Michael R Botchan

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/7400580/publications.pdf

Version: 2024-02-01

27 papers 9,689 citations

304743 22 h-index 27 g-index

43 all docs 43 docs citations

times ranked

43

9161 citing authors

#	Article	IF	CITATIONS
1	The Genome Sequence of <i>Drosophila melanogaster</i> . Science, 2000, 287, 2185-2195.	12.6	5,566
2	Inhibition of SV40 replication in simian cells by specific pBR322 DNA sequences. Nature, 1981, 293, 79-81.	27.8	769
3	Isolation of the Cdc45/Mcm2-7/GINS (CMG) complex, a candidate for the eukaryotic DNA replication fork helicase. Proceedings of the National Academy of Sciences of the United States of America, 2006, 103, 10236-10241.	7.1	615
4	Association of the Origin Recognition Complex with Heterochromatin and HP1 in Higher Eukaryotes. Cell, 1997, 91, 311-323.	28.9	388
5	Activation of BPV-1 replication in vitro by the transcription factor E2. Nature, 1991, 353, 628-632.	27.8	355
6	The structural basis for MCM2–7 helicase activation by GINS and Cdc45. Nature Structural and Molecular Biology, 2011, 18, 471-477.	8.2	290
7	DNA topology, not DNA sequence, is a critical determinant for Drosophila ORC–DNA binding. EMBO Journal, 2004, 23, 897-907.	7.8	221
8	Mechanisms for initiating cellular DNA replication. Science, 2017, 355, .	12.6	171
9	Expression of enhanced levels of small RNA polymerase III transcripts encoded by the B2 repeats in simian virus 40-transformed mouse cells. Nature, 1985, 314, 553-556.	27.8	149
10	Mechanisms and regulation of DNA replication initiation in eukaryotes. Critical Reviews in Biochemistry and Molecular Biology, 2017, 52, 107-144.	5.2	140
11	Distinct Cytoplasmic and Nuclear Fractions of <i>Drosophila</i> Heterochromatin Protein 1: Their Phosphorylation Levels and Associations with Origin Recognition Complex Proteins. Journal of Cell Biology, 1998, 142, 307-318.	5.2	115
12	CRISPR germline engineering—the community speaks. Nature Biotechnology, 2015, 33, 478-486.	17.5	110
13	Crystal structure of the eukaryotic origin recognition complex. Nature, 2015, 519, 321-326.	27.8	109
14	DNA binding polarity, dimerization, and ATPase ring remodeling in the CMG helicase of the eukaryotic replisome. ELife, 2014, 3, e03273.	6.0	103
15	A new class of disordered elements controls DNA replication through initiator self-assembly. ELife, 2019, 8, .	6.0	92
16	Cdc45 (cell division cycle protein 45) guards the gate of the Eukaryote Replisome helicase stabilizing leading strand engagement. Proceedings of the National Academy of Sciences of the United States of America, 2015, 112, E249-58.	7.1	78
17	Molecular Basis for ATP-Hydrolysis-Driven DNA Translocation by the CMG Helicase of the Eukaryotic Replisome. Cell Reports, 2019, 28, 2673-2688.e8.	6.4	74
18	Crystal Structure of the Human Papillomavirus Type 18 E2 Activation Domain. Science, 1999, 284, 1673-1677.	12.6	67

#	Article	IF	CITATIONS
19	ATP-dependent conformational dynamics underlie the functional asymmetry of the replicative helicase from a minimalist eukaryote. Proceedings of the National Academy of Sciences of the United States of America, 2012, 109, 11999-12004.	7.1	65
20	A Meier-Gorlin syndrome mutation in a conserved C-terminal helix of Orc6 impedes origin recognition complex formation. ELife, 2013, 2, e00882.	6.0	45
21	Conformational control and DNA-binding mechanism of the metazoan origin recognition complex. Proceedings of the National Academy of Sciences of the United States of America, 2018, 115, E5906-E5915.	7.1	34
22	CDK Phosphorylation Inhibits the DNA-binding and ATP-hydrolysis Activities of the Drosophila Origin Recognition Complex. Journal of Biological Chemistry, 2005, 280, 39740-39751.	3.4	32
23	Structural Mechanisms for Replicating DNA in Eukaryotes. Annual Review of Biochemistry, 2021, 90, 77-106.	11.1	29
24	DNA Replication: Making Two Forks from One Prereplication Complex. Molecular Cell, 2010, 40, 860-861.	9.7	21
25	Hitchhiking without Covalent Integration. Cell, 2004, 117, 280-281.	28.9	20
26	Chromatin reader L(3)mbt requires the Myb–MuvB/DREAM transcriptional regulatory complex for chromosomal recruitment. Proceedings of the National Academy of Sciences of the United States of America, 2014, 111, E4234-43.	7.1	17
27	Molecular determinants of phase separation for Drosophila DNA replication licensing factors. ELife, 2021, 10, .	6.0	11