Nikolay Skryabin

List of Publications by Year
in descending order

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25	334	1163117 8	839539 18
papers	citations	h-index	g-index

27	27	27	750
all docs	docs citations	times ranked	citing authors

#	Article	IF	Citations
1	Single gene microdeletions and microduplication of 3p26.3 in three unrelated families: CNTN6 as a new candidate gene for intellectual disability. Molecular Cytogenetics, 2014, 7, 97.	0.9	51
2	Array CGH analysis of a cohort of Russian patients with intellectual disability. Gene, 2014, 536, 145-150.	2.2	40
3	Karyotype of the blastocoel fluid demonstrates low concordance with both trophectoderm and inner cell mass. Fertility and Sterility, 2018, 109, 1127-1134.e1.	1.0	38
4	Genomic structural variations for cardiovascular and metabolic comorbidity. Scientific Reports, 2017, 7, 41268.	3.3	29
5	Phenotypic Drift as a Cause for Intratumoral Morphological Heterogeneity of Invasive Ductal Breast Carcinoma Not Otherwise Specified. BioResearch Open Access, 2013, 2, 148-154.	2.6	26
6	Clinically relevant morphological structures in breast cancer represent transcriptionally distinct tumor cell populations with varied degrees of epithelial-mesenchymal transition and CD44+CD24- stemness. Oncotarget, 2017, 8, 61163-61180.	1.8	22
7	Allele-Specific Biased Expression of the CNTN6 Gene in iPS Cell-Derived Neurons from a Patient with Intellectual Disability and 3p26.3 Microduplication Involving the CNTN6 Gene. Molecular Neurobiology, 2018, 55, 6533-6546.	4.0	18
8	Compound phenotype in a girl with r(22), concomitant microdeletion 22q13.32-q13.33 and mosaic monosomy 22. Molecular Cytogenetics, 2018, 11, 26.	0.9	15
9	A mosaic intragenic microduplication of LAMA1 and a constitutional 18p11.32 microduplication in a patient with keratosis pilaris and intellectual disability. American Journal of Medical Genetics, Part A, 2018, 176, 2395-2403.	1.2	9
10	Epigenetic effects of trisomy 16 in human placenta. Molecular Biology, 2013, 47, 373-381.	1.3	8
11	Relationship between morphological and cytogenetic heterogeneity in invasive micropapillary carcinoma of the breast: a report of one case. Journal of Clinical Pathology, 2015, 68, 758-762.	2.0	8
12	Epigenetic status of imprinted genes in placenta during recurrent pregnancy loss. Russian Journal of Genetics, 2017, 53, 376-387.	0.6	8
13	Delineation of Clinical Manifestations of the Inherited Xq24 Microdeletion Segregating with sXCI in Mothers: Two Novel Cases with Distinct Phenotypes Ranging from UBE2A Deficiency Syndrome to Recurrent Pregnancy Loss. Cytogenetic and Genome Research, 2020, 160, 245-254.	1.1	8
14	A de novo microtriplication at 4q21.21â€q21.22 in a patient with a vascular malignant hemangioma, elongated sigmoid colon, developmental delay, and absence of speech. American Journal of Medical Genetics, Part A, 2016, 170, 2089-2096.	1.2	7
15	DNA methylation profile in human placental tissues. Molecular Biology, 2011, 45, 493-499.	1.3	6
16	Somatic genome variations in vascular tissues and peripheral blood leukocytes in patients with atherosclerosis. Russian Journal of Genetics, 2014, 50, 870-878.	0.6	6
17	Gene Expression Profiling Revealed 2 Types of Bronchial Basal Cell Hyperplasia and Squamous Metaplasia With Different Progression Potentials. Applied Immunohistochemistry and Molecular Morphology, 2020, 28, 477-483.	1.2	6
18	Dynamics of aberrant methylation of functional groups of genes in progression of breast cancer. Molecular Biology, 2013, 47, 267-274.	1.3	5

#	Article	IF	Citations
19	Preimplantation genetic diagnosis by blastocentesis: Problems and perspectives. Russian Journal of Genetics, 2016, 52, 1-7.	0.6	5
20	Multilocus epimutations of imprintome in the pathology of human embryo development. Molecular Biology, 2012, 46, 183-191.	1.3	4
21	Molecular karyotyping of cell-free DNA from blastocoele fluid as a basis for noninvasive preimplantation genetic screening of aneuploidy. Russian Journal of Genetics, 2015, 51, 1123-1128.	0.6	4
22	NLRP7 variants in spontaneous abortions with multilocus imprinting disturbances from women with recurrent pregnancy loss. Journal of Assisted Reproduction and Genetics, 2021, 38, 2893-2908.	2.5	4
23	Epigenetic silencing of genomic structural variations. Russian Journal of Genetics, 2017, 53, 1072-1079.	0.6	3
24	Analysis of mutations spectrum in the ATP7B gene in patients with Wilson disease using massively parallel sequencing. Klinicheskaya Laboratornaya Diagnostika, 2022, 67, 250-256.	0.5	1
25	Runs of homozygosity in spontaneous abortions from families with recurrent pregnancy loss. Vavilovskii Zhurnal Genetiki I Selektsii, 2019, 23, 244-249.	1.1	0