

Johanna Lundin

List of Publications by Year in descending order

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14
papers

275
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1306789

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1058022

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times ranked

591
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#	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. <i>Genome Medicine</i> , 2019, 11, 68.	3.6	88
2	22q11.2 microduplication in two patients with bladder exstrophy and hearing impairment. <i>European Journal of Medical Genetics</i> , 2010, 53, 61-65.	0.7	42
3	Haploinsufficiency of ZNF462 is associated with craniofacial anomalies, corpus callosum dysgenesis, ptosis, and developmental delay. <i>European Journal of Human Genetics</i> , 2017, 25, 946-951.	1.4	33
4	Replicative and non-replicative mechanisms in the formation of clustered CNVs are indicated by whole genome characterization. <i>PLoS Genetics</i> , 2018, 14, e1007780.	1.5	28
5	WNT3 involvement in human bladder exstrophy and cloaca development in zebrafish. <i>Human Molecular Genetics</i> , 2015, 24, 5069-5078.	1.4	23
6	Rare copy number variants contribute pathogenic alleles in patients with intestinal malrotation. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e549.	0.6	12
7	Evaluation of the ISL1 gene in the pathogenesis of bladder exstrophy in a Swedish cohort. <i>Human Genome Variation</i> , 2018, 5, 18009.	0.4	9
8	Further support linking the 22q11.2 microduplication to an increased risk of bladder exstrophy and highlighting LZTR1 as a candidate gene. <i>Molecular Genetics & Genomic Medicine</i> , 2019, 7, e666.	0.6	9
9	<i>PHIP</i> - a novel candidate breast cancer susceptibility locus on 6q14.1. <i>Oncotarget</i> , 2017, 8, 102769-102782.	0.8	9
10	Flanking complex copy number variants in the same family formed through unequal crossing-over during meiosis. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2018, 812, 1-4.	0.4	6
11	A Case with Bladder Exstrophy and Unbalanced X Chromosome Rearrangement. <i>European Journal of Pediatric Surgery</i> , 2014, 24, 353-359.	0.7	5
12	NEIL1 is a candidate gene associated with common variable immunodeficiency in a patient with a chromosome 15q24 deletion. <i>Clinical Immunology</i> , 2017, 176, 71-76.	1.4	5
13	Two novel colorectal cancer risk loci in the region on chromosome 9q22.32. <i>Oncotarget</i> , 2018, 9, 11170-11179.	0.8	4
14	Study on genetic stability in human urothelial cells in vitro. <i>Journal of Tissue Engineering and Regenerative Medicine</i> , 2018, 12, e720-e726.	1.3	2