Johanna Lundin

List of Publications by Year
in descending order

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14 papers	275 citations	1307594 7 h-index	1058476 14 g-index
14 all docs	14 docs citations	14 times ranked	591 citing authors

#	Article	IF	Citations
1	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
2	22q11.2 microduplication in two patients with bladder exstrophy and hearing impairment. European Journal of Medical Genetics, 2010, 53, 61-65.	1.3	42
3	Haploinsufficiency of ZNF462 is associated with craniofacial anomalies, corpus callosum dysgenesis, ptosis, and developmental delay. European Journal of Human Genetics, 2017, 25, 946-951.	2.8	33
4	Replicative and non-replicative mechanisms in the formation of clustered CNVs are indicated by whole genome characterization. PLoS Genetics, 2018, 14, e1007780.	3.5	28
5	WNT3 involvement in human bladder exstrophy and cloaca development in zebrafish. Human Molecular Genetics, 2015, 24, 5069-5078.	2.9	23
6	Rare copy number variants contribute pathogenic alleles in patients with intestinal malrotation. Molecular Genetics & Genomic Medicine, 2019, 7, e549.	1.2	12
7	Evaluation of the ISL1 gene in the pathogenesis of bladder exstrophy in a Swedish cohort. Human Genome Variation, 2018, 5, 18009.	0.7	9
8	Further support linking the 22q11.2 microduplication to an increased risk of bladder exstrophy and highlighting LZTR1 as a candidate gene. Molecular Genetics & Genomic Medicine, 2019, 7, e666.	1.2	9
9	PHIP- a novel candidate breast cancer susceptibility locus on 6q14.1. Oncotarget, 2017, 8, 102769-102782.	1.8	9
10	Flanking complex copy number variants in the same family formed through unequal crossing-over during meiosis. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2018, 812, 1-4.	1.0	6
11	A Case with Bladder Exstrophy and Unbalanced X Chromosome Rearrangement. European Journal of Pediatric Surgery, 2014, 24, 353-359.	1.3	5
12	NEIL1 is a candidate gene associated with common variable immunodeficiency in a patient with a chromosome 15q24 deletion. Clinical Immunology, 2017, 176, 71-76.	3.2	5
13	Two novel colorectal cancer risk loci in the region on chromosome 9q22.32. Oncotarget, 2018, 9, 11170-11179.	1.8	4
14	Study on genetic stability in human urothelial cells in vitro. Journal of Tissue Engineering and Regenerative Medicine, 2018, 12, e720-e726.	2.7	2