

Edward L Bolt

List of Publications by Year in descending order

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49
papers

1,504
citations

279487

23
h-index

329751

37
g-index

51
all docs

51
docs citations

51
times ranked

1458
citing authors

#	ARTICLE	IF	CITATIONS
1	Gene dissection demonstrates that the <i>Escherichia coli cysG</i> gene encodes a multifunctional protein. <i>Biochemical Journal</i> , 1994, 302, 837-844.	1.7	109
2	Intrinsic sequence specificity of the Cas1 integrase directs new spacer acquisition. <i>ELife</i> , 2015, 4, .	2.8	104
3	Archaeal Hel308 helicase targets replication forks in vivo and in vitro and unwinds lagging strands. <i>Nucleic Acids Research</i> , 2005, 33, 3678-3690.	6.5	91
4	Interactions Between RuvA and RuvC at Holliday Junctions: Inhibition of Junction Cleavage and Formation of a RuvA-RuvC-DNA Complex. <i>Journal of Molecular Biology</i> , 1996, 264, 878-890.	2.0	86
5	Different genome stability proteins underpin primed and naïve adaptation in <i>E. coli</i> CRISPR-Cas immunity. <i>Nucleic Acids Research</i> , 2015, 43, 10821-10830.	6.5	86
6	Molecular insights into DNA interference by CRISPR-associated nuclease-helicase Cas3. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2014, 111, 16359-16364.	3.3	85
7	Substrate Specificity of RusA Resolvase Reveals the DNA Structures Targeted by RuvAB and RecG In Vivo. <i>Molecular Cell</i> , 2002, 10, 187-198.	4.5	68
8	Sequence Specificity and Biochemical Characterization of the RusA Holliday Junction Resolvase of <i>Escherichia coli</i> . <i>Journal of Biological Chemistry</i> , 1997, 272, 14873-14882.	1.6	65
9	Replication fork regression in vitro by the Werner syndrome protein (WRN): Holliday junction formation, the effect of leading arm structure and a potential role for WRN exonuclease activity. <i>Nucleic Acids Research</i> , 2007, 35, 5729-5747.	6.5	58
10	Helicase dissociation and annealing of RNA-DNA hybrids by <i>Escherichia coli</i> Cas3 protein. <i>Biochemical Journal</i> , 2011, 439, 85-95.	1.7	56
11	Interactions of RadB, a DNA Repair Protein in Archaea, with DNA and ATP. <i>Journal of Molecular Biology</i> , 2006, 358, 46-56.	2.0	38
12	Molecular biology of Hel308 helicase in archaea. <i>Biochemical Society Transactions</i> , 2009, 37, 74-78.	1.6	38
13	The role of Cas8 in type I CRISPR interference. <i>Bioscience Reports</i> , 2015, 35, .	1.1	37
14	Characterization of the <i>Rhodobacter sphaeroides</i> 5-aminolaevulinic acid synthase isoenzymes, HemA and HemT, isolated from recombinant <i>Escherichia coli</i> . <i>FEBS Journal</i> , 1999, 265, 290-299.	0.2	33
15	The Conjugative DNA Translocase TrwB Is a Structure-specific DNA-binding Protein. <i>Journal of Biological Chemistry</i> , 2010, 285, 17537-17544.	1.6	32
16	Cas3 is a limiting factor for CRISPR-Cas immunity in <i>Escherichia coli</i> cells lacking H-NS. <i>BMC Microbiology</i> , 2016, 16, 28.	1.3	30
17	A novel nuclease-ATPase (Nar71) from archaea is part of a proposed thermophilic DNA repair system. <i>Nucleic Acids Research</i> , 2004, 32, 6176-6186.	6.5	28
18	Tuning in to Interference: R-Loops and Cascade Complexes in CRISPR Immunity. <i>Journal of Molecular Biology</i> , 2012, 422, 607-616.	2.0	28

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19	Molecular Ageing of Alpha- and Beta-Synucleins: Protein Damage and Repair Mechanisms. PLoS ONE, 2013, 8, e61442.	1.1	28
20	Identification of three aspartic acid residues essential for catalysis by the RusA holliday junction resolvase 1 Edited by A. R. Fersht. Journal of Molecular Biology, 1999, 286, 403-415.	2.0	27
21	CRISPR-Cas adaptation in Escherichia coli requires RecBCD helicase but not nuclease activity, is independent of homologous recombination, and is antagonized by 5' ssDNA exonucleases. Nucleic Acids Research, 2018, 46, 10173-10183.	6.5	27
22	CRISPR-Cas immunity, DNA repair and genome stability. Bioscience Reports, 2018, 38, .	1.1	27
23	RusA proteins from the extreme thermophile Aquifex aeolicus and lactococcal phage ϕ t resolve Holliday junctions. Molecular Microbiology, 2002, 44, 549-559.	1.2	24
24	The Structure of Escherichia coli RusA Endonuclease Reveals a New Holliday Junction DNA Binding Fold. Structure, 2003, 11, 1557-1567.	1.6	24
25	Holliday Junction Binding and Resolution by the Rap Structure-specific Endonuclease of Phage ϕ t. Journal of Molecular Biology, 2004, 340, 739-751.	2.0	24
26	Archaeal Hel308 Domain V Couples DNA Binding to ATP Hydrolysis and Positions DNA for Unwinding Over the Helicase Ratchet. Journal of Molecular Biology, 2007, 374, 1139-1144.	2.0	21
27	Physical interaction between archaeal DNA repair helicase Hel308 and Replication Protein A (RPA). DNA Repair, 2011, 10, 306-313.	1.3	21
28	Cas3 Protein—A Review of a Multi-Tasking Machine. Genes, 2020, 11, 208.	1.0	21
29	RusA Holliday junction resolvase: DNA complex structure—insights into selectivity and specificity. Nucleic Acids Research, 2006, 34, 5577-5584.	6.5	19
30	Isoaspartate, carbamoyl phosphate synthase-1, and carbonic anhydrase-III as biomarkers of liver injury. Biochemical and Biophysical Research Communications, 2015, 458, 626-631.	1.0	19
31	DNA binding and unwinding by Hel308 helicase requires dual functions of a winged helix domain. DNA Repair, 2017, 57, 125-132.	1.3	16
32	Winged helix domains with unknown function in Hel308 and related helicases. Biochemical Society Transactions, 2011, 39, 140-144.	1.6	15
33	Cas3 stimulates runaway replication of a ColE1 plasmid in Escherichia coli and antagonises RNaseHI. RNA Biology, 2013, 10, 770-778.	1.5	13
34	DNA replication roadblocks caused by Cascade interference complexes are alleviated by RecG DNA repair helicase. RNA Biology, 2019, 16, 543-548.	1.5	13
35	Analysis of conserved basic residues associated with DNA binding (Arg69) and catalysis (Lys76) by the RusA holliday junction resolvase. Journal of Molecular Biology, 2000, 304, 165-176.	2.0	11
36	CRISPR-Cas adaptive immunity and the three Rs. Bioscience Reports, 2017, 37, .	1.1	11

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37	The HelQ human DNA repair helicase utilizes a PWI-like domain for DNA loading through interaction with RPA, triggering DNA unwinding by the HelQ helicase core. <i>NAR Cancer</i> , 2021, 3, zcaa043.	1.6	11
38	Genetic analysis of an archaeal Holliday junction resolvase in <i>Escherichia coli</i> 1 Edited by J. Karn. <i>Journal of Molecular Biology</i> , 2001, 310, 577-589.	2.0	10
39	A Gold Standard, CRISPR/Cas9-Based Complementation Strategy Reliant on 24 Nucleotide Bookmark Sequences. <i>Genes</i> , 2020, 11, 458.	1.0	10
40	Remodeling and Control of Homologous Recombination by DNA Helicases and Translocases that Target Recombinases and Synapsis. <i>Genes</i> , 2016, 7, 52.	1.0	8
41	Mechanistic insights into Lhr helicase function in DNA repair. <i>Biochemical Journal</i> , 2020, 477, 2935-2947.	1.7	7
42	Identification of <i>Escherichia coli</i> ygaQ and rpmG as novel mitomycin C resistance factors implicated in DNA repair. <i>Bioscience Reports</i> , 2016, 36, e00290.	1.1	5
43	Adaptation processes that build CRISPR immunity: creative destruction, updated. <i>Essays in Biochemistry</i> , 2019, 63, 227-235.	2.1	5
44	A Tryptophan "Gate"™ in the CRISPR-Cas3 Nuclease Controls ssDNA Entry into the Nuclease Site, That When Removed Results in Nuclease Hyperactivity. <i>International Journal of Molecular Sciences</i> , 2021, 22, 2848.	1.8	5
45	Helicases that interact with replication forks: new candidates from archaea. <i>Biochemical Society Transactions</i> , 2005, 33, 1471.	1.6	3
46	Homologous recombination in Archaea: new Holliday junction helicases. <i>Biochemical Society Transactions</i> , 2003, 31, 703-705.	1.6	2
47	Integration of diverse DNA substrates by a casposase can be targeted to R-loops <i>in vitro</i> by its fusion to Cas9. <i>Bioscience Reports</i> , 2021, 41, .	1.1	2
48	Crystallization of RusA Holliday junction resolvase from <i>Escherichia coli</i> . <i>Acta Crystallographica Section D: Biological Crystallography</i> , 2003, 59, 2262-2264.	2.5	1
49	Functions of Hel308 helicases in promoting genome stability in metazoans and archaea. <i>FASEB Journal</i> , 2006, 20, LB55.	0.2	0