	17
20	Missense Mutations of the Pro65 Residue of PCGF2 Cause a Recognizable Syndrome Associated with Craniofacial, Neurological, Cardiovascular, and Skeletal Features. American Journal of Human Genetics, 2018, 103, 786-793.	2.6	17
21	Hemizygous UBA5 missense mutation unmasks recessive disorder in a patient with infantile-onset encephalopathy, acquired microcephaly, small cerebellum, movement disorder and severe neurodevelopmental delay. European Journal of Medical Genetics, 2019, 62, 97-102.	0.7	15
22	Using Structural Analysis In Silico to Assess the Impact of Missense Variants in MEN1. Journal of the Endocrine Society, 2019, 3, 2258-2275.	0.1	14
23	Phenotype of CNTNAP1: a study of patients demonstrating a specific severe congenital hypomyelinating neuropathy with survival beyond infancy. European Journal of Human Genetics, 2018, 26, 796-807.	1.4	13
24	Homozygous Hypomorphic <i>HNF1A</i> Alleles Are a Novel Cause of Young-Onset Diabetes and Result in Sulfonylurea-Sensitive Diabetes. Diabetes Care, 2020, 43, 909-912.	4.3	13
25	<i><scp>SOS</scp>1</i> frameshift mutations cause pure mucosal neuroma syndrome, a clinical phenotype distinct from multiple endocrine neoplasia type 2B. Clinical Endocrinology, 2016, 84, 715-719.	1.2	11
26	Late-onset Pseudoxanthoma Elasticum Associated with a Hypomorphic ABCC6 Variant. American Journal of Ophthalmology, 2020, 218, 255-260.	1.7	11
27	A hypomorphic allele of SLC35D1 results in Schneckenbecken-like dysplasia. Human Molecular Genetics, 2019, 28, 3543-3551.	1.4	9
28	Copy number variation of <i>LINGO1</i> in familial dystonic tremor. Neurology: Genetics, 2019, 5, e307.	0.9	8
29	Refinement of the critical genomic region for congenital hyperinsulinismÂin the Chromosome 9p deletion syndrome. Wellcome Open Research, 2019, 4, 149.	0.9	5
30	An enhanced method for targeted next generation sequencing copy number variant detection using ExomeDepth. Wellcome Open Research, 0, 2, 49.	0.9	4
31	Refinement of the critical genomic region for hypoglycaemia in the Chromosome 9p deletion syndrome. Wellcome Open Research, 2019, 4, 149.	0.9	3
32	Mild MDPL in a patient with a novel de novo missense variant in the Cys-B region of POLD1. European Journal of Human Genetics, 2022, 30, 960-966.	1.4	2
33	Response to Comment on Misra et al. Homozygous Hypomorphic HNF1A Alleles Are a Novel Cause of Young-Onset Diabetes and Result in Sulfonylurea-Sensitive Diabetes. Diabetes Care 2020;43:909–912. Diabetes Care, 2020, 43, e155-e156.	4.3	0
34	Robinow syndrome in an extremely preterm infant: Novel homozygous ROR2 variant detected by rapid exome sequencing. American Journal of Medical Genetics, Part A, 2021, , .	0.7	0