

Susan T Lovett

List of Publications by Year in descending order

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95
papers

5,619
citations

76326

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82547

72
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112
all docs

112
docs citations

112
times ranked

5309
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|--|------|-----------|
| 1 | DnaA and SspA Regulation of the <i>iraD</i> gene of <i>E. coli</i> : an alternative DNA damage response independent of LexA/RecA. <i>Genetics</i> , 2022, , . | 2.9 | 3 |
| 2 | DNA damage-signaling, homologous recombination and genetic mutation induced by 5-azacytidine and DNA-protein crosslinks in <i>Escherichia coli</i> . <i>Mutation Research - Fundamental and Molecular Mechanisms of Mutagenesis</i> , 2021, 822, 111742. | 1.0 | 3 |
| 3 | The Role of Replication Clamp-Loader Protein HolC of <i>Escherichia coli</i> in Overcoming Replication/Transcription Conflicts. <i>MBio</i> , 2021, 12, . | 4.1 | 6 |
| 4 | Alternative complexes formed by the <i>Escherichia coli</i> clamp loader accessory protein HolC (x) with replication protein HolD (f) and repair protein YoaA. <i>DNA Repair</i> , 2021, 100, 103006. | 2.8 | 7 |
| 5 | DNA polymerase III protein, HolC, helps resolve replication/transcription conflicts. <i>Microbial Cell</i> , 2021, 8, 143-145. | 3.2 | 3 |
| 6 | Genetic Analysis of DinG Family Helicase YoaA and Its Interaction with Replication Clamp Loader Protein HolC in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2021, 203, e0022821. | 2.2 | 2 |
| 7 | Frequent template switching in postreplication gaps: suppression of deleterious consequences by the <i>Escherichia coli</i> Uup and RadD proteins. <i>Nucleic Acids Research</i> , 2020, 48, 212-230. | 14.5 | 12 |
| 8 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Infection and Immunity</i> , 2020, 88, . | 2.2 | 0 |
| 9 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Microbiology Spectrum</i> , 2020, 8, . | 3.0 | 0 |
| 10 | Identifying Small Molecules That Promote Quasipalindrome-Associated Template-Switch Mutations in <i>Escherichia coli</i> . <i>G3: Genes, Genomes, Genetics</i> , 2020, 10, 1809-1815. | 1.8 | 2 |
| 11 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Antimicrobial Agents and Chemotherapy</i> , 2020, 64, . | 3.2 | 0 |
| 12 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Journal of Virology</i> , 2020, 94, . | 3.4 | 0 |
| 13 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Journal of Bacteriology</i> , 2020, 202, . | 2.2 | 0 |
| 14 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Microbiology and Molecular Biology Reviews</i> , 2020, 84, . | 6.6 | 0 |
| 15 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Journal of Microbiology and Biology Education</i> , 2020, 21, . | 1.0 | 2 |
| 16 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>MSystems</i> , 2020, 5, . | 3.8 | 0 |
| 17 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Microbiology Resource Announcements</i> , 2020, 9, . | 0.6 | 0 |
| 18 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>MBio</i> , 2020, 11, . | 4.1 | 3 |

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| 19 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Journal of Clinical Microbiology</i> , 2020, 58, . | 3.9 | 1 |
| 20 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Applied and Environmental Microbiology</i> , 2020, 86, . | 3.1 | 1 |
| 21 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>MSphere</i> , 2020, 5, . | 2.9 | 1 |
| 22 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Molecular and Cellular Biology</i> , 2020, 40, . | 2.3 | 0 |
| 23 | The ASM Journals Committee Values the Contributions of Black Microbiologists. <i>Clinical Microbiology Reviews</i> , 2020, 33, . | 13.6 | 1 |
| 24 | Structure-Activity Relationship of Peptide-Conjugated Chloramphenicol for Inhibiting <i>Escherichia coli</i> . <i>Journal of Medicinal Chemistry</i> , 2019, 62, 10245-10257. | 6.4 | 7 |
| 25 | Stimulation of Replication Template-Switching by DNA-Protein Crosslinks. <i>Genes</i> , 2019, 10, 14. | 2.4 | 6 |
| 26 | Diglycine Enables Rapid Intrabacterial Hydrolysis for Activating Antibiotics against Gram-negative Bacteria. <i>Angewandte Chemie</i> , 2019, 131, 10741-10744. | 2.0 | 7 |
| 27 | Diglycine Enables Rapid Intrabacterial Hydrolysis for Activating Antibiotics against Gram-negative Bacteria. <i>Angewandte Chemie - International Edition</i> , 2019, 58, 10631-10634. | 13.8 | 24 |
| 28 | Between sisters: Watching replication-associated recombinational DNA repair. <i>Journal of Cell Biology</i> , 2018, 217, 2225-2227. | 5.2 | 0 |
| 29 | Template-switching during replication fork repair in bacteria. <i>DNA Repair</i> , 2017, 56, 118-128. | 2.8 | 43 |
| 30 | SSB recruitment of Exonuclease I aborts template-switching in <i>Escherichia coli</i> . <i>DNA Repair</i> , 2017, 57, 12-16. | 2.8 | 7 |
| 31 | Recombinational branch migration by the RadA/Sms paralog of RecA in <i>Escherichia coli</i> . <i>ELife</i> , 2016, 5, . | 6.0 | 44 |
| 32 | Connecting Replication and Repair: YoaA, a Helicase-Related Protein, Promotes Azidothymidine Tolerance through Association with Chi, an Accessory Clamp Loader Protein. <i>PLoS Genetics</i> , 2015, 11, e1005651. | 3.5 | 20 |
| 33 | Genetic analysis of <i>Escherichia coli</i> ...: functional motifs and genetic interactions. <i>Molecular Microbiology</i> , 2015, 95, 769-779. | 2.5 | 49 |
| 34 | Break-Induced DNA Replication. <i>Cold Spring Harbor Perspectives in Biology</i> , 2013, 5, a010397-a010397. | 5.5 | 191 |
| 35 | Azidothymidine and other chain terminators are mutagenic for template-switch-generated genetic mutations. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2012, 109, 6171-6174. | 7.1 | 22 |
| 36 | A glimpse of molecular competition. <i>Nature</i> , 2012, 491, 198-200. | 27.8 | 1 |

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|----|---|------|-----------|
| 37 | β-Galactosidase-instructed formation of molecular nanofibers and a hydrogel. <i>Nanoscale</i> , 2011, 3, 2859. | 5.6 | 34 |
| 38 | Phenotypic Landscape of a Bacterial Cell. <i>Cell</i> , 2011, 144, 143-156. | 28.9 | 623 |
| 39 | The DNA Exonucleases of <i>Escherichia coli</i> . <i>EcoSal Plus</i> , 2011, 4, . | 5.4 | 77 |
| 40 | Toxicity and tolerance mechanisms for azidothymidine, a replication gap-promoting agent, in <i>Escherichia coli</i> . <i>DNA Repair</i> , 2011, 10, 260-270. | 2.8 | 42 |
| 41 | Insights Into Mutagenesis Using <i>Escherichia coli</i> Chromosomal <i>lacZ</i> Strains That Enable Detection of a Wide Spectrum of Mutational Events. <i>Genetics</i> , 2011, 188, 247-262. | 2.9 | 31 |
| 42 | The 2011 Thomas Hunt Morgan Medal: James Haber. <i>Genetics</i> , 2011, 187, 987-989. | 2.9 | 0 |
| 43 | A Role for Nonessential Domain II of Initiator Protein, DnaA, in Replication Control. <i>Genetics</i> , 2009, 183, 39-49. | 2.9 | 25 |
| 44 | Growth Phase and (p)ppGpp Control of IraD, a Regulator of RpoS Stability, in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2009, 191, 7436-7446. | 2.2 | 52 |
| 45 | A DNA damage response in <i>Escherichia coli</i> involving the alternative sigma factor, RpoS. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 611-616. | 7.1 | 79 |
| 46 | The ObgE/CgtA GTPase influences the stringent response to amino acid starvation in <i>Escherichia coli</i> . <i>Molecular Microbiology</i> , 2009, 73, 253-266. | 2.5 | 67 |
| 47 | Cell cycle synchronization of <i>Escherichia coli</i> using the stringent response, with fluorescence labeling assays for DNA content and replication. <i>Methods</i> , 2009, 48, 8-13. | 3.8 | 99 |
| 48 | Reconstitution of initial steps of dsDNA break repair by the RecF pathway of <i>E. coli</i> . <i>Genes and Development</i> , 2009, 23, 1234-1245. | 5.9 | 138 |
| 49 | Mechanisms of Recombination: Lessons from <i>E. coli</i> . <i>Critical Reviews in Biochemistry and Molecular Biology</i> , 2008, 43, 347-370. | 5.2 | 91 |
| 50 | The Stringent Response and Cell Cycle Arrest in <i>Escherichia coli</i> . <i>PLoS Genetics</i> , 2008, 4, e1000300. | 3.5 | 119 |
| 51 | RecA-independent recombination is efficient but limited by exonucleases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2007, 104, 216-221. | 7.1 | 113 |
| 52 | Polymerase Switching in DNA Replication. <i>Molecular Cell</i> , 2007, 27, 523-526. | 9.7 | 23 |
| 53 | Chromosome segregation control by <i>Escherichia coli</i> ObgE GTPase. <i>Molecular Microbiology</i> , 2007, 65, 569-581. | 2.5 | 41 |
| 54 | Replication arrest-stimulated recombination: Dependence on the RecA paralog, RadA/Sms and translesion polymerase, DinB. <i>DNA Repair</i> , 2006, 5, 1421-1427. | 2.8 | 57 |

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| 55 | Cis and Trans-acting Effects on a Mutational Hotspot Involving a Replication Template Switch. <i>Journal of Molecular Biology</i> , 2006, 356, 300-311. | 4.2 | 45 |
| 56 | DNA Repeat Rearrangements Mediated by DnaK-Dependent Replication Fork Repair. <i>Molecular Cell</i> , 2006, 21, 595-604. | 9.7 | 88 |
| 57 | RecJ exonuclease: substrates, products and interaction with SSB. <i>Nucleic Acids Research</i> , 2006, 34, 1084-1091. | 14.5 | 112 |
| 58 | The role of replication initiation control in promoting survival of replication fork damage. <i>Molecular Microbiology</i> , 2006, 60, 229-239. | 2.5 | 43 |
| 59 | Resurrecting a broken genome. <i>Nature</i> , 2006, 443, 517-519. | 27.8 | 28 |
| 60 | A Bacterial G Protein-Mediated Response to Replication Arrest. <i>Molecular Cell</i> , 2005, 17, 549-560. | 9.7 | 77 |
| 61 | Filling the Gaps in Replication Restart Pathways. <i>Molecular Cell</i> , 2005, 17, 751-752. | 9.7 | 23 |
| 62 | Encoded errors: mutations and rearrangements mediated by misalignment at repetitive DNA sequences. <i>Molecular Microbiology</i> , 2004, 52, 1243-1253. | 2.5 | 232 |
| 63 | Connecting Replication and Recombination. <i>Molecular Cell</i> , 2003, 11, 554-556. | 9.7 | 20 |
| 64 | Stabilization of perfect and imperfect tandem repeats by single-strand DNA exonucleases. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2003, 100, 1134-1139. | 7.1 | 48 |
| 65 | Role for radA/sms in Recombination Intermediate Processing in <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2002, 184, 6836-6844. | 2.2 | 103 |
| 66 | Crossing Over Between Regions of Limited Homology in <i>Escherichia coli</i> : RecA-Dependent and RecA-Independent Pathways. <i>Genetics</i> , 2002, 160, 851-859. | 2.9 | 135 |
| 67 | Instability of repetitive DNA sequences: The role of replication in multiple mechanisms. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 8319-8325. | 7.1 | 313 |
| 68 | In vivo requirement for RecJ, ExoVII, ExoI, and ExoX in methyl-directed mismatch repair. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2001, 98, 6765-6770. | 7.1 | 192 |
| 69 | Redundant Exonuclease Involvement in <i>Escherichia coli</i> Methyl-directed Mismatch Repair. <i>Journal of Biological Chemistry</i> , 2001, 276, 31053-31058. | 3.4 | 114 |
| 70 | Evidence for Two Mechanisms of Palindrome-Stimulated Deletion in <i>Escherichia coli</i> : Single-Strand Annealing and Replication Slipped Mismatching. <i>Genetics</i> , 2001, 158, 527-540. | 2.9 | 72 |
| 71 | A Thermostable Single-Strand DNase from <i>Methanococcus jannaschii</i> Related to the RecJ Recombination and Repair Exonuclease from <i>Escherichia coli</i> . <i>Journal of Bacteriology</i> , 2000, 182, 607-612. | 2.2 | 24 |
| 72 | A novel mutational hotspot in a natural quasipalindrome in <i>Escherichia coli</i> . <i>Journal of Molecular Biology</i> , 2000, 302, 553-564. | 4.2 | 65 |

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|----|--|------|-----------|
| 73 | Exonuclease X of Escherichia coli. Journal of Biological Chemistry, 1999, 274, 30094-30100. | 3.4 | 63 |
| 74 | Expansion of DNA repeats in Escherichia coli : effects of recombination and replication functions 1 Edited by J. H. Miller. Journal of Molecular Biology, 1999, 289, 21-27. | 4.2 | 48 |
| 75 | Identification of RNase T as a High-Copy Suppressor of the UV Sensitivity Associated With Single-Strand DNA Exonuclease Deficiency in Escherichia coli. Genetics, 1999, 151, 929-934. | 2.9 | 25 |
| 76 | Tandem Repeat Recombination Induced by Replication Fork Defects in Escherichia coli Requires a Novel Factor, RadC. Genetics, 1999, 152, 5-13. | 2.9 | 60 |
| 77 | Mutational Analysis of the RecJ Exonuclease of Escherichia coli : Identification of Phosphoesterase Motifs. Journal of Bacteriology, 1999, 181, 6098-6102. | 2.2 | 37 |
| 78 | Slipped Misalignment Mechanisms of Deletion Formation: In Vivo Susceptibility to Nucleases. Journal of Bacteriology, 1999, 181, 477-482. | 2.2 | 55 |
| 79 | Slipped misalignment mechanisms of deletion formation: analysis of deletion endpoints. Journal of Molecular Biology, 1998, 276, 559-569. | 4.2 | 35 |
| 80 | Identification of a Potent DNase Activity Associated with RNase T of Escherichia coli. Journal of Biological Chemistry, 1998, 273, 35126-35131. | 3.4 | 30 |
| 81 | Single-Strand DNA-Specific Exonucleases in Escherichia coli: Roles in Repair and Mutation Avoidance. Genetics, 1998, 149, 7-16. | 2.9 | 129 |
| 82 | Crystal structures of Escherichia coli and Salmonella typhimurium 3-isopropylmalate dehydrogenase and comparison with their thermophilic counterpart from Thermus thermophilus. Journal of Molecular Biology, 1997, 266, 1016-1031. | 4.2 | 139 |
| 83 | Purification, catalytic properties and thermostability of 3-isopropylmalate dehydrogenase from Escherichia coli. BBA - Proteins and Proteomics, 1997, 1337, 105-112. | 2.1 | 24 |
| 84 | Enhanced Deletion Formation by Aberrant DNA Replication in Escherichia coli. Genetics, 1997, 146, 457-470. | 2.9 | 99 |
| 85 | Stabilization of diverged tandem repeats by mismatch repair: evidence for deletion formation via a misaligned replication intermediate.. Proceedings of the National Academy of Sciences of the United States of America, 1996, 93, 7120-7124. | 7.1 | 77 |
| 86 | Enhancement of RecA Strand-transfer Activity by the RecJ Exonuclease of Escherichia coli. Journal of Biological Chemistry, 1995, 270, 6881-6885. | 3.4 | 41 |
| 87 | Revision of the amino-acid sequence of 3-isopropylmalate dehydrogenase from Salmonella typhimurium by means of X-ray crystallography. Gene, 1995, 164, 85-87. | 2.2 | 6 |
| 88 | Recombination between repeats in Escherichia coli by a recA-independent, proximity-sensitive mechanism. Molecular Genetics and Genomics, 1994, 245, 294-300. | 2.4 | 88 |
| 89 | Release of 5'-terminal deoxyribose-phosphate residues from incised abasic sites in DNA by the Escherichia coli RecJ protein. Nucleic Acids Research, 1994, 22, 993-998. | 14.5 | 104 |
| 90 | Sequence of the RAD55 gene of Saccharomyces cerevisiae: similarity of RAD55 to prokaryotic RecA and other RecA-like proteins. Gene, 1994, 142, 103-106. | 2.2 | 104 |

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| 91 | Two related recombinases are required for site-specific recombination at dif and cer in <i>E. coli</i> K12. <i>Cell</i> , 1993, 75, 351-361. | 28.9 | 324 |
| 92 | Characterization of Null Mutants of the <i>rad55</i> Gene of <i>Saccharomyces cerevisiae</i> : Effects of Temperature, Osmotic Strength and Mating Type. <i>Genetics</i> , 1987, 116, 547-553. | 2.9 | 113 |
| 93 | Genetic Analysis of Regulation of the RecF Pathway of Recombination in <i>Escherichia coli</i> K-12. <i>Journal of Bacteriology</i> , 1983, 153, 1471-1478. | 2.2 | 81 |
| 94 | The DNA Damage Response. , 0, , 205-228. | | 12 |
| 95 | Misalignment-Mediated Mutations and Genetic Rearrangements at Repetitive DNA Sequences. , 0, , 449-464. | | 0 |