## Xu-Dong Zhu

## List of Publications by Citations

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27 1,617 11.3 4.24 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
24	Cell-cycle-regulated association of RAD50/MRE11/NBS1 with TRF2 and human telomeres. <i>Nature Genetics</i> , <b>2000</b> , 25, 347-52	36.3	509
23	ERCC1/XPF removes the 3Woverhang from uncapped telomeres and represses formation of telomeric DNA-containing double minute chromosomes. <i>Molecular Cell</i> , <b>2003</b> , 12, 1489-98	17.6	319
22	MRE11-RAD50-NBS1 and ATM function as co-mediators of TRF1 in telomere length control. <i>Nature Structural and Molecular Biology</i> , <b>2007</b> , 14, 832-40	17.6	92
21	ATM and ATR Signaling Regulate the Recruitment of Human Telomerase to Telomeres. <i>Cell Reports</i> , <b>2015</b> , 13, 1633-46	10.6	84
20	Arginine methylation regulates telomere length and stability. <i>Molecular and Cellular Biology</i> , <b>2009</b> , 29, 4918-34	4.8	51
19	Human XPF controls TRF2 and telomere length maintenance through distinctive mechanisms. <i>Mechanisms of Ageing and Development</i> , <b>2008</b> , 129, 602-10	5.6	44
18	Cockayne syndrome group B protein regulates DNA double-strand break repair and checkpoint activation. <i>EMBO Journal</i> , <b>2015</b> , 34, 1399-416	13	41
17	Post-translational modifications of TRF1 and TRF2 and their roles in telomere maintenance. <i>Mechanisms of Ageing and Development</i> , <b>2012</b> , 133, 421-34	5.6	41
16	Cockayne Syndrome group B protein interacts with TRF2 and regulates telomere length and stability. <i>Nucleic Acids Research</i> , <b>2012</b> , 40, 9661-74	20.1	35
15	XPF with mutations in its conserved nuclease domain is defective in DNA repair but functions in TRF2-mediated telomere shortening. <i>DNA Repair</i> , <b>2007</b> , 6, 157-66	4.3	34
14	ATM and CDK2 control chromatin remodeler CSB to inhibit RIF1 in DSB repair pathway choice. <i>Nature Communications</i> , <b>2017</b> , 8, 1921	17.4	33
13	ATM regulates proteasome-dependent subnuclear localization of TRF1, which is important for telomere maintenance. <i>Nucleic Acids Research</i> , <b>2012</b> , 40, 3975-89	20.1	30
12	Cyclin B-dependent kinase 1 regulates human TRF1 to modulate the resolution of sister telomeres. <i>Nature Communications</i> , <b>2011</b> , 2, 371	17.4	29
11	Cleavage-dependent ligation by the FLP recombinase. Characterization of a mutant FLP protein with an alteration in a catalytic amino acid. <i>Journal of Biological Chemistry</i> , <b>1995</b> , 270, 23044-54	5.4	26
10	Phosphorylated (pT371)TRF1 is recruited to sites of DNA damage to facilitate homologous recombination and checkpoint activation. <i>Nucleic Acids Research</i> , <b>2013</b> , 41, 10268-82	20.1	14
9	Methylated TRF2 associates with the nuclear matrix and serves as a potential biomarker for cellular senescence. <i>Aging</i> , <b>2014</b> , 6, 248-63	5.6	11
8	CSB interacts with BRCA1 in late S/G2 to promote MRN- and CtIP-mediated DNA end resection. <i>Nucleic Acids Research</i> , <b>2019</b> , 47, 10678-10692	20.1	10

## LIST OF PUBLICATIONS

7	Cdk-dependent phosphorylation regulates TRF1 recruitment to PML bodies and promotes C-circle production in ALT cells. <i>Journal of Cell Science</i> , <b>2016</b> , 129, 2559-72	5.3	10	
6	Biphasic recruitment of TRF2 to DNA damage sites promotes non-sister chromatid homologous recombination repair. <i>Journal of Cell Science</i> , <b>2018</b> , 131,	5.3	10	
5	CSB cooperates with SMARCAL1 to maintain telomere stability in ALT cells. <i>Journal of Cell Science</i> , <b>2020</b> , 133,	5.3	9	
4	TRF1 phosphorylation on T271 modulates telomerase-dependent telomere length maintenance as well as the formation of ALT-associated PML bodies. <i>Scientific Reports</i> , <b>2016</b> , 6, 36913	4.9	8	
3	Efficient UV repair requires disengagement of the CSB winged helix domain from the CSB ATPase domain. <i>DNA Repair</i> , <b>2018</b> , 68, 58-67	4.3	7	
2	The role of single-stranded DNA in Flp-mediated strand exchange. <i>Journal of Biological Chemistry</i> , <b>1998</b> , 273, 4921-7	5.4	1	
1	The Winged Helix Domain of CSB Regulates RNAPII Occupancy at Promoter Proximal Pause Sites. <i>International Journal of Molecular Sciences</i> , <b>2021</b> , 22,	6.3	1	