# Sergey V Razin

## List of Publications by Citations



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|---|---|---|---|
| 220 papers | 4,349 citations | 34 h-index | 56 g-index |
| 236 ext. papers | 5,061 ext. citations | 6.4 avg, IF | 5.74 L-index |

| # | Paper | IF | Citations |
|---|---|---|---|
| 220 | Single-nucleus Hi-C reveals unique chromatin reorganization at oocyte-to-zygote transition. *Nature*, **2017**, 544, 110-114 | 50.4 | 418 |
| 219 | Active chromatin and transcription play a key role in chromosome partitioning into topologically associating domains. *Genome Research*, **2016**, 26, 70-84 | 9.7 | 225 |
| 218 | Replication origins are attached to the nuclear skeleton. *Nucleic Acids Research*, **1986**, 14, 8189-207 | 20.1 | 157 |
| 217 | Cys2His2 zinc finger protein family: classification, functions, and major members. *Biochemistry (Moscow)*, **2012**, 77, 217-26 | 2.9 | 107 |
| 216 | Large-scale fragmentation of mammalian DNA in the course of apoptosis proceeds via excision of chromosomal DNA loops and their oligomers. *Journal of Biological Chemistry*, **1995**, 270, 20239-41 | 5.4 | 106 |
| 215 | Mechanisms of heat shock response in mammals. *Cellular and Molecular Life Sciences*, **2013**, 70, 4229-41 | 10.3 | 86 |
| 214 | Small molecule compounds that induce cellular senescence. *Aging Cell*, **2016**, 15, 999-1017 | 9.9 | 82 |
| 213 | Proteins tightly bound to DNA in the regions of DNA attachment to the skeletal structures of interphase nuclei and metaphase chromosomes. *Cell*, **1981**, 27, 65-73 | 56.2 | 81 |
| 212 | Control of human cytomegalovirus gene expression by differential histone modifications during lytic and latent infection of a monocytic cell line. *Gene*, **2006**, 384, 120-8 | 3.8 | 78 |
| 211 | Different topoisomerase II antitumor drugs direct similar specific long-range fragmentation of an amplified c-MYC gene locus in living cells and in high-salt-extracted nuclei. *Proceedings of the National Academy of Sciences of the United States of America*, **1995**, 92, 102-6 | 11.5 | 74 |
| 210 | Disclosure of a structural milieu for the proximity ligation reveals the elusive nature of an active chromatin hub. *Nucleic Acids Research*, **2013**, 41, 3563-75 | 20.1 | 72 |
| 209 | The channels model of nuclear matrix structure. *BioEssays*, **1995**, 17, 443-50 | 4.1 | 68 |
| 208 | Specificity and functional significance of DNA interaction with the nuclear matrix: new approaches to clarify the old questions. *International Review of Cytology*, **1995**, 162B, 405-48 | | 67 |
| 207 | Chromosome conformation capture (from 3C to 5C) and its ChIP-based modification. *Methods in Molecular Biology*, **2009**, 567, 171-88 | 1.4 | 63 |
| 206 | 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 | 3.4 | 62 |
| 205 | Nucleolus: A Central Hub for Nuclear Functions. *Trends in Cell Biology*, **2019**, 29, 647-659 | 18.3 | 61 |
| 204 | Chromatin domains and regulation of transcription. *Journal of Molecular Biology*, **2007**, 369, 597-607 | 6.5 | 61 |

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|---|---|---|---|
| 203 | Visualization of individual DNA loops and a map of loop domains in the human dystrophin gene. *Nucleic Acids Research*, **2004**, 32, 2079-86 | 20.1 | 60 |
| 202 | Nuclear lamina integrity is required for proper spatial organization of chromatin in Drosophila. *Nature Communications*, **2019**, 10, 1176 | 17.4 | 52 |
| 201 | Dual effect of heat shock on DNA replication and genome integrity. *Molecular Biology of the Cell*, **2012**, 23, 3450-60 | 3.5 | 50 |
| 200 | CTCF-dependent enhancer blockers at the upstream region of the chicken alpha-globin gene domain. *Nucleic Acids Research*, **2004**, 32, 1354-62 | 20.1 | 49 |
| 199 | In vivo formaldehyde cross-linking: it is time for black box analysis. *Briefings in Functional Genomics*, **2015**, 14, 163-5 | 4.9 | 48 |
| 198 | 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, 265-70 | 3.4 | 47 |
| 197 | Transcription factories in the context of the nuclear and genome organization. *Nucleic Acids Research*, **2011**, 39, 9085-92 | 20.1 | 46 |
| 196 | Spatial configuration of the chicken alpha-globin gene domain: immature and active chromatin hubs. *Nucleic Acids Research*, **2008**, 36, 4629-40 | 20.1 | 46 |
| 195 | Mechanism of heat stress-induced cellular senescence elucidates the exclusive vulnerability of early S-phase cells to mild genotoxic stress. *Nucleic Acids Research*, **2015**, 43, 6309-20 | 20.1 | 43 |
| 194 | Long-range fragmentation of the eukaryotic genome by exogenous and endogenous nucleases proceeds in a specific fashion via preferential DNA cleavage at matrix attachment sites. *Journal of Biological Chemistry*, **1995**, 270, 18685-90 | 5.4 | 42 |
| 193 | Genomic domains and regulatory elements operating at the domain level. *International Review of Cytology*, **2003**, 226, 63-125 | | 41 |
| 192 | Low ionic strength extraction of nuclease-treated nuclei destroys the attachment of transcriptionally active DNA to the nuclear skeleton. *Nucleic Acids Research*, **1985**, 13, 7427-44 | 20.1 | 41 |
| 191 | Synthetically Lethal Interactions of ATM, ATR, and DNA-PKcs. *Trends in Cancer*, **2018**, 4, 755-768 | 12.5 | 40 |
| 190 | Organization of the 3'-boundary of the chicken alpha globin gene domain and characterization of a CR 1-specific protein binding site. *Nucleic Acids Research*, **1990**, 18, 401-9 | 20.1 | 39 |
| 189 | A. E. Braunstein Plenary Lecture. Nuclear skeleton, DNA domains and control of replication and transcription. *FEBS Journal*, **1991**, 200, 613-24 | | 37 |
| 188 | Chromatin without the 30-nm fiber: constrained disorder instead of hierarchical folding. *Epigenetics* , **2014**, 9, 653-7 | 5.7 | 35 |
| 187 | Communication of genome regulatory elements in a folded chromosome. *FEBS Letters*, **2013**, 587, 1840-7 | 3.8 | 34 |
| 186 | A requiem to the nuclear matrix: from a controversial concept to 3D organization of the nucleus. *Chromosoma*, **2014**, 123, 217-24 | 2.8 | 33 |

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|---|---|---|---|
| 185 | Dynamics of double strand breaks and chromosomal translocations. *Molecular Cancer*, **2014**, 13, 249 | 42.1 | 33 |
| 184 | Single-cell Hi-C bridges microscopy and genome-wide sequencing approaches to study 3D chromatin organization. *BioEssays*, **2017**, 39, 1700104 | 4.1 | 32 |
| 183 | Chromosomal DNA loops may constitute basic units of the eukaryotic genome organization and evolution. *Critical Reviews in Eukaryotic Gene Expression*, **1999**, 9, 279-83 | 1.3 | 31 |
| 182 | Perinucleolar relocalization and nucleolin as crucial events in the transcriptional activation of key genes in mantle cell lymphoma. *Blood*, **2014**, 123, 2044-53 | 2.2 | 30 |
| 181 | DNA loop anchorage region colocalizes with the replication origin located downstream to the human gene encoding lamin B2. *Journal of Cellular Biochemistry*, **1998**, 69, 13-8 | 4.7 | 30 |
| 180 | A CTCF-dependent silencer located in the differentially methylated area may regulate expression of a housekeeping gene overlapping a tissue-specific gene domain. *Molecular and Cellular Biology*, **2006**, 26, 1589-97 | 4.8 | 30 |
| 179 | The 33 kb transcript of the chicken alpha-globin gene domain is part of the nuclear matrix. *Journal of Cellular Biochemistry*, **2004**, 92, 445-57 | 4.7 | 30 |
| 178 | Weak interactions in higher-order chromatin organization. *Nucleic Acids Research*, **2020**, 48, 4614-4626 | 20.1 | 30 |
| 177 | Non-clonability correlates with genomic instability: a case study of a unique DNA region. *Journal of Molecular Biology*, **2001**, 307, 481-6 | 6.5 | 29 |
| 176 | Induction of transcription within chromosomal DNA loops flanked by MAR elements causes an association of loop DNA with the nuclear matrix. *Nucleic Acids Research*, **2005**, 33, 4157-63 | 20.1 | 28 |
| 175 | Functional architecture of chromosomal DNA domains. *Critical Reviews in Eukaryotic Gene Expression*, **1996**, 6, 247-69 | 1.3 | 28 |
| 174 | Identification of c-Myb Target Genes in K562 Cells Reveals a Role for c-Myb as a Master Regulator. *Genes and Cancer*, **2011**, 2, 805-17 | 2.9 | 27 |
| 173 | Breakpoint cluster regions of the AML-1 and ETO genes contain MAR elements and are preferentially associated with the nuclear matrix in proliferating HEL cells. *Journal of Cell Science*, **2004**, 117, 4583-90 | 5.3 | 27 |
| 172 | Transcriptional regulation and spatial organisation of the human AML1/RUNX1 gene. *Journal of Cellular Biochemistry*, **2011**, 112, 1997-2005 | 4.7 | 25 |
| 171 | Nuclear compartments, genome folding, and enhancer-promoter communication. *International Review of Cell and Molecular Biology*, **2015**, 315, 183-244 | 6 | 24 |
| 170 | The anti-cancer drugs curaxins target spatial genome organization. *Nature Communications*, **2019**, 10, 1441 | 17.4 | 22 |
| 169 | Mapping of replication origins and termination sites in the Duchenne muscular dystrophy gene. *Genomics*, **1997**, 45, 24-30 | 4.3 | 22 |
| 168 | Mapping long-range chromatin organization within the chicken alpha-globin gene domain using oligonucleotide DNA arrays. *Genomics*, **2005**, 85, 143-51 | 4.3 | 22 |

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| 167 | The distribution of tightly bound proteins along the DNA chain reflects the type of cell differentiation. *Nucleic Acids Research*, **1988**, 16, 3617-33 | 20.1 | 22 |
| 166 | The Role of Liquid-Liquid Phase Separation in the Compartmentalization of Cell Nucleus and Spatial Genome Organization. *Biochemistry (Moscow)*, **2020**, 85, 643-650 | 2.9 | 21 |
| 165 | Quantitative analysis of genomic element interactions by molecular colony technique. *Nucleic Acids Research*, **2014**, 42, e36 | 20.1 | 21 |
| 164 | Breakpoint Clusters: Reason or Consequence?. *Critical Reviews in Eukaryotic Gene Expression*, **2004**, 14, 65-78 | 1.3 | 21 |
| 163 | Human cytomegalovirus proteins PP65 and IEP72 are targeted to distinct compartments in nuclei and nuclear matrices of infected human embryo fibroblasts. *Journal of Cellular Biochemistry*, **2003**, 90, 1056-67 | 4.7 | 20 |
| 162 | Extensive methylation of a part of the CpG island located 3.0-4.5 kbp upstream to the chicken alpha-globin gene cluster may contribute to silencing the globin genes in non-erythroid cells. *Journal of Molecular Biology*, **2000**, 299, 845-52 | 6.5 | 19 |
| 161 | Actual ligation frequencies in the chromosome conformation capture procedure. *PLoS ONE*, **2013**, 8, e60403 | 3.7 | 18 |
| 160 | Chromatin loops, illegitimate recombination, and genome evolution. *BioEssays*, **2009**, 31, 278-86 | 4.1 | 18 |
| 159 | Assembly of nuclear matrix-bound protein complexes involved in non-homologous end joining is induced by inhibition of DNA topoisomerase II. *Journal of Cellular Physiology*, **2006**, 207, 660-7 | 7 | 18 |
| 158 | The specificity of human lymphocyte nucleolar DNA long-range fragmentation by endogenous topoisomerase II and exogenous Bal 31 nuclease depends on cell proliferation status. *Biochemistry*, **1995**, 34, 4133-8 | 3.2 | 18 |
| 157 | Interaction in vivo between the two matrix attachment regions flanking a single chromatin loop. *Journal of Molecular Biology*, **2009**, 386, 929-37 | 6.5 | 17 |
| 156 | In the nucleus and cytoplasm of chicken erythroleukemic cells, prosomes containing the p23K subunit are found in centers of globin (pre-)mRNA processing and accumulation. *Experimental Cell Research*, **1999**, 250, 569-75 | 4.2 | 17 |
| 155 | The 3D Genome as a Target for Anticancer Therapy. *Trends in Molecular Medicine*, **2020**, 26, 141-149 | 11.5 | 17 |
| 154 | Early S-phase cell hypersensitivity to heat stress. *Cell Cycle*, **2016**, 15, 337-44 | 4.7 | 16 |
| 153 | Cell-cycle-dependent localization of human cytomegalovirus UL83 phosphoprotein in the nucleolus and modulation of viral gene expression in human embryo fibroblasts in vitro. *Journal of Cellular Biochemistry*, **2011**, 112, 307-17 | 4.7 | 16 |
| 152 | Chemotherapy-related secondary leukemias: A role for DNA repair by error-prone non-homologous end joining in topoisomerase II - Induced chromosomal rearrangements. *Gene*, **2007**, 391, 76-9 | 3.8 | 16 |
| 151 | Functional analysis of DNA sequences located within a cluster of DNase U hypersensitive sites colocalizing with a MAR element at the upstream border of the chicken ?-globin gene domain. *Journal of Cellular Biochemistry*, **1999**, 74, 38-49 | 4.7 | 16 |
| 150 | Domain organization of eukaryotic genome. *Cell Biology International Reports*, **1992**, 16, 697-708 | | 16 |

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| 149 | Initiated complexes of RNA polymerase II are concentrated in the nuclear skeleton associated DNA. *Experimental Cell Research*, **1985**, 158, 273-5 | 4.2 | 16 |
| 148 | Hypoosmotic stress induces R loop formation in nucleoli and ATR/ATM-dependent silencing of nucleolar transcription. *Nucleic Acids Research*, **2019**, 47, 6811-6825 | 20.1 | 15 |
| 147 | 5-hydroxymethylcytosine in DNA repair: A new player or a red herring?. *Cell Cycle*, **2017**, 16, 1499-1501 | 4.7 | 15 |
| 146 | Distinct distribution of ectopically expressed histone variants H2A.Bbd and MacroH2A in open and closed chromatin domains. *PLoS ONE*, **2012**, 7, e47157 | 3.7 | 15 |
| 145 | Mapping of the nuclear matrix-bound chromatin hubs by a new M3C experimental procedure. *Nucleic Acids Research*, **2010**, 38, 8051-60 | 20.1 | 15 |
| 144 | In chicken leukemia cells globin genes are fully transcribed but their rnas are retained in the perinucleolar area. *Experimental Cell Research*, **2001**, 270, 159-65 | 4.2 | 15 |
| 143 | Analysis of the replication direction through the domain of alpha-globin-encoding chicken genes. *Gene*, **1995**, 166, 255-9 | 3.8 | 15 |
| 142 | Organization of specific DNA sequence elements in the region of the replication origin and matrix attachment site in the chicken alpha-globin gene domain. *Molecular Genetics and Genomics*, **1992**, 235, 381-8 | | 15 |
| 141 | 3D genomics imposes evolution of the domain model of eukaryotic genome organization. *Chromosoma*, **2017**, 126, 59-69 | 2.8 | 14 |
| 140 | Studying RNA-DNA interactome by Red-C identifies noncoding RNAs associated with various chromatin types and reveals transcription dynamics. *Nucleic Acids Research*, **2020**, 48, 6699-6714 | 20.1 | 14 |
| 139 | Gene functioning and storage within a folded genome. *Cellular and Molecular Biology Letters*, **2017**, 22, 18 | 8.1 | 14 |
| 138 | TMEM8 - a non-globin gene entrapped in the globin web. *Nucleic Acids Research*, **2009**, 37, 7394-406 | 20.1 | 14 |
| 137 | Repositioning of ETO gene in cells treated with VP-16, an inhibitor of DNA-topoisomerase II. *Journal of Cellular Biochemistry*, **2008**, 104, 692-9 | 4.7 | 14 |
| 136 | RNA-dependent nuclear matrix contains a 33 kb globin full domain transcript as well as prosomes but no 26S proteasomes. *Journal of Cellular Biochemistry*, **2005**, 94, 529-39 | 4.7 | 14 |
| 135 | Specific radial positions of centromeres of human chromosomes X, 1, and 19 remain unchanged in chromatin-depleted nuclei of primary human fibroblasts: evidence for the organizing role of the nuclear matrix. *Journal of Cellular Biochemistry*, **2005**, 96, 850-7 | 4.7 | 14 |
| 134 | Rearrangement of chromatin domains in cancer and development. *Journal of Cellular Biochemistry*, **2000**, Suppl 35, 54-60 | 4.7 | 14 |
| 133 | The sequence-specific nuclear matrix binding factor F6 is a chicken GATA-like protein. *Molecular Genetics and Genomics*, **1993**, 238, 309-14 | | 14 |
| 132 | Transcriptional enhancer in the vicinity of a replication origin within the 5' region of the chicken alpha-globin gene domain. *Journal of Molecular Biology*, **1991**, 217, 595-8 | 6.5 | 14 |

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| 131 | Suppression of liquid-liquid phase separation by 1,6-hexanediol partially compromises the 3D genome organization in living cells. *Nucleic Acids Research*, **2021**, 49, 10524-10541 | 20.1 | 14 |
| 130 | The clustering of CpG islands may constitute an important determinant of the 3D organization of interphase chromosomes. *Epigenetics*, **2014**, 9, 951-63 | 5.7 | 13 |
| 129 | Spatial organization of the chicken beta-globin gene domain in erythroid cells of embryonic and adult lineages. *Epigenetics and Chromatin*, **2012**, 5, 16 | 5.8 | 13 |
| 128 | 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 alpha-globin gene domain. *Biochemical and Biophysical Research Communications*, **1991**, 179, 512-9 | 3.4 | 13 |
| 127 | Chromatin Domains and Territories: Flexibly Rigid. *Critical Reviews in Eukaryotic Gene Expression*, **2004**, 14, 79-88 | 1.3 | 13 |
| 126 | Acetylation of core histones causes the unfolding of 30 nm chromatin fiber: analysis by agarose gel electrophoresis. *Biochemical and Biophysical Research Communications*, **1993**, 196, 455-60 | 3.4 | 12 |
| 125 | Characterization of the chromatin structure in the upstream region of the chicken alpha-globin gene domain. *Molecular Genetics and Genomics*, **1994**, 242, 649-52 | | 12 |
| 124 | Quantitative differences in TAD border strength underly the TAD hierarchy in Drosophila chromosomes. *Journal of Cellular Biochemistry*, **2019**, 120, 4494-4503 | 4.7 | 12 |
| 123 | Activation of the alpha-globin gene expression correlates with dramatic upregulation of nearby non-globin genes and changes in local and large-scale chromatin spatial structure. *Epigenetics and Chromatin*, **2017**, 10, 35 | 5.8 | 11 |
| 122 | Unraveling the mechanisms of chromatin fibril packaging. *Nucleus*, **2016**, 7, 319-24 | 3.9 | 11 |
| 121 | Compartmentalization of the cell nucleus and spatial organization of the genome. *Molecular Biology*, **2015**, 49, 21-39 | 1.2 | 11 |
| 120 | Association of the mammalian transcriptional regulator kaiso with centrosomes and the midbody. *Cell Cycle*, **2009**, 8, 2303-4 | 4.7 | 11 |
| 119 | Early replication timing of the chicken alpha-globin gene domain correlates with its open chromatin state in cells of different lineages. *Genomics*, **2009**, 93, 481-6 | 4.3 | 11 |
| 118 | DNA-protein interactions and spatial organization of DNA. *Molecular Biology Reports*, **1993**, 18, 167-75 | 2.8 | 11 |
| 117 | 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, 1263-8 | 3.4 | 11 |
| 116 | Divide and Rule: Phase Separation in Eukaryotic Genome Functioning. *Cells*, **2020**, 9, | 7.9 | 11 |
| 115 | Topologically-associating domains: gene warehouses adapted to serve transcriptional regulation. *Transcription*, **2016**, 7, 84-90 | 4.8 | 11 |
| 114 | Order and stochasticity in the folding of individual Drosophila genomes. *Nature Communications*, **2021**, 12, 41 | 17.4 | 11 |

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|---|---|---|---|
| 113 | Domains of □and □globin genes in the context of the structural-functional organization of the eukaryotic genome. *Biochemistry (Moscow)*, **2012**, 77, 1409-23 | 2.9 | 10 |
| 112 | In embryonic chicken erythrocytes actively transcribed alpha globin genes are not associated with the nuclear matrix. *Journal of Cellular Biochemistry*, **2009**, 106, 170-8 | 4.7 | 10 |
| 111 | Sensitivity of human embryonic and induced pluripotent stem cells to a topoisomerase II poison etoposide. *Cell Cycle*, **2011**, 10, 2035-7 | 4.7 | 10 |
| 110 | Distribution of topoisomerase II-mediated cleavage sites and relation to structural and functional landmarks in 830 kb of Drosophila DNA. *Nucleic Acids Research*, **1997**, 25, 2041-6 | 20.1 | 10 |
| 109 | Changes in chromosome positioning may contribute to the development of diseases related to X-chromosome aneuploidy. *Journal of Cellular Physiology*, **2007**, 213, 278-83 | 7 | 10 |
| 108 | Breakpoint clusters: reason or consequence?. *Critical Reviews in Eukaryotic Gene Expression*, **2004**, 14, 65-77 | 1.3 | 9 |
| 107 | Distinct Patterns of Colocalization of the CCND1 and CMYC Genes With Their Potential Translocation Partner IGH at Successive Stages of B-Cell Differentiation. *Journal of Cellular Biochemistry*, **2016**, 117, 1506-10 | 4.7 | 8 |
| 106 | Nuclear matrix and structural and functional compartmentalization of the eucaryotic cell nucleus. *Biochemistry (Moscow)*, **2014**, 79, 608-18 | 2.9 | 8 |
| 105 | Modulatory effect of rRNA synthesis and ppUL83 nucleolar compartmentalization on human cytomegalovirus gene expression in vitro. *Journal of Cellular Biochemistry*, **2009**, 108, 415-23 | 4.7 | 8 |
| 104 | Joint cultivation of human erythroblastoid cells and mouse fibroblasts triggers release of a wide spectrum of cytotoxic factors. *Biochemical and Biophysical Research Communications*, **1997**, 234, 655-9 | 3.4 | 8 |
| 103 | Mechanisms controlling activation of the alpha-globin gene domain in chicken erythroid cells. *Biochemistry (Moscow)*, **2007**, 72, 467-70 | 2.9 | 8 |
| 102 | Transgenic Goats in the World Pharmaceutical Industry of the 21st Century. *Russian Journal of Genetics*, **2002**, 38, 1-14 | 0.6 | 8 |
| 101 | TGF-□ is the factor secreted by proliferative chondrocytes to inhibit neo-angiogenesis. *Journal of Cellular Biochemistry*, **2001**, 81, 79-88 | 4.7 | 8 |
| 100 | Excision of chromosomal DNA loops by treatment of permeabilised cells with Bal 31 nuclease. *Molecular Genetics and Genomics*, **1995**, 249, 253-6 | | 8 |
| 99 | A simple and reproducible method for analysis of chromatin condensation. *Biochemical and Biophysical Research Communications*, **1993**, 193, 113-8 | 3.4 | 8 |
| 98 | Inducing cellular senescence by using genetically encoded photosensitizers. *Aging*, **2016**, 8, 2449-2462 | 5.6 | 8 |
| 97 | C-TALE, a new cost-effective method for targeted enrichment of Hi-C/3C-seq libraries. *Methods*, **2020**, 170, 48-60 | 4.6 | 8 |
| 96 | Evolution of the Genome 3D Organization: Comparison of Fused and Segregated Globin Gene Clusters. *Molecular Biology and Evolution*, **2017**, 34, 1492-1504 | 8.3 | 7 |

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|---|---|---|---|
| 95 | Heat Stress-Induced Transcriptional Repression. *Biochemistry (Moscow)*, **2015**, 80, 990-3 | 2.9 | 7 |
| 94 | The Role of Crowding Forces in Juxtaposing β-Globin Gene Domain Remote Regulatory Elements in Mouse Erythroid Cells. *PLoS ONE*, **2015**, 10, e0139855 | 3.7 | 7 |
| 93 | Formaldehyde fixation of cells does not greatly reduce the ability to amplify cellular DNA. *Analytical Biochemistry*, **2009**, 390, 94-6 | 3.1 | 7 |
| 92 | Unusual compartmentalization of CTCF and other transcription factors in the course of terminal erythroid differentiation. *Biochimica Et Biophysica Acta - Molecular Cell Research*, **2007**, 1773, 924-33 | 4.9 | 7 |
| 91 | 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 | 4.7 | 7 |
| 90 | DNA-protein complexes of the nuclear matrix: visualization and partial characterization of the protein component. *Biochemical and Biophysical Research Communications*, **1989**, 162, 175-83 | 3.4 | 7 |
| 89 | Correlations of repetitive and AT-rich DNA segments within the chicken globin gene domains. *Molecular Biology Reports*, **1986**, 11, 177-87 | 2.8 | 7 |
| 88 | Structural-Functional Domains of the Eukaryotic Genome. *Biochemistry (Moscow)*, **2018**, 83, 302-312 | 2.9 | 6 |
| 87 | The broken MLL gene is frequently located outside the inherent chromosome territory in human lymphoid cells treated with DNA topoisomerase II poison etoposide. *PLoS ONE*, **2013**, 8, e75871 | 3.7 | 6 |
| 86 | The inactivation of the β-gene in chicken erythroblasts of adult lineage is not mediated by packaging of the embryonic part of the β-globin gene domain into a repressive heterochromatin-like structure. *Epigenetics*, **2011**, 6, 1481-8 | 5.7 | 6 |
| 85 | The upstream area of the chicken alpha-globin gene domain is transcribed in both directions in the same cells. *FEBS Letters*, **2005**, 579, 4746-50 | 3.8 | 6 |
| 84 | Spatial organization of the eukaryotic genome and the action of epigenetic mechanisms. *Russian Journal of Genetics*, **2006**, 42, 1353-1361 | 0.6 | 6 |
| 83 | Specific cleavage of chicken alpha A-globin and human c-Ha-ras genes by two molecular forms of calf thymus topoisomerase I. *Molecular and Cellular Biochemistry*, **1991**, 101, 115-24 | 4.2 | 6 |
| 82 | Studies on structure and function of chromatin. *Molecular and Cellular Biochemistry*, **1981**, 40, 29-48 | 4.2 | 6 |
| 81 | The IGH locus relocalizes to a "recombination compartment" in the perinucleolar region of differentiating B-lymphocytes. *Oncotarget*, **2017**, 8, 40079-40089 | 3.3 | 6 |
| 80 | Heat stress induces formation of cytoplasmic granules containing HSC70 protein. *Doklady Biochemistry and Biophysics*, **2015**, 463, 213-5 | 0.8 | 5 |
| 79 | Mammalian Diaphanous-related formin-1 restricts early phases of influenza A/NWS/33 virus (H1N1) infection in LLC-MK2 cells by affecting cytoskeleton dynamics. *Molecular and Cellular Biochemistry*, **2018**, 437, 185-201 | 4.2 | 5 |
| 78 | Fragment of intron 5.2 of the human RUNX1 gene important for transcription activation is neither enhancer nor MAR-element. *Doklady Biochemistry and Biophysics*, **2012**, 442, 26-9 | 0.8 | 5 |

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|---|---|---|---|
| 77 | Transcription factories and spatial organization of eukaryotic genomes. *Biochemistry (Moscow)*, **2010**, 75, 1307-15 | 2.9 | 5 |
| 76 | Expression of full-length human pro-urokinase in mammary glands of transgenic mice. *Transgenic Research*, **2009**, 18, 747-56 | 3.3 | 5 |
| 75 | Study of spatial organization of chicken alpha-globin gene domain by 3C technique. *Biochemistry (Moscow)*, **2008**, 73, 1192-9 | 2.9 | 5 |
| 74 | Inhibition of DNA topoisomerase II may trigger illegitimate recombination in living cells: experiments with a model system. *Journal of Cellular Biochemistry*, **2006**, 99, 598-608 | 4.7 | 5 |
| 73 | Histone Chaperone FACT and Curaxins: Effects on Genome Structure and Function. *Journal of Cancer Metastasis and Treatment*, **2019**, 5, | 3.8 | 5 |
| 72 | Suppression of liquid-liquid phase separation by 1,6-hexanediol partially compromises the 3D genome organization in living cells | | 5 |
| 71 | Non-coding RNAs in chromatin folding and nuclear organization. *Cellular and Molecular Life Sciences* , **2021**, 78, 5489-5504 | 10.3 | 5 |
| 70 | Chromatin domains and territories: flexibly rigid. *Critical Reviews in Eukaryotic Gene Expression*, **2004**, 14, 79-88 | 1.3 | 5 |
| 69 | Transcription-controlling regulatory elements of the eukaryotic genome. *Molecular Biology*, **2015**, 49, 185-194 | 1.2 | 4 |
| 68 | Single-cell genome-wide studies give new insight into nongenetic cell-to-cell variability in animals. *Histochemistry and Cell Biology*, **2016**, 146, 239-54 | 2.4 | 4 |
| 67 | Heat shock-induced dissociation of TRF2 from telomeres does not initiate a telomere-dependent DNA damage response. *Cell Biology International*, **2014**, 38, 675-81 | 4.5 | 4 |
| 66 | Transcription factor RUNX1. *Molecular Biology*, **2012**, 46, 755-767 | 1.2 | 4 |
| 65 | HP1β is not necessary for the structural maintenance of centromeric heterochromatin. *Epigenetics*, **2011**, 6, 380-7 | 5.7 | 4 |
| 64 | Chromatin and transcription regulation. *Molecular Biology*, **2007**, 41, 343-348 | 1.2 | 4 |
| 63 | An unusual extended DNA loop attachment region is located in the human dystrophin gene. *Journal of Cellular Physiology*, **2006**, 209, 515-21 | 7 | 4 |
| 62 | A modified protocol of Capture-C allows affordable and flexible high-resolution promoter interactome analysis. *Scientific Reports*, **2020**, 10, 15491 | 4.9 | 4 |
| 61 | Folded genome as a platform for the functional compartmentalization of the eukaryotic cell nucleus. *Biopolymers and Cell*, **2014**, 30, 83-89 | 0.3 | 3 |
| 60 | The two waves in single-cell 3D genomics. *Seminars in Cell and Developmental Biology*, **2021**, 121, 143-143 | 7.5 | 3 |

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| 59 | Treacle and TOPBP1 control replication stress response in the nucleolus. *Journal of Cell Biology*, **2021**, 220, | 7.3 | 3 |
| 58 | Manipulation of Cellular Processes via Nucleolus Hijaking in the Course of Viral Infection in Mammals. *Cells*, **2021**, 10, | 7.9 | 3 |
| 57 | 3D Genomics. *Molecular Biology*, **2019**, 53, 802-812 | 1.2 | 3 |
| 56 | RedChIP identifies noncoding RNAs associated with genomic sites occupied by Polycomb and CTCF proteins.. *Proceedings of the National Academy of Sciences of the United States of America*, **2022**, 119, | 11.5 | 3 |
| 55 | Benchmark of software tools for prokaryotic chromosomal interaction domain identification. *Bioinformatics*, **2020**, 36, 4560-4567 | 7.2 | 2 |
| 54 | Chromatin Trapping of Factors Involved in DNA Replication and Repair Underlies Heat-Induced Radio- and Chemosensitization. *Cells*, **2020**, 9, | 7.9 | 2 |
| 53 | Structural-Functional Organization of the Eukaryotic Cell Nucleus and Transcription Regulation: Introduction to This Special Issue of Biochemistry (Moscow). *Biochemistry (Moscow)*, **2018**, 83, 299-301 | 2.9 | 2 |
| 52 | Proteasomes raise the microtubule dynamics in influenza A (H1N1) virus-infected LLC-MK2 cells. *Cellular and Molecular Biology Letters*, **2015**, 20, 840-66 | 8.1 | 2 |
| 51 | Spatial organization of housekeeping genes in interphase nuclei. *Molecular Biology*, **2014**, 48, 886-895 | 1.2 | 2 |
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