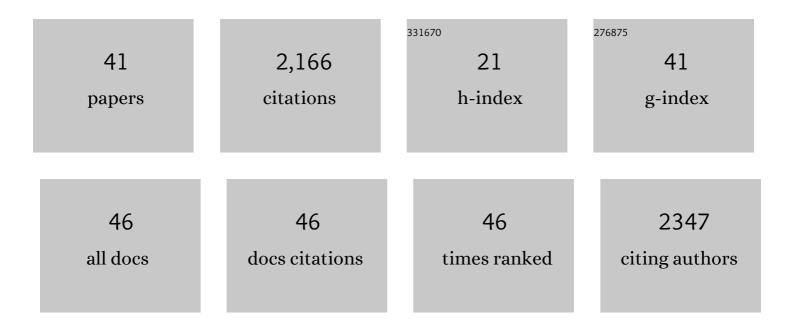
Tahir H Tahirov

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/2987825/publications.pdf Version: 2024-02-01



Τλμίο Η Τλμιρον

#	Article	IF	CITATIONS
1	Structural basis for transcription elongation by bacterial RNA polymerase. Nature, 2007, 448, 157-162.	27.8	380
2	Crystal structure of HIV-1 Tat complexed with human P-TEFb. Nature, 2010, 465, 747-751.	27.8	255
3	Structure of a T7 RNA polymerase elongation complex at 2.9 à resolution. Nature, 2002, 420, 43-50.	27.8	218
4	DNA Polymerase δ and ζ Switch by Sharing Accessory Subunits of DNA Polymerase δ. Journal of Biological Chemistry, 2012, 287, 17281-17287.	3.4	144
5	Structural basis for inhibition of DNA replication by aphidicolin. Nucleic Acids Research, 2014, 42, 14013-14021.	14.5	104
6	Evolution of DNA polymerases: an inactivated polymerase-exonuclease module in Pol Îμ and a chimeric origin of eukaryotic polymerases from two classes of archaeal ancestors. Biology Direct, 2009, 4, 11.	4.6	102
7	Mechanism of Concerted RNA-DNA Primer Synthesis by the Human Primosome. Journal of Biological Chemistry, 2016, 291, 10006-10020.	3.4	100
8	X-ray structure of the complex of regulatory subunits of human DNA polymerase delta. Cell Cycle, 2008, 7, 3026-3036.	2.6	81
9	Crystal Structure of the Human Primase. Journal of Biological Chemistry, 2015, 290, 5635-5646.	3.4	65
10	Insight into the Human DNA Primase Interaction with Template-Primer. Journal of Biological Chemistry, 2016, 291, 4793-4802.	3.4	60
11	Crystal structure of the C-terminal domain of human DNA primase large subunit. Cell Cycle, 2011, 10, 926-931.	2.6	55
12	Crystal Structure of the Human Pol α B Subunit in Complex with the C-terminal Domain of the Catalytic Subunit. Journal of Biological Chemistry, 2015, 290, 14328-14337.	3.4	53
13	Crystal structure of HIV-1 Tat complexed with human P-TEFb and AFF4. Cell Cycle, 2014, 13, 1788-1797.	2.6	51
14	Elaborated Action of the Human Primosome. Genes, 2017, 8, 62.	2.4	41
15	The C-terminal Domain of the DNA Polymerase Catalytic Subunit Regulates the Primase and Polymerase Activities of the Human DNA Polymerase α-Primase Complex. Journal of Biological Chemistry, 2014, 289, 22021-22034.	3.4	32
16	Iron–Sulfur Clusters in DNA Polymerases and Primases of Eukaryotes. Methods in Enzymology, 2018, 599, 1-20.	1.0	32
17	Crystal structure of the human Políµ B-subunit in complex with the C-terminal domain of the catalytic subunit. Journal of Biological Chemistry, 2017, 292, 15717-15730.	3.4	30
18	Modulation of mutagenesis in eukaryotes by DNA replication fork dynamics and quality of nucleotide pools. Environmental and Molecular Mutagenesis, 2012, 53, 699-724.	2.2	28

Τα<mark></mark>ΗΙΓΑΗΙΓΟν

#	Article	IF	CITATIONS
19	Activity and fidelity of human DNA polymerase α depend on primer structure. Journal of Biological Chemistry, 2018, 293, 6824-6843.	3.4	28
20	Structural basis of Ets1 cooperative binding to palindromic sequences on stromelysin-1 promoter DNA. Cell Cycle, 2010, 9, 3126-3134.	2.6	27
21	Replication protein A binds RNA and promotes R-loop formation. Journal of Biological Chemistry, 2020, 295, 14203-14213.	3.4	26
22	Inaccurate DNA Synthesis in Cell Extracts of Yeast Producing Active Human DNA Polymerase Iota. PLoS ONE, 2011, 6, e16612.	2.5	25
23	Crystal Structure of Mouse Elf3 C-terminal DNA-binding Domain in Complex with Type II TGF-Î ² Receptor Promoter DNA. Journal of Molecular Biology, 2010, 397, 278-289.	4.2	24
24	A novel variant of DNA polymerase ζ, Rev3ΔC, highlights differential regulation of Pol32 as a subunit of polymerase δversus ζ in Saccharomyces cerevisiae. DNA Repair, 2014, 24, 138-149.	2.8	22
25	High-Resolution Crystals of Methionine Aminopeptidase fromPyrococcus furiosusObtained by Water-Mediated Transformation. Journal of Structural Biology, 1998, 121, 68-72.	2.8	20
26	Structure and Function of Eukaryotic DNA Polymerase δ. Sub-Cellular Biochemistry, 2012, 62, 217-236.	2.4	19
27	Structural Basis of Ets1 Cooperative Binding to Widely Separated Sites on Promoter DNA. PLoS ONE, 2012, 7, e33698.	2.5	18
28	Divalent ions attenuate DNA synthesis by human DNA polymerase \hat{I}_{\pm} by changing the structure of the template/primer or by perturbing the polymerase reaction. DNA Repair, 2016, 43, 24-33.	2.8	16
29	Insight into RNA–DNA primer length counting by human primosome. Nucleic Acids Research, 2022, 50, 6264-6270.	14.5	15
30	Translesion activity of PrimPol on DNA with cisplatin and DNA–protein cross-links. Scientific Reports, 2021, 11, 17588.	3.3	14
31	Comment on "The [4Fe4S] cluster of human DNA primase functions as a redox switch using DNA charge transport― Science, 2017, 357, .	12.6	12
32	Functional mapping of the fission yeast DNA polymerase δB-subunit Cdc1 by site-directed and random pentapeptide insertion mutagenesis. BMC Molecular Biology, 2009, 10, 82.	3.0	10
33	Structure and Biophysics of CBFβ/RUNX and Its Translocation Products. Advances in Experimental Medicine and Biology, 2017, 962, 21-31.	1.6	10
34	Compact reduced thioredoxin structure from the thermophilic bacteria Thermus thermophilus. Proteins: Structure, Function and Bioinformatics, 2005, 61, 1032-1037.	2.6	9
35	Comparison of the kinetic parameters of the truncated catalytic subunit and holoenzyme of human DNA polymerase É>. DNA Repair, 2015, 29, 16-22.	2.8	9
36	Structural and functional insight into mismatch extension by human DNA polymerase α. Proceedings of the United States of America, 2022, 119, e2111744119.	7.1	8

TAHIR H TAHIROV

#	Article	IF	CITATIONS
37	Crystallization and preliminary X-ray diffraction analysis of human DNA primase. Acta Crystallographica Section F, Structural Biology Communications, 2014, 70, 206-210.	0.8	7
38	Crystal structure of a purine/pyrimidine phosphoribosyltransferase-related protein from Thermus thermophilus HB8. Proteins: Structure, Function and Bioinformatics, 2005, 61, 658-665.	2.6	4
39	Preliminary crystallographic analysis of mouse Elf3 C-terminal DNA-binding domain in complex with type II TGF-β receptor promoter DNA. Acta Crystallographica Section F: Structural Biology Communications, 2009, 65, 1261-1263.	0.7	4
40	Efficient discrimination against RNA-containing primers by human DNA polymerase ε. Scientific Reports, 2022, 12, .	3.3	4
41	Crystallization and preliminary crystallographic analysis of the complex of the second and third regulatory subunits of human Pol δ. Acta Crystallographica Section F: Structural Biology Communications, 2008, 64, 822-824.	0.7	3