

Lennart F Johansson

List of Publications by Year in descending order

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Version: 2024-02-01

15
papers

552
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1040056

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940533

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docs citations

16
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1954
citing authors

#	ARTICLE	IF	CITATIONS
1	Targeted Next-Generation Sequencing can Replace Sanger Sequencing in Clinical Diagnostics. <i>Human Mutation</i> , 2013, 34, 1035-1042.	2.5	248
2	CoNVaDING: Single Exon Variation Detection in Targeted NGS Data. <i>Human Mutation</i> , 2016, 37, 457-464.	2.5	79
3	Whole-exome sequencing is a powerful approach for establishing the etiological diagnosis in patients with intellectual disability and microcephaly. <i>BMC Medical Genomics</i> , 2015, 9, 7.	1.5	65
4	Next-generation sequencing-based genome diagnostics across clinical genetics centers: implementation choices and their effects. <i>European Journal of Human Genetics</i> , 2015, 23, 1142-1150.	2.8	56
5	A next-generation sequencing method for gene doping detection that distinguishes low levels of plasmid DNA against a background of genomic DNA. <i>Gene Therapy</i> , 2019, 26, 338-346.	4.5	27
6	Successful Noninvasive Trisomy 18 Detection Using Single Molecule Sequencing. <i>Clinical Chemistry</i> , 2013, 59, 705-709.	3.2	11
7	NIPTRIC: an online tool for clinical interpretation of non-invasive prenatal testing (NIPT) results. <i>Scientific Reports</i> , 2016, 6, 38359.	3.3	10
8	Diagnostic yield of targeted next generation sequencing in 2002 Dutch cardiomyopathy patients. <i>International Journal of Cardiology</i> , 2021, 332, 99-104.	1.7	9
9	NIPTeR: an R package for fast and accurate trisomy prediction in non-invasive prenatal testing. <i>BMC Bioinformatics</i> , 2018, 19, 531.	2.6	7
10	Clinical Value of EGFR Copy Number Gain Determined by Amplicon-Based Targeted Next Generation Sequencing in Patients with EGFR-Mutated NSCLC. <i>Targeted Oncology</i> , 2021, 16, 215-226.	3.6	7
11	Detection of Fusion Genes to Determine Minimal Residual Disease in Leukemia Using Next-Generation Sequencing. <i>Clinical Chemistry</i> , 2020, 66, 1084-1092.	3.2	6
12	The ELIXIR Human Copy Number Variations Community: building bioinformatics infrastructure for research. <i>F1000Research</i> , 2020, 9, 1229.	1.6	5
13	Genetic Screening Test to Detect Translocations in Acute Leukemias by Use of Targeted Locus Amplification. <i>Clinical Chemistry</i> , 2018, 64, 1096-1103.	3.2	3
14	Targeted RNA-Sequencing Enables Detection of Relevant Translocations and Single Nucleotide Variants and Provides a Method for Classification of Hematological Malignanciesâ€“RANKING. <i>Clinical Chemistry</i> , 2020, 66, 1521-1530.	3.2	2
15	Low Detection Rates of Genetic FH in Cohort of Patients With Severe Hypercholesterolemia in the United Arab Emirates. <i>Frontiers in Genetics</i> , 2021, 12, 809256.	2.3	1