Petr PeÄinka

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7789030/publications.pdf

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32	1,239	15	32
papers	citations	h-index	g-index
36	36	36	1261 citing authors
all docs	docs citations	times ranked	

#	Article	IF	CITATIONS
1	Searching for New Z-DNA/Z-RNA Binding Proteins Based on Structural Similarity to Experimentally Validated Zα Domain. International Journal of Molecular Sciences, 2022, 23, 768.	4.1	11
2	Are There Hidden Genes in DNA/RNA Vaccines?. Frontiers in Immunology, 2022, 13, 801915.	4.8	9
3	Unheeded SARS-CoV-2 proteins? A deep look into negative-sense RNA. Briefings in Bioinformatics, 2022, 23, .	6.5	15
4	Regulation of Phenolic Compound Production by Light Varying in Spectral Quality and Total Irradiance. International Journal of Molecular Sciences, 2022, 23, 6533.	4.1	13
5	Amino Acid Composition in Various Types of Nucleic Acid-Binding Proteins. International Journal of Molecular Sciences, 2021, 22, 922.	4.1	14
6	Tracing dsDNA Virus–Host Coevolution through Correlation of Their G-Quadruplex-Forming Sequences. International Journal of Molecular Sciences, 2021, 22, 3433.	4.1	11
7	Letter to the Editor: Significant mutation enrichment in inverted repeat sites of new SARS-CoV-2 strains. Briefings in Bioinformatics, 2021, 22, .	6.5	2
8	G-Quadruplex in Gene Encoding Large Subunit of Plant RNA Polymerase II: A Billion-Year-Old Story. International Journal of Molecular Sciences, 2021, 22, 7381.	4.1	13
9	Analyses of viral genomes for G-quadruplex forming sequences reveal their correlation with the type of infection. Biochimie, 2021, 186, 13-27.	2.6	33
10	The Changes in the p53 Protein across the Animal Kingdom Point to Its Involvement in Longevity. International Journal of Molecular Sciences, 2021, 22, 8512.	4.1	9
11	Searching for G-Quadruplex-Binding Proteins in Plants: New Insight into Possible G-Quadruplex Regulation. BioTech, 2021, 10, 20.	2.6	7
12	Transcriptomic and Proteomic Analysis of Drought Stress Response in Opium Poppy Plants during the First Week of Germination. Plants, 2021, 10, 1878.	3.5	9
13	SARS-CoV-2 hot-spot mutations are significantly enriched within inverted repeats and CpG island loci. Briefings in Bioinformatics, 2021, 22, 1338-1345.	6.5	20
14	Characterization of p53 Family Homologs in Evolutionary Remote Branches of Holozoa. International Journal of Molecular Sciences, 2020, 21, 6.	4.1	40
15	G-Quadruplexes in the Archaea Domain. Biomolecules, 2020, 10, 1349.	4.0	31
16	In-Depth Bioinformatic Analyses of Nidovirales Including Human SARS-CoV-2, SARS-CoV, MERS-CoV Viruses Suggest Important Roles of Non-canonical Nucleic Acid Structures in Their Lifecycles. Frontiers in Microbiology, 2020, 11, 1583.	3.5	57
17	p53 Binds Preferentially to Non-B DNA Structures Formed by the Pyrimidine-Rich Strands of GAA·TTC Trinucleotide Repeats Associated with Friedreich's Ataxia. Molecules, 2019, 24, 2078.	3.8	6
18	The Presence and Localization of G-Quadruplex Forming Sequences in the Domain of Bacteria. Molecules, 2019, 24, 1711.	3.8	75

#	Article	IF	CITATIONS
19	Liver regeneration during the associating liver partition and portal vein ligation for staged hepatectomy procedure in Susà ¿½ scrofa is positively modulated by stem cells. Oncology Letters, 2018, 15, 6309-6321.	1.8	2
20	The Amino Acid Composition of Quadruplex Binding Proteins Reveals a Shared Motif and Predicts New Potential Quadruplex Interactors. Molecules, 2018, 23, 2341.	3.8	51
21	Bioinformatics analyses and inÂvitro evidence for five and six stacked G-quadruplex forming sequences. Biochimie, 2018, 150, 70-75.	2.6	17
22	Electrochemical Activity of Wedelolactone and Probing its Interaction with DNA Using Voltammetry at a Carbon Electrode. Electroanalysis, 2015, 27, 2268-2271.	2.9	5
23	Impact of cadmium, cobalt and nickel on sequence-specific DNA binding of p63 and p73 in vitro and in cells. Biochemical and Biophysical Research Communications, 2015, 456, 29-34.	2.1	7
24	Preferential Binding of Hot Spot Mutant p53 Proteins to Supercoiled DNA In Vitro and in Cells. PLoS ONE, 2013, 8, e59567.	2.5	34
25	Selective binding of tumor suppressor p53 protein to topologically constrained DNA: Modulation by intercalative drugs. Biochemical and Biophysical Research Communications, 2010, 393, 894-899.	2.1	22
26	DNA topology influences p53 sequence-specific DNA binding through structural transitions within the target sites. Biochemical Journal, 2008, 412, 57-63.	3.7	33
27	DNA modification with cisplatin affects sequence-specific DNA binding of p53 and p73 proteins in a target site-dependent manner. FEBS Journal, 2006, 273, 4693-4706.	4.7	9
28	A Single-Surface Electrochemical Biosensor for the Detection of DNA Triplet Repeat Expansion. Electroanalysis, 2006, 18, 141-151.	2.9	33
29	Enhancement of p53 sequence-specific binding by DNA supercoiling. Oncogene, 2004, 23, 2119-2127.	5.9	37
30	Microanalysis of DNA by stripping transfer voltammetry. Bioelectrochemistry, 2004, 63, 249-252.	4.6	26
31	Role of tumor suppressor p53 domains in selective binding to supercoiled DNA. Nucleic Acids Research, 2002, 30, 4966-4974.	14.5	57
32	DNA tetraplex formation in the control region of c-myc. Nucleic Acids Research, 1998, 26, 1167-1172.	14.5	525