## Alexey S Ruzov

## List of Publications by Citations

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18 42 1,397 37 g-index h-index citations papers 6.3 1,571 43 3.99 avg, IF L-index ext. citations ext. papers

#	Paper	IF	Citations
42	The p120 catenin partner Kaiso is a DNA methylation-dependent transcriptional repressor. <i>Genes and Development</i> , <b>2001</b> , 15, 1613-8	12.6	359
41	Enzymatic approaches and bisulfite sequencing cannot distinguish between 5-methylcytosine and 5-hydroxymethylcytosine in DNA. <i>BioTechniques</i> , <b>2010</b> , 48, 317-9	2.5	173
40	Lineage-specific distribution of high levels of genomic 5-hydroxymethylcytosine in mammalian development. <i>Cell Research</i> , <b>2011</b> , 21, 1332-42	24.7	161
39	Kaiso is a genome-wide repressor of transcription that is essential for amphibian development. <i>Development (Cambridge)</i> , <b>2004</b> , 131, 6185-94	6.6	90
38	N-methyladenosine regulates the stability of RNA:DNA hybrids in human cells. <i>Nature Genetics</i> , <b>2020</b> , 52, 48-55	36.3	82
37	Transient accumulation of 5-carboxylcytosine indicates involvement of active demethylation in lineage specification of neural stem cells. <i>Cell Reports</i> , <b>2014</b> , 7, 1353-1361	10.6	70
36	xDnmt1 regulates transcriptional silencing in pre-MBT Xenopus embryos independently of its catalytic function. <i>Development (Cambridge)</i> , <b>2008</b> , 135, 1295-302	6.6	58
35	The interaction of xKaiso with xTcf3: a revised model for integration of epigenetic and Wnt signalling pathways. <i>Development (Cambridge)</i> , <b>2009</b> , 136, 723-7	6.6	45
34	5-hydroxymethyl-cytosine enrichment of non-committed cells is not a universal feature of vertebrate development. <i>Epigenetics</i> , <b>2012</b> , 7, 383-9	5.7	44
33	The non-methylated DNA-binding function of Kaiso is not required in early Xenopus laevis development. <i>Development (Cambridge)</i> , <b>2009</b> , 136, 729-38	6.6	40
32	Planarian MBD2/3 is required for adult stem cell pluripotency independently of DNA methylation. <i>Developmental Biology</i> , <b>2013</b> , 384, 141-53	3.1	27
31	MBD4 and MLH1 are required for apoptotic induction in xDNMT1-depleted embryos. <i>Development</i> (Cambridge), 2009, 136, 2277-86	6.6	27
30	5-Carboxylcytosine levels are elevated in human breast cancers and gliomas. <i>Clinical Epigenetics</i> , <b>2015</b> , 7, 88	7.7	25
29	Epigenetic silencing in embryogenesis. Experimental Cell Research, 2005, 309, 241-9	4.2	23
28	High constitutive level of NF-kappaB is crucial for viability of adenocarcinoma cells. <i>Cell Death and Differentiation</i> , <b>2001</b> , 8, 621-30	12.7	23
27	Semi-quantitative immunohistochemical detection of 5-hydroxymethyl-cytosine reveals conservation of its tissue distribution between amphibians and mammals. <i>Epigenetics</i> , <b>2012</b> , 7, 137-40	5.7	21
26	Dynamics of 5-carboxylcytosine during hepatic differentiation: Potential general role for active demethylation by DNA repair in lineage specification. <i>Epigenetics</i> , <b>2017</b> , 12, 277-286	5.7	18

## (2021-2018)

25	Molecular Mechanisms Governing the Stem Cell' Fate in Brain Cancer: Factors of Stemness and Quiescence. <i>Frontiers in Cellular Neuroscience</i> , <b>2018</b> , 12, 388	6.1	18	
24	Medulloblastoma and ependymoma cells display increased levels of 5-carboxylcytosine and elevated expression. <i>Clinical Epigenetics</i> , <b>2017</b> , 9, 18	7.7	12	
23	Population Epigenomics: Advancing Understanding of Phenotypic Plasticity, Acclimation, Adaptation and Diseases. <i>Population Genomics</i> , <b>2018</b> , 179-260	1.4	12	
22	5-Carboxylcytosine is localized to euchromatic regions in the nuclei of follicular cells in axolotl ovary. <i>Nucleus</i> , <b>2012</b> , 3, 565-9	3.9	11	
21	Developmental Functions of the Dynamic DNA Methylome and Hydroxymethylome in the Mouse and Zebrafish: Similarities and Differences. <i>Frontiers in Cell and Developmental Biology</i> , <b>2018</b> , 6, 27	5.7	8	
20	Detection of Modified Forms of Cytosine Using Sensitive Immunohistochemistry. <i>Journal of Visualized Experiments</i> , <b>2016</b> ,	1.6	8	
19	Cloning and developmental expression of MARK/Par-1/MELK-related protein kinase xMAK-V in Xenopus laevis. <i>Development Genes and Evolution</i> , <b>2004</b> , 214, 139-43	1.8	7	
18	White matter tract and glial-associated changes in 5-hydroxymethylcytosine following chronic cerebral hypoperfusion. <i>Brain Research</i> , <b>2014</b> , 1592, 82-100	3.7	6	
17	WilmsUTumor Protein 1 and Enzymatic Oxidation of 5-Methylcytosine in Brain Tumors: Potential Perspectives. <i>Frontiers in Cell and Developmental Biology</i> , <b>2018</b> , 6, 26	5.7	5	
16	Kaiso, a New Protein of the BTB/POZ Family, Specifically Binds to Methylated DNA Sequences. <i>Russian Journal of Genetics</i> , <b>2001</b> , 37, 603-609	0.6	4	
15	Mass spectrometry reveals the presence of specific set of epigenetic DNA modifications in the Norway spruce genome. <i>Scientific Reports</i> , <b>2019</b> , 9, 19314	4.9	4	
14	Evidence for Noncytosine Epigenetic DNA Modifications in Multicellular Eukaryotes: An Overview. <i>Methods in Molecular Biology</i> , <b>2021</b> , 2198, 15-25	1.4	3	
13	Immunohistochemical Detection of Oxidized Forms of 5-Methylcytosine in Embryonic and Adult Brain Tissue. <i>Neuromethods</i> , <b>2016</b> , 125-137	0.4	2	
12	Immunostaining for DNA Modifications: Computational Analysis of Confocal Images. <i>Journal of Visualized Experiments</i> , <b>2017</b> ,	1.6	2	
11	A B-cell targeting virus disrupts potentially protective genomic methylation patterns in lymphoid tissue by increasing global 5-hydroxmethylcytosine levels. <i>Veterinary Research</i> , <b>2014</b> , 45, 108	3.8	2	
10	5-formylcytosine and 5-hydroxymethyluracil as surrogate markers of TET2 and SF3B1 mutations in myelodysplastic syndrome, respectively. <i>Haematologica</i> , <b>2020</b> , 105, e213-e215	6.6	1	
9	A B-cell targeting virus disrupts potentially protective genomic methylation patterns in lymphoid tissue by increasing global 5-hydroxymethylcytosine levels. <i>Veterinary Research</i> , <b>2014</b> , 45, 108	3.8	1	
8	LINE-1 transcription in round spermatids is associated with accretion of 5-carboxylcytosine in their open reading frames. <i>Communications Biology</i> , <b>2021</b> , 4, 691	6.7	1	

7	SWI/SNF complexes as determinants of R-loop metabolism. <i>Nature Genetics</i> , <b>2021</b> , 53, 940-941	36.3	1
6	Modified Forms of Cytosine in Eukaryotes: DNA (De)methylation and Beyond. <i>Methods in Molecular Biology</i> , <b>2021</b> , 2198, 3-13	1.4	1
5	A high constitutive level of NF- <b>B</b> is necessary for the viability of mouse adenocarcinoma cells: A possible role of p53. <i>Molecular Biology</i> , <b>2000</b> , 34, 655-661	1.2	
4	Detection of Low-Abundance DNA Modifications Using Signal Amplification-Based Immunocytochemistry. <i>Methods in Molecular Biology</i> , <b>2021</b> , 2198, 169-181	1.4	
3	Analysis of 5-Carboxylcytosine Distribution Using DNA Immunoprecipitation. <i>Methods in Molecular Biology</i> , <b>2021</b> , 2198, 311-319	1.4	
2	Detection and Quantification of RNA Modifications on RNADNA Hybrids Using SID-UPLC-MS/MS. <i>Methods in Molecular Biology</i> , <b>2022</b> , 127-143	1.4	
1	Detecting and Mapping N6-Methyladenosine on RNA/DNA Hybrids. <i>Methods in Molecular Biology</i> , <b>2022</b> , 329-344	1.4	