Hans W Hombauer

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Ligation of newly replicated DNA controls the timing of DNA mismatch repair. Current Biology, 2021, 31, 1268-1276.e6.	1.8	19
2	Inactivation of folylpolyglutamate synthetase Met7 results in genome instability driven by an increased dUTP/dTTP ratio. Nucleic Acids Research, 2020, 48, 264-277.	6.5	7
3	Extensive 5′-surveillance guards against non-canonical NAD-caps of nuclear mRNAs in yeast. Nature Communications, 2020, 11, 5508.	5.8	28
4	Identification of MLH2/hPMS1 dominant mutations that prevent DNA mismatch repair function. Communications Biology, 2020, 3, 751.	2.0	5
5	A genetic screen pinpoints ribonucleotide reductase residues that sustain dNTP homeostasis and specifies a highly mutagenic type of dNTP imbalance. Nucleic Acids Research, 2019, 47, 237-252.	6.5	16
6	Alterations in cellular metabolism triggered by <i>URA7</i> or <i>GLN3</i> inactivation cause imbalanced dNTP pools and increased mutagenesis. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E4442-E4451.	3.3	30
7	Visualization of mismatch repair complexes using fluorescence microscopy. DNA Repair, 2016, 38, 58-67.	1.3	16
8	New insights into the mechanism of DNA mismatch repair. Chromosoma, 2015, 124, 443-462.	1.0	103
9	Mlh2 Is an Accessory Factor for DNA Mismatch Repair in Saccharomyces cerevisiae. PLoS Genetics, 2014, 10, e1004327.	1.5	36
10	PCNA and Msh2-Msh6 Activate an Mlh1-Pms1 Endonuclease Pathway Required for Exo1-Independent Mismatch Repair. Molecular Cell, 2014, 55, 291-304.	4.5	89
11	Checkpoint Kinases Regulate a Global Network of Transcription Factors in Response to DNA Damage. Cell Reports, 2013, 4, 174-188.	2.9	61
12	Dominant Mutations in S. cerevisiae PMS1 Identify the Mlh1-Pms1 Endonuclease Active Site and an Exonuclease 1-Independent Mismatch Repair Pathway. PLoS Genetics, 2013, 9, e1003869.	1.5	52
13	Visualization of Eukaryotic DNA Mismatch Repair Reveals Distinct Recognition and Repair Intermediates. Cell, 2011, 147, 1040-1053.	13.5	183
14	Mismatch Repair, But Not Heteroduplex Rejection, Is Temporally Coupled to DNA Replication. Science, 2011, 334, 1713-1716.	6.0	109
15	SRD5A3 Is Required for Converting Polyprenol to Dolichol and Is Mutated in a Congenital Glycosylation Disorder. Cell, 2010, 142, 203-217.	13.5	253
16	Cdc28/Cdk1 positively and negatively affects genome stability in S. cerevisiae. Journal of Cell Biology, 2009, 185, 423-437.	2.3	37
17	Generation of Active Protein Phosphatase 2A Is Coupled to Holoenzyme Assembly. PLoS Biology, 2007, 5, e155.	2.6	74
18	Identification of a Subunit of a Novel Kleisin-β/SMC Complex as a Potential Substrate of Protein Phosphatase 2A. Current Biology, 2003, 13, 2058-2064.	1.8	84

#	Article	IF	CITATIONS
19	A novel and essential mechanism determining specificity and activity of protein phosphatase 2A (PP2A) in vivo. Genes and Development, 2003, 17, 2138-2150.	2.7	89