## Irfahan Kassam

List of Publications by Year in descending order

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1040056 996975 15 374 9 15 citations h-index g-index papers 1144 16 16 16 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Long-range epigenetic regulation is conferred by genetic variation located at thousands of independent loci. Nature Communications, 2015, 6, 6326.	12.8	115
2	Association of the <i>APOE-<math>\hat{l}\mu</math>4</i> allele with outcome of traumatic brain injury in children and youth: a meta-analysis and meta-regression. Journal of Neurology, Neurosurgery and Psychiatry, 2016, 87, 433-440.	1.9	44
3	The effect of X-linked dosage compensation on complex trait variation. Nature Communications, 2019, 10, 3009.	12.8	44
4	Tissue-specific sex differences in human gene expression. Human Molecular Genetics, 2019, 28, 2976-2986.	2.9	41
5	Blood triglyceride levels are associated with DNA methylation at the serine metabolism gene PHGDH. Scientific Reports, 2017, 7, 11207.	3.3	32
6	Integrative Profiling of T790M-Negative EGFR-Mutated NSCLC Reveals Pervasive Lineage Transition and Therapeutic Opportunities. Clinical Cancer Research, 2021, 27, 5939-5950.	7.0	21
7	Impact of BMI and waist circumference on epigenome-wide DNA methylation and identification of epigenetic biomarkers in blood: an EWAS in multi-ethnic Asian individuals. Clinical Epigenetics, 2021, 13, 195.	4.1	17
8	Autosomal genetic control of human gene expression does not differ across the sexes. Genome Biology, 2016, 17, 248.	8.8	15
9	New Polygenic Risk Score to Predict High Myopia in Singapore Chinese Children. Translational Vision Science and Technology, 2021, 10, 26.	2.2	11
10	Genetic determinants of tissue factor pathway inhibitor plasma levels. Thrombosis and Haemostasis, 2015, 114, 245-257.	3.4	9
11	Evidence for mitochondrial genetic control of autosomal gene expression. Human Molecular Genetics, 2016, 25, ddw347.	2.9	6
12	Genome-wide identification of cis DNA methylation quantitative trait loci in three Southeast Asian Populations. Human Molecular Genetics, 2021, 30, 603-618.	2.9	5
13	Sex-specific effect of CPB2 Ala147Thr but not Thr325lle variants on the risk of venous thrombosis: A comprehensive meta-analysis. PLoS ONE, 2017, 12, e0177768.	2.5	5
14	The Potential of Current Polygenic Risk Scores to Predict High Myopia and Myopic Macular Degeneration in Multiethnic Singapore Adults. Ophthalmology, 2022, 129, 890-902.	5.2	5
15	The autosomal genetic control of sexually dimorphic traits in humans is largely the same across the sexes. Genome Biology, 2016, 17, 169.	8.8	4