

# Olga Jarinova

## List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/12058823/publications.pdf>

Version: 2024-02-01

21  
papers

1,068  
citations

840776

11  
h-index

839539

18  
g-index

22  
all docs

22  
docs citations

22  
times ranked

2594  
citing authors

#	ARTICLE	IF	CITATIONS
1	Interplay between probe design and test performance: overlap between genomic regions of interest, capture regions and high quality reference calls influence performance of WES-based assays. <i>Human Genetics</i> , 2021, 140, 289-297.	3.8	0
2	Harmonizing the Collection of Clinical Data on Genetic Testing Requisition Forms to Enhance Variant Interpretation in Hypertrophic Cardiomyopathy (HCM). <i>Journal of Molecular Diagnostics</i> , 2021, 23, 589-598.	2.8	5
3	Evidence-Based Assessment of Genes in Dilated Cardiomyopathy. <i>Circulation</i> , 2021, 144, 7-19.	1.6	213
4	Implementation of Epilepsy Multigene Panel Testing in Ontario, Canada. <i>Canadian Journal of Neurological Sciences</i> , 2020, 47, 61-68.	0.5	6
5	<i>ALU</i> transposition induces familial hypertrophic cardiomyopathy. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2020, 8, e951.	1.2	1
6	Adopting High-Resolution Allele Frequencies Substantially Expedites Variant Interpretation in Genetic Diagnostic Laboratories. <i>Journal of Molecular Diagnostics</i> , 2019, 21, 602-611.	2.8	0
7	Genetic Diagnostic Testing for Inherited Cardiomyopathies. <i>Journal of Molecular Diagnostics</i> , 2019, 21, 437-448.	2.8	7
8	Validation of bcbio-nextgen Pipeline Based on NextSeq500 Exome Sequencing. , 2019, , .		1
9	Leveraging the power of new molecular technologies in the clinical setting requires unprecedented awareness of limitations and drawbacks: experience of one diagnostic laboratory. <i>Journal of Medical Genetics</i> , 2019, 56, 408-412.	3.2	3
10	Data sharing as a national quality improvement program: reporting on BRCA1 and BRCA2 variant-interpretation comparisons through the Canadian Open Genetics Repository (COGR). <i>Genetics in Medicine</i> , 2018, 20, 294-302.	2.4	27
11	Reinterpretation of sequence variants: one diagnostic laboratory's experience, and the need for standard guidelines. <i>Genetics in Medicine</i> , 2018, 20, 365-368.	2.4	28
12	Scaling resolution of variant classification differences in ClinVar between 41 clinical laboratories through an outlier approach. <i>Human Mutation</i> , 2018, 39, 1641-1649.	2.5	50
13	Fragile X testing as a second-tier test. <i>Genetics in Medicine</i> , 2017, 19, 1380-1380.	2.4	7
14	Next-generation sequencing for diagnosis of rare diseases in the neonatal intensive care unit. <i>Cmaj</i> , 2016, 188, E254-E260.	2.0	86
15	Identification of a pathogenic <i>FTO</i> mutation by next-generation sequencing in a newborn with growth retardation and developmental delay. <i>Journal of Medical Genetics</i> , 2016, 53, 200-207.	3.2	50
16	Resolution of refractory hypotension and anuria in a premature newborn with loss of function of ACE. <i>American Journal of Medical Genetics, Part A</i> , 2015, 167, 1654-1658.	1.2	10
17	Regulatory variations in the era of next-generation sequencing: Implications for clinical molecular diagnostics. <i>Human Mutation</i> , 2012, 33, 1021-1030.	2.5	13
18	17p13.3 microduplications are associated with split-hand/foot malformation and long-bone deficiency (SHFLD). <i>European Journal of Human Genetics</i> , 2011, 19, 1144-1151.	2.8	32

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19	Functional Analysis of the Chromosome 9p21.3 Coronary Artery Disease Risk Locus. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2009, 29, 1671-1677.	2.4	350
20	Functional resolution of duplicated <i>hoxb5</i> genes in teleosts. <i>Development (Cambridge)</i> , 2008, 135, 3543-3553.	2.5	23
21	Regulatory Roles of Conserved Intergenic Domains in Vertebrate <i>Dlx</i> Bigene Clusters. <i>Genome Research</i> , 2003, 13, 533-543.	5.5	153