

# Marc-Phillip Mp Hitz

## List of Publications by Year in descending order

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Version: 2024-02-01

28  
papers

969  
citations

840776

11  
h-index

526287

27  
g-index

33  
all docs

33  
docs citations

33  
times ranked

2203  
citing authors

#	ARTICLE	IF	CITATIONS
1	Distinct genetic architectures for syndromic and nonsyndromic congenital heart defects identified by exome sequencing. <i>Nature Genetics</i> , 2016, 48, 1060-1065.	21.4	351
2	Rare Variants in NR2F2 Cause Congenital Heart Defects in Humans. <i>American Journal of Human Genetics</i> , 2014, 94, 574-585.	6.2	146
3	Rare Copy Number Variants Contribute to Congenital Left-Sided Heart Disease. <i>PLoS Genetics</i> , 2012, 8, e1002903.	3.5	119
4	Accurate and fast feature selection workflow for high-dimensional omics data. <i>PLoS ONE</i> , 2017, 12, e0189875.	2.5	60
5	Loss of ADAMTS19 causes progressive non-syndromic heart valve disease. <i>Nature Genetics</i> , 2020, 52, 40-47.	21.4	46
6	Functional Zebrafish Studies Based on Human Genotyping Point to Netrin-1 as a Link Between Aberrant Cardiovascular Development and Thyroid Dysgenesis. <i>Endocrinology</i> , 2015, 156, 377-388.	2.8	38
7	Extensive identification of genes involved in congenital and structural heart disorders and cardiomyopathy. , 2022, 1, 157-173.		22
8	Ventricular septal defect and restrictive cardiomyopathy in a paediatric TNNI3 mutation carrier. <i>Cardiology in the Young</i> , 2010, 20, 574-576.	0.8	20
9	Fetal Cardiac Troponin Isoforms Rescue the Increased Ca <sup>2+</sup> Sensitivity Produced by a Novel Double Deletion in Cardiac Troponin T Linked to Restrictive Cardiomyopathy. <i>Journal of Biological Chemistry</i> , 2011, 286, 20901-20912.	3.4	19
10	Cardiac specific expression of Xenopus Popeye-1. <i>Mechanisms of Development</i> , 2002, 115, 123-126.	1.7	17
11	Integrative analysis of genomic variants reveals new associations of candidate haploinsufficient genes with congenital heart disease. <i>PLoS Genetics</i> , 2021, 17, e1009679.	3.5	17
12	Systems genetics analysis identifies calcium-signaling defects as novel cause of congenital heart disease. <i>Genome Medicine</i> , 2020, 12, 76.	8.2	15
13	Common Genetic Variants Contribute to Risk of Transposition of the Great Arteries. <i>Circulation Research</i> , 2022, 130, 166-180.	4.5	15
14	Enabling Global Clinical Collaborations on Identifiable Patient Data: The Minerva Initiative. <i>Frontiers in Genetics</i> , 2019, 10, 611.	2.3	14
15	DNA methylation profiling allows for characterization of atrial and ventricular cardiac tissues and hiPSC-CMs. <i>Clinical Epigenetics</i> , 2019, 11, 89.	4.1	12
16	Levosimendan for bridging in a pediatric patient with Alström syndrome awaiting heart-lung transplantation. <i>Clinical Research in Cardiology</i> , 2008, 97, 846-848.	3.3	9
17	Germline variants in HEY2 functional domains lead to congenital heart defects and thoracic aortic aneurysms. <i>Genetics in Medicine</i> , 2021, 23, 103-110.	2.4	7
18	Rare variants in KDR, encoding VEGF Receptor 2, are associated with tetralogy of Fallot. <i>Genetics in Medicine</i> , 2021, 23, 1952-1960.	2.4	7

#	ARTICLE	IF	CITATIONS
19	Recent advances in congenital heart disease genomics. F1000Research, 2017, 6, 869.	1.6	7
20	Familial ventricular aneurysms and septal defects map to chromosome 10p15. European Heart Journal, 2011, 32, 568-573.	2.2	6
21	Hypoplastic Left Heart Syndrome With Left Ventricular Myocardial Sinusoids: Echocardiographic and Angiographic Findings in the First Neonate Surviving the Norwood I and II Procedure. Circulation, 2008, 117, e319-21.	1.6	4
22	The analysis of heterotaxy patients reveals new loss-of-function variants of GRK5. Scientific Reports, 2016, 6, 33231.	3.3	4
23	The omics discovery REST interface. Nucleic Acids Research, 2020, 48, W380-W384.	14.5	3
24	Race for healthy hearts. Nature, 2015, 520, 160-161.	27.8	2
25	Patients with congenital heart defect and their families support genetic heart research. Congenital Heart Disease, 2018, 13, 685-689.	0.2	2
26	Genetik der angeborenen Herzfehler. Medizinische Genetik, 2017, 29, 248-256.	0.2	1
27	Exome sequencing in individuals with cardiovascular laterality defects identifies potential candidate genes. European Journal of Human Genetics, 2022, , .	2.8	1
28	Genetic regulation of heart valve development: Clinical implications. Aswan Heart Centre Science & Practice Series, 2011, 2011, .	0.3	0