

Lars A Larsen

List of Publications by Year in descending order

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31
papers

1,480
citations

516215

16
h-index

414034

32
g-index

37
all docs

37
docs citations

37
times ranked

2811
citing authors

#	ARTICLE	IF	CITATIONS
1	Comparison of Outcome in Patients With Familial Versus Spontaneous Atrial Septal Defect. <i>American Journal of Cardiology</i> , 2022, 173, 128-131.	0.7	2
2	MCPH1: A Novel Case Report and a Review of the Literature. <i>Genes</i> , 2022, 13, 634.	1.0	5
3	Integrative analysis of genomic variants reveals new associations of candidate haploinsufficient genes with congenital heart disease. <i>PLoS Genetics</i> , 2021, 17, e1009679.	1.5	17
4	Mutation burden in patients with small unrepaired atrial septal defects. <i>International Journal of Cardiology Congenital Heart Disease</i> , 2021, 4, 100164.	0.2	6
5	RRP7A links primary microcephaly to dysfunction of ribosome biogenesis, resorption of primary cilia, and neurogenesis. <i>Nature Communications</i> , 2020, 11, 5816.	5.8	34
6	Systems genetics analysis identifies calcium-signaling defects as novel cause of congenital heart disease. <i>Genome Medicine</i> , 2020, 12, 76.	3.6	15
7	Model system identification of novel congenital heart disease gene candidates: focus on RPL13. <i>Human Molecular Genetics</i> , 2019, 28, 3954-3969.	1.4	19
8	Haploinsufficiency of ARHGAP42 is associated with hypertension. <i>European Journal of Human Genetics</i> , 2019, 27, 1296-1303.	1.4	12
9	CEP128 Localizes to the Subdistal Appendages of the Mother Centriole and Regulates TGF- β /BMP Signaling at the Primary Cilium. <i>Cell Reports</i> , 2018, 22, 2584-2592.	2.9	59
10	Familial co-occurrence of congenital heart defects follows distinct patterns. <i>European Heart Journal</i> , 2018, 39, 1015-1022.	1.0	32
11	IFT20 modulates ciliary PDGFR β signaling by regulating the stability of Cbl E3 ubiquitin ligases. <i>Journal of Cell Biology</i> , 2018, 217, 151-161.	2.3	54
12	The E3 ubiquitin ligase SMURF1 regulates cell-fate specification and outflow tract septation during mammalian heart development. <i>Scientific Reports</i> , 2018, 8, 9542.	1.6	20
13	Challenges for the Sustainability of University-Run Biobanks. <i>Biopreservation and Biobanking</i> , 2018, 16, 312-321.	0.5	12
14	Mutation of the Planar Cell Polarity Gene VANGL1 in Adolescent Idiopathic Scoliosis. <i>Spine</i> , 2017, 42, E702-E707.	1.0	16
15	A novel mutation in CDK5RAP2 gene causes primary microcephaly with speech impairment and sparse eyebrows in a consanguineous Pakistani family. <i>European Journal of Medical Genetics</i> , 2017, 60, 627-630.	0.7	10
16	Patient-specific three-dimensional explant spheroids derived from human nasal airway epithelium: a simple methodological approach for ex vivo studies of primary ciliary dyskinesia. <i>Cilia</i> , 2017, 6, 3.	1.8	16
17	How Suitable Are Registry Data for Recurrence Risk Calculations? Validation of Diagnoses on 1,593 Families With Congenital Heart Disease. <i>World Journal for Pediatric & Congenital Heart Surgery</i> , 2016, 7, 169-177.	0.3	5
18	Familial Atrial Septal Defect and Sudden Cardiac Death: Identification of a Novel <i>NKX2-5</i> Mutation and a Review of the Literature. <i>Congenital Heart Disease</i> , 2016, 11, 283-290.	0.0	81

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19	A novel splice site mutation in CEP135 is associated with primary microcephaly in a Pakistani family. <i>Journal of Human Genetics</i> , 2016, 61, 271-273.	1.1	16
20	Cilia and coordination of signaling networks during heart development. <i>Organogenesis</i> , 2014, 10, 108-125.	0.4	77
21	Of mice and men: molecular genetics of congenital heart disease. <i>Cellular and Molecular Life Sciences</i> , 2014, 71, 1327-1352.	2.4	159
22	TGF- β 2 Signaling Is Associated with Endocytosis at the Pocket Region of the Primary Cilium. <i>Cell Reports</i> , 2013, 3, 1806-1814.	2.9	248
23	Genetic and environmental risk factors in congenital heart disease functionally converge in protein networks driving heart development. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 14035-14040.	3.3	117
24	Screening of congenital heart disease patients using multiplex ligation-dependent probe amplification: Early diagnosis of syndromic patients. <i>American Journal of Medical Genetics, Part A</i> , 2012, 158A, 720-725.	0.7	27
25	Dissecting spatio-temporal protein networks driving human heart development and related disorders. <i>Molecular Systems Biology</i> , 2010, 6, 381.	3.2	80
26	Haploinsufficiency of TAB2 Causes Congenital Heart Defects in Humans. <i>American Journal of Human Genetics</i> , 2010, 86, 839-849.	2.6	97
27	Using Nucleofection of siRNA Constructs for Knockdown of Primary Cilia in P19.CL6 Cancer Stem Cell Differentiation into Cardiomyocytes. <i>Methods in Cell Biology</i> , 2009, 94, 181-197.	0.5	16
28	The primary cilium coordinates early cardiogenesis and hedgehog signaling in cardiomyocyte differentiation. <i>Journal of Cell Science</i> , 2009, 122, 3070-3082.	1.2	91
29	Molecular characterization of two patients with de novo interstitial deletions in 4q22-q24. <i>American Journal of Medical Genetics, Part A</i> , 2009, 149A, 1830-1833.	0.7	4
30	High frequency of submicroscopic genomic aberrations detected by tiling path array comparative genome hybridisation in patients with isolated congenital heart disease. <i>Journal of Medical Genetics</i> , 2008, 45, 704-709.	1.5	110
31	Single-strand conformation polymorphism analysis using capillary array electrophoresis for large-scale mutation detection. <i>Nature Protocols</i> , 2007, 2, 1458-1466.	5.5	20