

# Stéphane Zaffran

## List of Publications by Year in descending order

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93  
papers

6,994  
citations

117453

34  
h-index

60497

81  
g-index

107  
all docs

107  
docs citations

107  
times ranked

7097  
citing authors

#	ARTICLE	IF	CITATIONS
1	Clinical insights into a tertiary care center cohort of patients with bicuspid aortic valve. <i>International Journal of Cardiovascular Imaging</i> , 2022, 38, 51-59.	0.7	2
2	Identification of two variants in <i>AGRN</i> and <i>RPL3L</i> genes in a patient with catecholaminergic polymorphic ventricular tachycardia suggesting new candidate disease genes and digenic inheritance. <i>Clinical Case Reports (discontinued)</i> , 2022, 10, e05339.	0.2	7
3	Single Cell Approaches to Understand the Earliest Steps in Heart Development. <i>Current Cardiology Reports</i> , 2022, 24, 611-621.	1.3	2
4	Dilated-Left Ventricular Non-Compaction Cardiomyopathy in a Pediatric Case with SPEG Compound Heterozygous Variants. <i>International Journal of Molecular Sciences</i> , 2022, 23, 5205.	1.8	1
5	Identification of non-synonymous variations in <i>ROBO1</i> and <i>GATA5</i> genes in a family with bicuspid aortic valve disease. <i>Journal of Human Genetics</i> , 2022, , .	1.1	3
6	Molecular autopsy and clinical family screening in a case of sudden cardiac death reveals <i>ACTN2</i> mutation related to hypertrophic/dilated cardiomyopathy and a novel <i>LZTR1</i> variant associated with Noonan syndrome. <i>Molecular Genetics &amp; Genomic Medicine</i> , 2022, 10, .	0.6	5
7	<i>Mesp1</i> controls the chromatin and enhancer landscapes essential for spatiotemporal patterning of early cardiovascular progenitors. <i>Nature Cell Biology</i> , 2022, 24, 1114-1128.	4.6	11
8	Multiallelic rare variants support an oligogenic origin of sudden cardiac death in the young. <i>Herz</i> , 2021, 46, 94-102.	0.4	6
9	Side-dependent effect in the response of valve endothelial cells to bidirectional shear stress. <i>International Journal of Cardiology</i> , 2021, 323, 220-228.	0.8	6
10	Outflow Tract Formation—Embryonic Origins of Conotruncal Congenital Heart Disease. <i>Journal of Cardiovascular Development and Disease</i> , 2021, 8, 42.	0.8	18
11	A roadmap for the Human Developmental Cell Atlas. <i>Nature</i> , 2021, 597, 196-205.	13.7	114
12	<i>Msx1</i> haploinsufficiency modifies the <i>Pax9</i> -deficient cardiovascular phenotype. <i>BMC Developmental Biology</i> , 2021, 21, 14.	2.1	6
13	Identification of a peripheral blood gene signature predicting aortic valve calcification. <i>Physiological Genomics</i> , 2020, 52, 563-574.	1.0	11
14	<i>Piezo1</i> is required for outflow tract and aortic valve development.. <i>Journal of Molecular and Cellular Cardiology</i> , 2020, 143, 51-62.	0.9	44
15	<i>Hox</i> -dependent coordination of mouse cardiac progenitor cell patterning and differentiation. <i>ELife</i> , 2020, 9, .	2.8	41
16	<i>Krox20</i> Regulates Endothelial Nitric Oxide Signaling in Aortic Valve Development and Disease. <i>Journal of Cardiovascular Development and Disease</i> , 2019, 6, 39.	0.8	5
17	A severe clinical phenotype of Noonan syndrome with neonatal hypertrophic cardiomyopathy in the second case worldwide with <i>RAF1</i> S259Y neomutation. <i>Genetical Research</i> , 2019, 101, e6.	0.3	4
18	Human pre-valvular endocardial cells derived from pluripotent stem cells recapitulate cardiac pathophysiological valvulogenesis. <i>Nature Communications</i> , 2019, 10, 1929.	5.8	60

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19	Transcriptome analysis of mouse and human sinoatrial node cells reveals a conserved genetic program. <i>Development (Cambridge)</i> , 2019, 146, .	1.2	54
20	Krox20 defines a subpopulation of cardiac neural crest cells contributing to arterial valves and bicuspid aortic valve. <i>Development (Cambridge)</i> , 2018, 145, .	1.2	34
21	Asb21±â€“Filamin A Axis Is Essential for Actin Cytoskeleton Remodeling During Heart Development. <i>Circulation Research</i> , 2018, 122, e34-e48.	2.0	29
22	Giant congenital melanocytic nevus with vascular malformation and epidermal cysts associated with a somatic activating mutation in <i>BRAF</i> . <i>Pigment Cell and Melanoma Research</i> , 2018, 31, 437-441.	1.5	22
23	Myocardial Bmp2 gain causes ectopic EMT and promotes cardiomyocyte proliferation and immaturity. <i>Cell Death and Disease</i> , 2018, 9, 399.	2.7	24
24	Ectopic expression of <i>Hoxb1</i> induces cardiac and craniofacial malformations. <i>Genesis</i> , 2018, 56, e23221.	0.8	18
25	Hox and Tale transcription factors in heart development and disease. <i>International Journal of Developmental Biology</i> , 2018, 62, 837-846.	0.3	20
26	Bmp2 and Notch cooperate to pattern the embryonic endocardium. <i>Development (Cambridge)</i> , 2018, 145, .	1.2	30
27	Novel ALPK3 mutation in a Tunisian patient with pediatric cardiomyopathy and facio-thoraco-skeletal features. <i>Journal of Human Genetics</i> , 2018, 63, 1077-1082.	1.1	16
28	T-box genes and retinoic acid signaling regulate the segregation of arterial and venous pole progenitor cells in the murine second heart field. <i>Human Molecular Genetics</i> , 2018, 27, 3747-3760.	1.4	59
29	Analysis of HOXB1 gene in a cohort of patients with sporadic ventricular septal defect. <i>Molecular Biology Reports</i> , 2018, 45, 1507-1513.	1.0	0
30	Origines gÃ©nÃ©tique et dÃ©veloppementale de la bicuspidie aortique. <i>Archives Des Maladies Du Coeur Et Des Vaisseaux - Pratique</i> , 2017, 2017, 22-26.	0.0	0
31	The alternatively spliced LRRFIP1 Isoform-1 is a key regulator of the Wnt/Î²-catenin transcription pathway. <i>Biochimica Et Biophysica Acta - Molecular Cell Research</i> , 2017, 1864, 1142-1152.	1.9	13
32	Mechanisms of retinoic acid signaling during cardiogenesis. <i>Mechanisms of Development</i> , 2017, 143, 9-19.	1.7	74
33	Hoxa1 and Hoxb1 are required for pharyngeal arch artery development. <i>Mechanisms of Development</i> , 2017, 143, 1-8.	1.7	23
34	Reduced aggrecan expression affects cardiac outflow tract development in zebrafish and is associated with bicuspid aortic valve disease in humans. <i>International Journal of Cardiology</i> , 2017, 249, 340-343.	0.8	14
35	<i>FOXC1</i> haploinsufficiency due to 6p25 deletion in a patient with rapidly progressing aortic valve disease. <i>American Journal of Medical Genetics, Part A</i> , 2017, 173, 2489-2493.	0.7	7
36	Hox Genes in Cardiovascular Development and Diseases. <i>Journal of Developmental Biology</i> , 2016, 4, 14.	0.9	30

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37	Cardiac outflow morphogenesis depends on effects of retinoic acid signaling on multiple cell lineages. <i>Developmental Dynamics</i> , 2016, 245, 388-401.	0.8	30
38	0352: Decrease of Krox20 expression leads to aortic valve dysfunction and thickening of the valve leaflets. <i>Archives of Cardiovascular Diseases Supplements</i> , 2016, 8, 55.	0.0	0
39	Actionable Genes, Core Databases, and Locus-Specific Databases. <i>Human Mutation</i> , 2016, 37, 1299-1307.	1.1	6
40	WES/WGS Reporting of Mutations from Cardiovascular "Actionable" Genes in Clinical Practice: A Key Role for UMD Knowledgebases in the Era of Big Databases. <i>Human Mutation</i> , 2016, 37, 1308-1317.	1.1	5
41	Krox20 heterozygous mice: A model of aortic regurgitation associated with decreased expression of fibrillar collagen genes. <i>Archives of Cardiovascular Diseases</i> , 2016, 109, 188-198.	0.7	4
42	Disruption of CXCR4 signaling in pharyngeal neural crest cells causes DiGeorge syndrome-like malformations. <i>Development (Cambridge)</i> , 2016, 143, 582-8.	1.2	33
43	An uncommon cause of tricuspid regurgitation: three-dimensional echocardiographic incremental value, surgical and genetic insights. <i>European Journal of Cardio-thoracic Surgery</i> , 2016, 50, 180-182.	0.6	1
44	0268 : Involvement of LRRFip1 gene and canonical Wnt pathway in Mitral Valve Prolapse (MVP). <i>Archives of Cardiovascular Diseases Supplements</i> , 2015, 7, 204.	0.0	0
45	<i>scp</i> <i>M</i> <i>scx1</i> <i>cre</i> <i>ERT2</i> knock-in allele: A useful tool to target embryonic and adult cardiac valves. <i>Genesis</i> , 2015, 53, 337-345.	0.8	9
46	Hoxb1 regulates proliferation and differentiation of second heart field progenitors in pharyngeal mesoderm and genetically interacts with Hoxa1 during cardiac outflow tract development. <i>Developmental Biology</i> , 2015, 406, 247-258.	0.9	48
47	Loss of Krox20 results in aortic valve regurgitation and impaired transcriptional activation of fibrillar collagen genes. <i>Cardiovascular Research</i> , 2014, 104, 443-455.	1.8	19
48	Tbx1 Coordinates Addition of Posterior Second Heart Field Progenitor Cells to the Arterial and Venous Poles of the Heart. <i>Circulation Research</i> , 2014, 115, 790-799.	2.0	105
49	0174: LRRFip1 and Wnt pathway involvement in mitral valve prolapse. <i>Archives of Cardiovascular Diseases Supplements</i> , 2014, 6, 72.	0.0	0
50	Anterior Hox Genes in Cardiac Development and Great Artery Patterning. <i>Journal of Cardiovascular Development and Disease</i> , 2014, 1, 3-13.	0.8	5
51	Retinoids and Cardiac Development. <i>Journal of Developmental Biology</i> , 2014, 2, 50-71.	0.9	20
52	Genetic Lineage Tracing Analysis of Anterior Hox Expressing Cells. <i>Methods in Molecular Biology</i> , 2014, 1196, 37-48.	0.4	5
53	<i>Fibroblast growth factor 10</i> gene regulation in the second heart field by Tbx1, Nkx2-5, and Islet1 reveals a genetic switch for down-regulation in the myocardium. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 18273-18280.	3.3	109
54	Value of In Vivo T2 Measurement for Myocardial Fibrosis Assessment in Diabetic Mice at 11.75 T. <i>Investigative Radiology</i> , 2012, 47, 319-323.	3.5	34

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55	Early cardiac development: a view from stem cells to embryos. <i>Cardiovascular Research</i> , 2012, 96, 352-362.	1.8	115
56	ISL1 Directly Regulates FGF10 Transcription during Human Cardiac Outflow Formation. <i>PLoS ONE</i> , 2012, 7, e30677.	1.1	46
57	New developments in the second heart field. <i>Differentiation</i> , 2012, 84, 17-24.	1.0	77
58	Hox genes define distinct progenitor sub-domains within the second heart field. <i>Developmental Biology</i> , 2011, 353, 266-274.	0.9	144
59	A Retinoic Acid Responsive Hoxa3 Transgene Expressed in Embryonic Pharyngeal Endoderm, Cardiac Neural Crest and a Subdomain of the Second Heart Field. <i>PLoS ONE</i> , 2011, 6, e27624.	1.1	26
60	Expression of <i>Slit</i> and <i>Robo</i> genes in the developing mouse heart. <i>Developmental Dynamics</i> , 2010, 239, 3303-3311.	0.8	38
61	Decreased Levels of Embryonic Retinoic Acid Synthesis Accelerate Recovery From Arterial Growth Delay in a Mouse Model of DiGeorge Syndrome. <i>Circulation Research</i> , 2010, 106, 686-694.	2.0	82
62	Endogenous retinoic acid regulates cardiac progenitor differentiation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 9234-9239.	3.3	96
63	Conotruncal defects associated with anomalous pulmonary venous connections. <i>Archives of Cardiovascular Diseases</i> , 2009, 102, 105-110.	0.7	25
64	Genetics and embryological mechanisms of congenital heart diseases. <i>Archives of Cardiovascular Diseases</i> , 2009, 102, 59-63.	0.7	38
65	Atrial myocardium derives from the posterior region of the second heart field, which acquires left-right identity as <i>Pitx2c</i> is expressed. <i>Development (Cambridge)</i> , 2008, 135, 1157-1167.	1.2	134
66	Myocardium at the base of the aorta and pulmonary trunk is prefigured in the outflow tract of the heart and in subdomains of the second heart field. <i>Developmental Biology</i> , 2008, 313, 25-34.	0.9	62
67	Retinoic acid deficiency alters second heart field formation. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 2913-2918.	3.3	186
68	Correction: Pax3 and Pax7 have distinct and overlapping functions in adult muscle progenitor cells. <i>Journal of Cell Biology</i> , 2007, 176, 125-125.	2.3	0
69	The <i>Drosophila Hand</i> gene is required for remodeling of the developing adult heart and midgut during metamorphosis. <i>Developmental Biology</i> , 2007, 311, 287-296.	0.9	30
70	An <i>Nkx2-5/Bmp2/Smad1</i> Negative Feedback Loop Controls Heart Progenitor Specification and Proliferation. <i>Cell</i> , 2007, 128, 947-959.	13.5	470
71	Pax3 and Pax7 have distinct and overlapping functions in adult muscle progenitor cells. <i>Journal of Cell Biology</i> , 2006, 172, 91-102.	2.3	599
72	Myocardial cell lineages in the mammalian embryo: The second heart field. <i>Journal of Molecular and Cellular Cardiology</i> , 2006, 40, 992.	0.9	0

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73	Rotation of the Myocardial Wall of the Outflow Tract Is Implicated in the Normal Positioning of the Great Arteries. <i>Circulation Research</i> , 2006, 98, 421-428.	2.0	190
74	Congenital heart defects in <i>Fgfr2-IIIb</i> and <i>Fgf10</i> mutant mice. <i>Cardiovascular Research</i> , 2006, 71, 50-60.	1.8	86
75	Cardioblast-intrinsic Tinman activity controls proper diversification and differentiation of myocardial cells in <i>Drosophila</i> . <i>Development (Cambridge)</i> , 2006, 133, 4073-4083.	1.2	86
76	Development of the Larval Visceral Musculature. , 2006, , 62-78.		11
77	Building the mammalian heart from two sources of myocardial cells. <i>Nature Reviews Genetics</i> , 2005, 6, 826-835.	7.7	1,051
78	<i>Fgf10</i> expression identifies parabronchial smooth muscle cell progenitors and is required for their entry into the smooth muscle cell lineage. <i>Development (Cambridge)</i> , 2005, 132, 2157-2166.	1.2	168
79	Direct Isolation of Satellite Cells for Skeletal Muscle Regeneration. <i>Science</i> , 2005, 309, 2064-2067.	6.0	939
80	The homeodomain of Tinman mediates homo- and heterodimerization of NK proteins. <i>Biochemical and Biophysical Research Communications</i> , 2005, 334, 361-369.	1.0	17
81	Right Ventricular Myocardium Derives From the Anterior Heart Field. <i>Circulation Research</i> , 2004, 95, 261-268.	2.0	334
82	Cell history determines the maintenance of transcriptional differences between left and right ventricular cardiomyocytes in the developing mouse heart. <i>Journal of Cell Science</i> , 2003, 116, 5005-5013.	1.2	8
83	La Souris comme modèle d'étude de la morphogenèse du cœur chez les mammifères : origine des myocytes et études d'explants cardiaques. <i>Société De Biologie Journal</i> , 2003, 197, 187-194.	0.3	1
84	Early Signals in Cardiac Development. <i>Circulation Research</i> , 2002, 91, 457-469.	2.0	272
85	The $\beta$ 3 tubulin gene is a direct target of bagpipe and biniou in the visceral mesoderm of <i>Drosophila</i> . <i>Mechanisms of Development</i> , 2002, 114, 85-93.	1.7	16
86	Cardiogenesis in the <i>Drosophila</i> Model: Control Mechanisms during Early Induction and Diversification of Cardiac Progenitors. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2002, 67, 1-12.	2.0	24
87	Pericardin, a <i>Drosophila</i> type IV collagen-like protein is involved in the morphogenesis and maintenance of the heart epithelium during dorsal ectoderm closure. <i>Development (Cambridge)</i> , 2002, 129, 3241-53.	1.2	48
88	<i>biniou</i> ( <i>FoxF</i> ), a central component in a regulatory network controlling visceral mesoderm development and midgut morphogenesis in <i>Drosophila</i> . <i>Genes and Development</i> , 2001, 15, 2900-2915.	2.7	133
89	Molecular cloning and embryonic expression of dFKBP59, a novel <i>Drosophila</i> FK506-binding protein. <i>Gene</i> , 2000, 246, 103-109.	1.0	17
90	The NK-2 homeobox gene scarecrow ( <i>scro</i> ) is expressed in pharynx, ventral nerve cord and brain of <i>Drosophila</i> embryos. <i>Mechanisms of Development</i> , 2000, 94, 237-241.	1.7	58

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91	The Heterotrimeric Protein Go Is Required for the Formation of Heart Epithelium in Drosophila. Journal of Cell Biology, 1999, 145, 1063-1076.	2.3	71
92	Cellular interactions during heart morphogenesis in the Drosophila embryo. Biology of the Cell, 1995, 84, 13-24.	0.7	33
93	<i>Hox</i>-Dependent Coordination of Cardiac Cell Patterning and Differentiation. SSRN Electronic Journal, 0, , .	0.4	1