Nathalie Gaudreault

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/4957928/publications.pdf

Version: 2024-02-01

19 papers

831 citations

12 h-index 18 g-index

22 all docs 22 docs citations

times ranked

22

1752 citing authors

#	Article	IF	CITATIONS
1	The Null QO _{Ourém} Variant within a Copy-Neutral Loss-of-Heterozygosity Event Causing Alpha-1 Antitrypsin Deficiency. American Journal of Respiratory Cell and Molecular Biology, 2022, 66, 700-702.	1.4	O
2	Unravelling actionable biology using transcriptomic data to integrate mitotic index and Ki-67 in the management of lung neuroendocrine tumors. Oncotarget, 2021, 12, 209-220.	0.8	1
3	Enhancer-associated aortic valve stenosis risk locus 1p21.2 alters NFATC2 binding site and promotes fibrogenesis. IScience, 2021, 24, 102241.	1.9	9
4	Prioritization of candidate causal genes for asthma in susceptibility loci derived from UK Biobank. Communications Biology, 2021, 4, 700.	2.0	77
5	Lipoprotein Proteomics and Aortic Valve Transcriptomics Identify Biological Pathways Linking Lipoprotein(a) Levels to Aortic Stenosis. Metabolites, 2021, 11, 459.	1.3	14
6	Phenome-wide analyses establish a specific association between aortic valve PALMD expression and calcific aortic valve stenosis. Communications Biology, 2020, 3, 477.	2.0	12
7	Transcriptomic data helps refining classification of pulmonary carcinoid tumors with increased mitotic counts. Modern Pathology, 2020, 33, 1712-1721.	2.9	15
8	Genetic Association Analyses Highlight <i>IL6</i> , <i>ALPL</i> , and <i>NAV1</i> As 3 New Susceptibility Genes Underlying Calcific Aortic Valve Stenosis. Circulation Genomic and Precision Medicine, 2019, 12, e002617.	1.6	45
9	A transcriptome-wide association study identifies PALMD as a susceptibility gene for calcific aortic valve stenosis. Nature Communications, 2018, 9, 988.	5.8	93
10	DNA methylation of a PLPP3 MIR transposon-based enhancer promotes an osteogenic programme in calcific aortic valve disease. Cardiovascular Research, 2018, 114, 1525-1535.	1.8	27
11	Tumor-based gene expression biomarkers to predict survival following curative intent resection for stage I lung adenocarcinoma. PLoS ONE, 2018, 13, e0207513.	1.1	3
12	Transcriptomic Microenvironment of Lung Adenocarcinoma. Cancer Epidemiology Biomarkers and Prevention, 2017, 26, 389-396.	1.1	6
13	RNA expression profile of calcified bicuspid, tricuspid, and normal human aortic valves by RNA sequencing. Physiological Genomics, 2016, 48, 749-761.	1.0	52
14	Altered DNA Methylation of Long Noncoding RNA <i>H19</i> in Calcific Aortic Valve Disease Promotes Mineralization by Silencing <i>NOTCH1</i> . Circulation, 2016, 134, 1848-1862.	1.6	182
15	Identification of Gender-Specific Genetic Variants in Patients With Bicuspid Aortic Valve. American Journal of Cardiology, 2016, 117, 420-426.	0.7	53
16	Calcium Signaling Pathway Genes <i>RUNX2</i> and <i>CACNA1C</i> Are Associated With Calcific Aortic Valve Disease. Circulation: Cardiovascular Genetics, 2015, 8, 812-822.	5.1	51
17	NOTCH1 genetic variants in patients with tricuspid calcific aortic valve stenosis. Journal of Heart Valve Disease, 2013, 22, 142-9.	0.5	25
18	Molecular Signature of Smoking in Human Lung Tissues. Cancer Research, 2012, 72, 3753-3763.	0.4	111

#	Article	IF	CITATIONS
19	Replication of Genetic Association Studies in Aortic Stenosis in Adults. American Journal of Cardiology, 2011, 108, 1305-1310.	0.7	28