Sarah U Morton

List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

35 606 10 24 g-index

42 1,017 9.8 3.46 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
35	Genome-Wide De Novo Variants in Congenital Heart Disease Are Not Associated With Maternal Diabetes or Obesity <i>Circulation Genomic and Precision Medicine</i> , 2022 , CIRCGEN121003500	5.2	O
34	Transcription factor protein interactomes reveal genetic determinants in heart disease Cell, 2022,	56.2	3
33	Neither cardiac mitochondrial DNA variation or copy number contribute to congenital heart disease risk <i>American Journal of Human Genetics</i> , 2022 ,	11	1
32	An ancient founder mutation located between and is responsible for increased microtia risk in Amerindigenous populations <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022 , 119, e2203928119	11.5	O
31	Increased Breastfeeding Proportion Is Associated with Improved Gross Motor Skills at 35 Years of Age: A Pilot Study. <i>Nutrients</i> , 2022 , 14, 2215	6.7	
30	Reducing Benzodiazepine Exposure by Instituting a Guideline for Dexmedetomidine Usage in the NICU. <i>Pediatrics</i> , 2021 , 148,	7.4	2
29	microRNA-mRNA Profile of Skeletal Muscle Differentiation and Relevance to Congenital Myotonic Dystrophy. <i>International Journal of Molecular Sciences</i> , 2021 , 22,	6.3	2
28	Immune activation during brain infection in African infants with frequent cytomegalovirus co-infection. <i>IScience</i> , 2021 , 24, 102351	6.1	O
27	Quantification of magnetic resonance spectroscopy data using a combined reference: Application in typically developing infants. <i>NMR in Biomedicine</i> , 2021 , 34, e4520	4.4	1
26	Mechanisms of Congenital Heart Disease Caused by NAA15 Haploinsufficiency. <i>Circulation Research</i> , 2021 , 128, 1156-1169	15.7	2
25	Abnormal Right-Hemispheric Sulcal Patterns Correlate with Executive Function in Adolescents with Tetralogy of Fallot. <i>Cerebral Cortex</i> , 2021 , 31, 4670-4680	5.1	1
24	Association of Damaging Variants in Genes With Increased Cancer Risk Among Patients With Congenital Heart Disease. <i>JAMA Cardiology</i> , 2021 , 6, 457-462	16.2	12
23	Genomic frontiers in congenital heart disease. <i>Nature Reviews Cardiology</i> , 2021 ,	14.8	15
22	Congenital Heart Defects Due to Missense Variants. <i>Circulation Genomic and Precision Medicine</i> , 2020 , 13, e002843	5.2	3
21	Screening With Reticulocyte Hemoglobin Increased Iron Sufficiency Among NICU Patients. <i>Pediatric Quality & Safety</i> , 2020 , 5, e258	1	1
20	Genomic analyses implicate noncoding de novo variants in congenital heart disease. <i>Nature Genetics</i> , 2020 , 52, 769-777	36.3	33
19	De Novo Damaging Variants, Clinical Phenotypes, and Post-Operative Outcomes in Congenital Heart Disease. <i>Circulation Genomic and Precision Medicine</i> , 2020 , 13, e002836	5.2	15

18	EM-mosaic detects mosaic point mutations that contribute to congenital heart disease. <i>Genome Medicine</i> , 2020 , 12, 42	14.4	8
17	Rare genetic variation at transcription factor binding sites modulates local DNA methylation profiles. <i>PLoS Genetics</i> , 2020 , 16, e1009189	6	7
16	mutations in hiPSCs inform mechanisms for maldevelopment of the heart, pancreas, and diaphragm. <i>ELife</i> , 2020 , 9,	8.9	9
15	Maternal Dietary Intake of Omega-3 Fatty Acids Correlates Positively with Regional Brain Volumes in 1-Month-Old Term Infants. <i>Cerebral Cortex</i> , 2020 , 30, 2057-2069	5.1	8
14	infection with frequent viral coinfection contributes to postinfectious hydrocephalus in Ugandan infants. <i>Science Translational Medicine</i> , 2020 , 12,	17.5	13
13	Association of nucleated red blood cell count with mortality among neonatal intensive care unit patients. <i>Pediatrics and Neonatology</i> , 2020 , 61, 592-597	1.8	4
12	Abnormal Left-Hemispheric Sulcal Patterns Correlate with Neurodevelopmental Outcomes in Subjects with Single Ventricular Congenital Heart Disease. <i>Cerebral Cortex</i> , 2020 , 30, 476-487	5.1	11
11	Genome sequencing as a first-line genetic test in familial dilated cardiomyopathy. <i>Genetics in Medicine</i> , 2019 , 21, 650-662	8.1	36
10	Mammalian Hbs1L deficiency causes congenital anomalies and developmental delay associated with Pelota depletion and 80S monosome accumulation. <i>PLoS Genetics</i> , 2019 , 15, e1007917	6	4
9	Psychosocial Stress and Adversity: Effects from the Perinatal Period to Adulthood. <i>NeoReviews</i> , 2019 , 20, e686-e696	1.1	15
8	Response to Brodehl et al. <i>Genetics in Medicine</i> , 2019 , 21, 1248-1249	8.1	
7	Reducing time to initiation and advancement of enteral feeding in an all-referral neonatal intensive care unit. <i>Journal of Perinatology</i> , 2018 , 38, 936-943	3.1	4
6	Genome-Wide Association Study Identifies a Susceptibility Locus for Comitant Esotropia and Suggests a Parent-of-Origin Effect 2018 , 59, 4054-4064		10
5	Contribution of rare inherited and de novo variants in 2,871 congenital heart disease probands. <i>Nature Genetics</i> , 2017 , 49, 1593-1601	36.3	348
4	Separating Putative Pathogens from Background Contamination with Principal Orthogonal Decomposition: Evidence for Leptospira in the Ugandan Neonatal Septisome. <i>Frontiers in Medicine</i> , 2016 , 3, 22	4.9	4
3	Treatment options for apnoea of prematurity. <i>Archives of Disease in Childhood: Fetal and Neonatal Edition</i> , 2016 , 101, F352-6	4.7	26
2	Skeletal muscle microRNA and messenger RNA profiling in cofilin-2 deficient mice reveals cell cycle dysregulation hindering muscle regeneration. <i>PLoS ONE</i> , 2015 , 10, e0123829	3.7	4
1	Whole Genome De Novo Variant Identification with FreeBayes and Neural Network Approaches		3