Oksana Naumova

List of Publications by Year in Descending Order

35 papers	818 citations	13 h-index	28 g-index
40 ext. papers	933 ext. citations	2.7 avg, IF	3.54 L-index

#	Paper	IF	Citations
35	A Study of the Genomic Variations Associated with Autistic Spectrum Disorders in a Russian Cohort of Patients Using Whole-Exome Sequencing. *Genes*, **2022**, 13, 920	4.2	0
34	Neuroendocrine and autonomic stress systems activity in young adults raised by mothers with mental health and substance abuse problems: A prospective cohort study. *Developmental Psychobiology*, **2021**, 63, e22213	3	
33	Identification of Pathogenic CNVs in Unexplained Developmental Disabilities Using Exome Sequencing: A Family Trio Study. *Russian Journal of Genetics*, **2021**, 57, 1351-1355	0.6	1
32	Cohesion of Cortical Language Networks During Word Processing Is Predicted by a Common Polymorphism in the SETBP1 Gene. *New Directions for Child and Adolescent Development*, **2020**, 2020, 131-155	1.3	
31	Resting-State EEG Spectral Power in Children with Experience of Early Deprivation. *ksperimental?na Psihologi*, **2020**, 13, 115-124	0.6	
30	Next Generation Sequencing of 134 Children with Autism Spectrum Disorder and Regression. *Genes* , **2020**, 11,	4.2	7
29	Male pseudohermaphroditism: A case study of 46,XY disorder of sexual development using whole-exome sequencing. *Clinical Case Reports (discontinued)*, **2020**, 8, 2889-2894	0.7	
28	Genome-Wide Homozygosity Mapping Reveals Genes Associated With Cognitive Ability in Children From Saudi Arabia. *Frontiers in Genetics*, **2019**, 10, 888	4.5	4
27	Improved Educational Achievement as a Path to Desistance. *New Directions for Child and Adolescent Development*, **2019**, 2019, 111-135	1.3	6
26	Effects of early social deprivation on epigenetic statuses and adaptive behavior of young children: A study based on a cohort of institutionalized infants and toddlers. *PLoS ONE*, **2019**, 14, e0214285	3.7	14
25	Language Outcomes in Adults with a History of Institutionalization: Behavioral and Neurophysiological Characterization. *Scientific Reports*, **2019**, 9, 4252	4.9	9
24	A Study of the Association between Breastfeeding and DNA Methylation in Peripheral Blood Cells of Infants. *Russian Journal of Genetics*, **2019**, 55, 749-755	0.6	5
23	Negative parenting modulates the association between mother's DNA methylation profiles and adult offspring depression. *Developmental Psychobiology*, **2019**, 61, 304-310	3	4
22	DNA methylation alterations in the genome of a toddler with cri-du-chat syndrome. *Clinical Case Reports (discontinued)*, **2018**, 6, 14-17	0.7	4
21	Developmental dynamics of the epigenome: A longitudinal study of three toddlers. *Neurotoxicology and Teratology*, **2018**, 66, 125-131	3.9	3
20	Aberrant DNA methylation in lymphocytes of children with neurodevelopmental disorders. *Russian Journal of Genetics*, **2017**, 53, 1243-1258	0.6	1
19	*Russian Journal of Genetics*, **2017**, 1320-1337	0.8	

18	Epigenetic regulation of cognition: A circumscribed review of the field. *Development and Psychopathology*, **2016**, 28, 1285-1304	4.3	26
17	Health, Development and Epigenetic Characteristics of Institutionalized Children: A Preliminary Study based on a Small Cohort. *Procedia, Social and Behavioral Sciences*, **2016**, 233, 225-230		
16	Epigenetic Patterns Modulate the Connection Between Developmental Dynamics of Parenting and Offspring Psychosocial Adjustment. *Child Development*, **2016**, 87, 98-110	4.9	31
15	Gene expression in the human brain: the current state of the study of specificity and spatiotemporal dynamics. *Child Development*, **2013**, 84, 76-88	4.9	37
14	Comparison of whole-genome DNA methylation patterns in whole blood, saliva, and lymphoblastoid cell lines. *Behavior Genetics*, **2013**, 43, 168-76	3.2	55
13	A substantial prehistoric European ancestry amongst Ashkenazi maternal lineages. *Nature Communications*, **2013**, 4, 2543	17.4	62
12	Differential patterns of whole-genome DNA methylation in institutionalized children and children raised by their biological parents. *Development and Psychopathology*, **2012**, 24, 143-55	4.3	211
11	Age-related changes of gene expression in the neocortex: preliminary data on RNA-Seq of the transcriptome in three functionally distinct cortical areas. *Development and Psychopathology*, **2012**, 24, 1427-42	4.3	18
10	Childhood adversity and DNA methylation of genes involved in the hypothalamus-pituitary-adrenal axis and immune system: whole-genome and candidate-gene associations. *Development and Psychopathology*, **2012**, 24, 1417-25	4.3	97
9	10.1007/s11177-008-2015-1 **2010**, 44, 215		
8	Mitochondrial DNA diversity in Kazym Khanty. *Russian Journal of Genetics*, **2009**, 45, 756-760	0.6	2
7	Mitochondrial DNA variability in populations and ethnic groups of Tatars of the Tobol-Irtysh basin. *Russian Journal of Genetics*, **2009**, 45, 1107-1116	0.6	1
6	Analysis of Y chromosome STR haplotypes in the European part of Russia reveals high diversities but non-significant genetic distances between populations. *International Journal of Legal Medicine*, **2008**, 122, 219-23	3.1	48
5	Boundaries and clines in the West Eurasian Y-chromosome landscape: insights from the European part of Russia. *American Journal of Physical Anthropology*, **2008**, 137, 41-7	2.5	22
4	Mitochondrial DNA diversity in Siberian Tatars of the Tobol-Irtysh basin. *Russian Journal of Genetics*, **2008**, 44, 215-226	0.6	13
3	Mitochondrial DNA Polymorphism in Russian Population form Five Oblasts of the European Part of Russia. *Russian Journal of Genetics*, **2005**, 41, 1040-1045	0.6	7
2	Mitochondrial DNA and Y-chromosome variation in the caucasus. *Annals of Human Genetics*, **2004**, 68, 205-21	2.2	105
1	Genetic evidence concerning the origins of South and North Ossetians. *Annals of Human Genetics*, **2004**, 68, 588-99	2.2	22