

# Larisa M Haupt

## List of Publications by Year in descending order

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131  
papers

3,976  
citations

186265

28  
h-index

144013

57  
g-index

141  
all docs

141  
docs citations

141  
times ranked

6626  
citing authors

#	ARTICLE	IF	CITATIONS
1	Normal human mammary epithelial cells spontaneously escape senescence and acquire genomic changes. <i>Nature</i> , 2001, 409, 633-637.	27.8	604
2	Mitochondrial Genome Acquisition Restores Respiratory Function and Tumorigenic Potential of Cancer Cells without Mitochondrial DNA. <i>Cell Metabolism</i> , 2015, 21, 81-94.	16.2	582
3	Locked nucleic acid (LNA) single nucleotide polymorphism (SNP) genotype analysis and validation using real-time PCR. <i>Nucleic Acids Research</i> , 2004, 32, e55-e55.	14.5	143
4	Next Generation Sequencing Methods for Diagnosis of Epilepsy Syndromes. <i>Frontiers in Genetics</i> , 2018, 9, 20.	2.3	102
5	The osteogenic transcription factor Runx2 regulates components of the fibroblast growth factor/proteoglycan signaling axis in osteoblasts. <i>Journal of Cellular Biochemistry</i> , 2009, 107, 144-154.	2.6	87
6	Synergism between Wnt3a and Heparin Enhances Osteogenesis via a Phosphoinositide 3-Kinase/Akt/RUNX2 Pathway. <i>Journal of Biological Chemistry</i> , 2010, 285, 26233-26244.	3.4	86
7	Regulatory Mechanisms of Epigenetic miRNA Relationships in Human Cancer and Potential as Therapeutic Targets. <i>Cancers</i> , 2020, 12, 2922.	3.7	84
8	Review: Alternative Splicing (AS) of Genes As An Approach for Generating Protein Complexity. <i>Current Genomics</i> , 2013, 14, 182-194.	1.6	80
9	An epigenetic clock for human skeletal muscle. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2020, 11, 887-898.	7.3	70
10	The heparan sulfate proteoglycan (HSPG) glypican-3 mediates commitment of MC3T3-E1 cells toward osteogenesis. <i>Journal of Cellular Physiology</i> , 2009, 220, 780-791.	4.1	68
11	Human Mesenchymal Stem Cells Retain Multilineage Differentiation Capacity Including Neural Marker Expression after Extended In Vitro Expansion. <i>PLoS ONE</i> , 2015, 10, e0137255.	2.5	68
12	Genotypes of the MTHFR C677T and MTRR A66G genes act independently to reduce migraine disability in response to vitamin supplementation. <i>Pharmacogenetics and Genomics</i> , 2012, 22, 741-749.	1.5	65
13	Comparison of genomic DNA extraction techniques from whole blood samples: a time, cost and quality evaluation study. <i>Molecular Biology Reports</i> , 2012, 39, 5961-5966.	2.3	63
14	Cell surface heparan sulfate proteoglycans as novel markers of human neural stem cell fate determination. <i>Stem Cell Research</i> , 2016, 16, 92-104.	0.7	62
15	Sulfated glycosaminoglycans mediate the effects of FGF2 on the osteogenic potential of rat calvarial osteoprogenitor cells. <i>Journal of Cellular Physiology</i> , 2006, 209, 811-825.	4.1	57
16	Effect of Coffee Combining Green Coffee Bean Constituents with Typical Roasting Products on the Nrf2/ARE Pathway in Vitro and in Vivo. <i>Journal of Agricultural and Food Chemistry</i> , 2012, 60, 9631-9641.	5.2	51
17	Mesenchymal stem cells, neural lineage potential, heparan sulfate proteoglycans and the matrix. <i>Developmental Biology</i> , 2014, 388, 1-10.	2.0	49
18	A causal role for TRESK loss of function in migraine mechanisms. <i>Brain</i> , 2019, 142, 3852-3867.	7.6	49

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19	TNF and TNF receptor expression and insulin sensitivity in human omental and subcutaneous adipose tissue – influence of BMI and adipose distribution. <i>Diabetes and Vascular Disease Research</i> , 2006, 3, 26-33.	2.0	48
20	Exploiting Heparan Sulfate Proteoglycans in Human Neurogenesis – Controlling Lineage Specification and Fate. <i>Frontiers in Integrative Neuroscience</i> , 2017, 11, 28.	2.1	46
21	A Potential Epigenetic Marker Mediating Serum Folate and Vitamin B <sub>12</sub> Levels Contributes to the Risk of Ischemic Stroke. <i>BioMed Research International</i> , 2015, 2015, 1-4.	1.9	43
22	Osteogenic Differentiation of Murine Embryonic Stem Cells is Mediated by Fibroblast Growth Factor Receptors. <i>Stem Cells and Development</i> , 2007, 16, 305-318.	2.1	42
23	Heparan Sulfate Proteoglycans and Human Breast Cancer Epithelial Cell Tumorigenicity. <i>Journal of Cellular Biochemistry</i> , 2014, 115, 967-976.	2.6	42
24	Induction of antioxidative Nrf2 gene transcription by coffee in humans: depending on genotype?. <i>Molecular Biology Reports</i> , 2012, 39, 7155-7162.	2.3	41
25	Sustained release and osteogenic potential of heparan sulfate-doped fibrin glue scaffolds within a rat cranial model. <i>Journal of Molecular Histology</i> , 2007, 38, 425-433.	2.2	40
26	Analysis of the MTHFR C677T variant with migraine phenotypes. <i>BMC Research Notes</i> , 2010, 3, 213.	1.4	40
27	Loss of chromosomal integrity in human mammary epithelial cells subsequent to escape from senescence. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2001, 6, 235-243.	2.7	39
28	Meta-analysis of genome-wide DNA methylation and integrative omics of age in human skeletal muscle. <i>Journal of Cachexia, Sarcopenia and Muscle</i> , 2021, 12, 1064-1078.	7.3	37
29	Semaphorin – plexin signalling genes associated with human breast tumourigenesis. <i>Gene</i> , 2011, 489, 63-69.	2.2	28
30	Genetic association analysis of miRNA SNPs implicates MIR145 in breast cancer susceptibility. <i>BMC Medical Genetics</i> , 2015, 16, 107.	2.1	28
31	Effects of Dietary Folate Intake on Migraine Disability and Frequency. <i>Headache</i> , 2015, 55, 301-309.	3.9	28
32	Significant differences in gene expression of GABA receptors in peripheral blood leukocytes of migraineurs. <i>Gene</i> , 2011, 490, 32-36.	2.2	27
33	BDNF and TNF- $\alpha$ polymorphisms in memory. <i>Molecular Biology Reports</i> , 2013, 40, 5483-5490.	2.3	27
34	Methylome-wide association study of whole blood DNA in the Norfolk Island isolate identifies robust loci associated with age. <i>Aging</i> , 2017, 9, 753-768.	3.1	27
35	Stimulation of MMP-11 (stromelysin-3) expression in mouse fibroblasts by cytokines, collagen and co-culture with human breast cancer cell lines. <i>BMC Cancer</i> , 2004, 4, 40.	2.6	26
36	Investigation of Brain-Derived Neurotrophic Factor (BDNF) Gene Variants in Migraine. <i>Headache</i> , 2014, 54, 1184-1193.	3.9	26

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37	Association of a Notch 3 gene polymorphism with migraine susceptibility. <i>Cephalalgia</i> , 2011, 31, 264-270.	3.9	25
38	Association study of calcitonin gene-related polypeptide-alpha (CALCA) gene polymorphism with migraine. <i>Brain Research</i> , 2011, 1378, 119-124.	2.2	24
39	The human $\mu$ -opioid receptor gene polymorphism (A118G) is associated with head pain severity in a clinical cohort of female migraine with aura patients. <i>Journal of Headache and Pain</i> , 2012, 13, 513-519.	6.0	24
40	Association study of the calcitonin gene-related polypeptide-alpha (CALCA) and the receptor activity modifying 1 (RAMP1) genes with migraine. <i>Gene</i> , 2013, 515, 187-192.	2.2	24
41	Exome Sequencing Diagnoses X-Linked Moesin-Associated Immunodeficiency in a Primary Immunodeficiency Case. <i>Frontiers in Immunology</i> , 2018, 9, 420.	4.8	24
42	Genetic investigation of methylenetetrahydrofolate reductase (MTHFR) and catechol-O-methyl transferase (COMT) in multiple sclerosis. <i>Brain Research Bulletin</i> , 2006, 69, 327-331.	3.0	23
43	Association of a <i>GRIA3</i> Gene Polymorphism With Migraine in an Australian Case-Control Cohort. <i>Headache</i> , 2013, 53, 1245-1249.	3.9	22
44	Targeted next generation sequencing identifies novel NOTCH3 gene mutations in CADASIL diagnostics patients. <i>Human Genomics</i> , 2016, 10, 38.	2.9	21
45	Ion channelopathies and migraine pathogenesis. <i>Molecular Genetics and Genomics</i> , 2017, 292, 729-739.	2.1	21
46	Data defining markers of human neural stem cell lineage potential. <i>Data in Brief</i> , 2016, 7, 206-215.	1.0	20
47	Association of the microRNA-Single Nucleotide Polymorphism rs2910164 in miR146a with sporadic breast cancer susceptibility: A case control study. <i>Gene</i> , 2016, 576, 256-260.	2.2	20
48	Skeletal muscle methylome and transcriptome integration reveals profound sex differences related to muscle function and substrate metabolism. <i>Clinical Epigenetics</i> , 2021, 13, 202.	4.1	20
49	Development of an eight gene expression profile implicating human breast tumours of all grade. <i>Molecular Biology Reports</i> , 2012, 39, 3879-3892.	2.3	19
50	Epigenetics and migraine; complex mitochondrial interactions contributing to disease susceptibility. <i>Gene</i> , 2014, 543, 1-7.	2.2	19
51	BDNF Variants May Modulate Long-Term Visual Memory Performance in a Healthy Cohort. <i>International Journal of Molecular Sciences</i> , 2017, 18, 655.	4.1	19
52	The NRP1 migraine risk variant shows evidence of association with menstrual migraine. <i>Journal of Headache and Pain</i> , 2018, 19, 31.	6.0	19
53	Heritability and genome-wide linkage analysis of migraine in the genetic isolate of Norfolk Island. <i>Gene</i> , 2012, 494, 119-123.	2.2	18
54	Glycosaminoglycan and growth factor mediated murine calvarial cell proliferation. <i>Journal of Molecular Histology</i> , 2007, 38, 415-424.	2.2	17

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55	Association of heparan sulfate proteoglycans SDC1 and SDC4 polymorphisms with breast cancer in an Australian Caucasian population. <i>Tumor Biology</i> , 2015, 36, 1731-1738.	1.8	17
56	Dysregulated MicroRNA Expression Profiles and Potential Cellular, Circulating and Polymorphic Biomarkers in Non-Hodgkin Lymphoma. <i>Genes</i> , 2016, 7, 130.	2.4	17
57	Stromal fibroblasts regulate microvascular-like network architecture in a bioengineered breast tumour angiogenesis model. <i>Acta Biomaterialia</i> , 2020, 114, 256-269.	8.3	17
58	Comprehensive Exonic Sequencing of Hemiplegic Migraine-Related Genes in a Cohort of Suspected Proband Identifies Known and Potential Pathogenic Variants. <i>Cells</i> , 2020, 9, 2368.	4.1	17
59	Investigation of polymorphisms in genes involved in estrogen metabolism in menstrual migraine. <i>Gene</i> , 2017, 607, 36-40.	2.2	15
60	Genome wide association study of response to interval and continuous exercise training: the Predict-HIT study. <i>Journal of Biomedical Science</i> , 2021, 28, 37.	7.0	15
61	Analysis of chromosome 1 microsatellite markers and the FHM2-ATP1A2 gene mutations in migraine pedigrees. <i>Neurological Research</i> , 2005, 27, 647-652.	1.3	14
62	In vitro and in vivo MMP gene expression localisation by In Situ-RT-PCR in cell culture and paraffin embedded human breast cancer cell line xenografts. <i>BMC Cancer</i> , 2006, 6, 18.	2.6	14
63	Analysis of 3 common polymorphisms in the KCNK18 gene in an Australian Migraine Case-control cohort. <i>Gene</i> , 2013, 528, 343-346.	2.2	14
64	Genetic polymorphisms in miRNAs targeting the estrogen receptor and their effect on breast cancer risk. <i>Meta Gene</i> , 2014, 2, 226-236.	0.6	14
65	Genetic variants associated with exercise performance in both moderately trained and highly trained individuals. <i>Molecular Genetics and Genomics</i> , 2020, 295, 515-523.	2.1	14
66	Heparanase Promotes Syndecan-1 Expression to Mediate Fibrillar Collagen and Mammographic Density in Human Breast Tissue Cultured ex vivo. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 599.	3.7	14
67	Temporal and functional changes in glycosaminoglycan expression during osteogenesis. <i>Journal of Molecular Histology</i> , 2007, 38, 469-481.	2.2	13
68	Role of the apolipoprotein E and catechol-O-methyltransferase genes in prospective and retrospective memory traits. <i>Gene</i> , 2012, 506, 135-140.	2.2	13
69	Computational epigenetic profiling of CpG islets in MTHFR. <i>Molecular Biology Reports</i> , 2014, 41, 8285-8292.	2.3	13
70	Tiered analysis of whole-exome sequencing for epilepsy diagnosis. <i>Molecular Genetics and Genomics</i> , 2020, 295, 751-763.	2.1	13
71	Association Study of MTHFD1 Coding Polymorphisms R134K and R653Q With Migraine Susceptibility. <i>Headache</i> , 2014, 54, 1506-1514.	3.9	12
72	Genome-wide allele-specific methylation is enriched at gene regulatory regions in a multi-generation pedigree from the Norfolk Island isolate. <i>Epigenetics and Chromatin</i> , 2019, 12, 60.	3.9	12

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73	Investigating diagnostic sequencing techniques for CADASIL diagnosis. <i>Human Genomics</i> , 2020, 14, 2.	2.9	12
74	Syndecan-1 Facilitates the Human Mesenchymal Stem Cell Osteo-Adipogenic Balance. <i>International Journal of Molecular Sciences</i> , 2020, 21, 3884.	4.1	12
75	Ion torrent high throughput mitochondrial genome sequencing (HTMGS). <i>PLoS ONE</i> , 2019, 14, e0224847.	2.5	11
76	Long-Term Consumption of Anthocyanin-Rich Fruit Juice: Impact on Gut Microbiota and Antioxidant Markers in Lymphocytes of Healthy Males. <i>Antioxidants</i> , 2021, 10, 27.	5.1	11
77	Purification and characterization of heparan sulfate from human primary osteoblasts. <i>Journal of Cellular Biochemistry</i> , 2009, 108, 1132-1142.	2.6	10
78	Investigation of Two Wnt Signalling Pathway Single Nucleotide Polymorphisms in a Breast Cancer-Affected Australian Population. <i>Twin Research and Human Genetics</i> , 2011, 14, 562-567.	0.6	10
79	Heparan Sulfate Proteoglycans as Drivers of Neural Progenitors Derived From Human Mesenchymal Stem Cells. <i>Frontiers in Molecular Neuroscience</i> , 2018, 11, 134.	2.9	10
80	Three-Dimensional Models as a New Frontier for Studying the Role of Proteoglycans in the Normal and Malignant Breast Microenvironment. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 569454.	3.7	10
81	Using Monozygotic Twins to Dissect Common Genes in Posttraumatic Stress Disorder and Migraine. <i>Frontiers in Neuroscience</i> , 2021, 15, 678350.	2.8	10
82	Single Nucleotide Polymorphisms in MIR143 Contribute to Protection Against Non-Hodgkin Lymphoma (NHL) in Caucasian Populations. <i>Genes</i> , 2019, 10, 185.	2.4	9
83	Investigation of the 1758G>C and 2880A>G variants within the NCOA3 gene in a breast cancer affected Australian population. <i>Gene</i> , 2011, 482, 68-72.	2.2	8
84	Confirmation that Xq27 and Xq28 are susceptibility loci for migraine in independent pedigrees and a case-control cohort. <i>Neurogenetics</i> , 2012, 13, 97-101.	1.4	8
85	Investigation of lymphotoxin 1± genetic variants in migraine. <i>Gene</i> , 2013, 512, 527-531.	2.2	8
86	Genetic Variation in Cytokine-Related Genes and Migraine Susceptibility. <i>Twin Research and Human Genetics</i> , 2013, 16, 1079-1086.	0.6	8
87	Genetic Analysis of <sc>GRIA2</sc> and <sc>GRIA4</sc> Genes in Migraine. <i>Headache</i> , 2014, 54, 303-312.	3.9	8
88	In silico analyses reveal common cellular pathways affected by loss of heterozygosity (LOH) events in the lymphomagenesis of Non-Hodgkinâ€™s lymphoma (NHL). <i>BMC Genomics</i> , 2014, 15, 390.	2.8	8
89	Genetic and epigenetic variants in the MTHFR gene are not associated with non-Hodgkin lymphoma. <i>Meta Gene</i> , 2015, 6, 91-95.	0.6	8
90	An emerging role for epigenetic factors in relation to executive function. <i>Briefings in Functional Genomics</i> , 2018, 17, 170-180.	2.7	8

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91	Comprehensive Exonic Sequencing of Known Ataxia Genes in Episodic Ataxia. <i>Biomedicines</i> , 2020, 8, 134.	3.2	8
92	HSPGs glypicanâ€1 and glypicanâ€4 are human neuronal proteins characteristic of different neural phenotypes. <i>Journal of Neuroscience Research</i> , 2020, 98, 1619-1645.	2.9	8
93	Epigenetic Regulation of miR-92a and TET2 and Their Association in Non-Hodgkin Lymphoma. <i>Frontiers in Genetics</i> , 2021, 12, 768913.	2.3	8
94	Matrix metalloproteinase localisation by in situ-RT-PCR in archival human breast biopsy material. <i>Molecular and Cellular Probes</i> , 2008, 22, 83-89.	2.1	7
95	A CREB1 Gene Polymorphism (rs2253206) Is Associated with Prospective Memory in a Healthy Cohort. <i>Frontiers in Behavioral Neuroscience</i> , 2017, 11, 86.	2.0	7
96	Investigating the influence of mtDNA and nuclear encoded mitochondrial variants on high intensity interval training outcomes. <i>Scientific Reports</i> , 2020, 10, 11089.	3.3	7
97	Association of the SNP rs2623047 in the HSPG modification enzyme SULF1 with an Australian Caucasian Breast Cancer Cohort. <i>Gene</i> , 2014, 547, 50-54.	2.2	6
98	Case-control study of ADARB1 and ADARB2 gene variants in migraine. <i>Journal of Headache and Pain</i> , 2015, 16, 511.	6.0	6
99	Geneâ€centric analysis implicates nuclear encoded mitochondrial protein gene variants in migraine susceptibility. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2017, 5, 157-163.	1.2	6
100	Exploring Neuronal Vulnerability to Head Trauma Using a Whole Exome Approach. <i>Journal of Neurotrauma</i> , 2020, 37, 1870-1879.	3.4	6
101	Genetic Association Analysis Implicates Six MicroRNA-Related SNPs With Increased Risk of Breast Cancer in Australian Caucasian Women. <i>Clinical Breast Cancer</i> , 2021, 21, e694-e703.	2.4	6
102	Gene Expression Profiling in Human Breast Cancer - Toward Personalised Therapeutics?â€!2009-04-21~!2010-02-19~!2010-07-06~!. <i>Open Breast Cancer Journal</i> , 2010, 2, 46-59.	0.2	6
103	Current Understanding of DNA Methylation and Age-related Disease. , 2018, 2, 1-1.		6
104	Heparan Sulfate Proteoglycans, Tumour Progression and the Cancer Stem Cell Niche. <i>Current Cancer Therapy Reviews</i> , 2009, 5, 256-260.	0.3	5
105	Potential antioxidant response to coffee â€ A matter of genotype?. <i>Meta Gene</i> , 2014, 2, 525-539.	0.6	5
106	Methylenetetrahydrofolate Reductase CpG Islands: Epigenotyping. <i>Journal of Clinical Laboratory Analysis</i> , 2016, 30, 335-344.	2.1	5
107	Variant Call Formatâ€Diagnostic Annotation and Reporting Tool. <i>Journal of Molecular Diagnostics</i> , 2019, 21, 951-960.	2.8	5
108	Investigation of APOE isoforms and the association between APOE E3 and E4 with migraine in the Australian Caucasian population. <i>NeuroReport</i> , 2013, 24, 499-503.	1.2	4

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109	Saliva as a comparable-quality source of DNA for Whole Exome Sequencing on Ion platforms. <i>Genomics</i> , 2020, 112, 1437-1443.	2.9	4
110	An investigation of genetic polymorphisms in heparan sulfate proteoglycan core proteins and key modification enzymes in an Australian Caucasian multiple sclerosis population. <i>Human Genomics</i> , 2020, 14, 18.	2.9	4
111	Mini review: genome and transcriptome editing using CRISPR-cas systems for haematological malignancy gene therapy. <i>Transgenic Research</i> , 2021, 30, 129-141.	2.4	4
112	Three-Dimensional Human Neural Stem Cell Models to Mimic Heparan Sulfate Proteoglycans and the Neural Niche. <i>Seminars in Thrombosis and Hemostasis</i> , 2021, 47, 308-315.	2.7	4
113	A genome-wide methylation study of body fat traits in the Norfolk Island isolate. <i>Nutrition, Metabolism and Cardiovascular Diseases</i> , 2021, 31, 1556-1563.	2.6	4
114	Pedigree derived mutation rate across the entire mitochondrial genome of the Norfolk Island population. <i>Scientific Reports</i> , 2022, 12, 6827.	3.3	4
115	A combinatorial in silico approach for microRNA-target identification: Order out of chaos. <i>Biochimie</i> , 2021, 187, 121-130.	2.6	3
116	Investigation of Mitochondrial Related Variants in a Cerebral Small Vessel Disease Cohort. <i>Molecular Neurobiology</i> , 2022, 59, 5366-5378.	4.0	3
117	ISâ€RTâ€PCR assay detection of MTâ€MMP in a human breast cancer cell line. <i>IUBMB Life</i> , 1996, 39, 553-561.	3.4	2
118	Techniques for RNA extraction from cells cultured in starPEGâ€heparin hydrogels. <i>Open Biology</i> , 2021, 11, 200388.	3.6	2
119	Mechanical Pressure Driving Proteoglycan Expression in Mammographic Density: a Self-perpetuating Cycle?. <i>Journal of Mammary Gland Biology and Neoplasia</i> , 2021, 26, 277-296.	2.7	2
120	Label-free isolation and cultivation of patient-matched human mammary epithelial and stromal cells from normal breast tissue. <i>European Journal of Cell Biology</i> , 2021, 100, 151187.	3.6	2
121	Association of polymorphisms in <i>ARRB2</i> and clinical response to methadone for pain in advanced cancer. <i>Pharmacogenomics</i> , 2022, 23, 281-289.	1.3	2
122	Syndecan-1 and -4 influence Wnt signaling and cell migration in human breast cancers. <i>Biochimie</i> , 2022, 198, 60-75.	2.6	2
123	Expression QTL analysis of glaucoma endophenotypes in the Norfolk Island isolate provides evidence that immune-related genes are associated with optic disc size. <i>Journal of Human Genetics</i> , 2018, 63, 83-87.	2.3	1
124	Investigation of the <i>CADM2</i> polymorphism rs17518584 in memory and executive functions measures in a cohort of young healthy individuals. <i>Neurobiology of Learning and Memory</i> , 2018, 155, 330-336.	1.9	1
125	Development of an accurate genomic ancestry prediction strategy to enable the accounting of Australian and Japanese historical military remains. <i>Australian Journal of Forensic Sciences</i> , 2022, 54, 416-436.	1.2	1
126	Proteoglycans, Neurogenesis and Stem Cell Differentiation. <i>Biology of Extracellular Matrix</i> , 2021, , 111-152.	0.3	1



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127	The MinION as a cost-effective technology for diagnostic screening of the SCN1A gene in epilepsy patients. <i>Epilepsy Research</i> , 2021, 172, 106593.	1.6	1
128	Multi-phenotype genome-wide association studies of the Norfolk Island isolate implicate pleiotropic loci involved in chronic kidney disease. <i>Scientific Reports</i> , 2021, 11, 19425.	3.3	1
129	Novel compound heterozygous missense mutations in GDAP1 cause Charcot-Marie-Tooth type 4A. <i>Journal of Genetics</i> , 2021, 100, 1.	0.7	0
130	Evaluation of an ancestry prediction strategy for historical military remains using a World War II-era sample and pedigrees with family-level admixture. <i>Australian Journal of Forensic Sciences</i> , 0, , 1-18.	1.2	0
131	Discriminating head trauma outcomes using machine learning and genomics. <i>Journal of Molecular Medicine</i> , 2021, , 1.	3.9	0