Ashish Kapoor

List of Publications by Year
in descending order

Source: https://exaly.com/author-pdf/9085390/publications.pdf

Version: 2024-02-01

21	844	687363 13	752698 20
papers	citations	h-index	g-index

24	24	24	1219
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Molecular Genetic Anatomy and Risk Profile of Hirschsprungâ€™s Disease. New England Journal of Medicine, 2019, 380, 1421-1432.	27.0	131
2	Functional Loss of Semaphorin 3C and/or Semaphorin 3D and Their Epistatic Interaction with Ret Are Critical to Hirschsprung Disease Liability. American Journal of Human Genetics, 2015, 96, 581-596.	6.2	118
3	Enhancer Variants Synergistically Drive Dysfunction of a Gene Regulatory Network In Hirschsprung Disease. Cell, 2016, 167, 355-368.e10.	28.9	112
4	A Multicenter Study of BRD2 as a Risk Factor for Juvenile Myoclonic Epilepsy. Epilepsia, 2007, 48, 706-712.	5.1	76
5	An Enhancer Polymorphism at the Cardiomyocyte Intercalated Disc Protein NOS1AP Locus Is a Major Regulator of the QT Interval. American Journal of Human Genetics, 2014, 94, 854-869.	6.2	72
6	Population variation in total genetic risk of Hirschsprung disease from common RET, SEMA3 and NRG1 susceptibility polymorphisms. Human Molecular Genetics, 2015, 24, 2997-3003.	2.9	66
7	An idiopathic epilepsy syndrome linked to 3q13.3â€q21 and missense mutations in the extracellular calcium sensing receptor gene. Annals of Neurology, 2008, 64, 158-167.	5.3	65
8	Trans-ethnic meta-analysis of genome-wide association studies for Hirschsprung disease. Human Molecular Genetics, 2016, 25, ddw333.	2.9	38
9	Effects of RET and NRG1 polymorphisms in Indonesian patients with Hirschsprung disease. Journal of Pediatric Surgery, 2014, 49, 1614-1618.	1.6	37
10	Human cardiac cis-regulatory elements, their cognate transcription factors, and regulatory DNA sequence variants. Genome Research, 2018, 28, 1577-1588.	5.5	25
11	Multiple SCN5A variant enhancers modulate its cardiac gene expression and the QT interval. Proceedings of the National Academy of Sciences of the United States of America, 2019, 116, 10636-10645.	7.1	22
12	Mendelian Puzzles. Science, 2012, 335, 930-931.	12.6	17
13	Combined Genetic Effects of RET and NRG1 Susceptibility Variants on Multifactorial Hirschsprung Disease in Indonesia. Journal of Surgical Research, 2019, 233, 96-99.	1.6	16
14	A novel genetic locus for juvenile myoclonic epilepsy at chromosome 5q12â€“q14. Human Genetics, 2007, 121, 655-662.	3.8	12
15	A multi-enhancer RET regulatory code is disrupted in Hirschsprung disease. Genome Research, 2021, 31, 2199-2208.	5.5	10
16	Testing the Ret and Sema3d genetic interaction in mouse enteric nervous system development. Human Molecular Genetics, 2017, 26, 1811-1820.	2.9	8
17	Rare coding TTN variants are associated with electrocardiographic QT interval in the general population. Scientific Reports, 2016, 6, 28356.	3.3	6
18	Generation of a cre recombinase-conditional Nos1ap over-expression transgenic mouse. Biotechnology Letters, 2014, 36, 1179-1185.	2.2	3

#	Article	IF	Citations
19	Multiple, independent, common variants at RET, SEMA3 and NRG1 gut enhancers specify Hirschsprung disease risk in European ancestry subjects. Journal of Pediatric Surgery, 2021, 56, 2286-2294.	1.6	3
20	Sequence-based correction of barcode bias in massively parallel reporter assays. Genome Research, 2021, 31, 1638-1645.	5.5	3
21	Genetics and Genomics in Cardiovascular Gene Discovery. , 2012, , 231-259.		2