Deborah J G Mackay

List of Publications by Year in Descending Order

Source: https://exaly.com/author-pdf/8659403/deborah-j-g-mackay-publications-by-year.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

80 5,670 38 75 g-index

84 6,527 7.5 ext. citations avg, IF 5.09 L-index

#	Paper	IF	Citations
80	A patient with multilocus imprinting disturbance involving hypomethylation at 11p15 and 14q32, and phenotypic features of Beckwith-Wiedemann and Temple syndromes <i>American Journal of Medical Genetics, Part A</i> , 2022 ,	2.5	O
79	Experiences of adolescents living with Silver-Russell syndrome. <i>Archives of Disease in Childhood</i> , 2021 , 106, 1195-1201	2.2	1
78	Whole-genome analysis as a diagnostic tool for patients referred for diagnosis of Silver-Russell syndrome: a real-world study. <i>Journal of Medical Genetics</i> , 2021 ,	5.8	4
77	Phenotype of genetically confirmed Silver-Russell syndrome beyond childhood. <i>Journal of Medical Genetics</i> , 2020 , 57, 683-691	5.8	6
76	Growth disrupting mutations in epigenetic regulatory molecules are associated with abnormalities of epigenetic aging. <i>Genome Research</i> , 2019 , 29, 1057-1066	9.7	20
75	A Diagnosis for All Rare Genetic Diseases: The Horizon and the Next Frontiers. <i>Cell</i> , 2019 , 177, 32-37	56.2	53
74	Discrepant molecular and clinical diagnoses in Beckwith-Wiedemann and Silver-Russell syndromes. <i>Genetical Research</i> , 2019 , 101, e3	1.1	9
73	Genetic diagnosis of subfertility: the impact of meiosis and maternal effects. <i>Journal of Medical Genetics</i> , 2019 , 56, 271-282	5.8	10
7 2	Lived experience of Silver-Russell syndrome: implications for management during childhood and into adulthood. <i>Archives of Disease in Childhood</i> , 2019 , 104, 76-82	2.2	7
71	Genetics, molar pregnancies and medieval ideas of monstrous births: the lump of flesh in. <i>Medical Humanities</i> , 2019 , 45, 2-9	1.4	1
70	The phenotypic variations of multi-locus imprinting disturbances associated with maternal-effect variants of NLRP5 range from overt imprinting disorder to apparently healthy phenotype. <i>Clinical Epigenetics</i> , 2019 , 11, 190	7.7	13
69	Genomic imprinting disorders: lessons on how genome, epigenome and environment interact. <i>Nature Reviews Genetics</i> , 2019 , 20, 235-248	30.1	151
68	Revisiting Wilms tumour surveillance in Beckwith-Wiedemann syndrome with IC2 methylation loss, reply. <i>European Journal of Human Genetics</i> , 2018 , 26, 471-472	5.3	7
67	Expert consensus document: Clinical and molecular diagnosis, screening and management of Beckwith-Wiedemann syndrome: an international consensus statement. <i>Nature Reviews Endocrinology</i> , 2018 , 14, 229-249	15.2	234
66	Maternal variants in and other maternal effect proteins are associated with multilocus imprinting disturbance in offspring. <i>Journal of Medical Genetics</i> , 2018 , 55, 497-504	5.8	66
65	Growth Hormone Improves Short-Term Growth in Patients with Temple Syndrome. <i>Hormone Research in Paediatrics</i> , 2018 , 90, 407-413	3.3	7
64	Human imprinting disorders: Principles, practice, problems and progress. <i>European Journal of Medical Genetics</i> , 2017 , 60, 618-626	2.6	35

(2014-2017)

63	Diagnosis and management of Silver-Russell syndrome: first international consensus statement. <i>Nature Reviews Endocrinology</i> , 2017 , 13, 105-124	15.2	224
62	Recent Advances in Imprinting Disorders. <i>Clinical Genetics</i> , 2017 , 91, 3-13	4	62
61	Maternal uniparental disomy of chromosome 20: a novel imprinting disorder of growth failure. <i>Genetics in Medicine</i> , 2016 , 18, 309-15	8.1	52
60	Temple syndrome as a result of isolated hypomethylation of the 14q32 imprinted DLK1/MEG3 region. <i>American Journal of Medical Genetics, Part A</i> , 2016 , 170A, 170-5	2.5	22
59	Prenatal molecular testing for Beckwith-Wiedemann and Silver-Russell syndromes: a challenge for molecular analysis and genetic counseling. <i>European Journal of Human Genetics</i> , 2016 , 24, 784-93	5.3	34
58	EMQN best practice guidelines for the molecular genetic testing and reporting of chromosome 11p15 imprinting disorders: Silver-Russell and Beckwith-Wiedemann syndrome. <i>European Journal of Human Genetics</i> , 2016 , 24, 1377-87	5.3	54
57	The effect of early, comprehensive genomic testing on clinical care in neonatal diabetes: an international cohort study. <i>Lancet, The</i> , 2015 , 386, 957-63	40	186
56	Role of noninsulin therapies alone or in combination in chromosome 6q24-related transient neonatal diabetes: sulfonylurea improves but does not always normalize insulin secretion. <i>Diabetes Care</i> , 2015 , 38, e86-7	14.6	18
55	Beyond the gene roundtable discussion. <i>Textual Practice</i> , 2015 , 29, 415-432	0.1	1
54	Very small deletions within the NESP55 gene in pseudohypoparathyroidism type 1b. <i>European Journal of Human Genetics</i> , 2015 , 23, 494-9	5.3	18
53	Mutations in NLRP5 are associated with reproductive wastage and multilocus imprinting disorders in humans. <i>Nature Communications</i> , 2015 , 6, 8086	17.4	102
52	Multilocus methylation defects in imprinting disorders. <i>Biomolecular Concepts</i> , 2015 , 6, 47-57	3.7	31
51	Neonatal diabetes in Ukraine: incidence, genetics, clinical phenotype and treatment. <i>Journal of Pediatric Endocrinology and Metabolism</i> , 2015 , 28, 1279-86	1.6	25
50	Transient Neonatal Diabetes Mellitus followed by recurrent asymptomatic hypoglycaemia: a case report. <i>BMC Pediatrics</i> , 2015 , 15, 200	2.6	7
49	A statistical method for single sample analysis of HumanMethylation450 array data: genome-wide methylation analysis of patients with imprinting disorders. <i>Clinical Epigenetics</i> , 2015 , 7, 48	7.7	12
48	Imprinting disorders: a group of congenital disorders with overlapping patterns of molecular changes affecting imprinted loci. <i>Clinical Epigenetics</i> , 2015 , 7, 123	7.7	115
47	Report and review of described associations of Mayer-Rokitansky-Kater-Hauser syndrome and Silver-Russell syndrome. <i>Journal of Paediatrics and Child Health</i> , 2015 , 51, 555-560	1.3	6
46	Genome-wide DNA methylation analysis of patients with imprinting disorders identifies differentially methylated regions associated with novel candidate imprinted genes. <i>Journal of Medical Genetics</i> , 2014 , 51, 229-38	5.8	70

45	A familial disorder of altered DNA-methylation. <i>Journal of Medical Genetics</i> , 2014 , 51, 407-12	5.8	22
44	Chromosome 6q24 transient neonatal diabetes mellitus and protein sensitive hyperinsulinaemic hypoglycaemia. <i>Journal of Pediatric Endocrinology and Metabolism</i> , 2014 , 27, 1065-9	1.6	1
43	Temple syndrome: improving the recognition of an underdiagnosed chromosome 14 imprinting disorder: an analysis of 51 published cases. <i>Journal of Medical Genetics</i> , 2014 , 51, 495-501	5.8	136
42	Clinical presentation of 6q24 transient neonatal diabetes mellitus (6q24 TNDM) and genotype-phenotype correlation in an international cohort of patients. <i>Diabetologia</i> , 2013 , 56, 758-62	10.3	90
41	Targeted methylation testing of a patient cohort broadens the epigenetic and clinical description of imprinting disorders. <i>American Journal of Medical Genetics, Part A</i> , 2013 , 161A, 2174-82	2.5	65
40	Transient neonatal diabetes, ZFP57, and hypomethylation of multiple imprinted loci: a detailed follow-up. <i>Diabetes Care</i> , 2013 , 36, 505-12	14.6	48
39	Insights into the molecular mechanism for type 2 diabetes susceptibility at the KCNQ1 locus from temporal changes in imprinting status in human islets. <i>Diabetes</i> , 2013 , 62, 987-92	0.9	87
38	Hepatoblastoma in a child with a paternally-inherited ABCC8 mutation and mosaic paternal uniparental disomy 11p causing focal congenital hyperinsulinism. <i>European Journal of Medical Genetics</i> , 2013 , 56, 114-7	2.6	16
37	Transient neonatal diabetes mellitus in a Turkish patient with three novel homozygous variants in the ZFP57 gene. <i>JCRPE Journal of Clinical Research in Pediatric Endocrinology</i> , 2013 , 5, 125-8	1.9	5
36	Mutation of HERC2 causes developmental delay with Angelman-like features. <i>Journal of Medical Genetics</i> , 2013 , 50, 65-73	5.8	50
35	Evidence for anticipation in Beckwith-Wiedemann syndrome. <i>European Journal of Human Genetics</i> , 2013 , 21, 1344-8	5.3	23
34	Familial occurrence of neonatal diabetes with duplications in chromosome 6q24: treatment with sulfonylurea and 40-yr follow-up. <i>Pediatric Diabetes</i> , 2012 , 13, 155-62	3.6	15
33	Imprinting Disorders of Early Childhood 2012 , 137-160		1
32	Beckwith-Wiedemann syndrome caused by maternally inherited mutation of an OCT-binding motif in the IGF2/H19-imprinting control region, ICR1. <i>European Journal of Human Genetics</i> , 2012 , 20, 240-3	5.3	51
31	Deletions and rearrangements of the H19/IGF2 enhancer region in patients with Silver-Russell syndrome and growth retardation. <i>Journal of Medical Genetics</i> , 2011 , 48, 308-11	5.8	33
30	An atypical case of hypomethylation at multiple imprinted loci. <i>European Journal of Human Genetics</i> , 2011 , 19, 360-2	5.3	33
29	Methylation analysis of 79 patients with growth restriction reveals novel patterns of methylation change at imprinted loci. <i>European Journal of Human Genetics</i> , 2010 , 18, 648-55	5.3	80
28	Incidence of neonatal diabetes in Austria-calculation based on the Austrian Diabetes Register. <i>Pediatric Diabetes</i> , 2010 , 11, 18-23	3.6	42

(2005-2010)

27	Epigenotype-phenotype correlations in Silver-Russell syndrome. <i>Journal of Medical Genetics</i> , 2010 , 47, 760-8	5.8	98
26	Further refinement of the critical minimal genetic region for the imprinting disorder 6q24 transient neonatal diabetes. <i>Diabetologia</i> , 2010 , 53, 2347-51	10.3	27
25	DNA hypomethylation, transient neonatal diabetes, and prune belly sequence in one of two identical twins. <i>European Journal of Pediatrics</i> , 2010 , 169, 207-13	4.1	19
24	Investigation of 90 patients referred for molecular cytogenetic analysis using aCGH uncovers previously unsuspected anomalies of imprinting. <i>American Journal of Medical Genetics, Part A</i> , 2010 , 152A, 1990-3	2.5	6
23	Transient neonatal diabetes mellitus type 1. <i>American Journal of Medical Genetics, Part C: Seminars in Medical Genetics</i> , 2010 , 154C, 335-42	3.1	81
22	Concurrent course of transient neonatal diabetes with cholestasis and paucity of interlobular bile ducts: a case report. <i>Pediatric and Developmental Pathology</i> , 2009 , 12, 417-20	2.2	6
21	Hypomethylation at multiple maternally methylated imprinted regions including PLAGL1 and GNAS loci in Beckwith-Wiedemann syndrome. <i>European Journal of Human Genetics</i> , 2009 , 17, 611-9	5.3	176
20	Hypomethylation of multiple imprinted loci in individuals with transient neonatal diabetes is associated with mutations in ZFP57. <i>Nature Genetics</i> , 2008 , 40, 949-51	36.3	417
19	Clinical characterisation of the multiple maternal hypomethylation syndrome in siblings. <i>European Journal of Human Genetics</i> , 2008 , 16, 453-61	5.3	46
18	Mosaic paternal uniparental isodisomy and an ABCC8 gene mutation in a patient with permanent neonatal diabetes and hemihypertrophy. <i>Diabetes</i> , 2008 , 57, 255-8	0.9	12
17	Zellweger syndrome resulting from maternal isodisomy of chromosome 1. <i>American Journal of Medical Genetics, Part A</i> , 2007 , 143A, 2172-7	2.5	20
16	Mutations in ATP-sensitive K+ channel genes cause transient neonatal diabetes and permanent diabetes in childhood or adulthood. <i>Diabetes</i> , 2007 , 56, 1930-7	0.9	273
15	Isolated imprinting mutation of the DLK1/GTL2 locus associated with a clinical presentation of maternal uniparental disomy of chromosome 14. <i>Journal of Medical Genetics</i> , 2007 , 44, 637-40	5.8	54
14	Permanent neonatal diabetes caused by dominant, recessive, or compound heterozygous SUR1 mutations with opposite functional effects. <i>American Journal of Human Genetics</i> , 2007 , 81, 375-82	11	161
13	Assessment of the role of common genetic variation in the transient neonatal diabetes mellitus (TNDM) region in type 2 diabetes: a comparative genomic and tagging single nucleotide polymorphism approach. <i>Diabetes</i> , 2006 , 55, 2272-6	0.9	10
12	Epimutation of the TNDM locus and the Beckwith-Wiedemann syndrome centromeric locus in individuals with transient neonatal diabetes mellitus. <i>Human Genetics</i> , 2006 , 119, 179-84	6.3	51
11	A maternal hypomethylation syndrome presenting as transient neonatal diabetes mellitus. <i>Human Genetics</i> , 2006 , 120, 262-9	6.3	139
10	Bisulphite sequencing of the transient neonatal diabetes mellitus DMR facilitates a novel diagnostic test but reveals no methylation anomalies in patients of unknown aetiology. <i>Human Genetics</i> , 2005 , 116, 255-61	6.3	51

9	Monozygous triplets discordant for transient neonatal diabetes mellitus and for imprinting of the TNDM differentially methylated region. <i>Human Genetics</i> , 2005 , 117, 398-401	6.3	12
8	Relapsing diabetes can result from moderately activating mutations in KCNJ11. <i>Human Molecular Genetics</i> , 2005 , 14, 925-34	5.6	165
7	Activating mutations in the gene encoding the ATP-sensitive potassium-channel subunit Kir6.2 and permanent neonatal diabetes. <i>New England Journal of Medicine</i> , 2004 , 350, 1838-49	59.2	930
6	Relaxation of imprinted expression of ZAC and HYMAI in a patient with transient neonatal diabetes mellitus. <i>Human Genetics</i> , 2002 , 110, 139-44	6.3	70
5	Maturity onset diabetes of the young (MODY) and early onset Type II diabetes are not caused by loss of imprinting at the transient neonatal diabetes (TNDM) locus. <i>Diabetologia</i> , 2001 , 44, 924	10.3	5
4	Characterization of the methylation-sensitive promoter of the imprinted ZAC gene supports its role in transient neonatal diabetes mellitus. <i>Journal of Biological Chemistry</i> , 2001 , 276, 18653-6	5.4	52
3	Transient neonatal diabetes: widening the understanding of the etiopathogenesis of diabetes. <i>Diabetes</i> , 2000 , 49, 1359-66	0.9	214
2	The RhoS progress: a potential role during neuritogenesis for the Rho family of GTPases. <i>Trends in Neurosciences</i> , 1995 , 18, 496-501	13.3	110
1	Growth disrupting mutations in epigenetic regulatory molecules are associated with abnormalities of epigenetic aging		2