Elizabeth C Chao

List of Publications by Year
in descending order

12	1,098	933447 10	1199594 12
papers	citations	h-index	g-index

12	12	12	3597
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Enhanced utility of family-centered diagnostic exome sequencing with inheritance modelâ€“based analysis: results from 500 unselected families with undiagnosed genetic conditions. Genetics in Medicine, 2015, 17, 578-586.	2.4	401
2	Clinical laboratories collaborate to resolve differences in variant interpretations submitted to ClinVar. Genetics in Medicine, 2017, 19, 1096-1104.	2.4	200
3	Patient decisions for disclosure of secondary findings among the first 200 individuals undergoing clinical diagnostic exome sequencing. Genetics in Medicine, 2014, 16, 395-399.	2.4	99
4	Hereditary predisposition to ovarian cancer, looking beyond BRCA1/BRCA2. Gynecologic Oncology, 2015, 137, 86-92.	1.4	97
5	Update on molecular diagnosis of hereditary hemorrhagic telangiectasia. Human Genetics, 2010, 128, 61-77.	3.8	74
6	ELP2 is a novel gene implicated in neurodevelopmental disabilities. American Journal of Medical Genetics, Part A, 2015, 167, 1391-1395.	1.2	61
7	Congenital lethal motor neuron disease with a novel defect in ribosome biogenesis. Neurology, 2014, 82, 1322-1330.	1.1	50
8	Candidate-gene criteria for clinical reporting: diagnostic exome sequencing identifies altered candidate genes among 8% of patients with undiagnosed diseases. Genetics in Medicine, 2017, 19, 224-235.	2.4	47
9	Breast cancer risk is similar for CHEK2 founder and non-founder mutation carriers. Cancer Genetics, 2016, 209, 403-407.	0.4	41
10	Incidental detection of acquired variants in germline genetic and genomic testing: a points to consider statement of the American College of Medical Genetics and Genomics (ACMG). Genetics in Medicine, 2021, 23, 1179-1184.	2.4	13
11	The importance of proper bioinformatics analysis and clinical interpretation of tumor genomic profiling: a case study of undifferentiated sarcoma and a constitutional pathogenic BRCA2 mutation and an MLH1 variant of uncertain significance. Familial Cancer, 2015, 14, 481-485.	1.9	8
12	Clinical germline diagnostic exome sequencing for hereditary cancer: Findings within novel candidate genes are prevalent. Cancer Genetics, 2018, 224-225, 12-20.	0.4	7