

# Mathew J Wallis

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/5000867/publications.pdf>

Version: 2024-02-01

30  
papers

964  
citations

687363

13  
h-index

501196

28  
g-index

30  
all docs

30  
docs citations

30  
times ranked

2246  
citing authors

#	ARTICLE	IF	CITATIONS
1	Standardized practices for RNA diagnostics using clinically accessible specimens reclassifies 75% of putative splicing variants. <i>Genetics in Medicine</i> , 2022, 24, 130-145.	2.4	45
2	Any symptom, in any organ, at any age: A case report of multiple genetic diagnoses mimicking mitochondrial disease in an adult with kidney disease. <i>Nephrology</i> , 2022, , .	1.6	1
3	Population DNA screening for medically actionable disease risk in adults. <i>Medical Journal of Australia</i> , 2022, 216, 278-280.	1.7	10
4	Germline variants in tumor suppressor FBXW7 lead to impaired ubiquitination and a neurodevelopmental syndrome. <i>American Journal of Human Genetics</i> , 2022, 109, 601-617.	6.2	16
5	Heterozygous <i>PNPT1</i> Variants Cause Spinocerebellar Ataxia Type 25. <i>Annals of Neurology</i> , 2022, 92, 122-137.	5.3	8
6	Lymphedema distichiasis syndrome may be caused by <i>FOXC2</i> promoter-enhancer dissociation and disruption of a topological associated domain. <i>American Journal of Medical Genetics, Part A</i> , 2021, 185, 150-156.	1.2	4
7	Clinical impact of genomic testing in patients with suspected monogenic kidney disease. <i>Genetics in Medicine</i> , 2021, 23, 183-191.	2.4	70
8	The clinical utility of exome sequencing and extended bioinformatic analyses in adolescents and adults with a broad range of neurological phenotypes: an Australian perspective. <i>Journal of the Neurological Sciences</i> , 2021, 420, 117260.	0.6	16
9	Paediatric genomic testing: Navigating medicare rebatable genomic testing. <i>Journal of Paediatrics and Child Health</i> , 2021, 57, 477-483.	0.8	8
10	Cryopyrin-associated periodic syndrome: a treatable genetic inflammatory condition. <i>Practical Neurology</i> , 2021, 21, 424-426.	1.1	1
11	Small interstitial 9p24.3 deletions principally involving <i>KANK1</i> are likely benign copy number variants. <i>European Journal of Medical Genetics</i> , 2020, 63, 103618.	1.3	2
12	Fine-mapping of 150 breast cancer risk regions identifies 191 likely target genes. <i>Nature Genetics</i> , 2020, 52, 56-73.	21.4	120
13	Germline and Mosaic Variants in <i>PRKACA</i> and <i>PRKACB</i> Cause a Multiple Congenital Malformation Syndrome. <i>American Journal of Human Genetics</i> , 2020, 107, 977-988.	6.2	33
14	Genetic Kidney Disease in Southern Tasmania. <i>Kidney International Reports</i> , 2020, 5, 534-537.	0.8	1
15	Rapid Diagnosis of Spinocerebellar Ataxia 36 in a Three-Generation Family Using Short-Read Whole-Genome Sequencing Data. <i>Movement Disorders</i> , 2020, 35, 1675-1679.	3.9	12
16	Prospective Evaluation of the Utility of Whole Exome Sequencing in Dilated Cardiomyopathy. <i>Journal of the American Heart Association</i> , 2020, 9, e013346.	3.7	28
17	Homologous recombination DNA repair defects in <i>PALB2</i> -associated breast cancers. <i>Npj Breast Cancer</i> , 2019, 5, 23.	5.2	39
18	Pathogenic <i>WDFY3</i> variants cause neurodevelopmental disorders and opposing effects on brain size. <i>Brain</i> , 2019, 142, 2617-2630.	7.6	31

#	ARTICLE	IF	CITATIONS
19	A novel approach to offering additional genomic findingsâ€”A protocol to test a twoâ€”step approach in the healthcare system. <i>Journal of Genetic Counseling</i> , 2019, 28, 388-397.	1.6	14
20	Loss of TOP3B leads to increased R-loop formation and genome instability. <i>Open Biology</i> , 2019, 9, 190222.	3.6	46
21	Heritable DNA methylation marks associated with susceptibility to breast cancer. <i>Nature Communications</i> , 2018, 9, 867.	12.8	76
22	Surprisingly good outcome in antenatal diagnosis of severe hydrocephalus related to CCDC88C deficiency. <i>European Journal of Medical Genetics</i> , 2018, 61, 189-196.	1.3	9
23	A craniosynostosis massively parallel sequencing panel study in 309 Australian and New Zealand patients: findings and recommendations. <i>Genetics in Medicine</i> , 2018, 20, 1061-1068.	2.4	37
24	Blake's pouch cyst in 13q deletion syndrome: Posterior fossa malformations may occur due to disruption of multiple genes. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2442-2445.	1.2	8
25	A balanced paternal interchromosomal reciprocal insertion between 5q12.1q13.2 and 20p12.3p12.1 resulting in separate genetic conditions in two siblings. <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 1930-1933.	1.2	2
26	Fanconi anemia in 55â€”yearâ€”old identical twins first presenting as fatal postâ€”chemotherapy pancytopenia. <i>American Journal of Hematology</i> , 2016, 91, 1273-1276.	4.1	2
27	Dual genetic diagnoses: Atypical handâ€”footâ€”genital syndrome and developmental delay due to de novo mutations in <i>HOXA13</i> and <i>NRXN1</i> . <i>American Journal of Medical Genetics, Part A</i> , 2016, 170, 717-724.	1.2	11
28	COMMD1 is linked to the WASH complex and regulates endosomal trafficking of the copper transporter ATP7A. <i>Molecular Biology of the Cell</i> , 2015, 26, 91-103.	2.1	200
29	A severe phenotype of Gitelman syndrome with increased prostaglandin excretion and favorable response to indomethacin. <i>CKJ: Clinical Kidney Journal</i> , 2014, 7, 306-310.	2.9	13
30	CCDC22 deficiency in humans blunts activation of proinflammatory NF- $\kappa$ B signaling. <i>Journal of Clinical Investigation</i> , 2013, 123, 2244-2256.	8.2	101