Yegor S Vassetzky

List of Publications by Year
in descending order

147 papers	3,507 citations	109321 35 h-index	182427 51 g-index
154 all docs	154 docs citations	154 times ranked	4267 citing authors

#	Article	IF	Citations
1	Mitotic Remodeling of the Replicon and Chromosome Structure. Cell, 2005, 123, 787-801.	28.9	175
2	Nucleolus: A Central Hub for Nuclear Functions. Trends in Cell Biology, 2019, 29, 647-659.	7.9	119
3	Myoblasts from affected and nonâ€affected FSHD muscles exhibit morphological differentiation defects. Journal of Cellular and Molecular Medicine, 2010, 14, 275-289.	3.6	112
4	Functional muscle impairment in facioscapulohumeral muscular dystrophy is correlated with oxidative stress and mitochondrial dysfunction. Free Radical Biology and Medicine, 2012, 53, 1068-1079.	2.9	106
5	Increased levels of adenine nucleotide translocator 1 protein and response to oxidative stress are early events in facioscapulohumeral muscular dystrophy muscle. Journal of Molecular Medicine, 2005, 83, 216-224.	3.9	98
6	mTORC1 pathway in DNA damage response. Biochimica Et Biophysica Acta - Molecular Cell Research, 2018, 1865, 1293-1311.	4.1	97
7	The Epigenetic Landscape of Mammary Gland Development and Functional Differentiation. Journal of Mammary Gland Biology and Neoplasia, 2010, 15, 85-100.	2.7	88
8	Chromatin loop domain organization within the 4q35 locus in facioscapulohumeral dystrophy patients versus normal human myoblasts. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 6982-6987.	7.1	82
9	Simultaneous miRNA and mRNA transcriptome profiling of human myoblasts reveals a novel set of myogenic differentiation-associated miRNAs and their target genes. BMC Genomics, 2013, 14, 265.	2.8	75
10	DUX4-induced constitutive DNA damage and oxidative stress contribute to aberrant differentiation of myoblasts from FSHD patients. Free Radical Biology and Medicine, 2016, 99, 244-258.	2.9	73
11	DNA topoisomerase II mutations and resistance to anti-tumor drugs. BioEssays, 1995, 17, 767-774.	2.5	70
12	Chromosome Conformation Capture (from 3C to 5C) and Its ChIP-Based Modification. Methods in Molecular Biology, 2009, 567, 171-188.	0.9	70
13	Chromatin Domains and Regulation of Transcription. Journal of Molecular Biology, 2007, 369, 597-607.	4.2	69
14	Characterization of DNA pattern in the site of permanent attachment to the nuclear matrix located in the vicinity of replication origin. Biochemical and Biophysical Research Communications, 1990, 168, 9-15.	2.1	68
15	HIV-1 Tat protein induces DNA damage in human peripheral blood B-lymphocytes via mitochondrial ROS production. Redox Biology, 2018, 15, 97-108.	9.0	62
16	A nuclear matrix attachment site in the 4q35 locus has an enhancer-blocking activity in vivo: Implications for the facio-scapulo-humeral dystrophy. Genome Research, 2008, 18, 39-45.	5.5	61
17	Defective Regulation of MicroRNA Target Genes in Myoblasts from Facioscapulohumeral Dystrophy Patients. Journal of Biological Chemistry, 2013, 288, 34989-35002.	3.4	61
18	High Resolution Genome-Wide Analysis of Chromosomal Alterations in Burkitt's Lymphoma. PLoS ONE, 2009, 4, e7089.	2.5	60

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19	Chromatin remodelling and DNA replication: from nucleosomes to loop domains. Oncogene, 2001, 20, 3086-3093.	5.9	57
20	Transcription factories in the context of the nuclear and genome organization. Nucleic Acids Research, 2011, 39, 9085-9092.	14.5	53
21	Rearrangement of chromatin domains during development in<i>Xenopus</i>. Genes and Development, 2000, 14, 1541-1552.	5.9	53
22	Topoisomerase II forms multimers in vitro: effects of metals, beta-glycerophosphate, and phosphorylation of its C-terminal domain.. Molecular and Cellular Biology, 1994, 14, 6962-6974.	2.3	51
23	Antagonistic functional duality of cancer genes. Gene, 2013, 529, 199-207.	2.2	51
24	Nuclear matrix attachment regions and topoisomerase II binding and reaction sites in the vicinity of a chicken DNA replication origin. Biochemical and Biophysical Research Communications, 1991, 177, 265-270.	2.1	50
25	Order and stochasticity in the folding of individual Drosophila genomes. Nature Communications, 2021, 12, 41.	12.8	49
26	A requiem to the nuclear matrix: from a controversial concept to 3D organization of the nucleus. Chromosoma, 2014, 123, 217-224.	2.2	47
27	Nuclear skeleton, DNA domains and control of replication and transcription. FEBS Journal, 1991, 200, 613-624.	0.2	45
28	MiR-34a is up-regulated in response to low dose, low energy X-ray induced DNA damage in breast cells. Radiation Oncology, 2013, 8, 231.	2.7	45
29	A Functional Role for 4qA/B in the Structural Rearrangement of the 4q35 Region and in the Regulation of FRG1 and ANT1 in Facioscapulohumeral Dystrophy. PLoS ONE, 2008, 3, e3389.	2.5	44
30	HIV Tat induces a prolonged MYC relocalization next to IGH in circulating B-cells. Leukemia, 2017, 31, 2515-2522.	7.2	44
31	Oncogenic Properties of the EBV ZEBRA Protein. Cancers, 2020, 12, 1479.	3.7	43
32	Dynamics of double strand breaks and chromosomal translocations. Molecular Cancer, 2014, 13, 249.	19.2	42
33	Temozolomide promotes genomic and phenotypic changes in glioblastoma cells. Cancer Cell International, 2016, 16, 36.	4.1	40
34	Eukaryotic enhancers: common features, regulation, and participation in diseases. Cellular and Molecular Life Sciences, 2015, 72, 2361-2375.	5.4	39
35	A Comparison of Techniques to Evaluate the Effectiveness of Genome Editing. Trends in Biotechnology, 2018, 36, 147-159.	9.3	38
36	DNA replication initiates at domains overlapping with nuclear matrix attachment regions in the xenopus and mouse c-myc promoter. Gene, 2004, 332, 129-138.	2.2	37

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37	Perinucleolar relocalization and nucleolin as crucial events in the transcriptional activation of key genes in mantle cell lymphoma. Blood, 2014, 123, 2044-2053.	1.4	37
38	HIVâ€1, HAART and cancer: A complex relationship. International Journal of Cancer, 2020, 146, 2666-2679.	5.1	37
39	Expression of SARS-CoV-2 entry factors in lung epithelial stem cells and its potential implications for COVID-19. Scientific Reports, 2020, 10, 17772.	3.3	37
40	Pearls in the junk: Dissecting the molecular pathogenesis of facioscapulohumeral muscular dystrophy. Neuromuscular Disorders, 2009, 19, 17-20.	0.6	30
41	Bradykinin antagonists and thiazolidinone derivatives as new potential anti-cancer compounds. Bioorganic and Medicinal Chemistry, 2014, 22, 3815-3823.	3.0	27
42	Step-wise and punctuated genome evolution drive phenotype changes of tumor cells. Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis, 2015, 771, 56-69.	1.0	27
43	Ectopic expression of inactive forms of yeast DNA topoisomerase II confers resistance to the anti-tumour drug, etoposide. British Journal of Cancer, 1996, 73, 1201-1209.	6.4	25
44	Functional roles of HIV-1 Tat protein in the nucleus. Cellular and Molecular Life Sciences, 2016, 73, 589-601.	5.4	25
45	Dux4 controls migration of mesenchymal stem cells through the Cxcr4-Sdf1 axis. Oncotarget, 2016, 7, 65090-65108.	1.8	24
46	Facioscapulohumeral dystrophy myoblasts efficiently repair moderate levels of oxidative DNA damage. Histochemistry and Cell Biology, 2016, 145, 475-483.	1.7	24
47	Modulation of mTORC1 Signaling Pathway by HIV-1. Cells, 2020, 9, 1090.	4.1	24
48	A transcription-dependent DNase I â€" hypersensitive site in a far upstream segment of the chicken Î±-globin gene domain coincides with a matrix attachment region. Biochemical and Biophysical Research Communications, 1992, 184, 1226-1234.	2.1	23
49	Control of gene expression inXenopus early development. Genesis, 1998, 22, 122-131.	2.1	23
50	Mapping long-range chromatin organization within the chicken Î±-globin gene domain using oligonucleotide DNA arrays. Genomics, 2005, 85, 143-151.	2.9	23
51	The distribution of tightly bound proteins along the DNA chain reflects the type of cell differentiation. Nucleic Acids Research, 1988, 16, 3617-3633.	14.5	22
52	The KrÃ¼ppel-like Factor 15 as a Molecular Link between Myogenic Factors and a Chromosome 4q Transcriptional Enhancer Implicated in Facioscapulohumeral Dystrophy*. Journal of Biological Chemistry, 2011, 286, 44620-44631.	3.4	21
53	Distinct Distribution of Ectopically Expressed Histone Variants H2A.Bbd and MacroH2A in Open and Closed Chromatin Domains. PLoS ONE, 2012, 7, e47157.	2.5	20
54	Specification of Chromatin Domains and Regulation of Replication and Transcription During Development. Critical Reviews in Eukaryotic Gene Expression, 2000, 10, 8.	0.9	20

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55	Topologically-associating domains: gene warehouses adapted to serve transcriptional regulation. Transcription, 2016, 7, 84-90.	3.1	19
56	Interaction in vivo between the Two Matrix Attachment Regions Flanking a Single Chromatin Loop. Journal of Molecular Biology, 2009, 386, 929-937.	4.2	18
57	Topoisomerase II Forms Multimers In Vitro: Effects of Metals, Î²-Glycerophosphate, and Phosphorylation of Its C-Terminal Domain. Molecular and Cellular Biology, 1994, 14, 6962-6974.	2.3	18
58	SETDB1 fuels the lung cancer phenotype by modulating epigenome, 3D genome organization and chromatin mechanical properties. Nucleic Acids Research, 2022, 50, 4389-4413.	14.5	18
59	Transcriptional enhancer in the vicinity of a replication origin within the 5â€² region of the chicken Î±-globin gene domain. Journal of Molecular Biology, 1991, 217, 595-598.	4.2	17
60	Domain organization of eukaryotic genome. Cell Biology International Reports, 1992, 16, 697-708.	0.6	17
61	Tat basic domain: A â€œSwiss army knifeâ€•of HIVâ€1 Tat?. Reviews in Medical Virology, 2019, 29, e2031.	8.3	17
62	The sequence-specific nuclear matrix binding factor F6 is a chicken GATA-like protein. Molecular Genetics and Genomics, 1993, 238, 309-314.	2.4	16
63	HIVâ€1 Tat protein induces aberrant activation of AICDA in human Bâ€lymphocytes from peripheral blood. Journal of Cellular Physiology, 2019, 234, 15678-15685.	4.1	16
64	Chromatin Domains and Territories: Flexibly Rigid. Critical Reviews in Eukaryotic Gene Expression, 2004, 14, 79-88.	0.9	16
65	Rearrangement of chromatin domains in cancer and development. Journal of Cellular Biochemistry, 2000, 79, 54-60.	2.6	15
66	Selective matrix attachment regions in T helper cell subsets support loop conformation in the Ifng gene. Genes and Immunity, 2007, 8, 35-43.	4.1	15
67	The microRNA-205-5p is correlated to metastatic potential of 21T series: A breast cancer progression model. PLoS ONE, 2017, 12, e0173756.	2.5	15
68	The role of Alu-derived RNAs in Alzheimerâ€™s and other neurodegenerative conditions. Medical Hypotheses, 2018, 115, 29-34.	1.5	15
69	DUX4 Pathological Expression: Causes and Consequences in Cancer. Trends in Cancer, 2019, 5, 268-271.	7.4	15
70	The presence of sequence-specific protein binding sites correlate with replication activity and matrix binding in a 1.7 Kb-long DNA fragment of the chicken â^•globin gene domain. Biochemical and Biophysical Research Communications, 1991, 179, 512-519.	2.1	14
71	DNA polymorphism and epigenetic marks modulate the affinity of a scaffold/matrix attachment region to the nuclear matrix. European Journal of Human Genetics, 2014, 22, 1117-1123.	2.8	14
72	3D genomics imposes evolution of the domain model of eukaryotic genome organization. Chromosoma, 2017, 126, 59-69.	2.2	14

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73	A One-Step PCR-Based Assay to Evaluate the Efficiency and Precision of Genomic DNA-Editing Tools. Molecular Therapy - Methods and Clinical Development, 2017, 5, 43-50.	4.1	14
74	Nuclear matrix and structural and functional compartmentalization of the eucaryotic cell nucleus. Biochemistry (Moscow), 2014, 79, 608-618.	1.5	13
75	Control of DNA integrity in skeletal muscle under physiological and pathological conditions. Cellular and Molecular Life Sciences, 2017, 74, 3439-3449.	5.4	13
76	Genetic and Epigenetic Mechanisms of Î²-Globin Gene Switching. Biochemistry (Moscow), 2018, 83, 381-392.	1.5	13
77	DNA fragments which specifically bind to isolated nuclear matrix in vitro interact with matrix-associated DNA topoisomerase II. Biochemical and Biophysical Research Communications, 1989, 159, 1263-1268.	2.1	12
78	In embryonic chicken erythrocytes actively transcribed alpha globin genes are not associated with the nuclear matrix. Journal of Cellular Biochemistry, 2009, 106, 170-178.	2.6	12
79	Metal ions modify DNA-protecting and mutagen-scavenging capacities of the AV-153 1,4-dihydropyridine. Mutation Research - Genetic Toxicology and Environmental Mutagenesis, 2019, 845, 403077.	1.7	12
80	Analysis of genes regulated by DUX4 via oxidative stress reveals potential therapeutic targets for treatment of facioscapulohumeral dystrophy. Redox Biology, 2021, 43, 102008.	9.0	12
81	MUC1 Story: Great Expectations, Disappointments and the Renaissance. Current Medicinal Chemistry, 2019, 26, 554-563.	2.4	12
82	Early replication timing of the chicken Î±-globin gene domain correlates with its open chromatin state in cells of different lineages. Genomics, 2009, 93, 481-486.	2.9	11
83	Cancerâ€related genes in the transcription signature of facioscapulohumeral dystrophy myoblasts and myotubes. Journal of Cellular and Molecular Medicine, 2014, 18, 208-217.	3.6	11
84	Easy and robust electrotransfection protocol for efficient ectopic gene expression and genome editing in human B cells. Gene Therapy, 2023, 30, 167-171.	4.5	11
85	Epigenetic Modifications, Chromatin Distribution and <i>TP53</i> Transcription in a Model of Breast Cancer Progression. Journal of Cellular Biochemistry, 2015, 116, 533-541.	2.6	10
86	Distinct Patterns of Colocalization of the <i>CCND1</i> and <i>CMYC</i> Genes With Their Potential Translocation Partner <i>IGH</i> at Successive Stages of Bâ€Cell Differentiation. Journal of Cellular Biochemistry, 2016, 117, 1506-1510.	2.6	10
87	Evolution of the Genome 3D Organization: Comparison of Fused and Segregated Globin Gene Clusters. Molecular Biology and Evolution, 2017, 34, 1492-1504.	8.9	10
88	Analysis of the chicken DNA fragments that contain structural sites of attachment to the nuclear matrix: DNA-matrix interactions and replication. Journal of Cellular Biochemistry, 2000, 79, 1-14.	2.6	9
89	Histone deacetylase inhibitor abexinostat affects chromatin organization and gene transcription in normal B cells and in mantle cell lymphoma. Gene, 2016, 580, 134-143.	2.2	9
90	Correction of the FSHD myoblast differentiation defect by fusion with healthy myoblasts. Journal of Cellular Physiology, 2016, 231, 62-71.	4.1	9

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91	Dual Role of the Extracellular Domain of Human Mucin MUC1 in Metastasis. Journal of Cellular Biochemistry, 2017, 118, 4002-4011.	2.6	8
92	RNA-dependent disassembly of nuclear bodies. Journal of Cell Science, 2016, 129, 4509-4520.	2.0	7
93	Control of DUX4 Expression in Facioscapulohumeral Muscular Dystrophy and Cancer. Trends in Molecular Medicine, 2021, 27, 588-601.	6.7	7
94	The<i>IGH</i>locus relocalizes to a â€œrecombination compartmentâ€•in the perinucleolar region of differentiating B-lymphocytes. Oncotarget, 2017, 8, 40079-40089.	1.8	7
95	The upstream area of the chicken Î±-globin gene domain is transcribed in both directions in the same cells. FEBS Letters, 2005, 579, 4746-4750.	2.8	6
96	Recruitment of RNA Polymerase II in the Ifng Gene Promoter Correlates with the Nuclear Matrix Association in Activated T Helper Cells. Journal of Molecular Biology, 2007, 371, 317-322.	4.2	6
97	Analysis of telomeric DNA: Current approaches and methods. Russian Journal of Developmental Biology, 2009, 40, 125-144.	0.5	6
98	Ring-Like Distribution of Constitutive Heterochromatin in Bovine Senescent Cells. PLoS ONE, 2011, 6, e26844.	2.5	6
99	Tightly bound to DNA proteins: Possible universal substrates for intranuclear processes. Gene, 2012, 492, 54-64.	2.2	6
100	HIV-1 Tat Activates Akt/mTORC1 Pathway and AICDA Expression by Downregulating Its Transcriptional Inhibitors in B Cells. International Journal of Molecular Sciences, 2021, 22, 1588.	4.1	6
101	Chromatin domains and territories: flexibly rigid. Critical Reviews in Eukaryotic Gene Expression, 2004, 14, 79-88.	0.9	6
102	An unusual extended DNA loop attachment region is located in the human dystrophin gene. Journal of Cellular Physiology, 2006, 209, 515-521.	4.1	5
103	Loop domain organization of the p53 locus in normal and breast cancer cells correlates with the transcriptional status of the TP53 and the neighboring genes. Journal of Cellular Biochemistry, 2011, 112, 2072-2081.	2.6	5
104	Genome- and cell-based strategies in therapy of muscular dystrophies. Biochemistry (Moscow), 2016, 81, 678-690.	1.5	5
105	Uncoupling of oxidative phosphorylation and antioxidants affect fusion of primary human myoblasts in vitro. Biopolymers and Cell, 2016, 32, 111-117.	0.4	5
106	Molecular Coevolution of Nuclear and Nucleolar Localization Signals inside the Basic Domain of HIV-1 Tat. Journal of Virology, 2022, 96, JVI0150521.	3.4	5
107	Ca2+ Transportome and the Interorganelle Communication in Hepatocellular Carcinoma. Cells, 2022, 11, 815.	4.1	5
108	Determination of the chromatin domain structure in arrayed repeat regions: Organization of the somatic 5S RNA domain during embryogenesis inXenopus laevis. Journal of Cellular Biochemistry, 2007, 102, 1140-1148.	2.6	4

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109	Transcription- and Apoptosis-Dependent Long-Range Distribution of Tight DNAâ€“Protein Complexes in the Chicken Î±-Globin Gene. DNA and Cell Biology, 2008, 27, 615-621.	1.9	4
110	Development-dependent changes in the tight DNA-protein complexes of barley on chromosome and gene level. BMC Plant Biology, 2009, 9, 56.	3.6	4
111	Proteins tightly bound to DNA: New data and old problems. Biochemistry (Moscow), 2010, 75, 1240-1251.	1.5	4
112	Effect of Environmental Factors on Nuclear Organization and Transformation of Human B Lymphocytes. Biochemistry (Moscow), 2018, 83, 402-410.	1.5	4
113	Treatment of lymphoid cells with the topoisomerase II poison etoposide leads to an increased juxtaposition of AML1 and ETO genes on the surface of nucleoli. Biopolymers and Cell, 2011, 27, 398-403.	0.4	4
114	Translocations affecting human immunoglobulin heavy chain locus. Biopolymers and Cell, 2014, 30, 90-95.	0.4	3
115	Mobility of Nuclear Components and Genome Functioning. Biochemistry (Moscow), 2018, 83, 690-700.	1.5	3
116	HIV: implication in Burkitt lymphoma.. Biopolymers and Cell, 2012, 28, 285-287.	0.4	3
117	FSHD myoblasts fail to downregulate intermediate filament protein vimentin during myogenic differentiation. Biopolymers and Cell, 2011, 27, 359-363.	0.4	3
118	Nuclear matrix-associated DNA fragments enhance autonomous replication of plasmids in chicken cells. Biochimie, 1995, 77, 880-887.	2.6	2
119	T-antigen interactions with chromatin and p53 during the cell cycle in extracts from Xenopus eggs. Journal of Cellular Biochemistry, 1999, 75, 288-299.	2.6	2
120	A set of vectors for introduction of antibiotic resistance genes by in vitro Cre-mediated recombination. BMC Research Notes, 2008, 1, 135.	1.4	2
121	Evolution of Î±- and Î²-Globin genes and their regulatory systems in light of the hypothesis of domain organization of the genome. Biochemistry (Moscow), 2014, 79, 1141-1150.	1.5	2
122	Nuclear localization of translocation partners in differentiating B-cells. Doklady Biochemistry and Biophysics, 2015, 464, 312-314.	0.9	2
123	Histone deacetylase inhibitors and epigenetic regulation in lymphoid malignancies. Investigational New Drugs, 2015, 33, 1280-1291.	2.6	2
124	Heterochromatin restricts the mobility of nuclear bodies. Chromosoma, 2018, 127, 529-537.	2.2	2
125	DUX4, a Zygotic Genome Activator, Is Involved in Oncogenesis and Genetic Diseases. Russian Journal of Developmental Biology, 2020, 51, 176-182.	0.5	2
126	Live-Cell Imaging and Analysis of Nuclear Body Mobility. Methods in Molecular Biology, 2020, 2175, 1-9.	0.9	2

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127	Interaction between mesenchymal stem cells and myoblasts in the context of facioscapulohumeral muscular dystrophy contributes to the disease phenotype. Journal of Cellular Physiology, 0, , .	4.1	2
128	Topoisomerase I is associated with the regulatory region of transcriptionally active SV 40 minichromosomes. Molecular and Cellular Biochemistry, 1990, 95, 95-106.	3.1	1
129	Title is missing!. Russian Journal of Genetics, 2003, 39, 147-151.	0.6	1
130	Effect of DNA loop anchorage regions(LARs) and microinjection timing on expression ofÎ²-galactosidase gene injected into one-cell rabbit embryos. Journal of Cellular Biochemistry, 2004, 92, 1171-1179.	2.6	1
131	Academy reform needs a reality check. Nature, 2013, 499, 284-284.	27.8	1
132	Role of the Nucleolus in Rearrangements of the IGH Locus. Molecular Biology, 2018, 52, 182-189.	1.3	1
133	From an increase in the number of tandem repeats through the decrease of sialylation to the downregulation of MUC1 expression level. Journal of Cellular Biochemistry, 2019, 120, 4472-4484.	2.6	1
134	Rearrangement of chromatin domains in cancer and development. Journal of Cellular Biochemistry, 2000, 79, 54-60.	2.6	1
135	Differences in Transcription Patterns between Induced Pluripotent Stem Cells Produced from the Same Germ Layer Are Erased upon Differentiation. PLoS ONE, 2013, 8, e53033.	2.5	1
136	Growth suppression activity of bradykinin antagonists in glioma cells. Biopolymers and Cell, 2014, 30, 77-79.	0.4	1
137	MARs Wars: heterogeneity and clustering of DNA-binding domains in the nuclear matrix. Biopolymers and Cell, 2009, 25, 451-456.	0.4	1
138	Intranuclear localization of transcription factories and immunoglobulin heavy chain gene alleles during human B-cell maturation. Biopolymers and Cell, 2016, 32, 179-183.	0.4	1
139	Title is missing!. Russian Journal of Developmental Biology, 2003, 34, 213-217.	0.5	0
140	Spatial Organization of the Chicken Î±-Globin Gene Domain in Cells of Different Origins. Molecular Biology, 2005, 39, 851-856.	1.3	0
141	Basic science in Russia under threat. Nature, 2010, 467, 789-789.	27.8	0
142	miR-205 is involved in metastatic potential of 21T series, a breast cancer progression model. BMC Proceedings, 2013, 7, .	1.6	0
143	Structure and function of oncogene-transfected immortal cells. Biopolymers and Cell, 2014, 30, 25-28.	0.4	0
144	Nuclear skeleton, DNA domains and control of replication and transcription. , 1991, , 137-148.		0

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145	Cisplatin treatment of C6 rat glioma in vivo did not influence copy number alterations and growth pattern of tumor-derived resistant cells. Biopolymers and Cell, 2015, 31, 209-217.	0.4	0
146	Control of gene expression in Xenopus early development. Genesis, 1998, 22, 122-131.	2.1	0
147	Tâ€antigen interactions with chromatin and p53 during the cell cycle in extracts from Xenopus eggs. Journal of Cellular Biochemistry, 1999, 75, 288-299.	2.6	0