

Olga Fedorova

List of Publications by Year in descending order

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158
papers

2,660
citations

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times ranked

1433
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#	ARTICLE	IF	CITATIONS
1	The Role of Natural Polymorphic Variants of DNA Polymerase $\hat{\text{I}}^2$ in DNA Repair. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2390.	1.8	8
2	Comparative Analysis of Exo- and Endonuclease Activities of APE1-like Enzymes. <i>International Journal of Molecular Sciences</i> , 2022, 23, 2869.	1.8	3
3	Insights into Mechanisms of Damage Recognition and Catalysis by APE1-like Enzymes. <i>International Journal of Molecular Sciences</i> , 2022, 23, 4361.	1.8	5
4	Structural and Molecular Kinetic Features of Activities of DNA Polymerases. <i>International Journal of Molecular Sciences</i> , 2022, 23, 6373.	1.8	13
5	Pre-steady-state kinetic and mutational insights into mechanisms of endo- and exonuclease DNA processing by mutant forms of human AP endonuclease. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2022, 1866, 130198.	1.1	1
6	The Enigma of Substrate Recognition and Catalytic Efficiency of APE1-Like Enzymes. <i>Frontiers in Cell and Developmental Biology</i> , 2021, 9, 617161.	1.8	6
7	Mutational and Kinetic Analysis of APE1 Endoribonuclease Activity. <i>Molecular Biology</i> , 2021, 55, 211-224.	0.4	7
8	Comparative Analysis of the Activity of the Polymorphic Variants of Human Uracil-DNA-Glycosylases SMUG1 and MBD4. <i>Molecular Biology</i> , 2021, 55, 241-251.	0.4	1
9	Initial stages of DNA Base Excision Repair in Nucleosomes. <i>Molecular Biology</i> , 2021, 55, 167-181.	0.4	2
10	Common Kinetic Mechanism of Abasic Site Recognition by Structurally Different Apurinic/Apyrimidinic Endonucleases. <i>International Journal of Molecular Sciences</i> , 2021, 22, 8874.	1.8	6
11	Kinetic Analysis of the Interaction of Nicking Endonuclease BspD6I with DNA. <i>Biomolecules</i> , 2021, 11, 1420.	1.8	0
12	DNA Demethylation in the Processes of Repair and Epigenetic Regulation Performed by 2-Ketoglutarate-Dependent DNA Dioxygenases. <i>International Journal of Molecular Sciences</i> , 2021, 22, 10540.	1.8	6
13	Pre-Steady-State Kinetics of the SARS-CoV-2 Main Protease as a Powerful Tool for Antiviral Drug Discovery. <i>Frontiers in Pharmacology</i> , 2021, 12, 773198.	1.6	5
14	Efficiency of RNA Hydrolysis by Binase from <i>Bacillus pumilus</i> : The Impact of Substrate Structure, Metal Ions, and Low Molecular Weight Nucleotide Compounds. <i>Molecular Biology</i> , 2020, 54, 769-776.	0.4	2
15	Activity of Human Apurinic/Apyrimidinic Endonuclease APE1 Toward Damaged DNA and Native RNA With Non-canonical Structures. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 590848.	1.8	18
16	The role of active-site amino acid residues in the cleavage of DNA and RNA substrates by human apurinic/aprimidinic endonuclease APE1. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2020, 1864, 129718.	1.1	14
17	The Role of Active-Site Plasticity in Damaged-Nucleotide Recognition by Human Apurinic/Apyrimidinic Endonuclease APE1. <i>Molecules</i> , 2020, 25, 3940.	1.7	11
18	Lesion Recognition and Cleavage of Damage-Containing Quadruplexes and Bulged Structures by DNA Glycosylases. <i>Frontiers in Cell and Developmental Biology</i> , 2020, 8, 595687.	1.8	12

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19	An Assay for the Activity of Base Excision Repair Enzymes in Cellular Extracts Using Fluorescent DNA Probes. <i>Biochemistry (Moscow)</i> , 2020, 85, 480-489.	0.7	4
20	Kinetic Milestones of Damage Recognition by DNA Glycosylases of the Helix-Hairpin-Helix Structural Superfamily. <i>Advances in Experimental Medicine and Biology</i> , 2020, 1241, 1-18.	0.8	11
21	Effect of the Substrate Structure and Metal Ions on the Hydrolysis of Undamaged RNA by Human AP Endonuclease APE1. <i>Acta Naturae</i> , 2020, 12, 74-85.	1.7	10
22	Role of Arg243 and His239 Residues in the Recognition of Damaged Nucleotides by Human Uracil-DNA Glycosylase SMUG1. <i>Biochemistry (Moscow)</i> , 2020, 85, 594-603.	0.7	1
23	Modulation of the Apurinic/Apyrimidinic Endonuclease Activity of Human APE1 and of Its Natural Polymorphic Variants by Base Excision Repair Proteins. <i>International Journal of Molecular Sciences</i> , 2020, 21, 7147.	1.8	12
24	Role of Ionizing Amino Acid Residues in the Process of DNA Binding by Human AP Endonuclease 1 and in Its Catalysis. <i>Journal of Physical Chemistry B</i> , 2019, 123, 9546-9556.	1.2	16
25	The Role of Active-Site Residues Phe98, His239, and Arg243 in DNA Binding and in the Catalysis of Human Uracil-DNA Glycosylase SMUG1. <i>Molecules</i> , 2019, 24, 3133.	1.7	2
26	Conformational Dynamics of Damage Processing by Human DNA Glycosylase NEIL1. <i>Journal of Molecular Biology</i> , 2019, 431, 1098-1112.	2.0	17
27	Roles of Active-Site Amino Acid Residues in Specific Recognition of DNA Lesions by Human 8-Oxoguanine-DNA Glycosylase (OGG1). <i>Journal of Physical Chemistry B</i> , 2019, 123, 4878-4887.	1.2	8
28	The impact of single-nucleotide polymorphisms of human apurinic/aprimidinic endonuclease 1 on specific DNA binding and catalysis. <i>Biochimie</i> , 2019, 163, 73-83.	1.3	15
29	New Fluorescent Analogs of Nucleotides Based on 3-Hydroxychromone for Recording Conformational Changes of DNA. <i>Russian Journal of Bioorganic Chemistry</i> , 2019, 45, 599-607.	0.3	4
30	Comparative Analysis of Nucleotide Fluorescent Analogs for Registration of DNA Conformational Changes Induced by Interaction with Formamidopyrimidine-DNA Glycosylase Fpg. <i>Russian Journal of Bioorganic Chemistry</i> , 2019, 45, 591-598.	0.3	6
31	Conformational Dynamics of Dioxygenase AlkB and DNA in the Course of Catalytically Active Enzyme-Substrate Complex Formation. <i>Russian Journal of Bioorganic Chemistry</i> , 2019, 45, 630-640.	0.3	4
32	A Single-Turnover Kinetic Study of DNA Demethylation Catalyzed by Fe(II)/ α -Ketoglutarate-Dependent Dioxygenase AlkB. <i>Molecules</i> , 2019, 24, 4576.	1.7	7
33	Thermodynamics of the DNA Repair Process by Endonuclease VIII. <i>Acta Naturae</i> , 2019, 11, 29-37.	1.7	10
34	Thermodynamics of the DNA Repair Process by Endonuclease VIII. <i>Acta Naturae</i> , 2019, 11, 29-37.	1.7	3
35	The role of the N-terminal domain of human apurinic/aprimidinic endonuclease 1, APE1, in DNA glycosylase stimulation. <i>DNA Repair</i> , 2018, 64, 10-25.	1.3	30
36	Substrate specificity of human apurinic/aprimidinic endonuclease APE1 in the nucleotide incision repair pathway. <i>Nucleic Acids Research</i> , 2018, 46, 11454-11465.	6.5	38

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37	Data on PAGE analysis and MD simulation for the interaction of endonuclease Apn1 from <i>Saccharomyces cerevisiae</i> with DNA substrates containing 5,6-dihydrouracyl and 2-aminopurine. <i>Data in Brief</i> , 2018, 20, 1515-1524.	0.5	0
38	Apurinic/aprimidinic endonuclease Apn1 from <i>Saccharomyces cerevisiae</i> is recruited to the nucleotide incision repair pathway: Kinetic and structural features. <i>Biochimie</i> , 2018, 152, 53-62.	1.3	7
39	Kinetics and Thermodynamics of DNA Processing by Wild Type DNA-Glycosylase Endo III and Its Catalytically Inactive Mutant Forms. <i>Genes</i> , 2018, 9, 190.	1.0	19
40	Kinetic Basis of the Bifunctionality of SsoII DNA Methyltransferase. <i>Molecules</i> , 2018, 23, 1192.	1.7	0
41	Kinetic Features of 3'→5' Exonuclease Activity of Human AP-Endonuclease APE1. <i>Molecules</i> , 2018, 23, 2101.	1.7	29
42	The formation of catalytically competent enzyme-substrate complex is not a bottleneck in lesion excision by human alkyladenine DNA glycosylase. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017, 35, 950-967.	2.0	13
43	Interaction features of adenine DNA glycosylase MutY from <i>E. coli</i> with DNA substrates. <i>Russian Journal of Bioorganic Chemistry</i> , 2017, 43, 13-22.	0.3	4
44	The kinetic analysis of recognition of the damaged nucleotides by mutant forms of the 8-oxoguanine DNA glycosylase hOGG1. <i>Russian Journal of Bioorganic Chemistry</i> , 2017, 43, 1-12.	0.3	8
45	Pre-steady-state kinetic analysis of damage recognition by human single-strand selective monofunctional uracil-DNA glycosylase SMUG1. <i>Molecular BioSystems</i> , 2017, 13, 2638-2649.	2.9	26
46	Evolution of inhibitor-resistant natural mutant forms of HIV-1 protease probed by pre-steady state kinetic analysis. <i>Biochimie</i> , 2017, 142, 125-134.	1.3	2
47	Global DNA dynamics of 8-oxoguanine repair by human OGG1 revealed by stopped-flow kinetics and molecular dynamics simulation. <i>Molecular BioSystems</i> , 2017, 13, 1954-1966.	2.9	12
48	A real-time study of the interaction of TBP with a TATA box-containing duplex identical to an ancestral or minor allele of human gene LEP or TPI. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017, 35, 3070-3081.	2.0	16
49	Pre-steady state kinetics of DNA binding and abasic site hydrolysis by tyrosyl-DNA phosphodiesterase 1. <i>Journal of Biomolecular Structure and Dynamics</i> , 2017, 35, 2314-2327.	2.0	6
50	Mutational and Kinetic Analysis of Lesion Recognition by <i>Escherichia coli</i> Endonuclease VIII. <i>Genes</i> , 2017, 8, 140.	1.0	21
51	Search for Modified DNA Sites with the Human Methyl-CpG-Binding Enzyme MBD4. <i>Acta Naturae</i> , 2017, 9, 88-98.	1.7	25
52	Search for Modified DNA Sites with the Human Methyl-CpG-Binding Enzyme MBD4. <i>Acta Naturae</i> , 2017, 9, 88-98.	1.7	7
53	Effects of mono- and divalent metal ions on DNA binding and catalysis of human apurinic/aprimidinic endonuclease 1. <i>Molecular BioSystems</i> , 2016, 12, 1527-1539.	2.9	54
54	A kinetic mechanism of repair of DNA containing β -anomeric deoxyadenosine by human apurinic/aprimidinic endonuclease 1. <i>Molecular BioSystems</i> , 2016, 12, 3435-3446.	2.9	8

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55	Thermodynamic analysis of fast stages of specific lesion recognition by DNA repair enzymes. <i>Biochemistry (Moscow)</i> , 2016, 81, 1136-1152.	0.7	8
56	New oligonucleotide derivatives as unreactive substrate analogues and potential inhibitors of human apurinic/apyrimidinic endonuclease APE1. <i>Molecular BioSystems</i> , 2016, 12, 67-75.	2.9	9
57	Thermodynamics of Damaged DNA Binding and Catalysis by Human AP Endonuclease 1. <i>Acta Naturae</i> , 2016, 8, 103-110.	1.7	28
58	Thermodynamics of Damaged DNA Binding and Catalysis by Human AP Endonuclease 1. <i>Acta Naturae</i> , 2016, 8, 103-10.	1.7	14
59	Pre-steady-state Kinetic and Structural Analysis of Interaction of Methionine S-Methyltransferase from <i>Citrobacter freundii</i> with Inhibitors. <i>Journal of Biological Chemistry</i> , 2015, 290, 671-681.	1.6	19
60	Conformational Dynamics of DNA Repair by <i>Escherichia coli</i> Endonuclease III. <i>Journal of Biological Chemistry</i> , 2015, 290, 14338-14349.	1.6	42
61	The role of His-83 of yeast apurinic/apyrimidinic endonuclease Ape1 in catalytic incision of abasic sites in DNA. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2015, 1850, 1297-1309.	1.1	3
62	Active destabilization of base pairs by a DNA glycosylase wedge initiates damage recognition. <i>Nucleic Acids Research</i> , 2015, 43, 272-281.	6.5	49
63	Thermodynamics of the DNA Damage Repair Steps of Human 8-Oxoguanine DNA Glycosylase. <i>PLoS ONE</i> , 2014, 9, e98495.	1.1	36
64	The role of Asn-212 in the catalytic mechanism of human endonuclease APE1: Stopped-flow kinetic study of incision activity on a natural AP site and a tetrahydrofuran analogue. <i>DNA Repair</i> , 2014, 21, 43-54.	1.3	14
65	Role of the light-chain constant-domain switch in the structure and functionality of A17 reactibody. <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2014, 70, 708-719.	2.5	22
66	Effect of Some Substituents Increasing the Solubility of Zn(II) and Al(III) Phthalocyanines on Their Photophysical Properties. <i>Bioinorganic Chemistry and Applications</i> , 2014, 2014, 1-7.	1.8	8
67	Step-by-step mechanism of DNA damage recognition by human 8-oxoguanine DNA glycosylase. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2014, 1840, 387-395.	1.1	43
68	Pre-steady-state fluorescence analysis of damaged DNA transfer from human DNA glycosylases to AP endonuclease APE1. <i>Biochimica Et Biophysica Acta - General Subjects</i> , 2014, 1840, 3042-3051.	1.1	30
69	New Environment-Sensitive Multichannel DNA Fluorescent Label for Investigation of the Protein-DNA Interactions. <i>PLoS ONE</i> , 2014, 9, e100007.	1.1	44
70	Real-Time Interaction between TVR and the TATA Box of the Human Triosephosphate Isomerase Gene Promoter in the Norm and Pathology. <i>Acta Naturae</i> , 2014, 6, 36-40.	1.7	17
71	Structural Features of the Interaction between Human 8-Oxoguanine DNA Glycosylase hOGG1 and DNA. <i>Acta Naturae</i> , 2014, 6, 52-65.	1.7	9
72	Real-Time Interaction between TBP and the TATA Box of the Human Triosephosphate Isomerase Gene Promoter in the Norm and Pathology. <i>Acta Naturae</i> , 2014, 6, 36-40.	1.7	16

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73	Structural Features of the Interaction between Human 8-Oxoguanine DNA Glycosylase hOGG1 and DNA. <i>Acta Naturae</i> , 2014, 6, 52-65.	1.7	3
74	Reaction of 4-hydroxycoumarin with 2-acetyloxiranes. <i>Russian Journal of Organic Chemistry</i> , 2013, 49, 1497-1501.	0.3	2
75	DNA Damage Processing by Human 8-Oxoguanine-DNA Glycosylase Mutants with the Occluded Active Site. <i>Journal of Biological Chemistry</i> , 2013, 288, 28936-28947.	1.6	25
76	Effect of Complexation with Arabinogalactan on Pharmacokinetics of Guest Drugs in Rats: For Example, Warfarin. <i>BioMed Research International</i> , 2013, 2013, 1-4.	0.9	6
77	Direct DNA Lesion Reversal and Excision Repair in <i>Escherichia coli</i> . <i>EcoSal Plus</i> , 2013, 5, .	2.1	6
78	Pulsed Electron Double Resonance in Structural Studies of Spin-Labeled Nucleic Acids. <i>Acta Naturae</i> , 2013, 5, 9-32.	1.7	6
79	Pulsed electron double resonance in structural studies of spin-labeled nucleic acids. <i>Acta Naturae</i> , 2013, 5, 9-32.	1.7	2
80	Thermodynamics of the multi-stage DNA lesion recognition and repair by formamidopyrimidine-DNA glycosylase using pyrrolocytosine fluorescence stopped-flow pre-steady-state kinetics. <i>Nucleic Acids Research</i> , 2012, 40, 7384-7392.	6.5	57
81	Conformational dynamics of the interaction of <i>Escherichia coli</i> endonuclease VIII with DNA substrates. <i>DNA Repair</i> , 2012, 11, 884-891.	1.3	29
82	Kinetic mechanism of the interaction of <i>Saccharomyces cerevisiae</i> AP-endonuclease 1 with DNA substrates. <i>Biochemistry (Moscow)</i> , 2012, 77, 1162-1171.	0.7	4
83	Conformational Dynamics of Abasic DNA upon Interactions with AP Endonuclease 1 Revealed by Stopped-Flow Fluorescence Analysis. <i>Biochemistry</i> , 2012, 51, 1306-1321.	1.2	29
84	Highly Mutagenic Exocyclic DNA Adducts Are Substrates for the Human Nucleotide Incision Repair Pathway. <i>PLoS ONE</i> , 2012, 7, e51776.	1.1	29
85	PELDOR analysis of enzyme-induced structural changes in damaged DNA duplexes. <i>Molecular BioSystems</i> , 2011, 7, 2670.	2.9	21
86	¹ H CIDNP study of the kinetics and mechanism of the reversible photoinduced oxidation of tryptophyl-tryptophan dipeptide in aqueous solutions. <i>Russian Chemical Bulletin</i> , 2011, 60, 2579-2587.	0.4	8
87	Lys98 Substitution in Human AP Endonuclease 1 Affects the Kinetic Mechanism of Enzyme Action in Base Excision and Nucleotide Incision Repair Pathways. <i>PLoS ONE</i> , 2011, 6, e24063.	1.1	16
88	Mechanism of recognition and repair of damaged DNA by human 8-oxoguanine DNA glycosylase hOGG1. <i>Biochemistry (Moscow)</i> , 2011, 76, 118-130.	0.7	20
89	Kinetic mechanism of human apurinic/apyrimidinic endonuclease action in nucleotide incision repair. <i>Biochemistry (Moscow)</i> , 2011, 76, 273-281.	0.7	8
90	Pre-steady-state kinetics of interaction of wild-type and multiple drug-resistant HIV protease with first and second generation inhibitory drugs. <i>Doklady Biochemistry and Biophysics</i> , 2011, 440, 239-243.	0.3	1

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91	Deprotonation of Transient Guanosyl Cation Radical Catalyzed by Buffer in Aqueous Solution: TR-CIDNP Study. <i>Applied Magnetic Resonance</i> , 2011, 41, 239-250.	0.6	9
92	Mechanism of Antisense Oligonucleotide Interaction with Natural RNAs. <i>Journal of Biomolecular Structure and Dynamics</i> , 2011, 29, 27-50.	2.0	9
93	Dimeric Fe-Co Phthalocyanine Complex as a Reagent for the Selective Damage of Nucleic Acids. <i>Macrocyclic Chemistry</i> , 2011, 4, 135-137.	0.9	2
94	Real-time studies of conformational dynamics of the repair enzyme E. coli formamidopyrimidine-DNA glycosylase and its DNA complexes during catalytic cycle. <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2010, 685, 3-10.	0.4	39
95	Conformational dynamics and pre-steady-state kinetics of DNA glycosylases. <i>Biochemistry (Moscow)</i> , 2010, 75, 1225-1239.	0.7	7
96	Conformational Transitions in Human AP Endonuclease 1 and Its Active Site Mutant during Abasic Site Repair. <i>Biochemistry</i> , 2010, 49, 6451-6461.	1.2	42
97	Biophysical and X-ray Crystallographic Analysis of Mps1 Kinase Inhibitor Complexes. <i>Biochemistry</i> , 2010, 49, 1689-1701.	1.2	35
98	Genetic and Biochemical Characterization of Human AP Endonuclease 1 Mutants Deficient in Nucleotide Incision Repair Activity. <i>PLoS ONE</i> , 2010, 5, e12241.	1.1	37
99	Fe(II) phthalocyanine catalyzed oxidation of dGMP by molecular oxygen. <i>Bioorganic and Medicinal Chemistry Letters</i> , 2009, 19, 4335-4338.	1.0	6
100	Oxidation of DNA and its components with reactive oxygen species. <i>Russian Chemical Reviews</i> , 2009, 78, 659-678.	2.5	26
101	Reversible Chemical Step and Rate-Limiting Enzyme Regeneration in the Reaction Catalyzed by Formamidopyrimidine-DNA Glycosylase. <i>Biochemistry</i> , 2009, 48, 11335-11343.	1.2	46
102	Conformational Dynamics of Human AP Endonuclease in Base Excision and Nucleotide Incision Repair Pathways. <i>Journal of Biomolecular Structure and Dynamics</i> , 2009, 26, 637-652.	2.0	47
103	PELDOR study of conformations of double-spin-labeled single- and double-stranded DNA with non-nucleotide inserts. <i>Physical Chemistry Chemical Physics</i> , 2009, 11, 6826.	1.3	46
104	Substrate Recognition of Anthrax Lethal Factor Examined by Combinatorial and Pre-steady-state Kinetic Approaches. <i>Journal of Biological Chemistry</i> , 2009, 284, 17902-17913.	1.6	18
105	DNA-binding and Oxidative Properties of Cationic Phthalocyanines and Their Dimeric Complexes with Anionic Phthalocyanines Covalently Linked to Oligonucleotides. <i>Journal of Biomolecular Structure and Dynamics</i> , 2008, 26, 307-319.	2.0	16
106	Kinetic Conformational Analysis of Human 8-Oxoguanine-DNA Glycosylase. <i>Journal of Biological Chemistry</i> , 2007, 282, 1029-1038.	1.6	69
107	Quantitative surface-enhanced resonance Raman scattering of phthalocyanine-labelled oligonucleotides. <i>Nucleic Acids Research</i> , 2007, 35, e42-e42.	6.5	19
108	Fluorescence Spectroscopic and ¹⁹ F NMR Studies of Human Thymidylate Synthase with its Cognate RNA. <i>Journal of Biomolecular Structure and Dynamics</i> , 2007, 25, 253-269.	2.0	6

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109	Pre-Steady-State Kinetic Study of Substrate Specificity of Escherichia coli Formamidopyrimidine-DNA Glycosylase. <i>Biochemistry</i> , 2007, 46, 424-435.	1.2	68
110	Conjugates of Phthalocyanines with Oligonucleotides as Reagents for Sensitized or Catalytic DNA Modification. <i>Bioinorganic Chemistry and Applications</i> , 2006, 2006, 1-8.	1.8	8
111	Kinetic Study of DNA Modification by Phthalocyanine Derivative of the Oligonucleotide. <i>Bioinorganic Chemistry and Applications</i> , 2006, 2006, 1-10.	1.8	5
112	Effect of DNA and bivalent metal ions on the interaction of thermostable DNA polymerase Tte with dNTPs. <i>Russian Chemical Bulletin</i> , 2005, 54, 1306-1310.	0.4	0
113	Kinetics of substrate recognition and cleavage by human 8-oxoguanine-DNA glycosylase. <i>Nucleic Acids Research</i> , 2005, 33, 3919-3931.	6.5	108
114	Pre-steady-state kinetics shows differences in processing of various DNA lesions by Escherichia coli formamidopyrimidine-DNA glycosylase. <i>Nucleic Acids Research</i> , 2004, 32, 926-935.	6.5	57
115	Thermodynamics of Interaction of Phthalocyanine-Oligonucleotide Conjugates with Single- and Double-Stranded DNA. <i>Nucleosides, Nucleotides and Nucleic Acids</i> , 2004, 23, 983-987.	0.4	5
116	Quantitative parameters of cooperative interactions of oligonucleotides within tandem complexes. <i>Russian Chemical Bulletin</i> , 2003, 52, 2507-2516.	0.4	1
117	Thermodynamic, Kinetic, and Structural Basis for Recognition and Repair of 8-Oxoguanine in DNA by Fpg Protein from Escherichia coli. <i>Biochemistry</i> , 2002, 41, 7540-7548.	1.2	46
118	Stopped-Flow Kinetic Studies of the Interaction between Escherichia coli Fpg Protein and DNA Substrates. <i>Biochemistry</i> , 2002, 41, 1520-1528.	1.2	58
119	BINDING OF A PORPHYRIN CONJUGATE OF HOECHST 33258 TO DNA. I. UV-VISIBLE AND MELTING STUDIES DETECT MULTIPLE BINDING MODES TO A 12-MER NONSELF-COMPLEMENTARY DUPLEX. <i>Nucleosides, Nucleotides and Nucleic Acids</i> , 2001, 20, 131-143.	0.4	8
120	BINDING OF A DESMETALLO-PORPHYRIN CONJUGATE OF HOECHST 33258 TO DNA. III. STRONG BINDING TO SINGLE-STRAND OLIGONUCLEOTIDES. <i>Nucleosides, Nucleotides and Nucleic Acids</i> , 2001, 20, 157-168.	0.4	5
121	PHOTOSENSITIZED AND CATALYTIC OXIDATION OF DNA BY METALLOPHthalocyanine-OLIGONUCLEOTIDE CONJUGATES. <i>Nucleosides, Nucleotides and Nucleic Acids</i> , 2001, 20, 1259-1262.	0.4	15
122	CD and Melting Curves Structural Studies of the Tandem DNA Complex Formed with Oligonucleotides Carrying Photoactive and Sensitizing Groups in the Nick Region. <i>Journal of Biomolecular Structure and Dynamics</i> , 2001, 19, 515-526.	2.0	2
123	Title is missing!. <i>Molecular Biology</i> , 2000, 34, 814-822.	0.4	3
124	The synthesis of a cobalt(II) tetracarboxyphthalocyanine-deoxyribooligonucleotide conjugate as a reagent for the directed DNA modification. <i>Russian Journal of Bioorganic Chemistry</i> , 2000, 26, 104-110.	0.3	5
125	Real-Time Oligonucleotide Hybridization Kinetics Monitored by Resonant Mirror Technique. <i>IUBMB Life</i> , 1999, 48, 317-320.	1.5	15
126	Synthesis of New Oligonucleotide Derivatives with Porphyrins and Phthalocyanins. <i>Nucleosides & Nucleotides</i> , 1999, 18, 1515-1516.	0.5	4

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127	Structural Requirements of Double and Single Stranded DNA Substrates and Inhibitors, Including a Photoaffinity Label, of Fpg Protein From Escherichia Coli. <i>Journal of Biomolecular Structure and Dynamics</i> , 1999, 17, 301-310.	2.0	15
128	Real-time Oligonucleotide Hybridization Kinetics Monitored by Resonant Mirror Technique. <i>IUBMB Life</i> , 1999, 48, 317-320.	1.5	19
129	Cooperative Binding of Oligonucleotides to Adjacent Sites of Single-Stranded DNA: Sequence Composition Dependence at the Junction. <i>Journal of Biomolecular Structure and Dynamics</i> , 1999, 17, 259-265.	2.0	9
130	Quantitative Parameters of Cooperative Interactions of the Oligodeoxyribonucleotides on the Complementary Template. <i>Nucleosides & Nucleotides</i> , 1998, 17, 1705-1708.	0.5	0
131	Cooperative Interactions of the Oligodeoxyribonucleotides on the Complementary Template. The Influence of Chemical Groups and Mismatched Nucleotides at the 5'- and 3'-ends of Oligonucleotides on the Parameters of Cooperativity. <i>Journal of Biomolecular Structure and Dynamics</i> , 1997, 15, 369-380.	2.0	6
132	Kinetic Approach for Determination of Affinity Properties of Reactive Oligonucleotide Derivatives to Complementary Regions in Nucleic Acids. <i>Nucleosides & Nucleotides</i> , 1997, 16, 1807-1808.	0.5	0
133	Complexation of Photochromic Crown Ether Styryl Dyes with Mg ²⁺ As Probed by Surface-Enhanced Raman Scattering Spectroscopy. <i>Journal of Physical Chemistry B</i> , 1997, 101, 4077-4084.	1.2	13
134	Site-Specific Photomodification of DNA by Porphyrin-Oligonucleotide Conjugates Synthesized via a Solid Phase H-Phosphonate Approach. <i>Bioconjugate Chemistry</i> , 1997, 8, 49-56.	1.8	34
135	A series of meso-tris(N-methyl-pyridiniumyl)-(4-alkylamidophenyl) porphyrins: Synthesis, interaction with DNA and antibacterial activity. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1997, 1354, 252-260.	2.4	50
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