## Adrien Georges

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/9196128/publications.pdf

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567281 610901 1,147 23 15 24 citations h-index g-index papers 30 30 30 1783 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Association of the PHACTR1/EDN1 Genetic Locus With Spontaneous Coronary Artery Dissection. Journal of the American College of Cardiology, 2019, 73, 58-66.	2.8	147
2	Transcription factors: specific DNA binding and specific gene regulation. Trends in Genetics, 2014, 30, 211-219.	6.7	145
3	Kinesin's cover-neck bundle folds forward to generate force. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 19247-19252.	7.1	132
4	FOXL2: a central transcription factor of the ovary. Journal of Molecular Endocrinology, 2014, 52, R17-R33.	2.5	125
5	The transcription factor FOXL2 mobilizes estrogen signaling to maintain the identity of ovarian granulosa cells. ELife, 2014, 3, .	6.0	96
6	The transcription factor FOXL2: At the crossroads of ovarian physiology and pathology. Molecular and Cellular Endocrinology, 2012, 356, 55-64.	3.2	67
7	Functional Exploration of the Adult Ovarian Granulosa Cell Tumor-Associated Somatic FOXL2 Mutation p.Cys134Trp (c.402C>G). PLoS ONE, 2010, 5, e8789.	2.5	67
8	FOXL2, GATA4, and SMAD3 Co-Operatively Modulate Gene Expression, Cell Viability and Apoptosis in Ovarian Granulosa Cell Tumor Cells. PLoS ONE, 2014, 9, e85545.	2.5	55
9	Discovery of novel protein partners of the transcription factor FOXL2 provides insights into its physiopathological roles. Human Molecular Genetics, 2012, 21, 3264-3274.	2.9	41
10	Genomics of Fibromuscular Dysplasia. International Journal of Molecular Sciences, 2018, 19, 1526.	4.1	35
11	Genetic investigation of fibromuscular dysplasia identifies risk loci and shared genetics with common cardiovascular diseases. Nature Communications, 2021, 12, 6031.	12.8	34
12	Genome-Wide Association Study–Driven Gene-Set Analyses, Genetic, and Functional Follow-Up Suggest <i>GLIS1</i> as a Susceptibility Gene for Mitral Valve Prolapse. Circulation Genomic and Precision Medicine, 2019, 12, e002497.	3.6	31
13	A plasma proteogenomic signature for fibromuscular dysplasia. Cardiovascular Research, 2020, 116, 63-77.	3.8	27
14	Genome-wide association study reveals novel genetic loci: a new polygenic risk score for mitral valve prolapse. European Heart Journal, 2022, 43, 1668-1680.	2.2	25
15	SUMOylation of the Forkhead Transcription Factor FOXL2 Promotes Its Stabilization/Activation through Transient Recruitment to PML Bodies. PLoS ONE, 2011, 6, e25463.	2.5	24
16	Rare loss-of-function mutations of <i>PTGIR</i> are enriched in fibromuscular dysplasia. Cardiovascular Research, 2021, 117, 1154-1165.	3.8	20
17	Functional interplay between Mediator and RNA polymerase II in Rad2/XPG loading to the chromatin. Nucleic Acids Research, 2019, 47, 8988-9004.	14.5	17
18	Interaction with the Yesâ€associated protein (YAP) allows TEAD1 to positively regulate NAIP expression. FEBS Letters, 2013, 587, 3216-3223.	2.8	11

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#	Article	lF	CITATIONS
19	Transcriptome Analysis of IncRNAs in Pheochromocytomas and Paragangliomas. Journal of Clinical Endocrinology and Metabolism, 2020, 105, 898-907.	3.6	11
20	Chromatin Accessibility of Human Mitral Valves and Functional Assessment of MVP Risk Loci. Circulation Research, 2021, 128, e84-e101.	4.5	10
21	Genetic Study of <i>PHACTR1</i> and Fibromuscular Dysplasia, Meta-Analysis and Effects on Clinical Features of Patients. Hypertension, 2020, 76, e4-e7.	2.7	9
22	Computational estimates of annular diameter reveal genetic determinants of mitral valve function and disease. JCI Insight, 2022, 7, .	5.0	9
23	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. International Journal of Cardiology, 2021, 344, 213-219.	1.7	3