Janet F Partridge

List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	NSD1 mediates antagonism between SWI/SNF and polycomb complexes and is required for transcriptional activation upon EZH2 inhibition. Molecular Cell, 2022, 82, 2472-2489.e8.	9.7	18
2	Surprising phenotypic diversity of cancer-associated mutations of Gly 34 in the histone H3 tail. ELife, 2021, 10, .	6.0	22
3	Subtelomeric Chromatin in the Fission Yeast S. pombe. Microorganisms, 2021, 9, 1977.	3.6	2
4	Histone H3 Mutations: An Updated View of Their Role in Chromatin Deregulation and Cancer. Cancers, 2019, 11, 660.	3.7	105
5	Vitamin A differentially regulates cytokine expression in respiratory epithelial and macrophage cell lines. Cytokine, 2017, 91, 1-5.	3.2	21
6	Histone H3G34R mutation causes replication stress, homologous recombination defects and genomic instability in S. pombe. ELife, 2017, 6, .	6.0	36
7	SHREC Silences Heterochromatin via Distinct Remodeling and Deacetylation Modules. Molecular Cell, 2016, 62, 207-221.	9.7	45
8	Abo1, a conserved bromodomain <scp>AAA</scp> ― <scp>ATP</scp> ase, maintains global nucleosome occupancy and organisation. EMBO Reports, 2016, 17, 79-93.	4.5	22
9	Hotspots for Vitamin–Steroid–Thyroid Hormone Response Elements Within Switch Regions of Immunoglobulin Heavy Chain Loci Predict a Direct Influence of Vitamins and Hormones on B Cell Class Switch Recombination. Viral Immunology, 2016, 29, 132-136.	1.3	23
10	DNA Damage Response Checkpoint Activation Drives KP1019 Dependent Pre-Anaphase Cell Cycle Delay in S. cerevisiae. PLoS ONE, 2015, 10, e0138085.	2.5	8
11	Cancer-Associated Mutants of RNA Helicase DDX3X Are Defective in RNA-Stimulated ATP Hydrolysis. Journal of Molecular Biology, 2015, 427, 1779-1796.	4.2	66
12	Histone H3 mutations—a special role for H3.3 in tumorigenesis?. Chromosoma, 2015, 124, 177-189.	2.2	77
13	The landscape of somatic mutations in epigenetic regulators across 1,000 paediatric cancer genomes. Nature Communications, 2014, 5, 3630.	12.8	342
14	The Mi-2 Homolog Mit1 Actively Positions Nucleosomes within Heterochromatin To Suppress Transcription. Molecular and Cellular Biology, 2014, 34, 2046-2061.	2.3	29
15	Sir2 is required for Clr4 to initiate centromeric heterochromatin assembly in fission yeast. EMBO Journal, 2013, 32, 2321-2335.	7.8	68
16	Centromeric heterochromatin assembly in fission yeast—balancing transcription, RNA interference and chromatin modification. Chromosome Research, 2012, 20, 521-534.	2.2	28
17	Should I Stay or Should I Go? Chromodomain Proteins Seal the Fate of Heterochromatic Transcripts in Fission Yeast. Molecular Cell, 2012, 47, 153-155.	9.7	1
18	The Chp1–Tas3 core is a multifunctional platform critical for gene silencing by RITS. Nature Structural and Molecular Biology, 2011, 18, 1351-1357.	8.2	38

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19	RITS—connecting transcription, RNA interference, and heterochromatin assembly in fission yeast. Wiley Interdisciplinary Reviews RNA, 2011, 2, 632-646.	6.4	38
20	Cdk1 phosphorylation of the kinetochore protein Nsk1 prevents error-prone chromosome segregation. Journal of Cell Biology, 2011, 195, 583-593.	5.2	12
21	H3K9me-Independent Gene Silencing in Fission Yeast Heterochromatin by Clr5 and Histone Deacetylases. PLoS Genetics, 2011, 7, e1001268.	3.5	28
22	Continuous Requirement for the Clr4 Complex But Not RNAi for Centromeric Heterochromatin Assembly in Fission Yeast Harboring a Disrupted RITS Complex. PLoS Genetics, 2010, 6, e1001174.	3.5	24
23	Spreading the Silence. Developmental Cell, 2009, 16, 630-632.	7.0	1
24	High-Affinity Binding of Chp1 Chromodomain to K9 Methylated Histone H3 Is Required to Establish Centromeric Heterochromatin. Molecular Cell, 2009, 34, 36-46.	9.7	103
25	Chp1-Tas3 Interaction Is Required To Recruit RITS to Fission Yeast Centromeres and for Maintenance of Centromeric Heterochromatin. Molecular and Cellular Biology, 2008, 28, 2154-2166.	2.3	42
26	Centromeric chromatin in fission yeast. Frontiers in Bioscience - Landmark, 2008, Volume, 3896.	3.0	8
27	Plasticity of Fission Yeast CENP-A Chromatin Driven by Relative Levels of Histone H3 and H4. PLoS Genetics, 2007, 3, e121.	3.5	78
28	Functional Separation of the Requirements for Establishment and Maintenance of Centromeric Heterochromatin. Molecular Cell, 2007, 26, 593-602.	9.7	74
29	Characterization of Dicer-deficient murine embryonic stem cells. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 12135-12140.	7.1	742
30	RNA Interference (RNAi)-Dependent and RNAi-Independent Association of the Chp1 Chromodomain Protein with Distinct Heterochromatic Loci in Fission Yeast. Molecular and Cellular Biology, 2005, 25, 2331-2346.	2.3	80
31	Centromere Silencing and Function in Fission Yeast Is Governed by the Amino Terminus of Histone H3. Current Biology, 2003, 13, 1748-1757.	3.9	123
32	Schizosaccharomyces pombe Git7p, a Member of the Saccharomyces cerevisiae Sgt1p Family, Is Required for Glucose and Cyclic AMP Signaling, Cell Wall Integrity, and Septation. Eukaryotic Cell, 2002, 1, 558-567.	3.4	35
33	cis-Acting DNA from Fission Yeast Centromeres Mediates Histone H3 Methylation and Recruitment of Silencing Factors and Cohesin to an Ectopic Site. Current Biology, 2002, 12, 1652-1660.	3.9	165
34	Selective recognition of methylated lysine 9 on histone H3 by the HP1 chromo domain. Nature, 2001, 410, 120-124.	27.8	2,535
35	Requirement of Heterochromatin for Cohesion at Centromeres. Science, 2001, 294, 2539-2542.	12.6	583
36	Dimerisation of a chromo shadow domain and distinctions from the chromodomain as revealed by structural analysis. Current Biology, 2000, 10, 517-525.	3.9	228

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37	Distinct protein interaction domains and protein spreading in a complex centromere. Genes and Development, 2000, 14, 783-791.	5.9	219
38	Genetic characterisation of hda1+, a putative fission yeast histone deacetylase gene. Nucleic Acids Research, 1998, 26, 3247-3254.	14.5	25
39	Cell Cycle-dependent Transcription of CLN1 Involves Swi4 Binding to MCB-like Elements. Journal of Biological Chemistry, 1997, 272, 9071-9077.	3.4	52
40	A new component of the transcription factor DRTF1/E2F. Nature, 1993, 362, 83-87.	27.8	265