

Youngho Kwon

List of Publications by Year in descending order

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2,657
citations

201674

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docs citations

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#	ARTICLE	IF	CITATIONS
1	Rad54 and Rdh54 prevent Srs2-mediated disruption of Rad51 presynaptic filaments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	7.1	11
2	Bloom helicase mediates formation of large single-stranded DNA loops during DNA end processing. <i>Nature Communications</i> , 2022, 13, 2248.	12.8	11
3	Single-molecule visualization of human RECQ5 interactions with single-stranded DNA recombination intermediates. <i>Nucleic Acids Research</i> , 2021, 49, 285-305.	14.5	15
4	The Rad51 paralog complex Rad55-Rad57 acts as a molecular chaperone during homologous recombination. <i>Molecular Cell</i> , 2021, 81, 1043-1057.e8.	9.7	45
5	Biochemical Analysis of D-Loop Extension and DNA Strand Displacement Synthesis. <i>Methods in Molecular Biology</i> , 2021, 2153, 87-99.	0.9	3
6	Single-molecule studies of yeast Rad51 paralogs. <i>Methods in Enzymology</i> , 2021, 661, 343-362.	1.0	0
7	Rad54 Drives ATP Hydrolysis-Dependent DNA Sequence Alignment during Homologous Recombination. <i>Cell</i> , 2020, 181, 1380-1394.e18.	28.9	77
8	The ZGRF1 Helicase Promotes Recombinational Repair of Replication-Blocking DNA Damage in Human Cells. <i>Cell Reports</i> , 2020, 32, 107849.	6.4	9
9	Rad54 and Rdh54 occupy spatially and functionally distinct sites within the Rad51-associated DNA presynaptic complex. <i>EMBO Journal</i> , 2020, 39, e105705.	7.8	17
10	Defining the influence of Rad51 and Dmc1 lineage-specific amino acids on genetic recombination. <i>Genes and Development</i> , 2019, 33, 1191-1207.	5.9	38
11	Single-molecule visualization of human BLM helicase as it acts upon double- and single-stranded DNA substrates. <i>Nucleic Acids Research</i> , 2019, 47, 11225-11237.	14.5	32
12	Guidelines for DNA recombination and repair studies: Mechanistic assays of DNA repair processes. <i>Microbial Cell</i> , 2019, 6, 65-101.	3.2	10
13	Regulatory control of Sgs1 and Dna2 during eukaryotic DNA end resection. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2019, 116, 6091-6100.	7.1	35
14	The RecQ helicase Sgs1 drives ATP-dependent disruption of Rad51 filaments. <i>Nucleic Acids Research</i> , 2019, 47, 4694-4706.	14.5	26
15	The BRCA Tumor Suppressor Network in Chromosome Damage Repair by Homologous Recombination. <i>Annual Review of Biochemistry</i> , 2019, 88, 221-245.	11.1	104
16	Dynamic interactions of the homologous pairing 2 (Hop2) meiotic nuclear divisions 1 (Mnd1) protein complex with meiotic presynaptic filaments in budding yeast. <i>Journal of Biological Chemistry</i> , 2019, 294, 490-501.	3.4	19
17	Regulation of Hed1 and Rad54 binding during maturation of the meiosis-specific presynaptic complex. <i>EMBO Journal</i> , 2018, 37, .	7.8	33
18	Spontaneous self-segregation of Rad51 and Dmc1 DNA recombinases within mixed recombinase filaments. <i>Journal of Biological Chemistry</i> , 2018, 293, 4191-4200.	3.4	24

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19	Meiosis-specific recombinase Dmc1 is a potent inhibitor of the Srs2 antirecombinase. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2018, 115, E10041-E10048.	7.1	29
20	A DNA nick at Ku-blocked double-strand break ends serves as an entry site for exonuclease 1 (Exo1) or Sgs1â€“Dna2 in long-range DNA end resection. <i>Journal of Biological Chemistry</i> , 2018, 293, 17061-17069.	3.4	19
21	Sequence imperfections and base triplet recognition by the Rad51/RecA family of recombinases. <i>Journal of Biological Chemistry</i> , 2017, 292, 11125-11135.	3.4	26
22	Human RAD52 interactions with replication protein A and the RAD51 presynaptic complex. <i>Journal of Biological Chemistry</i> , 2017, 292, 11702-11713.	3.4	47
23	Reconstituted System for the Examination of Repair DNA Synthesis in Homologous Recombination. <i>Methods in Enzymology</i> , 2017, 591, 307-325.	1.0	8
24	BRCA1â€“BARD1 promotes RAD51-mediated homologous DNA pairing. <i>Nature</i> , 2017, 550, 360-365.	27.8	262
25	Yeast Srs2 Helicase Promotes Redistribution of Single-Stranded DNA-Bound RPA and Rad52 in Homologous Recombination Regulation. <i>Cell Reports</i> , 2017, 21, 570-577.	6.4	36
26	Dissociation of Rad51 Presynaptic Complexes and Heteroduplex DNA Joints by Tandem Assemblies of Srs2. <i>Cell Reports</i> , 2017, 21, 3166-3177.	6.4	43
27	Role of the Pif1-PCNA Complex in Pol Î´-Dependent Strand Displacement DNA Synthesis and Break-Induced Replication. <i>Cell Reports</i> , 2017, 21, 1707-1714.	6.4	62
28	Rad52, Maestro of Inverse Strand Exchange. <i>Molecular Cell</i> , 2017, 67, 1-3.	9.7	30
29	Plasticity of the Mre11â€“Rad50â€“Xrs2â€“Sae2 nuclease ensemble in the processing of DNA-bound obstacles. <i>Genes and Development</i> , 2017, 31, 2331-2336.	5.9	96
30	Protein dynamics of human RPA and RAD51 on ssDNA during assembly and disassembly of the RAD51 filament. <i>Nucleic Acids Research</i> , 2017, 45, 749-761.	14.5	120
31	Enrichment of Cdk1-cyclins at DNA double-strand breaks stimulates Fun30 phosphorylation and DNA end resection. <i>Nucleic Acids Research</i> , 2016, 44, 2742-2753.	14.5	39
32	Mek1 Down Regulates Rad51 Activity during Yeast Meiosis by Phosphorylation of Hed1. <i>PLoS Genetics</i> , 2016, 12, e1006226.	3.5	76
33	DNA Sequence Alignment by Microhomology Sampling during Homologous Recombination. <i>Cell</i> , 2015, 160, 856-869.	28.9	182
34	Promotion of BRCA2-Dependent Homologous Recombination by DSS1 via RPA Targeting and DNA Mimicry. <i>Molecular Cell</i> , 2015, 59, 176-187.	9.7	141
35	Base triplet stepping by the Rad51/RecA family of recombinases. <i>Science</i> , 2015, 349, 977-981.	12.6	145
36	Restriction of Replication Fork Regression Activities by a Conserved SMC Complex. <i>Molecular Cell</i> , 2014, 56, 436-445.	9.7	60

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37	The Fanconi Anemia Proteins FANCD2 and FANCI Interact and Regulate Each Other's Chromatin Localization. <i>Journal of Biological Chemistry</i> , 2014, 289, 25774-25782.	3.4	17
38	Protein dynamics during presynaptic-complex assembly on individual single-stranded DNA molecules. <i>Nature Structural and Molecular Biology</i> , 2014, 21, 893-900.	8.2	81
39	Regulation of DNA Pairing in Homologous Recombination. <i>Cold Spring Harbor Perspectives in Biology</i> , 2014, 6, a017954-a017954.	5.5	82
40	Concentration-Dependent Exchange of Replication Protein A on Single-Stranded DNA Revealed by Single-Molecule Imaging. <i>PLoS ONE</i> , 2014, 9, e87922.	2.5	176
41	Pif1 helicase and Pol δ promote recombination-coupled DNA synthesis via bubble migration. <i>Nature</i> , 2013, 502, 393-396.	27.8	265
42	Investigations of homologous recombination pathways and their regulation. <i>Yale Journal of Biology and Medicine</i> , 2013, 86, 453-61.	0.2	40
43	Biochemical Studies on Human Rad51-Mediated Homologous Recombination. <i>Methods in Molecular Biology</i> , 2011, 745, 421-435.	0.9	6
44	ATP-dependent Chromatin Remodeling by the <i>Saccharomyces cerevisiae</i> Homologous Recombination Factor Rdh54. <i>Journal of Biological Chemistry</i> , 2008, 283, 10445-10452.	3.4	36
45	Synergistic action of the <i>Saccharomyces cerevisiae</i> homologous recombination factors Rad54 and Rad51 in chromatin remodeling. <i>DNA Repair</i> , 2007, 6, 1496-1506.	2.8	23