Jayashri Ghosh

List of Publications by Year
in descending order

12	480	933447 10	1199594 12
papers	citations	h-index	g-index

12	12	12	747
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	DNA methylation differences between in vitro- and in vivo-conceived children are associated with ART procedures rather than infertility. Clinical Epigenetics, 2015, 7, 41.	4.1	94
2	Global DNA methylation levels are altered by modifiable clinical manipulations in assisted reproductive technologies. Clinical Epigenetics, 2017, 9, 14.	4.1	88
3	Epigenetic changes and assisted reproductive technologies. Epigenetics, 2020, 15, 12-25.	2.7	75
4	Genomeâ€Wideâ€Associated Variants in Migraine Susceptibility: A Replication Study From North <scp>I</scp>ndia. Headache, 2013, 53, 1583-1594.	3.9	43
5	Outlier DNA methylation levels as an indicator of environmental exposure and risk of undesirable birth outcome. Human Molecular Genetics, 2016, 25, 123-129.	2.9	34
6	Investigation of TNFA 308GÂ>ÂA and TNFB 252GÂ>ÂA polymorphisms in genetic susceptibility to migraine. Journal of Neurology, 2010, 257, 898-904.	3.6	33
7	Identification of a Novel ANKK1 and Other Dopaminergic (DRD2 and DBH) Gene Variants in Migraine Susceptibility. NeuroMolecular Medicine, 2013, 15, 61-73.	3.4	31
8	Epigenetic changes in preterm birth placenta suggest a role for ADAMTS genes in spontaneous preterm birth. Human Molecular Genetics, 2019, 28, 84-95.	2.9	24
9	Potential Role of Aromatase over Estrogen Receptor Gene Polymorphisms in Migraine Susceptibility: A Case Control Study from North India. PLoS ONE, 2012, 7, e34828.	2.5	23
10	Role of Dopaminergic Gene Polymorphisms (DBH 19 bp Indel and DRD2Nco I) in Genetic Susceptibility to Migraine in North Indian Population: Table 1. Pain Medicine, 2011, 12, 1109-1111.	1.9	19
11	Highly variant DNA methylation in normal tissues identifies a distinct subclass of cancer patients. Advances in Cancer Research, 2019, 142, 1-22.	5.0	8
12	Embryo cryopreservation leads to sex-specific DNA methylation perturbations in both human and mouse placentas. Human Molecular Genetics, 2022, 31, 3855-3872.	2.9	8