

# Stephen C West

## List of Publications by Year in descending order

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185  
papers

23,937  
citations

5126

86  
h-index

8878

150  
g-index

187  
all docs

187  
docs citations

187  
times ranked

15695  
citing authors

#	ARTICLE	IF	CITATIONS
1	Integrated genome and transcriptome analyses reveal the mechanism of genome instability in ataxia with oculomotor apraxia 2. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, .	3.3	16
2	Generation of double Holliday junction DNAs and their dissolution/resolution within a chromatin context. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2022, 119, e2123420119.	3.3	6
3	Defective ALC1 nucleosome remodeling confers PARPi sensitization and synthetic lethality with HRD. <i>Molecular Cell</i> , 2021, 81, 767-783.e11.	4.5	72
4	Charles M. Radding: A love of science and art. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2021, 118, e2025935118.	3.3	1
5	Coordinated roles of SLX4 and MutS $\hat{\nu}$ 2 in DNA repair and the maintenance of genome stability. <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 2021, 56, 157-177.	2.3	16
6	Targeting the nucleotide salvage factor DNPH1 sensitizes <i>BRCA</i> -deficient cells to PARP inhibitors. <i>Science</i> , 2021, 372, 156-165.	6.0	68
7	Tackling PARP inhibitor resistance. <i>Trends in Cancer</i> , 2021, 7, 1102-1118.	3.8	23
8	Repeat expansions confer WRN dependence in microsatellite-unstable cancers. <i>Nature</i> , 2020, 586, 292-298.	13.7	95
9	MutS $\hat{\nu}$ 2 Stimulates Holliday Junction Resolution by the SMX Complex. <i>Cell Reports</i> , 2020, 33, 108289.	2.9	23
10	Unresolved recombination intermediates lead to ultra-fine anaphase bridges, chromosome breaks and aberrations. <i>Nature Cell Biology</i> , 2018, 20, 92-103.	4.6	149
11	A new class of ultrafine anaphase bridges generated by homologous recombination. <i>Cell Cycle</i> , 2018, 17, 2101-2109.	1.3	29
12	GEN1 Endonuclease: Purification and Nuclease Assays. <i>Methods in Enzymology</i> , 2018, 600, 527-542.	0.4	4
13	Preparation and Resolution of Holliday Junction DNA Recombination Intermediates. <i>Methods in Enzymology</i> , 2018, 600, 569-590.	0.4	3
14	The SMX DNA Repair Tri-nuclease. <i>Molecular Cell</i> , 2017, 65, 848-860.e11.	4.5	98
15	Resolution of single and double Holliday junction recombination intermediates by GEN1. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, 443-450.	3.3	32
16	Genome Instability as a Consequence of Defects in the Resolution of Recombination Intermediates. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2017, 82, 207-212.	2.0	15
17	Analysis of Structure-Selective Endonuclease Activities From Yeast and Human Extracts. <i>Methods in Enzymology</i> , 2017, 591, 271-286.	0.4	4
18	SMX makes the cut in genome stability. <i>Oncotarget</i> , 2017, 8, 102765-102766.	0.8	0

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19	Keeping homologous recombination in check. <i>Cell Research</i> , 2016, 26, 397-398.	5.7	6
20	GEN1 promotes Holliday junction resolution by a coordinated nick and counter-nick mechanism. <i>Nucleic Acids Research</i> , 2015, 43, 10882-10892.	6.5	32
21	Structural and Mechanistic Analysis of the Slx1-Slx4 Endonuclease. <i>Cell Reports</i> , 2015, 10, 1467-1476.	2.9	28
22	Resolution of Recombination Intermediates: Mechanisms and Regulation. <i>Cold Spring Harbor Symposia on Quantitative Biology</i> , 2015, 80, 103-109.	2.0	95
23	Substrate specificity of the MUS81-EME2 structure selective endonuclease. <i>Nucleic Acids Research</i> , 2014, 42, 3833-3845.	6.5	50
24	Spatial control of the GEN1 Holliday junction resolvase ensures genome stability. <i>Nature Communications</i> , 2014, 5, 4844.	5.8	80
25	Roles of SLX1-SLX4, MUS81-EME1, and GEN1 in avoiding genome instability and mitotic catastrophe. <i>Genes and Development</i> , 2014, 28, 1124-1136.	2.7	106
26	Holliday junction processing enzymes as guardians of genome stability. <i>Trends in Biochemical Sciences</i> , 2014, 39, 409-419.	3.7	85
27	Holliday Junction Resolvases. <i>Cold Spring Harbor Perspectives in Biology</i> , 2014, 6, a023192-a023192.	2.3	165
28	Structure and mechanism of action of the BRCA2 breast cancer tumor suppressor. <i>Nature Structural and Molecular Biology</i> , 2014, 21, 962-968.	3.6	95
29	MUS81-EME2 Promotes Replication Fork Restart. <i>Cell Reports</i> , 2014, 7, 1048-1055.	2.9	119
30	Dual Control of Yen1 Nuclease Activity and Cellular Localization by Cdk and Cdc14 Prevents Genome Instability. <i>Molecular Cell</i> , 2014, 54, 94-106.	4.5	108
31	Holliday junction resolution: Regulation in space and time. <i>DNA Repair</i> , 2014, 19, 176-181.	1.3	124
32	Cell-Cycle Kinases Coordinate the Resolution of Recombination Intermediates with Chromosome Segregation. <i>Cell Reports</i> , 2013, 4, 76-86.	2.9	77
33	Coordinated Actions of SLX1-SLX4 and MUS81-EME1 for Holliday Junction Resolution in Human Cells. <i>Molecular Cell</i> , 2013, 52, 234-247.	4.5	252
34	DNA-dependent SUMO modification of PARP-1. <i>DNA Repair</i> , 2013, 12, 761-773.	1.3	24
35	Architecture and DNA Recognition Elements of the Fanconi Anemia FANCM-FAAP24 Complex. <i>Structure</i> , 2013, 21, 1648-1658.	1.6	26
36	Senataxin, Defective in the Neurodegenerative Disorder Ataxia with Oculomotor Apraxia 2, Lies at the Interface of Transcription and the DNA Damage Response. <i>Molecular and Cellular Biology</i> , 2013, 33, 406-417.	1.1	163

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37	Distinct Roles of Mus81, Yen1, Slx1-Slx4, and Rad1 Nucleases in the Repair of Replication-Born Double-Strand Breaks by Sister Chromatid Exchange. <i>Molecular and Cellular Biology</i> , 2012, 32, 1592-1603.	1.1	58
38	The DNA translocase activity of FANCM protects stalled replication forks. <i>Human Molecular Genetics</i> , 2012, 21, 2005-2016.	1.4	71
39	Regulatory Control of the Resolution of DNA Recombination Intermediates during Meiosis and Mitosis. <i>Cell</i> , 2011, 147, 158-172.	13.5	263
40	DNA interstrand crosslink repair and cancer. <i>Nature Reviews Cancer</i> , 2011, 11, 467-480.	12.8	847
41	Aberrant chromosome morphology in human cells defective for Holliday junction resolution. <i>Nature</i> , 2011, 471, 642-646.	13.7	190
42	Functional overlap between the structure-specific nucleases Yen1 and Mus81-Mms4 for DNA-damage repair in <i>S. cerevisiae</i> . <i>DNA Repair</i> , 2010, 9, 394-402.	1.3	86
43	Solution structures of the two PBZ domains from human APLF and their interaction with poly(ADP-ribose). <i>Nature Structural and Molecular Biology</i> , 2010, 17, 241-243.	3.6	89
44	The breast cancer tumor suppressor BRCA2 promotes the specific targeting of RAD51 to single-stranded DNA. <i>Nature Structural and Molecular Biology</i> , 2010, 17, 1263-1265.	3.6	217
45	InTERTpreting telomerase structure and function. <i>Nucleic Acids Research</i> , 2010, 38, 5609-5622.	6.5	146
46	The human Holliday junction resolvase GEN1 rescues the meiotic phenotype of a <i>Schizosaccharomyces pombe</i> mus81 mutant. <i>Nucleic Acids Research</i> , 2010, 38, 1866-1873.	6.5	51
47	RTEL-1 Enforces Meiotic Crossover Interference and Homeostasis. <i>Science</i> , 2010, 327, 1254-1258.	6.0	155
48	Identification of KIAA1018/FAN1, a DNA Repair Nuclease Recruited to DNA Damage by Monoubiquitinated FANCD2. <i>Cell</i> , 2010, 142, 65-76.	13.5	284
49	Mechanism of Holliday junction resolution by the human GEN1 protein. <i>Genes and Development</i> , 2010, 24, 1559-1569.	2.7	128
50	TRF2 promotes, remodels and protects telomeric Holliday junctions. <i>EMBO Journal</i> , 2009, 28, 641-651.	3.5	99
51	FANCM Connects the Genome Instability Disorders Bloom's Syndrome and Fanconi Anemia. <i>Molecular Cell</i> , 2009, 36, 943-953.	4.5	221
52	Poly(ADP-ribose)-Dependent Regulation of DNA Repair by the Chromatin Remodeling Enzyme ALC1. <i>Science</i> , 2009, 325, 1240-1243.	6.0	504
53	The search for a human Holliday junction resolvase. <i>Biochemical Society Transactions</i> , 2009, 37, 519-526.	1.6	47
54	Poly(ADP-ribose)-binding zinc finger motifs in DNA repair/checkpoint proteins. <i>Nature</i> , 2008, 451, 81-85.	13.7	367

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55	Single-stranded DNA-binding protein hSSB1 is critical for genomic stability. <i>Nature</i> , 2008, 453, 677-681.	13.7	220
56	Identification of Holliday junction resolvases from humans and yeast. <i>Nature</i> , 2008, 456, 357-361.	13.7	345
57	DNA Repair Synthesis Facilitates RAD52-Mediated Second-End Capture during DSB Repair. <i>Molecular Cell</i> , 2008, 29, 510-516.	4.5	109
58	FANCM and FAAP24 Function in ATR-Mediated Checkpoint Signaling Independently of the Fanconi Anemia Core Complex. <i>Molecular Cell</i> , 2008, 32, 313-324.	4.5	187
59	RTEL1 Maintains Genomic Stability by Suppressing Homologous Recombination. <i>Cell</i> , 2008, 135, 261-271.	13.5	315
60	Molecular Mechanism of DNA Deadenylation by the Neurological Disease Protein Aprataxin. <i>Journal of Biological Chemistry</i> , 2008, 283, 33994-34001.	1.6	33
61	More complexity to the Bloom's syndrome complex. <i>Genes and Development</i> , 2008, 22, 2737-2742.	2.7	33
62	Structural and Functional Relationships of the XPF/MUS81 Family of Proteins. <i>Annual Review of Biochemistry</i> , 2008, 77, 259-287.	5.0	244
63	RIDDLE immunodeficiency syndrome is linked to defects in 53BP1-mediated DNA damage signaling. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 16910-16915.	3.3	152
64	Actions of Aprataxin in Multiple DNA Repair Pathways. <i>Journal of Biological Chemistry</i> , 2007, 282, 9469-9474.	1.6	78
65	RAD51C deficiency in mice results in early prophase I arrest in males and sister chromatid separation at metaphase II in females. <i>Journal of Cell Biology</i> , 2007, 176, 581-592.	2.3	118
66	Role of RAD51C and XRCC3 in Genetic Recombination and DNA Repair. <i>Journal of Biological Chemistry</i> , 2007, 282, 1973-1979.	1.6	101
67	Defective DNA Repair and Neurodegenerative Disease. <i>Cell</i> , 2007, 130, 991-1004.	13.5	295
68	Identification of FAAP24, a Fanconi Anemia Core Complex Protein that Interacts with FANCM. <i>Molecular Cell</i> , 2007, 25, 331-343.	4.5	264
69	Interactions between human BRCA2 protein and the meiosis-specific recombinase DMC1. <i>EMBO Journal</i> , 2007, 26, 2915-2922.	3.5	104
70	Stabilization of RAD51 nucleoprotein filaments by the C-terminal region of BRCA2. <i>Nature Structural and Molecular Biology</i> , 2007, 14, 468-474.	3.6	240
71	The neurodegenerative disease protein aprataxin resolves abortive DNA ligation intermediates. <i>Nature</i> , 2006, 443, 713-716.	13.7	348
72	Interplay between human DNA repair proteins at a unique double-strand break in vivo. <i>EMBO Journal</i> , 2006, 25, 222-231.	3.5	172

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73	Synthetic Junctions as Tools to Identify and Characterize Holliday Junction Resolvases. <i>Methods in Enzymology</i> , 2006, 408, 485-501.	0.4	25
74	CDK-dependent phosphorylation of BRCA2 as a regulatory mechanism for recombinational repair. <i>Nature</i> , 2005, 434, 598-604.	13.7	428
75	BRCA2 BRC motifs bind RAD51-DNA filaments. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2005, 102, 8537-8542.	3.3	130
76	Recombination at Mammalian Telomeres: An Alternative Mechanism for Telomere Protection and Elongation. <i>Cell Cycle</i> , 2005, 4, 672-674.	1.3	30
77	Human DNA Polymerase $\delta$ Promotes DNA Synthesis from Strand Invasion Intermediates of Homologous Recombination. <i>Molecular Cell</i> , 2005, 20, 783-792.	4.5	287
78	RAD51 localization and activation following DNA damage. <i>Philosophical Transactions of the Royal Society B: Biological Sciences</i> , 2004, 359, 87-93.	1.8	91
79	Direct interaction of FANCD2 with BRCA2 in DNA damage response pathways. <i>Human Molecular Genetics</i> , 2004, 13, 1241-1248.	1.4	190
80	RAD51C Is Required for Holliday Junction Processing in Mammalian Cells. <i>Science</i> , 2004, 303, 243-246.	6.0	289
81	Happy Hollidays: 40th anniversary of the Holliday junction. <i>Nature Reviews Molecular Cell Biology</i> , 2004, 5, 937-944.	16.1	152
82	Human RECQ5 $\delta$ , a protein with DNA helicase and strand-annealing activities in a single polypeptide. <i>EMBO Journal</i> , 2004, 23, 2882-2891.	3.5	184
83	Holliday Junction Branch Migration and Resolution Assays. , 2004, 262, 239-254.		8
84	Conformational Changes Modulate the Activity of Human RAD51 Protein. <i>Journal of Molecular Biology</i> , 2004, 337, 817-827.	2.0	53
85	Telomere Maintenance Requires the RAD51D Recombination/Repair Protein. <i>Cell</i> , 2004, 117, 337-347.	13.5	204
86	BRCA2-dependent and independent formation of RAD51 nuclear foci. <i>Oncogene</i> , 2003, 22, 1115-1123.	2.6	173
87	Molecular views of recombination proteins and their control. <i>Nature Reviews Molecular Cell Biology</i> , 2003, 4, 435-445.	16.1	860
88	Eme1 is involved in DNA damage processing and maintenance of genomic stability in mammalian cells. <i>EMBO Journal</i> , 2003, 22, 6137-6147.	3.5	118
89	XRCC3 and Rad51 Modulate Replication Fork Progression on Damaged Vertebrate Chromosomes. <i>Molecular Cell</i> , 2003, 11, 1109-1117.	4.5	148
90	Cross-links between Fanconi anaemia and BRCA2. <i>DNA Repair</i> , 2003, 2, 231-234.	1.3	3

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91	Identification and Characterization of the Human Mus81-Eme1 Endonuclease. <i>Journal of Biological Chemistry</i> , 2003, 278, 25172-25178.	1.6	189
92	Structure of the single-strand annealing domain of human RAD52 protein. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 13492-13497.	3.3	211
93	Role of Mammalian RAD51L2 (RAD51C) in Recombination and Genetic Stability. <i>Journal of Biological Chemistry</i> , 2002, 277, 19322-19330.	1.6	88
94	Involvement of human polynucleotide kinase in double-strand break repair by non-homologous end joining. <i>EMBO Journal</i> , 2002, 21, 2827-2832.	3.5	234
95	Specific interaction of IP6 with human Ku70/80, the DNA-binding subunit of DNA-PK. <i>EMBO Journal</i> , 2002, 21, 2038-2044.	3.5	90
96	Holliday junction resolution in human cells: two junction endonucleases with distinct substrate specificities. <i>EMBO Journal</i> , 2002, 21, 5577-5585.	3.5	151
97	Distinct functions of BRCA1 and BRCA2 in double-strand break repair. <i>Breast Cancer Research</i> , 2001, 4, 9-13.	2.2	69
98	The efficiency of strand invasion by <i>Escherichia coli</i> RecA is dependent upon the length and polarity of ssDNA tails. <i>Journal of Molecular Biology</i> , 2001, 305, 23-31.	2.0	17
99	Role of BRCA2 in Control of the RAD51 Recombination and DNA Repair Protein. <i>Molecular Cell</i> , 2001, 7, 273-282.	4.5	617
100	Branch Migration and Holliday Junction Resolution Catalyzed by Activities from Mammalian Cells. <i>Cell</i> , 2001, 104, 259-268.	13.5	140
101	Visualization of recombination intermediates produced by RAD52-mediated single-strand annealing. <i>EMBO Reports</i> , 2001, 2, 905-909.	2.0	69
102	The Rad51 and Dmc1 recombinases: a non-identical twin relationship. <i>Trends in Biochemical Sciences</i> , 2001, 26, 131-136.	3.7	158
103	Identification and purification of two distinct complexes containing the five RAD51 paralogs. <i>Genes and Development</i> , 2001, 15, 3296-3307.	2.7	323
104	RuvAB-mediated branch migration does not involve extensive DNA opening within the RuvB hexamer. <i>Current Biology</i> , 2000, 10, 103-106.	1.8	19
105	The human Rad52 protein exists as a heptameric ring. <i>Current Biology</i> , 2000, 10, 337-340.	1.8	196
106	Werner's syndrome protein (WRN) migrates Holliday junctions and co-localizes with RPA upon replication arrest. <i>EMBO Reports</i> , 2000, 1, 80-84.	2.0	378
107	Precise binding of single-stranded DNA termini by human RAD52 protein. <i>EMBO Journal</i> , 2000, 19, 4175-4181.	3.5	77
108	Cleavage of Holliday Junctions by the <i>Escherichia coli</i> RuvABC Complex. <i>Journal of Biological Chemistry</i> , 2000, 275, 26467-26476.	1.6	43

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109	Functional interactions of Mycobacterium leprae RuvA with Escherichia coli RuvB and RuvC on holliday junctions 1 1Edited by M. Yaniv. Journal of Molecular Biology, 2000, 301, 839-850.	2.0	6
110	Reconstitution of the strand invasion step of double-strand break repair using human Rad51 Rad52 and RPA proteins. Journal of Molecular Biology, 2000, 304, 151-164.	2.0	104
111	Binding of Inositol Phosphate to DNA-PK and Stimulation of Double-Strand Break Repair. Cell, 2000, 102, 721-729.	13.5	231
112	Gross chromosomal rearrangements and genetic exchange between nonhomologous chromosomes following BRCA2 inactivation. Genes and Development, 2000, 14, 1400-1406.	2.7	267
113	Binding of double-strand breaks in DNA by human Rad52 protein. Nature, 1999, 398, 728-731.	13.7	279
114	The meiosis-specific recombinase hDmc1 forms ring structures and interacts with hRad51. EMBO Journal, 1999, 18, 6552-6560.	3.5	123
115	Heteroduplex Formation by Human Rad51 Protein: Effects of DNA End-structure, hRP-A and hRad52. Journal of Molecular Biology, 1999, 291, 363-374.	2.0	57
116	Helicase-defective RuvB D113E promotes RuvAB-mediated branch migration in Vitro 1 1Edited by J. Karn. Journal of Molecular Biology, 1999, 293, 505-519.	2.0	14
117	Binding of double-strand breaks in DNA by human Rad52 protein. Nature, 1999, 401, 403-403.	13.7	3
118	Title is missing!. Nature, 1999, 401, 403-403.	13.7	0
119	Synergistic actions of Rad51 and Rad52 in recombination and DNA repair. Nature, 1998, 391, 401-404.	13.7	371
120	Formation of RuvABC“Holliday junction complexes in vitro. Current Biology, 1998, 8, 725-727.	1.8	50
121	Role of the human RAD51 protein in homologous recombination and double-stranded-break repair. Trends in Biochemical Sciences, 1998, 23, 247-251.	3.7	492
122	RuvA Gets X-Rayed on Holliday. Cell, 1998, 94, 699-701.	13.5	16
123	Sequence-specificity of holliday junction resolution: identification of RuvC mutants defective in metal binding and target site recognition. Journal of Molecular Biology, 1998, 281, 17-29.	2.0	17
124	Coordinated actions of RuvABC in Holliday junction processing 1 1Edited by J. Karn. Journal of Molecular Biology, 1998, 281, 621-630.	2.0	69
125	Visualisation of human rad52 protein and its complexes with hrad51 and DNA. Journal of Molecular Biology, 1998, 284, 1027-1038.	2.0	109
126	PROCESSING OF RECOMBINATION INTERMEDIATES BY THE RuvABC PROTEINS. Annual Review of Genetics, 1997, 31, 213-244.	3.2	439



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127	Structure and subunit composition of the RuvAB-holliday junction complex 1 Edited by P. E. Wright. Journal of Molecular Biology, 1997, 266, 217-222.	2.0	85
128	Effect of DNA topology on holliday junction resolution by Escherichia coli RuvC and bacteriophage T7 endonuclease I. Journal of Molecular Biology, 1997, 270, 663-673.	2.0	24
129	Biochemical properties of RuvBD113N: a mutation in helicase motif II of the RuvB hexamer affects DNA binding and ATPase activities. Journal of Molecular Biology, 1997, 271, 704-717.	2.0	43
130	Purification of human Rad51 protein by selective spermidine precipitation. Mutation Research DNA Repair, 1997, 384, 65-72.	3.8	64
131	In Vitro Reconstitution of the Late Steps of Genetic Recombination in E. coli. Cell, 1997, 89, 607-617.	13.5	122
132	Bypass of DNA Heterologies During RuvAB-mediated Three- and Four-strand Branch Migration. Journal of Molecular Biology, 1996, 263, 582-596.	2.0	47
133	DNA Helicases: New Breeds of Translocating Motors and Molecular Pumps. Cell, 1996, 86, 177-180.	13.5	141
134	Human Rad51 Protein Promotes ATP-Dependent Homologous Pairing and Strand Transfer Reactions In Vitro. Cell, 1996, 87, 757-766.	13.5	630
135	Exchanging partners: recombination in E. coli. Trends in Genetics, 1996, 12, 20-26.	2.9	74
136	The directionality of RuvAB-mediated branch migration: in vitro studies with three-armed junctions. Genes To Cells, 1996, 1, 443-451.	0.5	19
137	DNA helicases get physical. Nature, 1996, 384, 316-317.	13.7	3
138	Role of RuvA in Branch Migration Reactions Catalyzed by the RuvA and RuvB Proteins of Escherichia coli. Journal of Biological Chemistry, 1996, 271, 19497-19502.	1.6	19
139	Characterisation of RuvAB-Holliday junction complexes by glycerol gradient sedimentation. Nucleic Acids Research, 1995, 23, 3621-3626.	6.5	10
140	Holliday junctions cleaved by Rad1?. Nature, 1995, 373, 27-28.	13.7	24
141	Structure of a multisubunit complex that promotes DNA branch migration. Nature, 1995, 374, 375-378.	13.7	182
142	Role of the Rad1 and Rad10 Proteins in Nucleotide Excision Repair and Recombination. Journal of Biological Chemistry, 1995, 270, 24638-24641.	1.6	123
143	Relaxing and unwinding on Holliday: DNA helicase-mediated branch migration. Mutation Research DNA Repair, 1995, 337, 149-159.	3.8	10
144	Branch migration during homologous recombination: assembly of a RuvAB-holliday junction complex in vitro. Cell, 1995, 80, 787-793.	13.5	72

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145	Unwinding of Closed Circular DNA by the Escherichia coli RuvA and RuvB Recombination/Repair Proteins. <i>Journal of Molecular Biology</i> , 1995, 247, 404-417.	2.0	34
146	Structural Analysis of the RuvC-Holliday Junction Complex Reveals an Unfolded Junction. <i>Journal of Molecular Biology</i> , 1995, 252, 213-226.	2.0	118
147	Formation, translocation and resolution of Holliday junctions during homologous genetic recombination. , 1995, , 17-21.		0
148	Activation of RuvC Holliday junction resolvase in vitro. <i>Nucleic Acids Research</i> , 1994, 22, 2490-2497.	6.5	50
149	XPG endonuclease makes the 3' incision in human DNA nucleotide excision repair. <i>Nature</i> , 1994, 371, 432-435.	13.7	450
150	The processing of recombination intermediates: Mechanistic insights from studies of bacterial proteins. <i>Cell</i> , 1994, 76, 9-15.	13.5	118
151	Processing of Holliday Junctions by RuvABC? An Overview. <i>Annals of the New York Academy of Sciences</i> , 1994, 726, 156-164.	1.8	5
152	Hexameric Rings of Escherichia coli RuvB Protein.. <i>Journal of Molecular Biology</i> , 1994, 243, 208-215.	2.0	76
153	An Assay for In Vitro Recombination Between Duplex DNA Molecules. , 1994, 30, 413-424.		2
154	Recombination genes and proteins. <i>Current Opinion in Genetics and Development</i> , 1994, 4, 221-228.	1.5	41
155	Formation of a RuvAB-Holliday Junction Complex in Vitro. <i>Journal of Molecular Biology</i> , 1993, 232, 397-405.	2.0	108
156	Resolution of holliday junctions by RuvC resolvase: Cleavage specificity and DNA distortion. <i>Cell</i> , 1993, 74, 1021-1031.	13.5	190
157	Enzymes and Molecular Mechanisms of Genetic Recombination. <i>Annual Review of Biochemistry</i> , 1992, 61, 603-640.	5.0	389
158	ATP-dependent branch migration of holliday junctions promoted by the RuvA and RuvB proteins of E. coli. <i>Cell</i> , 1992, 69, 1171-1180.	13.5	262
159	Purification and properties of the RuvA and RuvB proteins of Escherichia coli. <i>Molecular Genetics and Genomics</i> , 1992, 235, 1-10.	2.4	105
160	Biological roles of the Escherichia coli RuvA, RuvB and RuvC proteins revealed. <i>Molecular Microbiology</i> , 1992, 6, 2755-2759.	1.2	43
161	Formation and resolution of recombination intermediates by E. coli RecA and RuvC proteins. <i>Nature</i> , 1991, 354, 506-510.	13.7	243
162	Three-stranded DNA helices as intermediates in genetic recombination. <i>BioEssays</i> , 1991, 13, 37-38.	1.2	4

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163	T7 endonuclease I resolves Holliday junctions formed in vitro by RecA protein. <i>Nucleic Acids Research</i> , 1990, 18, 5633-5636.	6.5	20
164	Processing of recombination intermediates in vitro. <i>BioEssays</i> , 1990, 12, 151-154.	1.2	14
165	Specificity of binding to four-way junctions in DNA by bacteriophage T7 endonuclease I. <i>Nucleic Acids Research</i> , 1990, 18, 4377-4384.	6.5	42
166	Enzymatic formation and resolution of Holliday junctions in vitro. <i>Cell</i> , 1990, 60, 329-336.	13.5	56
167	Cleavage specificity of bacteriophage T4 endonuclease VII and bacteriophage T7 endonuclease I on synthetic branch migratable holliday junctions. <i>Journal of Molecular Biology</i> , 1990, 212, 723-735.	2.0	58
168	Homologous pairing and the formation of nascent synaptic intermediates between regions of duplex DNA by RecA protein. <i>Cell</i> , 1989, 56, 987-995.	13.5	59
169	Proteins from Yeast and Human Cells Specific for Model Holliday Junctions in DNA. , 1989, , 233-243.		1
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173	Role of RecA protein spiral filaments in genetic recombination. <i>Nature</i> , 1984, 309, 215-220.	13.7	389
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176	Enzymatic formation of biparental figure-eight molecules from plasmid DNA and their resolution in <i>E. coli</i> . <i>Cell</i> , 1983, 32, 817-829.	13.5	67
177	Role of SSB protein in RecA promoted branch migration reactions. <i>Molecular Genetics and Genomics</i> , 1982, 186, 333-338.	2.4	55
178	Postreplication repair in <i>E. coli</i> : Strand exchange reactions of gapped DNA by RecA protein. <i>Molecular Genetics and Genomics</i> , 1982, 187, 209-217.	2.4	98
179	Homologous pairing can occur before DNA strand separation in general genetic recombination. <i>Nature</i> , 1981, 290, 29-33.	13.7	49
180	Mechanism of <i>E. coli</i> RecA protein directed strand exchanges in post-replication repair of DNA. <i>Nature</i> , 1981, 294, 659-662.	13.7	89

#	ARTICLE	IF	CITATIONS
181	Genetic recombination: recA protein promotes homologous pairing between duplex DNA molecules without strand unwinding. <i>Nucleic Acids Research</i> , 1981, 9, 4201-4210.	6.5	15
182	INDUCTION AND REPRESSION OF THE <i>recA</i> GENE OF <i>ESCHERICHIA COLI</i> . , 1978, , 367-370.		0
183	Induction of protein synthesis in <i>Escherichia coli</i> following UV-or $\hat{\text{I}}^3$ -irradiation, mitomycin C treatment or <i>tif</i> expression. <i>Molecular Genetics and Genomics</i> , 1977, 151, 57-67.	2.4	50
184	Identification of protein X of <i>Escherichia coli</i> as the <i>recA</i> + <i>tif</i> + gene product. <i>Molecular Genetics and Genomics</i> , 1977, 155, 77-85.	2.4	153
185	<i>recA</i> +-dependent inactivation of the lambda repressor in <i>Escherichia coli</i> lysogens by $\hat{\text{I}}^3$ -radiation and by <i>tif</i> expression. <i>Molecular Genetics and Genomics</i> , 1975, 141, 1-8.	2.4	22