

Elika Sahlin

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3623514/publications.pdf>

Version: 2024-02-01

13
papers

395
citations

1039406

9
h-index

1125271

13
g-index

13
all docs

13
docs citations

13
times ranked

908
citing authors

#	ARTICLE	IF	CITATIONS
1	Integration of whole genome sequencing into a healthcare setting: high diagnostic rates across multiple clinical entities in 3219 rare disease patients. <i>Genome Medicine</i> , 2021, 13, 40.	3.6	116
2	Massive parallel sequencing in a family with rectal cancer. <i>Hereditary Cancer in Clinical Practice</i> , 2021, 19, 23.	0.6	3
3	Fetal HLA-G mediated immune tolerance and interferon response in preeclampsia. <i>EBioMedicine</i> , 2020, 59, 102872.	2.7	25
4	Pathogenic copy number variants are detected in a subset of patients with gastrointestinal malformations. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1084.	0.6	5
5	From cytogenetics to cytogenomics: whole-genome sequencing as a first-line test comprehensively captures the diverse spectrum of disease-causing genetic variation underlying intellectual disability. <i>Genome Medicine</i> , 2019, 11, 68.	3.6	88
6	Identification of putative pathogenic single nucleotide variants (SNVs) in genes associated with heart disease in 290 cases of stillbirth. <i>PLoS ONE</i> , 2019, 14, e0210017.	1.1	24
7	Knowledge and Attitudes Regarding Non-Invasive Prenatal Testing (NIPT) and Preferences for Risk Information among High School Students in Sweden. <i>Journal of Genetic Counseling</i> , 2017, 26, 447-454.	0.9	10
8	Positive Attitudes towards Non-Invasive Prenatal Testing (NIPT) in a Swedish Cohort of 1,003 Pregnant Women. <i>PLoS ONE</i> , 2016, 11, e0156088.	1.1	28
9	Fetal Calcifications Are Associated with Chromosomal Abnormalities. <i>PLoS ONE</i> , 2015, 10, e0123343.	1.1	9
10	Rare copy number variants are common in young children with autism spectrum disorder. <i>Acta Paediatrica, International Journal of Paediatrics</i> , 2015, 104, 610-618.	0.7	17
11	Amplification-free sequencing of cell-free DNA for prenatal non-invasive diagnosis of chromosomal aberrations. <i>Genomics</i> , 2015, 105, 150-158.	1.3	26
12	Mutation Screening and Array Comparative Genomic Hybridization Using a 180K Oligonucleotide Array in VACTERL Association. <i>PLoS ONE</i> , 2014, 9, e85313.	1.1	22
13	Molecular and Cytogenetic Analysis in Stillbirth: Results from 481 Consecutive Cases. <i>Fetal Diagnosis and Therapy</i> , 2014, 36, 326-332.	0.6	22