

Marie-Claude Gingras

List of Publications by Year in descending order

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Version: 2024-02-01

27
papers

6,960
citations

567144

15
h-index

552653

26
g-index

29
all docs

29
docs citations

29
times ranked

13816
citing authors

#	ARTICLE	IF	CITATIONS
1	Germline mutation in POLR2A: a heterogeneous, multi-systemic developmental disorder characterized by transcriptional dysregulation. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100014.	1.0	10
2	DNA methylation patterns identify subgroups of pancreatic neuroendocrine tumors with clinical association. <i>Communications Biology</i> , 2021, 4, 155.	2.0	26
3	Sequencing of a central nervous system tumor demonstrates cancer transmission in an organ transplant. <i>Life Science Alliance</i> , 2021, 4, e202000941.	1.3	1
4	Genetic testing in ambulatory cardiology clinics reveals high rate of findings with clinical management implications. <i>Genetics in Medicine</i> , 2021, 23, 2404-2414.	1.1	14
5	Transmission event of SARS-CoV-2 delta variant reveals multiple vaccine breakthrough infections. <i>BMC Medicine</i> , 2021, 19, 255.	2.3	137
6	NF- κ B and STAT3 co-operation enhances high glucose induced aggressiveness of cholangiocarcinoma cells. <i>Life Sciences</i> , 2020, 262, 118548.	2.0	9
7	Phenotypic expansion in <i>KIF1A</i> -related dominant disorders: A description of novel variants and review of published cases. <i>Human Mutation</i> , 2020, 41, 2094-2104.	1.1	8
8	Community-based recruitment and exome sequencing indicates high diagnostic yield in adults with intellectual disability. <i>Molecular Genetics & Genomic Medicine</i> , 2020, 8, e1439.	0.6	6
9	Whole-genome landscape of pancreatic neuroendocrine tumours. <i>Nature</i> , 2017, 543, 65-71.	13.7	716
10	Integrative Genomic Analysis of Cholangiocarcinoma Identifies Distinct IDH-Mutant Molecular Profiles. <i>Cell Reports</i> , 2017, 18, 2780-2794.	2.9	416
11	Functional annotation of rare gene aberration drivers of pancreatic cancer. <i>Nature Communications</i> , 2016, 7, 10500.	5.8	58
12	An open access pilot freely sharing cancer genomic data from participants in Texas. <i>Scientific Data</i> , 2016, 3, 160010.	2.4	19
13	Ampullary Cancers Harbor ELF3 Tumor Suppressor Gene Mutations and Exhibit Frequent WNT Dysregulation. <i>Cell Reports</i> , 2016, 14, 907-919.	2.9	107
14	Genomic analyses identify molecular subtypes of pancreatic cancer. <i>Nature</i> , 2016, 531, 47-52.	13.7	2,700
15	Genomic profiling guides the choice of molecular targeted therapy of pancreatic cancer. <i>Cancer Letters</i> , 2015, 363, 1-6.	3.2	21
16	Trans-ancestry mutational landscape of hepatocellular carcinoma genomes. <i>Nature Genetics</i> , 2014, 46, 1267-1273.	9.4	655
17	Pancreatic cancer genomes reveal aberrations in axon guidance pathway genes. <i>Nature</i> , 2012, 491, 399-405.	13.7	1,741
18	A novel therapeutic strategy for pancreatic neoplasia using a novel RNAi platform targeting PDX-1. <i>Nature Precedings</i> , 2011, , .	0.1	0

#	ARTICLE	IF	CITATIONS
19	A Primer on a Hepatocellular Carcinoma Bioresource Bank Using the Cancer Genome Atlas Guidelines: Practical Issues and Pitfalls. <i>World Journal of Surgery</i> , 2011, 35, 1732-1737.	0.8	4
20	Building a Comprehensive Genomic Program for Hepatocellular Carcinoma. <i>World Journal of Surgery</i> , 2011, 35, 1746-1750.	0.8	15
21	<i>PDX</i> . <i>Cancer</i> , 2011, 117, 723-733.	2.0	42
22	Basic Principles and Technologies for Deciphering the Genetic Map of Cancer. <i>World Journal of Surgery</i> , 2009, 33, 615-629.	0.8	13
23	Sequencing the Full Length of the Phosphatase and Tensin Homolog (PTEN) Gene in Hepatocellular Carcinoma (HCC) Using the 454 GS20 and Illumina GA DNA Sequencing Platforms. <i>World Journal of Surgery</i> , 2009, 33, 647-652.	0.8	6
24	Single nucleotide polymorphism in RECQL and survival in resectable pancreatic adenocarcinoma. <i>Hpb</i> , 2009, 11, 435-444.	0.1	16
25	TREM-1, MDL-1, and DAP12 expression is associated with a mature stage of myeloid development. <i>Molecular Immunology</i> , 2002, 38, 817-824.	1.0	124
26	CFFM4: a new member of the CD20/Fc μ R1 ² family. <i>Immunogenetics</i> , 2001, 53, 468-476.	1.2	21
27	Transendothelial migration induces rapid expression on neutrophils of granule-release VLA6 used for tissue infiltration. <i>Journal of Leukocyte Biology</i> , 1997, 62, 356-362.	1.5	31