

# Vidu Garg

## List of Publications by Year in descending order

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96  
papers

6,824  
citations

87723

38  
h-index

62479

80  
g-index

107  
all docs

107  
docs citations

107  
times ranked

7898  
citing authors

#	ARTICLE	IF	CITATIONS
1	Gender Differences in Physical Activity Engagement Among Adolescents With Congenital Heart Disease. <i>Journal of Pediatric Psychology</i> , 2022, 47, 859-869.	1.1	4
2	OUP accepted manuscript. <i>European Heart Journal</i> , 2022, , .	1.0	5
3	Probing single ventricle heart defects with <sc>patientâ€derived</sc> induced pluripotent stem cells and emerging technologies. <i>Birth Defects Research</i> , 2022, , .	0.8	3
4	Single-Cell RNA Sequencing Reveals Novel Genes Regulated by Hypoxia in the Lung Vasculature. <i>Journal of Vascular Research</i> , 2022, 59, 163-175.	0.6	4
5	Use of machine learning to classify high-risk variants of uncertain significance in lamin A/C cardiac disease. <i>Heart Rhythm</i> , 2022, 19, 676-685.	0.3	3
6	Inhibition of BKCa channels protects neonatal hearts against myocardial ischemia and reperfusion injury. <i>Cell Death Discovery</i> , 2022, 8, 175.	2.0	3
7	Exome sequencing in multiplex families with left-sided cardiac defects has high yield for disease gene discovery. <i>PLoS Genetics</i> , 2022, 18, e1010236.	1.5	8
8	Molecular genetic mechanisms of congenital heart disease. <i>Current Opinion in Genetics and Development</i> , 2022, 75, 101949.	1.5	10
9	In Vivo and In Vitro Genetic Models of Congenital Heart Disease. <i>Cold Spring Harbor Perspectives in Biology</i> , 2021, 13, a036764.	2.3	23
10	Galectin-3 and sST2 as Prognosticators for Heart Failure Requiring Extracorporeal Life Support: Jack nâ€™™ Jill. <i>Biomolecules</i> , 2021, 11, 166.	1.8	6
11	Nitric oxide prevents aortic valve calcification by S-nitrosylation of USP9X to activate NOTCH signaling. <i>Science Advances</i> , 2021, 7, .	4.7	43
12	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
13	Common deletion variants causing protocadherin-14 deficiency contribute to the complex genetics of BAV and left-sided congenital heart disease. <i>Human Genetics and Genomics Advances</i> , 2021, 2, 100037.	1.0	7
14	A Multi-Omics Approach Using a Mouse Model of Cardiac Malformations for Prioritization of Human Congenital Heart Disease Contributing Genes. <i>Frontiers in Cardiovascular Medicine</i> , 2021, 8, 683074.	1.1	2
15	Human Stem Cell Models of SARS-CoV-2 Infection in the Cardiovascular System. <i>Stem Cell Reviews and Reports</i> , 2021, 17, 2107-2119.	1.7	0
16	miRâ€145 transgenic mice develop cardiopulmonary complications leading to postnatal death. <i>Physiological Reports</i> , 2021, 9, e15013.	0.7	1
17	Genetics of congenital heart disease: a narrative review of recent advances and clinical implications. <i>Translational Pediatrics</i> , 2021, 10, 2366-2386.	0.5	26
18	Impact of maternal hyperglycemia on cardiac development: Insights from animal models. <i>Genesis</i> , 2021, 59, e23449.	0.8	4

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19	Decoding Genetics of Congenital Heart Disease Using Patient-Derived Induced Pluripotent Stem Cells (iPSCs). <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 630069.	1.8	17
20	A Randomized Clinical Trial Demonstrating Feasibility and Preliminary Efficacy of a Videoconference-Delivered Physical Activity Lifestyle Intervention Among Adolescents With a Congenital Heart Defect. <i>Annals of Behavioral Medicine</i> , 2021, , .	1.7	1
21	Cardiomyocyte Proliferation and Maturation: Two Sides of the Same Coin for Heart Regeneration. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 594226.	1.8	50
22	Novel frameshift variant in MYL2 reveals molecular differences between dominant and recessive forms of hypertrophic cardiomyopathy. <i>PLoS Genetics</i> , 2020, 16, e1008639.	1.5	16
23	Subtype-specific cardiomyocytes for precision medicine: Where are we now?. <i>Stem Cells</i> , 2020, 38, 822-833.	1.4	24
24	Abstract 436: Maternal Pregestational Diabetes Alters Fetal Gene Regulatory Program to Cause Spectrum of Congenital Heart Defects. <i>Circulation Research</i> , 2020, 127, .	2.0	0
25	Abstract 276: NOTCH1 is Essential for Ventricular Cardiomyocyte Differentiation of Human Induced Pluripotent Stem Cells. <i>Circulation Research</i> , 2020, 127, .	2.0	0
26	Abstract 16552: Chromosomal Microarray Abnormalities in More Than 1300 Patients With Congenital Heart Disease Provide Novel Clinical and Etiological Insights. <i>Circulation</i> , 2020, 142, .	1.6	0
27	Reclassification of Variants of Uncertain Significance in Children with Inherited Arrhythmia Syndromes is Predicted by Clinical Factors. <i>Pediatric Cardiology</i> , 2019, 40, 1679-1687.	0.6	24
28	Developmental origins for semilunar valve stenosis identified in mice harboring congenital heart disease-associated <i>GATA4</i> mutation. <i>DMM Disease Models and Mechanisms</i> , 2019, 12, .	1.2	17
29	Shaping the future heart: transgenerational outcomes of maternal metabolic syndrome. <i>American Journal of Physiology - Heart and Circulatory Physiology</i> , 2019, 316, H1141-H1143.	1.5	1
30	Another Notch in the Genetic Puzzle of Tetralogy of Fallot. <i>Circulation Research</i> , 2019, 124, 462-464.	2.0	7
31	Abstract 650: Dose-response Effect of Hyperglycemia in Maternal Diabetes Mediated Congenital Heart Defects. <i>Circulation Research</i> , 2019, 125, .	2.0	0
32	Nestin expression is dynamically regulated in cardiomyocytes during embryogenesis. <i>Journal of Cellular Physiology</i> , 2018, 233, 3218-3229.	2.0	21
33	The Role of Lipoprotein(a) in Calcific Aortic Valve Disease. <i>JAMA Cardiology</i> , 2018, 3, 24.	3.0	3
34	Maternal hyperglycemia and fetal cardiac development: Clinical impact and underlying mechanisms. <i>Birth Defects Research</i> , 2018, 110, 1504-1516.	0.8	72
35	Genetic Basis for Congenital Heart Disease: Revisited: A Scientific Statement From the American Heart Association. <i>Circulation</i> , 2018, 138, e653-e711.	1.6	387
36	Genetic basis of aortic valvular disease. <i>Current Opinion in Cardiology</i> , 2017, 32, 239-245.	0.8	22

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37	UTILITY OF GENETIC TESTING IN INFANTS WITH ISOLATED CONGENITAL HEART DISEASE. Journal of the American College of Cardiology, 2017, 69, 635.	1.2	0
38	Abnormal Longitudinal Growth of the Aorta in Children with Familial Bicuspid Aortic Valve. Pediatric Cardiology, 2017, 38, 1709-1715.	0.6	4
39	Assessment of large copy number variants in patients with apparently isolated congenital left-sided cardiac lesions reveals clinically relevant genomic events. American Journal of Medical Genetics, Part A, 2017, 173, 2176-2188.	0.7	17
40	Dose-dependent effect of Hyperglycemia during Cardiac Development. Journal of Molecular and Cellular Cardiology, 2017, 112, 153.	0.9	0
41	Epigenetic mechanisms underlying maternal diabetes-associated risk of congenital heart disease. JCI Insight, 2017, 2, .	2.3	59
42	Notch1 haploinsufficiency causes ascending aortic aneurysms in mice. JCI Insight, 2017, 2, .	2.3	44
43	Measuring genetic knowledge: a brief survey instrument for adolescents and adults. Clinical Genetics, 2016, 89, 235-243.	1.0	33
44	Endothelial Notch1 Is Required for Proper Development of the Semilunar Valves and Cardiac Outflow Tract. Journal of the American Heart Association, 2016, 5, .	1.6	55
45	Percutaneous Patent Ductus Arteriosus (PDA) Closure in Very Preterm Infants: Feasibility and Complications. Journal of the American Heart Association, 2016, 5, .	1.6	100
46	Enhancing Literacy in Cardiovascular Genetics: A Scientific Statement From the American Heart Association. Circulation: Cardiovascular Genetics, 2016, 9, 448-467.	5.1	64
47	Utilization of Whole Exome Sequencing to Identify Causative Mutations in Familial Congenital Heart Disease. Circulation: Cardiovascular Genetics, 2016, 9, 320-329.	5.1	71
48	A genome-wide association study of congenital cardiovascular left-sided lesions shows association with a locus on chromosome 20. Human Molecular Genetics, 2016, 25, 2331-2341.	1.4	31
49	Notch Signaling in Aortic Valve Development and Disease. , 2016, , 371-376.		13
50	Rationale for the Cytogenomics of Cardiovascular Malformations Consortium: A Phenotype Intensive Registry Based Approach. Journal of Cardiovascular Development and Disease, 2015, 2, 76-92.	0.8	7
51	Dynamic Heterogeneity of the Heart Valve Interstitial Cell Population in Mitral Valve Health and Disease. Journal of Cardiovascular Development and Disease, 2015, 2, 214-232.	0.8	26
52	Evidence of Aortopathy in Mice with Haploinsufficiency of Notch1 in Nos3-Null Background. Journal of Cardiovascular Development and Disease, 2015, 2, 17-30.	0.8	28
53	Lifetime Prevalence of Sexual Intercourse and Contraception Use at Last Sex Among Adolescents and Young Adults With Congenital Heart Disease. Journal of Adolescent Health, 2015, 56, 396-401.	1.2	22
54	Early versus delayed umbilical cord clamping in infants with congenital heart disease: a pilot, randomized, controlled trial. Journal of Perinatology, 2015, 35, 826-831.	0.9	28

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55	MicroRNA miR145 Regulates TGFBR2 Expression and Matrix Synthesis in Vascular Smooth Muscle Cells. <i>Circulation Research</i> , 2015, 116, 23-34.	2.0	72
56	Rare GATA5 sequence variants identified in individuals with bicuspid aortic valve. <i>Pediatric Research</i> , 2014, 76, 211-216.	1.1	74
57	Disruption of myocardial Gata4 and Tbx5 results in defects in cardiomyocyte proliferation and atrioventricular septation. <i>Human Molecular Genetics</i> , 2014, 23, 5025-5035.	1.4	50
58	Use of a targeted, combinatorial next-generation sequencing approach for the study of bicuspid aortic valve. <i>BMC Medical Genomics</i> , 2014, 7, 56.	0.7	50
59	Pharmacological Inhibitor of Notch Signaling Stabilizes the Progression of Small Abdominal Aortic Aneurysm in a Mouse Model. <i>Journal of the American Heart Association</i> , 2014, 3, e001064.	1.6	50
60	Genetics of Valvular Heart Disease. <i>Current Cardiology Reports</i> , 2014, 16, 487.	1.3	57
61	A unified test of linkage analysis and rare-variant association for analysis of pedigree sequence data. <i>Nature Biotechnology</i> , 2014, 32, 663-669.	9.4	93
62	Beyond genetics: focusing on maternal environment for congenital heart disease prevention. <i>Evidence-Based Medicine</i> , 2014, 19, e8-e8.	0.6	1
63	Etiology of Valvular Heart Disease. <i>Circulation Journal</i> , 2014, 78, 1801-1807.	0.7	45
64	Endothelial nitric oxide signaling regulates Notch1 in aortic valve disease. <i>Journal of Molecular and Cellular Cardiology</i> , 2013, 60, 27-35.	0.9	142
65	Compacting the heart with Notch. <i>Nature Medicine</i> , 2013, 19, 133-134.	15.2	10
66	Genetic Abnormalities in <i>FOXP1</i> Are Associated with Congenital Heart Defects. <i>Human Mutation</i> , 2013, 34, 1226-1230.	1.1	39
67	The Congenital Heart Disease Genetic Network Study. <i>Circulation Research</i> , 2013, 112, 698-706.	2.0	142
68	Congenital Heart Disease—Causing Gata4 Mutation Displays Functional Deficits In Vivo. <i>PLoS Genetics</i> , 2012, 8, e1002690.	1.5	77
69	Inhibition of Notch1 Signaling Reduces Abdominal Aortic Aneurysm in Mice by Attenuating Macrophage-Mediated Inflammation. <i>Arteriosclerosis, Thrombosis, and Vascular Biology</i> , 2012, 32, 3012-3023.	1.1	58
70	Submicroscopic Chromosomal Copy Number Variations Identified in Children With Hypoplastic Left Heart Syndrome. <i>Pediatric Cardiology</i> , 2012, 33, 757-763.	0.6	35
71	Molecular Basis of Cardiac Development and Congenital Heart Disease. , 2012, , 317-339.		2
72	Growth of the Normal Human Heart. , 2012, , 1305-1316.		0

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73	Inhibitory Role of Notch1 in Calcific Aortic Valve Disease. PLoS ONE, 2011, 6, e27743.	1.1	96
74	Human balanced translocation and mouse gene inactivation implicate Basonuclin 2 in distal urethral development. European Journal of Human Genetics, 2011, 19, 540-546.	1.4	28
75	Chromosomal Haplotypes by Genetic Phasing of Human Families. American Journal of Human Genetics, 2011, 89, 382-397.	2.6	63
76	Heredity of bicuspid aortic valve: is family screening indicated?. Heart, 2011, 97, 1193-1195.	1.2	19
77	Genetics of Congenital Heart Disease. Current Cardiology Reviews, 2010, 6, 91-97.	0.6	153
78	Impact of Mendelian inheritance in cardiovascular disease. Annals of the New York Academy of Sciences, 2010, 1214, 122-137.	1.8	13
79	Fetal and postnatal lung defects reveal a novel and required role for Fgf8 in lung development. Developmental Biology, 2010, 347, 92-108.	0.9	19
80	Identification of GATA6 Sequence Variants in Patients With Congenital Heart Defects. Pediatric Research, 2010, 68, 281-285.	1.1	105
81	Interaction of Gata4 and Gata6 with Tbx5 is critical for normal cardiac development. Developmental Biology, 2009, 326, 368-377.	0.9	168
82	EXPRESSION PATTERNS OF BASONUCLIN 2 IN PENILE TISSUE. Journal of Urology, 2009, 181, 253-253.	0.2	0
83	A novel mutation in LAMIN A/C is associated with isolated early-onset atrial fibrillation and progressive atrioventricular block followed by cardiomyopathy and sudden cardiac death. Heart Rhythm, 2009, 6, 707-710.	0.3	44
84	Cryptic Chromosomal Abnormalities Identified in Children With Congenital Heart Disease. Pediatric Research, 2008, 64, 358-363.	1.1	91
85	A Rare Human Sequence Variant Reveals Myocardin Autoinhibition. Journal of Biological Chemistry, 2008, 283, 35845-35852.	1.6	15
86	GATA4 sequence variants in patients with congenital heart disease. Journal of Medical Genetics, 2007, 44, 779-783.	1.5	180
87	Spectrum of heart disease associated with murine and human GATA4 mutation. Journal of Molecular and Cellular Cardiology, 2007, 43, 677-685.	0.9	218
88	Screening and biochemical analysis of GATA4 sequence variations identified in patients with congenital heart disease. American Journal of Medical Genetics, Part A, 2007, 143A, 817-823.	0.7	59
89	Molecular genetics of aortic valve disease. Current Opinion in Cardiology, 2006, 21, 180-184.	0.8	104
90	Insights into the genetic basis of congenital heart disease. Cellular and Molecular Life Sciences, 2006, 63, 1141-1148.	2.4	76

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91	Mutations in NOTCH1 cause aortic valve disease. Nature, 2005, 437, 270-274.	13.7	1,274
92	Genetic Underpinnings of Cardiogenesis and Congenital Heart Disease. , 2005, , 155-164.		0
93	GATA4 mutations cause human congenital heart defects and reveal an interaction with TBX5. Nature, 2003, 424, 443-447.	13.7	1,086
94	Tbx1, a DiGeorge Syndrome Candidate Gene, Is Regulated by Sonic Hedgehog during Pharyngeal Arch Development. Developmental Biology, 2001, 235, 62-73.	0.9	282
95	A common cis-acting sequence in the DiGeorge critical region regulates bi-directional transcription of UFD1L and CDC45L. Mechanisms of Development, 2001, 108, 81-92.	1.7	9
96	A Molecular Pathway Revealing a Genetic Basis for Human Cardiac and Craniofacial Defects. Science, 1999, 283, 1158-1161.	6.0	280