

Erfan Aref-Eshghi

List of Publications by Year in descending order

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Version: 2024-02-01

42
papers

1,686
citations

331670

21
h-index

302126

39
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44
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44
docs citations

44
times ranked

2385
citing authors

#	ARTICLE	IF	CITATIONS
1	Evaluation of DNA Methylation Episignatures for Diagnosis and Phenotype Correlations in 42 Mendelian Neurodevelopmental Disorders. <i>American Journal of Human Genetics</i> , 2020, 106, 356-370.	6.2	171
2	Genomic DNA Methylation Signatures Enable Concurrent Diagnosis and Clinical Genetic Variant Classification in Neurodevelopmental Syndromes. <i>American Journal of Human Genetics</i> , 2018, 102, 156-174.	6.2	135
3	Diagnostic Utility of Genome-wide DNA Methylation Testing in Genetically Unsolved Individuals with Suspected Hereditary Conditions. <i>American Journal of Human Genetics</i> , 2019, 104, 685-700.	6.2	125
4	Clinical Validation of Copy Number Variant Detection from Targeted Next-Generation Sequencing Panels. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 905-920.	2.8	104
5	Classification of osteoarthritis phenotypes by metabolomics analysis. <i>BMJ Open</i> , 2014, 4, e006286.	1.9	90
6	Clinical epigenomics: genome-wide DNA methylation analysis for the diagnosis of Mendelian disorders. <i>Genetics in Medicine</i> , 2021, 23, 1065-1074.	2.4	88
7	BAFopathies™ DNA methylation epi-signatures demonstrate diagnostic utility and functional continuum of Coffin-Siris and Nicolaides-Baraitser syndromes. <i>Nature Communications</i> , 2018, 9, 4885.	12.8	83
8	The defining DNA methylation signature of Kabuki syndrome enables functional assessment of genetic variants of unknown clinical significance. <i>Epigenetics</i> , 2017, 12, 923-933.	2.7	79
9	Gene domain-specific DNA methylation episignatures highlight distinct molecular entities of ADNP syndrome. <i>Clinical Epigenetics</i> , 2019, 11, 64.	4.1	71
10	Screening for genes that accelerate the epigenetic aging clock in humans reveals a role for the H3K36 methyltransferase NSD1. <i>Genome Biology</i> , 2019, 20, 146.	8.8	66
11	Peripheral blood epi-signature of Claes-Jensen syndrome enables sensitive and specific identification of patients and healthy carriers with pathogenic mutations in KDM5C. <i>Clinical Epigenetics</i> , 2018, 10, 21.	4.1	58
12	Relationship Between Blood Plasma and Synovial Fluid Metabolite Concentrations in Patients with Osteoarthritis. <i>Journal of Rheumatology</i> , 2015, 42, 859-865.	2.0	45
13	DNA methylation signatures in mendelian developmental disorders as a diagnostic bridge between genotype and phenotype. <i>Epigenomics</i> , 2019, 11, 563-575.	2.1	42
14	Genome-wide DNA methylation study of hip and knee cartilage reveals embryonic organ and skeletal system morphogenesis as major pathways involved in osteoarthritis. <i>BMC Musculoskeletal Disorders</i> , 2015, 16, 287.	1.9	41
15	Frameshift mutations at the C-terminus of HIST1H1E result in a specific DNA hypomethylation signature. <i>Clinical Epigenetics</i> , 2020, 12, 7.	4.1	40
16	Clinical Validation of a Genome-Wide DNA Methylation Assay for Molecular Diagnosis of Imprinting Disorders. <i>Journal of Molecular Diagnostics</i> , 2017, 19, 848-856.	2.8	39
17	Genomic DNA Methylation-Derived Algorithm Enables Accurate Detection of Malignant Prostate Tissues. <i>Frontiers in Oncology</i> , 2018, 8, 100.	2.8	38
18	Overexpression of MMP13 in human osteoarthritic cartilage is associated with the SMAD-independent TGF- β 2 signalling pathway. <i>Arthritis Research and Therapy</i> , 2015, 17, 264.	3.5	37

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19	Metabolomic analysis of human synovial fluid and plasma reveals that phosphatidylcholine metabolism is associated with both osteoarthritis and diabetes mellitus. <i>Metabolomics</i> , 2016, 12, 1.	3.0	37
20	De novo SMARCA2 variants clustered outside the helicase domain cause a new recognizable syndrome with intellectual disability and blepharophimosis distinct from Nicolaides-Baraitser syndrome. <i>Genetics in Medicine</i> , 2020, 22, 1838-1850.	2.4	31
21	Genome-wide DNA methylation and RNA analyses enable reclassification of two variants of uncertain significance in a patient with clinical Kabuki syndrome. <i>Human Mutation</i> , 2019, 40, 1684-1689.	2.5	27
22	Diagnostic Utility of Genome-Wide DNA Methylation Analysis in Mendelian Neurodevelopmental Disorders. <i>International Journal of Molecular Sciences</i> , 2020, 21, 9303.	4.1	23
23	Functional annotation of genomic variation: DNA methylation epigraphures in neurodevelopmental Mendelian disorders. <i>Human Molecular Genetics</i> , 2020, 29, R27-R32.	2.9	23
24	Identification of Dyslipidemic Patients Attending Primary Care Clinics Using Electronic Medical Record (EMR) Data from the Canadian Primary Care Sentinel Surveillance Network (CPCSSN) Database. <i>Journal of Medical Systems</i> , 2017, 41, 45.	3.6	22
25	Does the Prevalence of Dyslipidemias Differ between Newfoundland and the Rest of Canada? Findings from the Electronic Medical Records of the Canadian Primary Care Sentinel Surveillance Network. <i>Frontiers in Cardiovascular Medicine</i> , 2015, 2, 1.	2.4	21
26	Implementation of an NGS-based sequencing and gene fusion panel for clinical screening of patients with suspected hematologic malignancies. <i>European Journal of Haematology</i> , 2019, 103, 178-189.	2.2	21
27	Using Electronic Medical Record to Identify Patients With Dyslipidemia in Primary Care Settings: International Classification of Disease Code Matters From One Region to a National Database. <i>Biomedical Informatics Insights</i> , 2017, 9, 117822261668588.	4.6	18
28	SMAD3 Is Associated with the Total Burden of Radiographic Osteoarthritis: The Chingford Study. <i>PLoS ONE</i> , 2014, 9, e97786.	2.5	17
29	Single and mixed dyslipidaemia in Canadian primary care settings: findings from the Canadian primary care sentinel surveillance network database. <i>BMJ Open</i> , 2015, 5, e007954.	1.9	13
30	Low density lipoprotein cholesterol control status among Canadians at risk for cardiovascular disease: findings from the Canadian Primary Care Sentinel Surveillance Network Database. <i>Lipids in Health and Disease</i> , 2015, 14, 60.	3.0	11
31	SMAD3 Is Upregulated in Human Osteoarthritic Cartilage Independent of the Promoter DNA Methylation. <i>Journal of Rheumatology</i> , 2016, 43, 388-394.	2.0	10
32	Genetic and epigenetic profiling of BRCA1/2 in ovarian tumors reveals additive diagnostic yield and evidence of a genomic BRCA1/2 DNA methylation signature. <i>Journal of Human Genetics</i> , 2020, 65, 865-873.	2.3	10
33	Glucose-induced, duration-dependent genome-wide DNA methylation changes in human endothelial cells. <i>American Journal of Physiology - Cell Physiology</i> , 2020, 319, C268-C276.	4.6	10
34	Detection of a DNA Methylation Signature for the Intellectual Developmental Disorder, X-Linked, Syndromic, Armfield Type. <i>International Journal of Molecular Sciences</i> , 2021, 22, 1111.	4.1	10
35	Six-year time-trend analysis of dyslipidemia among adults in Newfoundland and Labrador: findings from the laboratory information system between 2009 and 2014. <i>Lipids in Health and Disease</i> , 2018, 17, 99.	3.0	5
36	The oncogenic roles of NTRK fusions and methods of molecular diagnosis. <i>Cancer Genetics</i> , 2021, 258-259, 110-119.	0.4	5

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37	Advanced approach for comprehensive mtDNA genome testing in mitochondrial disease. <i>Molecular Genetics and Metabolism</i> , 2022, 135, 93-101.	1.1	5
38	Epigenomic Mechanisms of Human Developmental Disorders. , 2018, , 837-859.		4
39	Genetic associations in community context: a mixed model approach identifies a functional variant in the RBP4 gene associated with HDL-C dyslipidemia. <i>BMC Medical Genetics</i> , 2018, 19, 205.	2.1	3
40	Clinical and technical assessment of MedExome vs. NGS panels in patients with suspected genetic disorders in Southwestern Ontario. <i>Journal of Human Genetics</i> , 2021, 66, 451-464.	2.3	2
41	A Novel TP53 Tandem Duplication in a Child with Li-Fraumeni Syndrome. <i>Journal of Physical Education and Sports Management</i> , 2022, , mcs.a006181.	1.2	2
42	Multiparametric magnetic resonance imaging of multifocal prostate cancer to reveal intra-prostatic genomic heterogeneity and novel radio-genomic correlates: Results of the Smarter Prostate Interventions and Therapeutics (SPIRIT) study.. <i>Journal of Clinical Oncology</i> , 2020, 38, 20-20.	1.6	0