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List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/7183040/publications.pdf>

Version: 2024-02-01

23
papers

525
citations

759233

12
h-index

677142

22
g-index

23
all docs

23
docs citations

23
times ranked

939
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparison of gene fusion detection methods in salivary gland tumors. <i>Human Pathology</i> , 2022, 123, 1-10.	2.0	3
2	A Next-Generation Sequencing Test for Severe Congenital Neutropenia. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 200-211.	2.8	2
3	Clinical Implications of a Targeted RNA-Sequencing Panel in the Detection of Gene Fusions in Solid Tumors. <i>Journal of Molecular Diagnostics</i> , 2021, 23, 1749-1760.	2.8	5
4	BRAF mutations may identify a clinically distinct subset of glioblastoma. <i>Scientific Reports</i> , 2021, 11, 19999.	3.3	15
5	Utility of copy number variants in the classification of intracranial ependymoma. <i>Cancer Genetics</i> , 2020, 240, 66-72.	0.4	0
6	Pediatric meningioma: a clinicopathologic and molecular study with potential grading implications. <i>Brain Pathology</i> , 2020, 30, 1134-1143.	4.1	17
7	Impact of Reducing DNA Input on Next-Generation Sequencing Library Complexity and Variant Detection. <i>Journal of Molecular Diagnostics</i> , 2020, 22, 720-727.	2.8	9
8	Beyond Panel-Based Testing: Exome Analysis Increases Sensitivity for Diagnosis of Genetic Kidney Disease. <i>Kidney360</i> , 2020, 1, 772-780.	2.1	7
9	Optimization of Population Frequency Cutoffs for Filtering Common Germline Polymorphisms from Tumor-Only Next-Generation Sequencing Data. <i>Journal of Molecular Diagnostics</i> , 2019, 21, 903-912.	2.8	13
10	Diagnostic Utility of Next-Generation Sequencing for Disorders of Somatic Mosaicism: A Five-Year Cumulative Cohort. <i>American Journal of Human Genetics</i> , 2019, 105, 734-746.	6.2	23
11	Beyond sequence variation: assessment of copy number variation in adult glioblastoma through targeted tumor somatic profiling. <i>Human Pathology</i> , 2019, 86, 170-181.	2.0	24
12	Genomic heterogeneity of ALK fusion breakpoints in non-small-cell lung cancer. <i>Modern Pathology</i> , 2018, 31, 791-808.	5.5	79
13	Analysis of point mutations and copy number variation in Grade II and III meningioma. <i>Experimental and Molecular Pathology</i> , 2018, 105, 328-333.	2.1	18
14	Discriminating a common somatic ASXL1 mutation (c.1934dup; p.G646Wfs*12) from artifact in myeloid malignancies using NGS. <i>Leukemia</i> , 2018, 32, 1874-1878.	7.2	18
15	Direct experimental manipulation of intestinal cells in <i>Ascaris suum</i> , with minor influences on the global transcriptome. <i>International Journal for Parasitology</i> , 2017, 47, 271-279.	3.1	6
16	Helminth.net: expansions to Nematode.net and an introduction to Trematode.net. <i>Nucleic Acids Research</i> , 2015, 43, D698-D706.	14.5	58
17	Inter and intra-specific diversity of parasites that cause lymphatic filariasis. <i>Infection, Genetics and Evolution</i> , 2013, 14, 137-146.	2.3	34
18	Localization of Wolbachia-like gene transcripts and peptides in adult <i>Onchocerca flexuosa</i> worms indicates tissue specific expression. <i>Parasites and Vectors</i> , 2013, 6, 2.	2.5	15

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19	Absence of Wolbachia Endobacteria in <i>Chandlerella quisqualis</i> , an Avian Filarial Parasite. <i>Journal of Parasitology</i> , 2012, 98, 382-387.	0.7	9
20	Comparing the mitochondrial genomes of Wolbachia-dependent and independent filarial nematode species. <i>BMC Genomics</i> , 2012, 13, 145.	2.8	39
21	Transcriptomic and Proteomic Analyses of a Wolbachia-Free Filarial Parasite Provide Evidence of Trans-Kingdom Horizontal Gene Transfer. <i>PLoS ONE</i> , 2012, 7, e45777.	2.5	20
22	Endosymbiont DNA in Endobacteria-Free Filarial Nematodes Indicates Ancient Horizontal Genetic Transfer. <i>PLoS ONE</i> , 2010, 5, e11029.	2.5	105
23	<i>Brugia malayi</i> : Whole genome amplification for genomic characterization of filarial parasites. <i>Experimental Parasitology</i> , 2008, 119, 256-263.	1.2	6