## Georgia Vasileiou

List of Publications by Year in descending order

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840776 888059 17 670 11 17 citations h-index g-index papers 19 19 19 1491 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Manifestation of epilepsy in a patient with <scp><i>EED</i></scp> â€related overgrowth ( <scp>Cohen–Gibson</scp> syndrome). American Journal of Medical Genetics, Part A, 2022, 188, 292-297.	1.2	3
2	Loss-of-function variants in SRRM2 cause a neurodevelopmental disorder. Genetics in Medicine, 2022, 24, 1774-1780.	2.4	16
3	<scp><i>ZMYND11</i></scp> variants are a novel cause of centrotemporal and generalised epilepsies with neurodevelopmental disorder. Clinical Genetics, 2021, 100, 412-429.	2.0	5
4	BDV Syndrome: an Emerging Syndrome With Profound Obesity and Neurodevelopmental Delay Resembling Prader-Willi Syndrome. Journal of Clinical Endocrinology and Metabolism, 2021, 106, 3413-3427.	3.6	9
5	Targeted sequencing of FH-deficient uterine leiomyomas reveals biallelic inactivating somatic fumarase variants and allows characterization of missense variants. Modern Pathology, 2020, 33, 2341-2353.	5.5	19
6	Breast MRI texture analysis for prediction of BRCA-associated genetic risk. BMC Medical Imaging, 2020, 20, 86.	2.7	8
7	A biallelic truncating <i>AEBP1</i> variant causes connective tissue disorder in two siblings. American Journal of Medical Genetics, Part A, 2019, 179, 50-56.	1.2	11
8	CTCF variants in 39 individuals with a variable neurodevelopmental disorder broaden the mutational and clinical spectrum. Genetics in Medicine, 2019, 21, 2723-2733.	2.4	48
9	Prenatal diagnosis of <i>HNF1B</i> \$â€associated renal cysts: Is there a need to differentiate intragenic variants from 17q12 microdeletion syndrome?. Prenatal Diagnosis, 2019, 39, 1136-1147.	2.3	16
10	<i>TRIM28</i> haploinsufficiency predisposes to Wilms tumor. International Journal of Cancer, 2019, 145, 941-951.	5.1	45
11	Mutations in the BAF-Complex Subunit DPF2 Are Associated with Coffin-Siris Syndrome. American Journal of Human Genetics, 2018, 102, 468-479.	6.2	63
12	BRCA mutations and their influence on pathological complete response and prognosis in a clinical cohort of neoadjuvantly treated breast cancer patients. Breast Cancer Research and Treatment, 2018, 171, 85-94.	2.5	56
13	Addition of triple negativity of breast cancer as an indicator for germline mutations in predisposing genes increases sensitivity of clinical selection criteria. BMC Cancer, 2018, 18, 926.	2.6	16
14	Risk, Prediction and Prevention of Hereditary Breast Cancer – Large-Scale Genomic Studies in Times of Big and Smart Data. Geburtshilfe Und Frauenheilkunde, 2018, 78, 481-492.	1.8	38
15	Gene panel sequencing in familial breast/ovarian cancer patients identifies multiple novel mutations also in genes others than BRCA1/2. International Journal of Cancer, 2017, 140, 95-102.	5.1	99
16	Chromatin-Remodeling-Factor ARID1B Represses Wnt/ $\hat{l}^2$ -Catenin Signaling. American Journal of Human Genetics, 2015, 97, 445-456.	6.2	67
17	De Novo Mutations in the Genome Organizer CTCF Cause Intellectual Disability. American Journal of Human Genetics, 2013, 93, 124-131.	6.2	151