Erik J Sontheimer

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84 11,798 38 103 h-index g-index citations papers 6.85 13,538 21.1 103 L-index ext. citations avg, IF ext. papers

| # | Paper | IF | Citations |
|----|---|-------------------|-----------|
| 84 | Origins and Mechanisms of miRNAs and siRNAs. <i>Cell</i> , 2009 , 136, 642-55 | 56.2 | 3659 |
| 83 | CRISPR interference limits horizontal gene transfer in staphylococci by targeting DNA. <i>Science</i> , 2008 , 322, 1843-5 | 33.3 | 1181 |
| 82 | Distinct roles for Drosophila Dicer-1 and Dicer-2 in the siRNA/miRNA silencing pathways. <i>Cell</i> , 2004 , 117, 69-81 | 56.2 | 1016 |
| 81 | CRISPR interference: RNA-directed adaptive immunity in bacteria and archaea. <i>Nature Reviews Genetics</i> , 2010 , 11, 181-90 | 30.1 | 711 |
| 80 | Efficient genome engineering in human pluripotent stem cells using Cas9 from Neisseria meningitidis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013 , 110, 15644-9 | 11.5 | 508 |
| 79 | Self versus non-self discrimination during CRISPR RNA-directed immunity. <i>Nature</i> , 2010 , 463, 568-71 | 50.4 | 444 |
| 78 | A Dicer-2-dependent 80s complex cleaves targeted mRNAs during RNAi in Drosophila. <i>Cell</i> , 2004 , 117, 83-94 | 56.2 | 348 |
| 77 | Assembly and function of RNA silencing complexes. <i>Nature Reviews Molecular Cell Biology</i> , 2005 , 6, 127- | -3488.7 | 330 |
| 76 | Silencing by small RNAs is linked to endosomal trafficking. <i>Nature Cell Biology</i> , 2009 , 11, 1150-6 | 23.4 | 279 |
| 75 | Naturally Occurring Off-Switches for CRISPR-Cas9. <i>Cell</i> , 2016 , 167, 1829-1838.e9 | 56.2 | 260 |
| 74 | Silence from within: endogenous siRNAs and miRNAs. <i>Cell</i> , 2005 , 122, 9-12 | 56.2 | 236 |
| 73 | Adenovirus-Mediated Somatic Genome Editing of Pten by CRISPR/Cas9 in Mouse Liver in Spite of Cas9-Specific Immune Responses. <i>Human Gene Therapy</i> , 2015 , 26, 432-42 | 4.8 | 226 |
| 72 | Processing-independent CRISPR RNAs limit natural transformation in Neisseria meningitidis. <i>Molecular Cell</i> , 2013 , 50, 488-503 | 17.6 | 206 |
| 71 | Metal ion catalysis during splicing of premessenger RNA. <i>Nature</i> , 1997 , 388, 801-5 | 50.4 | 155 |
| 70 | A Broad-Spectrum Inhibitor of CRISPR-Cas9. <i>Cell</i> , 2017 , 170, 1224-1233.e15 | 56.2 | 145 |
| 69 | Autoantibodies against a serine tRNA-protein complex implicated in cotranslational selenocysteine insertion. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1992 , 89, 973 | 9 ¹ 43 | 126 |
| 68 | An inside job for siRNAs. <i>Molecular Cell</i> , 2008 , 31, 309-12 | 17.6 | 108 |

(2004-2017)

| 67 | CRISPR/Cas9-mediated genome editing induces exon skipping by alternative splicing or exon deletion. <i>Genome Biology</i> , 2017 , 18, 108 | 18.3 | 103 |
|----|---|------|-----|
| 66 | Metal ion catalysis during the exon-ligation step of nuclear pre-mRNA splicing: extending the parallels between the spliceosome and group II introns. <i>Rna</i> , 2000 , 6, 199-205 | 5.8 | 98 |
| 65 | A role for ubiquitin in the spliceosome assembly pathway. <i>Nature Structural and Molecular Biology</i> , 2008 , 15, 444-51 | 17.6 | 93 |
| 64 | RNAi: RISC gets loaded. <i>Cell</i> , 2005 , 123, 543-5 | 56.2 | 87 |
| 63 | A Compact, High-Accuracy Cas9 with a Dinucleotide PAM for InIVivo Genome Editing. <i>Molecular Cell</i> , 2019 , 73, 714-726.e4 | 17.6 | 85 |
| 62 | Kinetic characterization of the second step of group II intron splicing: role of metal ions and the cleavage site 2FOH in catalysis. <i>Biochemistry</i> , 2000 , 39, 12939-52 | 3.2 | 72 |
| 61 | C-BERST: defining subnuclear proteomic landscapes at genomic elements with dCas9-APEX2. <i>Nature Methods</i> , 2018 , 15, 433-436 | 21.6 | 67 |
| 60 | Site-specific RNA crosslinking with 4-thiouridine. <i>Molecular Biology Reports</i> , 1994 , 20, 35-44 | 2.8 | 64 |
| 59 | Ubiquitin binding by a variant Jab1/MPN domain in the essential pre-mRNA splicing factor Prp8p. <i>Rna</i> , 2006 , 12, 292-302 | 5.8 | 63 |
| 58 | NmeCas9 is an intrinsically high-fidelity genome-editing platform. <i>Genome Biology</i> , 2018 , 19, 214 | 18.3 | 60 |
| 57 | Molecular requirements for RNA-induced silencing complex assembly in the Drosophila RNA interference pathway. <i>Journal of Biological Chemistry</i> , 2005 , 280, 39278-83 | 5.4 | 58 |
| 56 | All-in-one adeno-associated virus delivery and genome editing by Neisseria meningitidis Cas9 in vivo. <i>Genome Biology</i> , 2018 , 19, 137 | 18.3 | 58 |
| 55 | Type II-C CRISPR-Cas9 Biology, Mechanism, and Application. ACS Chemical Biology, 2018, 13, 357-365 | 4.9 | 57 |
| 54 | The Bacterial Origins of the CRISPR Genome-Editing Revolution. Human Gene Therapy, 2015 , 26, 413-24 | 4.8 | 56 |
| 53 | Short interfering RNA strand selection is independent of dsRNA processing polarity during RNAi in Drosophila. <i>Current Biology</i> , 2006 , 16, 530-5 | 6.3 | 53 |
| 52 | An engineered ScCas9 with broad PAM range and high specificity and activity. <i>Nature Biotechnology</i> , 2020 , 38, 1154-1158 | 44.5 | 51 |
| 51 | Potent Cas9 Inhibition in Bacterial and Human Cells by AcrIIC4 and AcrIIC5 Anti-CRISPR Proteins. <i>MBio</i> , 2018 , 9, | 7.8 | 51 |
| 50 | Molecular biology. Argonaute journeys into the heart of RISC. <i>Science</i> , 2004 , 305, 1409-10 | 33.3 | 46 |

| 49 | DNase H Activity of Neisseria meningitidis Cas9. <i>Molecular Cell</i> , 2015 , 60, 242-55 | 17.6 | 45 |
|----|--|---------------|----|
| 48 | Improved prime editors enable pathogenic allele correction and cancer modelling in adult mice. Nature Communications, 2021, 12, 2121 | 17.4 | 45 |
| 47 | Heavily and fully modified RNAs guide efficient SpyCas9-mediated genome editing. <i>Nature Communications</i> , 2018 , 9, 2641 | 17.4 | 44 |
| 46 | A Cas9 with PAM recognition for adenine dinucleotides. <i>Nature Communications</i> , 2020 , 11, 2474 | 17.4 | 38 |
| 45 | Anti-CRISPRs: Protein Inhibitors of CRISPR-Cas Systems. <i>Annual Review of Biochemistry</i> , 2020 , 89, 309-3 | 32 9.1 | 37 |
| 44 | Tissue-restricted genome editing in vivo specified by microRNA-repressible anti-CRISPR proteins. <i>Rna</i> , 2019 , 25, 1421-1431 | 5.8 | 35 |
| 43 | Structures of Neisseria meningitidis Cas9 Complexes in Catalytically Poised and Anti-CRISPR-Inhibited States. <i>Molecular Cell</i> , 2019 , 76, 938-952.e5 | 17.6 | 35 |
| 42 | Inhibition of CRISPR-Cas9 ribonucleoprotein complex assembly by anti-CRISPR AcrIIC2. <i>Nature Communications</i> , 2019 , 10, 2806 | 17.4 | 30 |
| 41 | Blanks, a nuclear siRNA/dsRNA-binding complex component, is required for Drosophila spermiogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2011 , 108, 3204-9 | 11.5 | 24 |
| 40 | Inhibition of CRISPR-Cas systems by mobile genetic elements. <i>Current Opinion in Microbiology</i> , 2017 , 37, 120-127 | 7.9 | 23 |
| 39 | The NIH Somatic Cell Genome Editing program. <i>Nature</i> , 2021 , 592, 195-204 | 50.4 | 21 |
| 38 | Anti-CRISPR AcrilA5 Potently inhibits All Cas9 Homologs Used for Genome Editing. <i>Cell Reports</i> , 2019 , 29, 1739-1746.e5 | 10.6 | 20 |
| 37 | Invasive DNA, chopped and in the CRISPR. Structure, 2009, 17, 786-8 | 5.2 | 20 |
| 36 | Orthogonal Cas9-Cas9 chimeras provide a versatile platform for genome editing. <i>Nature Communications</i> , 2018 , 9, 4856 | 17.4 | 19 |
| 35 | ATP modulates siRNA interactions with an endogenous human Dicer complex. <i>Rna</i> , 2005 , 11, 1719-24 | 5.8 | 18 |
| 34 | A Hyperthermophilic Phage Decoration Protein Suggests Common Evolutionary Origin with Herpesvirus Triplex Proteins and an Anti-CRISPR Protein. <i>Structure</i> , 2018 , 26, 936-947.e3 | 5.2 | 16 |
| 33 | Meiosis-induced alterations in transcript architecture and noncoding RNA expression in S. cerevisiae. <i>Rna</i> , 2012 , 18, 1142-53 | 5.8 | 15 |
| 32 | Thermodynamic and structural characterization of 2Fnitrogen-modified RNA duplexes. <i>Nucleic Acids Research</i> , 2004 , 32, 3446-55 | 20.1 | 14 |

(2019-2015)

| 31 | Primary processing of CRISPR RNA by the endonuclease Cas6 in Staphylococcus epidermidis. <i>FEBS Letters</i> , 2015 , 589, 3197-204 | 3.8 | 10 |
|----|---|------|----|
| 30 | "siRNAs and miRNAs": a meeting report on RNA silencing. <i>Rna</i> , 2004 , 10, 1165-73 | 5.8 | 10 |
| 29 | 5? Modifications Improve Potency and Efficacy of DNA Donors for Precision Genome Editing | | 8 |
| 28 | Efficient Homology-directed Repair with Circular ssDNA Donors | | 7 |
| 27 | Proteomics identification of Drosophila small interfering RNA-associated factors. <i>Molecular and Cellular Proteomics</i> , 2010 , 9, 1866-72 | 7.6 | 6 |
| 26 | R2D2 leads the silencing trigger to mRNATs death star. <i>Cell</i> , 2003 , 115, 132-3 | 56.2 | 6 |
| 25 | YAP1 Withdrawal in Hepatoblastoma Drives Therapeutic Differentiation of Tumor Cells to Functional Hepatocyte-Like Cells. <i>Hepatology</i> , 2021 , 73, 1011-1027 | 11.2 | 6 |
| 24 | SPO24 is a transcriptionally dynamic, small ORF-encoding locus required for efficient sporulation in Saccharomyces cerevisiae. <i>PLoS ONE</i> , 2014 , 9, e105058 | 3.7 | 5 |
| 23 | Bridging sulfur substitutions in the analysis of pre-mRNA splicing. <i>Methods</i> , 1999 , 18, 29-37 | 4.6 | 5 |
| 22 | Self-inactivating, all-in-one AAV vectors for precision Cas9 genome editing via homology-directed repair in vivo. <i>Nature Communications</i> , 2021 , 12, 6267 | 17.4 | 5 |
| 21 | CRISPR-enhanced human adipocyte browning as cell therapy for metabolic disease. <i>Nature Communications</i> , 2021 , 12, 6931 | 17.4 | 4 |
| 20 | 5TModifications improve potency and efficacy of DNA donors for precision genome editing. <i>ELife</i> , 2021 , 10, | 8.9 | 4 |
| 19 | NmeCas9 is an intrinsically high-fidelity genome editing platform | | 4 |
| 18 | Structural biology. Cascading into focus. <i>Science</i> , 2014 , 345, 1452-3 | 33.3 | 3 |
| 17 | Cas9 gets a classmate. <i>Nature Biotechnology</i> , 2015 , 33, 1240-1241 | 44.5 | 3 |
| 16 | Separation of Drosophila RNA silencing complexes by native gel electrophoresis. <i>Methods in Molecular Biology</i> , 2005 , 309, 11-6 | 1.4 | 2 |
| 15 | Adapting dCas9-APEX2 for subnuclear proteomic profiling. <i>Methods in Enzymology</i> , 2019 , 616, 365-383 | 1.7 | 1 |
| 14 | X-Tracting a New CRISPR-Cas Genome-Editing Platform from Metagenomic Data Sets. <i>CRISPR Journal</i> , 2019 , 2, 148-150 | 2.5 | 1 |

| 13 | Small RNAs of opposite signIbut same absolute value. <i>Cell</i> , 2012 , 151, 1157-8 | 56.2 | 1 |
|----|---|----------|---|
| 12 | Genome-wide detection of CRISPR editing in vivo using GUIDE-tag <i>Nature Communications</i> , 2022 , 13, 437 | 17.4 | 1 |
| 11 | Heavily and Fully Modified RNAs Guide Efficient SpyCas9-Mediated Genome Editing | | 1 |
| 10 | One Anti-CRISPR to Rule Them All: Potent Inhibition of Cas9 Homologs Used for Genome Editing. SSRN Electronic Journal, | 1 | 1 |
| 9 | Orthogonal CRISPR-Cas genome editing and efficient inhibition with anti-CRISPRs in zebrafish embryos | 3 | 1 |
| 8 | Potent Cas9 inhibition in bacterial and human cells by new anti-CRISPR protein families | | 1 |
| 7 | Tissue-specific Genome Editing in vivo by MicroRNA-repressible Anti-CRISPR Proteins | | 1 |
| 6 | Shutting down RNA-targeting CRISPR. <i>Science</i> , 2020 , 369, 31-32 | 33.3 | Ο |
| 5 | Quit stalling or you T l be silenced. <i>Cell</i> , 2013 , 152, 938-9 | 56.2 | 0 |
| 4 | CRISPR Shields: Fending Off Diverse Cas Nucleases with Nucleus-like Structures. <i>Molecular Cell</i> , 2020 , 77, 934-936 | 17.6 | |
| 3 | CRISPRs from scratch. <i>Nature Microbiology</i> , 2018 , 3, 261-262 | 26.6 | |
| 2 | RNA. CRISPR goes retro. <i>Science</i> , 2016 , 351, 920-1 | 33.3 | |
| 1 | Accelerating expansion. <i>Rna</i> , 2015 , 21, 510 | 5.8 | |