Ellika Sahlin

List of Publications by Year
in descending order

13 papers	395 citations	1040056 9 h-index	1125743 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. Genome Medicine, 2021, 13, 40.	8.2	116
2	Massive parallel sequencing in a family with rectal cancer. Hereditary Cancer in Clinical Practice, 2021, 19, 23.	1.5	3
3	Fetal HLA-G mediated immune tolerance and interferon response in preeclampsia. EBioMedicine, 2020, 59, 102872.	6.1	25
4	Pathogenic copy number variants are detected in a subset of patients with gastrointestinal malformations. Molecular Genetics & Genomic Medicine, 2020, 8, e1084.	1.2	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. Genome Medicine, 2019, 11, 68.	8.2	88
6	Identification of putative pathogenic single nucleotide variants (SNVs) in genes associated with heart disease in 290 cases of stillbirth. PLoS ONE, 2019, 14, e0210017.	2.5	24
7	Knowledge and Attitudes Regarding Nonâ€Invasive Prenatal Testing (NIPT) and Preferences for Risk Information among High School Students in Sweden. Journal of Genetic Counseling, 2017, 26, 447-454.	1.6	10
8	Positive Attitudes towards Non-Invasive Prenatal Testing (NIPT) in a Swedish Cohort of 1,003 Pregnant Women. PLoS ONE, 2016, 11, e0156088.	2.5	28
9	Fetal Calcifications Are Associated with Chromosomal Abnormalities. PLoS ONE, 2015, 10, e0123343.	2.5	9
10	Rare copy number variants are common in young children with autism spectrum disorder. Acta Paediatrica, International Journal of Paediatrics, 2015, 104, 610-618.	1.5	17
11	Amplification-free sequencing of cell-free DNA for prenatal non-invasive diagnosis of chromosomal aberrations. Genomics, 2015, 105, 150-158.	2.9	26
12	Mutation Screening and Array Comparative Genomic Hybridization Using a 180K Oligonucleotide Array in VACTERL Association. PLoS ONE, 2014, 9, e85313.	2.5	22
13	Molecular and Cytogenetic Analysis in Stillbirth: Results from 481 Consecutive Cases. Fetal Diagnosis and Therapy, 2014, 36, 326-332.	1.4	22