

# Adrien Georges

## List of Publications by Year in descending order

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Version: 2024-02-01

23  
papers

1,147  
citations

567281

15  
h-index

610901

24  
g-index

30  
all docs

30  
docs citations

30  
times ranked

1783  
citing authors

#	ARTICLE	IF	CITATIONS
1	Genome-wide association study reveals novel genetic loci: a new polygenic risk score for mitral valve prolapse. <i>European Heart Journal</i> , 2022, 43, 1668-1680.	2.2	25
2	Computational estimates of annular diameter reveal genetic determinants of mitral valve function and disease. <i>JCI Insight</i> , 2022, 7, .	5.0	9
3	Rare loss-of-function mutations of <i>PTGIR</i> are enriched in fibromuscular dysplasia. <i>Cardiovascular Research</i> , 2021, 117, 1154-1165.	3.8	20
4	Chromatin Accessibility of Human Mitral Valves and Functional Assessment of MVP Risk Loci. <i>Circulation Research</i> , 2021, 128, e84-e101.	4.5	10
5	Plasma and genetic determinants of soluble TREM-1 and major adverse cardiovascular events in a prospective cohort of acute myocardial infarction patients. Results from the FAST-MI 2010 study. <i>International Journal of Cardiology</i> , 2021, 344, 213-219.	1.7	3
6	Genetic investigation of fibromuscular dysplasia identifies risk loci and shared genetics with common cardiovascular diseases. <i>Nature Communications</i> , 2021, 12, 6031.	12.8	34
7	A plasma proteogenomic signature for fibromuscular dysplasia. <i>Cardiovascular Research</i> , 2020, 116, 63-77.	3.8	27
8	Transcriptome Analysis of lncRNAs in Pheochromocytomas and Paragangliomas. <i>Journal of Clinical Endocrinology and Metabolism</i> , 2020, 105, 898-907.	3.6	11
9	Genetic Study of <i>PHACTR1</i> and Fibromuscular Dysplasia, Meta-Analysis and Effects on Clinical Features of Patients. <i>Hypertension</i> , 2020, 76, e4-e7.	2.7	9
10	Functional interplay between Mediator and RNA polymerase II in Rad2/XPG loading to the chromatin. <i>Nucleic Acids Research</i> , 2019, 47, 8988-9004.	14.5	17
11	Genome-Wide Association Studyâ€Driven Gene-Set Analyses, Genetic, and Functional Follow-Up Suggest <i>GLIS1</i> as a Susceptibility Gene for Mitral Valve Prolapse. <i>Circulation Genomic and Precision Medicine</i> , 2019, 12, e002497.	3.6	31
12	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. <i>Journal of the American College of Cardiology</i> , 2019, 73, 58-66.	2.8	147
13	Genomics of Fibromuscular Dysplasia. <i>International Journal of Molecular Sciences</i> , 2018, 19, 1526.	4.1	35
14	FOXL2: a central transcription factor of the ovary. <i>Journal of Molecular Endocrinology</i> , 2014, 52, R17-R33.	2.5	125
15	Transcription factors: specific DNA binding and specific gene regulation. <i>Trends in Genetics</i> , 2014, 30, 211-219.	6.7	145
16	FOXL2, GATA4, and SMAD3 Co-Operatively Modulate Gene Expression, Cell Viability and Apoptosis in Ovarian Granulosa Cell Tumor Cells. <i>PLoS ONE</i> , 2014, 9, e85545.	2.5	55
17	The transcription factor FOXL2 mobilizes estrogen signaling to maintain the identity of ovarian granulosa cells. <i>ELife</i> , 2014, 3, .	6.0	96
18	Interaction with the Yes-associated protein (YAP) allows TEAD1 to positively regulate NAIP expression. <i>FEBS Letters</i> , 2013, 587, 3216-3223.	2.8	11

#	ARTICLE	IF	CITATIONS
19	Discovery of novel protein partners of the transcription factor FOXL2 provides insights into its physiopathological roles. <i>Human Molecular Genetics</i> , 2012, 21, 3264-3274.	2.9	41
20	The transcription factor FOXL2: At the crossroads of ovarian physiology and pathology. <i>Molecular and Cellular Endocrinology</i> , 2012, 356, 55-64.	3.2	67
21	SUMOylation of the Forkhead Transcription Factor FOXL2 Promotes Its Stabilization/Activation through Transient Recruitment to PML Bodies. <i>PLoS ONE</i> , 2011, 6, e25463.	2.5	24
22	Functional Exploration of the Adult Ovarian Granulosa Cell Tumor-Associated Somatic FOXL2 Mutation p.Cys134Trp (c.402C>G). <i>PLoS ONE</i> , 2010, 5, e8789.	2.5	67
23	Kinesin's cover-neck bundle folds forward to generate force. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 19247-19252.	7.1	132