

Takashi Kubota

List of Publications by Year in descending order

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687363

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1206
citing authors

#	ARTICLE	IF	CITATIONS
1	SWI/SNF and the histone chaperone Rtt106 drive expression of the Pleiotropic Drug Resistance network genes. <i>Nature Communications</i> , 2022, 13, 1968.	12.8	3
2	Ligation of newly replicated DNA controls the timing of DNA mismatch repair. <i>Current Biology</i> , 2021, 31, 1268-1276.e6.	3.9	19
3	Effective mismatch repair depends on timely control of PCNA retention on DNA by the Elg1 complex. <i>Nucleic Acids Research</i> , 2019, 47, 6826-6841.	14.5	20
4	Identification of Elg1 interaction partners and effects on post-replication chromatin re-formation. <i>PLoS Genetics</i> , 2018, 14, e1007783.	3.5	15
5	PCNA Retention on DNA into G2/M Phase Causes Genome Instability in Cells Lacking Elg1. <i>Cell Reports</i> , 2016, 16, 684-695.	6.4	65
6	Definition of the transcription factor TdIF1 consensus binding sequence through genome-wide mapping of its binding sites. <i>Genes To Cells</i> , 2015, 20, 242-254.	1.2	7
7	Replication-Coupled PCNA Unloading by the Elg1 Complex Occurs Genome-wide and Requires Okazaki Fragment Ligation. <i>Cell Reports</i> , 2015, 12, 774-787.	6.4	100
8	Rif1 controls DNA replication by directing Protein Phosphatase 1 to reverse Cdc7-mediated phosphorylation of the MCM complex. <i>Genes and Development</i> , 2014, 28, 372-383.	5.9	217
9	The Elg1 Replication Factor C-like Complex Functions in PCNA Unloading during DNA Replication. <i>Molecular Cell</i> , 2013, 50, 273-280.	9.7	230
10	Is PCNA unloading the central function of the Elg1/ATAD5 replication factor C-like complex?. <i>Cell Cycle</i> , 2013, 12, 2570-2579.	2.6	37
11	TdIF1 Recognizes a Specific DNA Sequence through Its Helix-Turn-Helix and AT-Hook Motifs to Regulate Gene Transcription. <i>PLoS ONE</i> , 2013, 8, e66710.	2.5	6
12	Quantitative proteomic analysis of yeast DNA replication proteins. <i>Methods</i> , 2012, 57, 196-202.	3.8	20
13	Quantitative Proteomic Analysis of Chromatin Reveals that Ctf18 Acts in the DNA Replication Checkpoint. <i>Molecular and Cellular Proteomics</i> , 2011, 10, M110.005561.	3.8	60
14	TdT interacting factor 1 enhances TdT ubiquitylation through recruitment of BPOZ2 into nucleus from cytoplasm. <i>Genes To Cells</i> , 2009, 14, 1415-1427.	1.2	5
15	Bood POZ containing gene type 2 is a human counterpart of yeast Btb3p and promotes the degradation of terminal deoxynucleotidyltransferase. <i>Genes To Cells</i> , 2008, 13, 439-457.	1.2	10
16	Identification of functional domains in TdIF1 and its inhibitory mechanism for TdT activity. <i>Genes To Cells</i> , 2007, 12, 941-959.	1.2	15
17	UDP-glucuronosyltransferase1A1 directly binds to albumin. <i>Hepatology Research</i> , 2005, 31, 241-245.	3.4	6
18	DNA polymerase lambda directly binds to proliferating cell nuclear antigen through its confined C-terminal region. <i>Genes To Cells</i> , 2005, 10, 705-715.	1.2	26

#	ARTICLE	IF	CITATIONS
19	Direct binding of TRpP-132 with TdT results in reduction of TdT activity. <i>Genes To Cells</i> , 2005, 11, 47-57.	1.2	12
20	Terminal deoxynucleotidyltransferase forms a ternary complex with a novel chromatin remodeling protein with 82 kDa and core histone. <i>Genes To Cells</i> , 2003, 8, 559-571.	1.2	15
21	Structures of Ganoderic Acid A and B, Two New Lanostane Type Bitter Triterpenes from <i>Ganoderma lucidum</i> (FR.) KARST.. <i>Helvetica Chimica Acta</i> , 1982, 65, 611-619.	1.6	159
22	Fatty Acid Composition of Mesocarp Oils of <i>Lindera strychnifolia</i> (Sieb. et Zucc.) F. Vill, <i>Neolitsea aciculata</i> (Blume) Koidz., and <i>Neolitsea sericea</i> (Blume) Koidz.. <i>Journal of Japan Oil Chemists Society</i> , 1980, 29, 426-427.	0.1	0
23	The Constituents of the Essential Oil from <i>Litsea japonica</i> (Thunb.) Juss., FRUIT. <i>Agricultural and Biological Chemistry</i> , 1978, 42, 1601-1603.	0.3	0