

Takuro Nakagawa

List of Publications by Year in descending order

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Version: 2024-02-01

34
papers

1,751
citations

257429

24
h-index

361001

35
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37
all docs

37
docs citations

37
times ranked

1810
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | Genome-wide localization of pre-RC sites and identification of replication origins in fission yeast. <i>EMBO Journal</i> , 2007, 26, 1327-1339. | 7.8 | 163 |
| 2 | The heterochromatin protein Swi6/HP1 activates replication origins at the pericentromeric region and silent mating-type locus. <i>Nature Cell Biology</i> , 2009, 11, 357-362. | 10.3 | 141 |
| 3 | The <i>Saccharomyces cerevisiae</i> MER3 gene, encoding a novel helicase-like protein, is required for crossover control in meiosis. <i>EMBO Journal</i> , 1999, 18, 5714-5723. | 7.8 | 128 |
| 4 | <i>Saccharomyces cerevisiae</i> Mer3 Helicase Stimulates 3' to 5' Heteroduplex Extension by Rad51. <i>Cell</i> , 2004, 117, 47-56. | 28.9 | 111 |
| 5 | Mcm10 plays an essential role in origin DNA unwinding after loading of the CMG components. <i>EMBO Journal</i> , 2012, 31, 2182-2194. | 7.8 | 97 |
| 6 | Ordered assembly of Sld3, GINS and Cdc45 is distinctly regulated by DDK and CDK for activation of replication origins. <i>EMBO Journal</i> , 2006, 25, 4663-4674. | 7.8 | 84 |
| 7 | Auxin-inducible protein depletion system in fission yeast. <i>BMC Cell Biology</i> , 2011, 12, 8. | 3.0 | 79 |
| 8 | Functions of the yeast meiotic recombination genes, MRE11 and MRE2. <i>Advances in Biophysics</i> , 1995, 31, 67-76. | 0.5 | 72 |
| 9 | Telomere-binding protein Taz1 controls global replication timing through its localization near late replication origins in fission yeast. <i>Genes and Development</i> , 2012, 26, 2050-2062. | 5.9 | 68 |
| 10 | Multiple functions of MutS- and MutL-related heterocomplexes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1999, 96, 14186-14188. | 7.1 | 60 |
| 11 | <i>Saccharomyces cerevisiae</i> Mer3 Is a DNA Helicase Involved in Meiotic Crossing Over. <i>Molecular and Cellular Biology</i> , 2002, 22, 3281-3291. | 2.3 | 59 |
| 12 | Rad51 suppresses gross chromosomal rearrangement at centromere in <i>Schizosaccharomyces pombe</i> . <i>EMBO Journal</i> , 2008, 27, 3036-3046. | 7.8 | 58 |
| 13 | The MER3 Helicase Involved in Meiotic Crossing Over Is Stimulated by Single-stranded DNA-binding Proteins and Unwinds DNA in the 3' to 5' Direction. <i>Journal of Biological Chemistry</i> , 2001, 276, 31487-31493. | 3.4 | 54 |
| 14 | Involvement of the MRE2 gene of yeast in formation of meiosis-specific double-strand breaks and crossover recombination through RNA splicing. <i>Genes To Cells</i> , 1997, 2, 65-79. | 1.2 | 50 |
| 15 | The Prereplication Complex Recruits XEco2 to Chromatin to Promote Cohesin Acetylation in <i>Xenopus</i> Egg Extracts. <i>Current Biology</i> , 2012, 22, 977-988. | 3.9 | 50 |
| 16 | DNA polymerization-independent functions of DNA polymerase epsilon in assembly and progression of the replisome in fission yeast. <i>Molecular Biology of the Cell</i> , 2012, 23, 3240-3253. | 2.1 | 41 |
| 17 | The MER3 DNA Helicase Catalyzes the Unwinding of Holliday Junctions. <i>Journal of Biological Chemistry</i> , 2002, 277, 28019-28024. | 3.4 | 38 |
| 18 | A novel allele of fission yeast rad11 that causes defects in DNA repair and telomere length regulation. <i>Nucleic Acids Research</i> , 2003, 31, 7141-7149. | 14.5 | 38 |

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 19 | MutS± maintains the mismatch repair capability by inhibiting PCNA unloading. <i>ELife</i> , 2016, 5, . | 6.0 | 37 |
| 20 | Nucleosomes around a mismatched base pair are excluded via an Msh2-dependent reaction with the aid of SNF2 family ATPase Smarcd1. <i>Genes and Development</i> , 2018, 32, 806-821. | 5.9 | 35 |
| 21 | CDK promotes interactions of Sld3 and Drc1 with Cut5 for initiation of DNA replication in fission yeast. <i>Molecular Biology of the Cell</i> , 2011, 22, 2620-2633. | 2.1 | 34 |
| 22 | Rad51 and Rad54 promote noncrossover recombination between centromere repeats on the same chromatid to prevent isochromosome formation. <i>Nucleic Acids Research</i> , 2016, 44, 10744-10757. | 14.5 | 30 |
| 23 | The DNA damage checkpoint pathway promotes extensive resection and nucleotide synthesis to facilitate homologous recombination repair and genome stability in fission yeast. <i>Nucleic Acids Research</i> , 2014, 42, 5644-5656. | 14.5 | 27 |
| 24 | A Novel Intermediate in Initiation Complex Assembly for Fission Yeast DNA Replication. <i>Molecular Biology of the Cell</i> , 2004, 15, 3740-3750. | 2.1 | 26 |
| 25 | Regulation of mitotic recombination between DNA repeats in centromeres. <i>Nucleic Acids Research</i> , 2017, 45, 11222-11235. | 14.5 | 26 |
| 26 | Mcm4 C-terminal domain of MCM helicase prevents excessive formation of single-stranded DNA at stalled replication forks. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2008, 105, 12973-12978. | 7.1 | 25 |
| 27 | Regulation of DNA Replication Machinery by Mrc1 in Fission Yeast. <i>Genetics</i> , 2006, 174, 155-165. | 2.9 | 24 |
| 28 | Heterochromatin suppresses gross chromosomal rearrangements at centromeres by repressing Tfs1/TFIIS-dependent transcription. <i>Communications Biology</i> , 2019, 2, 17. | 4.4 | 24 |
| 29 | Transcriptional silencing of centromere repeats by heterochromatin safeguards chromosome integrity. <i>Current Genetics</i> , 2019, 65, 1089-1098. | 1.7 | 20 |
| 30 | Abundance of Prereplicative Complexes (Pre-RCs) Facilitates Recombinational Repair under Replication Stress in Fission Yeast. <i>Journal of Biological Chemistry</i> , 2011, 286, 41701-41710. | 3.4 | 13 |
| 31 | DNA replication machinery prevents Rad52-dependent single-strand annealing that leads to gross chromosomal rearrangements at centromeres. <i>Communications Biology</i> , 2020, 3, 202. | 4.4 | 13 |
| 32 | Shelterin promotes tethering of late replication origins to telomeres for replication timing control. <i>EMBO Journal</i> , 2018, 37, . | 7.8 | 11 |
| 33 | Genome-wide localization of pre-RC sites and identification of replication origins in fission yeast. <i>EMBO Journal</i> , 2007, 26, 2821-2821. | 7.8 | 7 |
| 34 | Fission yeast Rad8/HLTF facilitates Rad52-dependent chromosomal rearrangements through PCNA lysine 107 ubiquitination. <i>PLoS Genetics</i> , 2021, 17, e1009671. | 3.5 | 5 |