

# Aurelien Trimouille

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4051090/publications.pdf>

Version: 2024-02-01

22  
papers

479  
citations

840776

11  
h-index

752698

20  
g-index

25  
all docs

25  
docs citations

25  
times ranked

955  
citing authors

#	ARTICLE	IF	CITATIONS
1	Phenotype and genotype of 87 patients with Mowat-Wilson syndrome and recommendations for care. <i>Genetics in Medicine</i> , 2018, 20, 965-975.	2.4	67
2	Solve-RD: systematic pan-European data sharing and collaborative analysis to solve rare diseases. <i>European Journal of Human Genetics</i> , 2021, 29, 1325-1331.	2.8	49
3	Neuroimaging findings in Mowat-Wilson syndrome: a study of 54 patients. <i>Genetics in Medicine</i> , 2017, 19, 691-700.	2.4	45
4	BLOC1S5 pathogenic variants cause a new type of Hermansky-Pudlak syndrome. <i>Genetics in Medicine</i> , 2020, 22, 1613-1622.	2.4	44
5	Major intra-familial phenotypic heterogeneity and incomplete penetrance due to a CACNA1A pathogenic variant. <i>European Journal of Medical Genetics</i> , 2019, 62, 103530.	1.3	34
6	Solving patients with rare diseases through programmatic reanalysis of genome-phenome data. <i>European Journal of Human Genetics</i> , 2021, 29, 1337-1347.	2.8	34
7	Further delineation of the phenotype caused by biallelic variants in the <i>WDR4</i> gene. <i>Clinical Genetics</i> , 2018, 93, 374-377.	2.0	33
8	<i>IL11RA</i> -related Crouzon-like autosomal recessive craniosynostosis in 10 new patients: Resemblances and differences. <i>Clinical Genetics</i> , 2018, 94, 373-380.	2.0	29
9	Functional and genetic analyses of <i>ZYG11B</i> provide evidences for its involvement in OAVS. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2020, 8, e1375.	1.2	21
10	Oculo-auriculo-vertebral spectrum: new genes and literature review on a complex disease. <i>Journal of Medical Genetics</i> , 2022, 59, 417-427.	3.2	20
11	A recurrent missense variant in <i>EYA3</i> gene is associated with oculo-auriculo-vertebral spectrum. <i>Human Genetics</i> , 2021, 140, 933-944.	3.8	14
12	Implication of folate deficiency in <i>CYP2U1</i> loss of function. <i>Journal of Experimental Medicine</i> , 2021, 218, .	8.5	13
13	<i>De novo</i> coding variants in the <i>AGO1</i> gene cause a neurodevelopmental disorder with intellectual disability. <i>Journal of Medical Genetics</i> , 2022, 59, 965-975.	3.2	13
14	Hereditary Mucoepithelial Dysplasia Results from Heterozygous Variants at p.Arg557 Mutational Hotspot in <i>SREBF1</i> , Encoding a Transcription Factor Involved in Cholesterol Homeostasis. <i>Journal of Investigative Dermatology</i> , 2020, 140, 1289-1292.e2.	0.7	12
15	Description of a family with X-linked oculo-auriculo-vertebral spectrum associated with polyalanine tract expansion in <i>ZIC3</i> . <i>Clinical Genetics</i> , 2020, 98, 384-389.	2.0	11
16	Rare variants in the <i>GABA<sub>A</sub></i> receptor subunit $\mu$ identified in patients with a wide spectrum of epileptic phenotypes. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2020, 8, e1388.	1.2	8
17	19p13 microduplications encompassing <i>NFIX</i> are responsible for intellectual disability, short stature and small head circumference. <i>European Journal of Human Genetics</i> , 2018, 26, 85-93.	2.8	7
18	A <i>MT-TL1</i> variant identified by whole exome sequencing in an individual with intellectual disability, epilepsy, and spastic tetraparesis. <i>European Journal of Human Genetics</i> , 2021, 29, 1359-1368.	2.8	7

#	ARTICLE	IF	CITATIONS
19	An in-frame deletion in BICD2 associated with a non-progressive form of SMALED. Clinical Neurology and Neurosurgery, 2018, 166, 1-3.	1.4	6
20	PADDAS syndrome associated with hair dysplasia caused by a de novo missense variant of PUM1. American Journal of Medical Genetics, Part A, 2019, 179, 1030-1033.	1.2	6
21	Deletion in 2q35 excluding the IHH gene leads to fetal severe limb anomalies and suggests a disruption of chromatin architecture. European Journal of Human Genetics, 2019, 27, 384-388.	2.8	3
22	Deletion of the transcription factor <i>SOX4</i> is implicated in syndromic nephroblastoma. Clinical Genetics, 2017, 92, 449-450.	2.0	2