

# Elizabeth C Chao

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/4350469/publications.pdf>

Version: 2024-02-01

12  
papers

1,098  
citations

933447

10  
h-index

1199594

12  
g-index

12  
all docs

12  
docs citations

12  
times ranked

3597  
citing authors

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