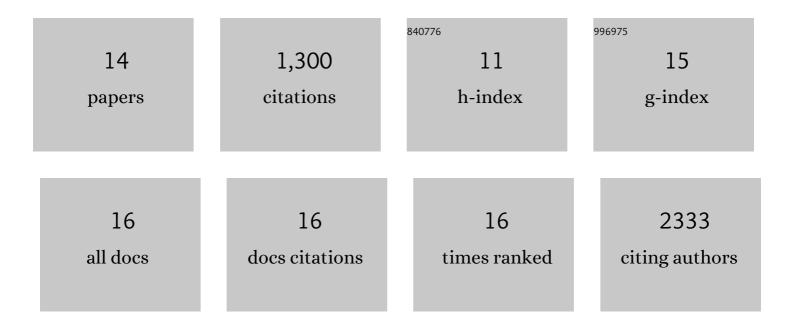
Claire S Leblond

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7199837/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Genetic correlates of phenotypic heterogeneity in autism. Nature Genetics, 2022, 54, 1293-1304.	21.4	51
2	Decreased phenol sulfotransferase activities associated with hyperserotonemia in autism spectrum disorders. Translational Psychiatry, 2021, 11, 23.	4.8	11
3	Operative list of genes associated with autism and neurodevelopmental disorders based on database review. Molecular and Cellular Neurosciences, 2021, 113, 103623.	2.2	51
4	Oligogenicity, C9orf72 expansion, and variant severity in ALS. Neurogenetics, 2020, 21, 227-242.	1.4	13
5	Social and non-social autism symptoms and trait domains are genetically dissociable. Communications Biology, 2019, 2, 328.	4.4	57
6	Both rare and common genetic variants contribute to autism in the Faroe Islands. Npj Genomic Medicine, 2019, 4, 1.	3.8	72
7	Somatic expansion of the C9orf72 hexanucleotide repeat does not occur in ALS spinal cord tissues. Neurology: Genetics, 2019, 5, e317.	1.9	8
8	Altered spinogenesis in iPSC-derived cortical neurons from patients with autism carrying de novo SHANK3 mutations. Scientific Reports, 2019, 9, 94.	3.3	51
9	Homozygous 2p11.2 deletion supports the implication of ELMOD3 in hearing loss and reveals the potential association of CAPG with ASD/ID etiology. Journal of Applied Genetics, 2019, 60, 49-56.	1.9	11
10	Heritability of the melatonin synthesis variability in autism spectrum disorders. Scientific Reports, 2017, 7, 17746.	3.3	28
11	A framework to identify contributing genes in patients with Phelan-McDermid syndrome. Npj Genomic Medicine, 2017, 2, 32.	3.8	58
12	11q24.2â€25 microâ€rearrangements in autism spectrum disorders: Relation to brain structures. American Journal of Medical Genetics, Part A, 2015, 167, 3019-3030.	1.2	25
13	Meta-analysis of SHANK Mutations in Autism Spectrum Disorders: A Gradient of Severity in Cognitive Impairments. PLoS Genetics, 2014, 10, e1004580.	3.5	501
14	Genetic and Functional Analyses of SHANK2 Mutations Suggest a Multiple Hit Model of Autism Spectrum Disorders. PLoS Genetics, 2012, 8, e1002521.	3.5	358