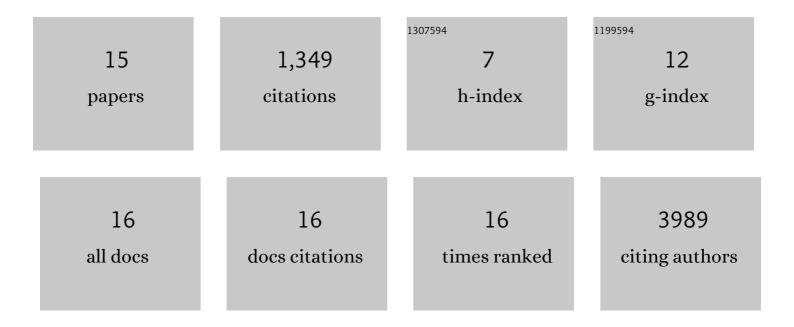
Jean McGowan-Jordan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/8977566/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Porokeratotic eccrine ostial and dermal duct nevus associated with an 11 megabase 3p deletion. Pediatric Dermatology, 2022, 39, 107-111.	0.9	0
2	Adopting High-Resolution Allele Frequencies Substantially Expedites Variant Interpretation in Genetic Diagnostic Laboratories. Journal of Molecular Diagnostics, 2019, 21, 602-611.	2.8	0
3	Genetic Diagnostic Testing for Inherited Cardiomyopathies. Journal of Molecular Diagnostics, 2019, 21, 437-448.	2.8	7
4	Atypical Hepatic Mesenchymal Hamartoma: Histologic Appearance, Immunophenotype, and Molecular Findings. Pediatric and Developmental Pathology, 2019, 22, 365-369.	1.0	3
5	Leveraging the power of new molecular technologies in the clinical setting requires unprecedented awareness of limitations and drawbacks: experience of one diagnostic laboratory. Journal of Medical Genetics, 2019, 56, 408-412.	3.2	3
6	Use of multicolor fluorescence in situ hybridization to detect deletions in clinical tissue sections. Laboratory Investigation, 2018, 98, 403-413.	3.7	10
7	Reinterpretation of sequence variants: one diagnostic laboratory's experience, and the need for standard guidelines. Genetics in Medicine, 2018, 20, 365-368.	2.4	28
8	Update on molecular findings in rhabdomyosarcoma. Pathology, 2017, 49, 238-246.	0.6	40
9	Variable developmental delays and characteristic facial features—A novel 7p22.3p22.2 microdeletion syndrome?. American Journal of Medical Genetics, Part A, 2017, 173, 1593-1600.	1.2	9
10	MDS with isolated del(5q) and internuclear bridging. Blood, 2017, 129, 2333-2333.	1.4	0
11	A rare case of pediatric lipoma with t(9;12)(p22;q14) and evidence of HMGA2-NFIB gene fusion. Cancer Genetics, 2017, 216-217, 100-104.	0.4	7
12	Genotype–phenotype characterization in 13 individuals with chromosome Xp11.22 duplications. American Journal of Medical Genetics, Part A, 2016, 170, 967-977.	1.2	11
13	CML with complex chromosome rearrangements and dysplastic megakaryocytes. Blood, 2016, 128, 604-604.	1.4	0
14	HGVS Recommendations for the Description of Sequence Variants: 2016 Update. Human Mutation, 2016, 37, 564-569.	2.5	1,194
15	Molecular Analysis of Cystinosis: Probable Irish Origin of the Most Common French Canadian Mutation. European Journal of Human Genetics, 1999, 7, 671-678.	2.8	35