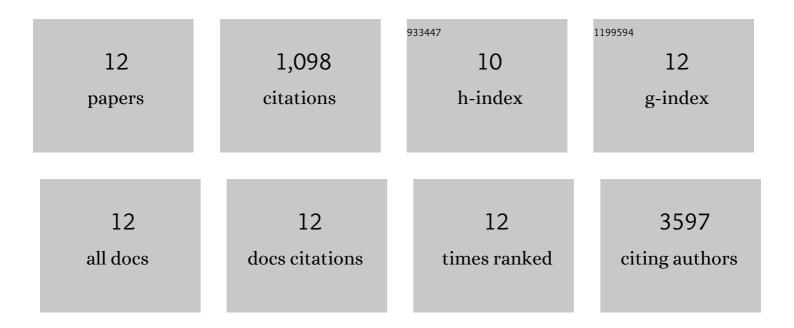
Elizabeth C Chao

List of Publications by Year in descending order

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| # | Article | IF | CITATIONS |
|----|---|-----|-----------|
| 1 | Enhanced utility of family-centered diagnostic exome sequencing with inheritance model–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 | <i>ELP2</i> 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 |