

# Erik J Sontheimer

## List of Publications by Year in Descending Order

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The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

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

| #  | Paper  | IF   | Citations |
|----|--|------|-----------|
| 84 | Genome-wide detection of CRISPR editing in vivo using GUIDE-tag.. <i>Nature Communications</i> , <b>2022</b> , 13, 437   | 17.4 | 1         |
| 83 | CRISPR-enhanced human adipocyte browning as cell therapy for metabolic disease. <i>Nature Communications</i> , <b>2021</b> , 12, 6931                                    | 17.4 | 4         |
| 82 | 5TModifications improve potency and efficacy of DNA donors for precision genome editing. <i>ELife</i> , <b>2021</b> , 10,  | 8.9  | 4         |
| 81 | Self-inactivating, all-in-one AAV vectors for precision Cas9 genome editing via homology-directed repair in vivo. <i>Nature Communications</i> , <b>2021</b> , 12, 6267  | 17.4 | 5         |
| 80 | Improved prime editors enable pathogenic allele correction and cancer modelling in adult mice. <i>Nature Communications</i> , <b>2021</b> , 12, 2121                     | 17.4 | 45        |
| 79 | The NIH Somatic Cell Genome Editing program. <i>Nature</i> , <b>2021</b> , 592, 195-204  | 50.4 | 21        |
| 78 | YAP1 Withdrawal in Hepatoblastoma Drives Therapeutic Differentiation of Tumor Cells to Functional Hepatocyte-Like Cells. <i>Hepatology</i> , <b>2021</b> , 73, 1011-1027 | 11.2 | 6         |
| 77 | A Cas9 with PAM recognition for adenine dinucleotides. <i>Nature Communications</i> , <b>2020</b> , 11, 2474   | 17.4 | 38        |
| 76 | An engineered ScCas9 with broad PAM range and high specificity and activity. <i>Nature Biotechnology</i> , <b>2020</b> , 38, 1154-1158                                   | 44.5 | 51        |
| 75 | Anti-CRISPRs: Protein Inhibitors of CRISPR-Cas Systems. <i>Annual Review of Biochemistry</i> , <b>2020</b> , 89, 309-332   | 29.1 | 37        |
| 74 | CRISPR Shields: Fending Off Diverse Cas Nucleases with Nucleus-like Structures. <i>Molecular Cell</i> , <b>2020</b> , 77, 934-936  | 17.6 |           |
| 73 | Shutting down RNA-targeting CRISPR. <i>Science</i> , <b>2020</b> , 369, 31-32  | 33.3 | 0         |
| 72 | Tissue-restricted genome editing in vivo specified by microRNA-repressible anti-CRISPR proteins. <i>Rna</i> , <b>2019</b> , 25, 1421-1431                                | 5.8  | 35        |
| 71 | Adapting dCas9-APEX2 for subnuclear proteomic profiling. <i>Methods in Enzymology</i> , <b>2019</b> , 616, 365-383   | 1.7  | 1         |
| 70 | Inhibition of CRISPR-Cas9 ribonucleoprotein complex assembly by anti-CRISPR AcrIIC2. <i>Nature Communications</i> , <b>2019</b> , 10, 2806                               | 17.4 | 30        |
| 69 | X-Tracting a New CRISPR-Cas Genome-Editing Platform from Metagenomic Data Sets. <i>CRISPR Journal</i> , <b>2019</b> , 2, 148-150   | 2.5  | 1         |
| 68 | Structures of Neisseria meningitidis Cas9 Complexes in Catalytically Poised and Anti-CRISPR-Inhibited States. <i>Molecular Cell</i> , <b>2019</b> , 76, 938-952.e5       | 17.6 | 35        |

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| 67 | Anti-CRISPR AcrIIA5 Potently Inhibits All Cas9 Homologs Used for Genome Editing. <i>Cell Reports</i> , <b>2019</b> , 29, 1739-1746.e5  | 10.6 | 20  |
| 66 | A Compact, High-Accuracy Cas9 with a Dinucleotide PAM for In Vivo Genome Editing. <i>Molecular Cell</i> , <b>2019</b> , 73, 714-726.e4   | 17.6 | 85  |
| 65 | CRISPRs from scratch. <i>Nature Microbiology</i> , <b>2018</b> , 3, 261-262  | 26.6 |     |
| 64 | Heavily and fully modified RNAs guide efficient SpyCas9-mediated genome editing. <i>Nature Communications</i> , <b>2018</b> , 9, 2641  | 17.4 | 44  |
| 63 | C-BERST: defining subnuclear proteomic landscapes at genomic elements with dCas9-APEX2. <i>Nature Methods</i> , <b>2018</b> , 15, 433-436  | 21.6 | 67  |
| 62 | Type II-C CRISPR-Cas9 Biology, Mechanism, and Application. <i>ACS Chemical Biology</i> , <b>2018</b> , 13, 357-365   | 4.9  | 57  |
| 61 | Orthogonal Cas9-Cas9 chimeras provide a versatile platform for genome editing. <i>Nature Communications</i> , <b>2018</b> , 9, 4856  | 17.4 | 19  |
| 60 | Potent Cas9 Inhibition in Bacterial and Human Cells by AcrIIC4 and AcrIIC5 Anti-CRISPR Proteins. <i>MBio</i> , <b>2018</b> , 9,  | 7.8  | 51  |
| 59 | NmeCas9 is an intrinsically high-fidelity genome-editing platform. <i>Genome Biology</i> , <b>2018</b> , 19, 214   | 18.3 | 60  |
| 58 | All-in-one adeno-associated virus delivery and genome editing by <i>Neisseria meningitidis</i> Cas9 in vivo. <i>Genome Biology</i> , <b>2018</b> , 19, 137                                     | 18.3 | 58  |
| 57 | A Hyperthermophilic Phage Decoration Protein Suggests Common Evolutionary Origin with Herpesvirus Triplex Proteins and an Anti-CRISPR Protein. <i>Structure</i> , <b>2018</b> , 26, 936-947.e3 | 5.2  | 16  |
| 56 | CRISPR/Cas9-mediated genome editing induces exon skipping by alternative splicing or exon deletion. <i>Genome Biology</i> , <b>2017</b> , 18, 108  | 18.3 | 103 |
| 55 | A Broad-Spectrum Inhibitor of CRISPR-Cas9. <i>Cell</i> , <b>2017</b> , 170, 1224-1233.e15  | 56.2 | 145 |
| 54 | Inhibition of CRISPR-Cas systems by mobile genetic elements. <i>Current Opinion in Microbiology</i> , <b>2017</b> , 37, 120-127  | 7.9  | 23  |
| 53 | RNA. CRISPR goes retro. <i>Science</i> , <b>2016</b> , 351, 920-1  | 33.3 |     |
| 52 | Naturally Occurring Off-Switches for CRISPR-Cas9. <i>Cell</i> , <b>2016</b> , 167, 1829-1838.e9  | 56.2 | 260 |
| 51 | 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> , <b>2015</b> , 26, 432-42              | 4.8  | 226 |
| 50 | The Bacterial Origins of the CRISPR Genome-Editing Revolution. <i>Human Gene Therapy</i> , <b>2015</b> , 26, 413-24  | 4.8  | 56  |

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|----|--|------|------|
| 49 | DNase H Activity of <i>Neisseria meningitidis</i> Cas9. <i>Molecular Cell</i> , <b>2015</b> , 60, 242-55   | 17.6 | 45   |
| 48 | Primary processing of CRISPR RNA by the endonuclease Cas6 in <i>Staphylococcus epidermidis</i> . <i>FEBS Letters</i> , <b>2015</b> , 589, 3197-204   | 3.8  | 10   |
| 47 | Accelerating expansion. <i>Rna</i> , <b>2015</b> , 21, 510   | 5.8  |      |
| 46 | Cas9 gets a classmate. <i>Nature Biotechnology</i> , <b>2015</b> , 33, 1240-1241   | 44.5 | 3    |
| 45 | Structural biology. Cascading into focus. <i>Science</i> , <b>2014</b> , 345, 1452-3   | 33.3 | 3    |
| 44 | SPO24 is a transcriptionally dynamic, small ORF-encoding locus required for efficient sporulation in <i>Saccharomyces cerevisiae</i> . <i>PLoS ONE</i> , <b>2014</b> , 9, e105058  | 3.7  | 5    |
| 43 | Quit stalling or you'll be silenced. <i>Cell</i> , <b>2013</b> , 152, 938-9  | 56.2 | 0    |
| 42 | Processing-independent CRISPR RNAs limit natural transformation in <i>Neisseria meningitidis</i> . <i>Molecular Cell</i> , <b>2013</b> , 50, 488-503   | 17.6 | 206  |
| 41 | Efficient genome engineering in human pluripotent stem cells using Cas9 from <i>Neisseria meningitidis</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2013</b> , 110, 15644-9 | 11.5 | 508  |
| 40 | Small RNAs of opposite sign but same absolute value. <i>Cell</i> , <b>2012</b> , 151, 1157-8   | 56.2 | 1    |
| 39 | Meiosis-induced alterations in transcript architecture and noncoding RNA expression in <i>S. cerevisiae</i> . <i>Rna</i> , <b>2012</b> , 18, 1142-53   | 5.8  | 15   |
| 38 | Blanks, a nuclear siRNA/dsRNA-binding complex component, is required for <i>Drosophila</i> spermiogenesis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , <b>2011</b> , 108, 3204-9    | 11.5 | 24   |
| 37 | Self versus non-self discrimination during CRISPR RNA-directed immunity. <i>Nature</i> , <b>2010</b> , 463, 568-71   | 50.4 | 444  |
| 36 | CRISPR interference: RNA-directed adaptive immunity in bacteria and archaea. <i>Nature Reviews Genetics</i> , <b>2010</b> , 11, 181-90   | 30.1 | 711  |
| 35 | Proteomics identification of <i>Drosophila</i> small interfering RNA-associated factors. <i>Molecular and Cellular Proteomics</i> , <b>2010</b> , 9, 1866-72   | 7.6  | 6    |
| 34 | Invasive DNA, chopped and in the CRISPR. <i>Structure</i> , <b>2009</b> , 17, 786-8  | 5.2  | 20   |
| 33 | Silencing by small RNAs is linked to endosomal trafficking. <i>Nature Cell Biology</i> , <b>2009</b> , 11, 1150-6  | 23.4 | 279  |
| 32 | Origins and Mechanisms of miRNAs and siRNAs. <i>Cell</i> , <b>2009</b> , 136, 642-55   | 56.2 | 3659 |

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|----|---|------|------|
| 31 | A role for ubiquitin in the spliceosome assembly pathway. <i>Nature Structural and Molecular Biology</i> , <b>2008</b> , 15, 444-51   | 17.6 | 93   |
| 30 | CRISPR interference limits horizontal gene transfer in staphylococci by targeting DNA. <i>Science</i> , <b>2008</b> , 322, 1843-5   | 33.3 | 1181 |
| 29 | An inside job for siRNAs. <i>Molecular Cell</i> , <b>2008</b> , 31, 309-12  | 17.6 | 108  |
| 28 | Short interfering RNA strand selection is independent of dsRNA processing polarity during RNAi in <i>Drosophila</i> . <i>Current Biology</i> , <b>2006</b> , 16, 530-5                      | 6.3  | 53   |
| 27 | Ubiquitin binding by a variant Jab1/MPN domain in the essential pre-mRNA splicing factor Prp8p. <i>Rna</i> , <b>2006</b> , 12, 292-302  | 5.8  | 63   |
| 26 | Molecular requirements for RNA-induced silencing complex assembly in the <i>Drosophila</i> RNA interference pathway. <i>Journal of Biological Chemistry</i> , <b>2005</b> , 280, 39278-83   | 5.4  | 58   |
| 25 | Silence from within: endogenous siRNAs and miRNAs. <i>Cell</i> , <b>2005</b> , 122, 9-12  | 56.2 | 236  |
| 24 | RNAi: RISC gets loaded. <i>Cell</i> , <b>2005</b> , 123, 543-5  | 56.2 | 87   |
| 23 | Assembly and function of RNA silencing complexes. <i>Nature Reviews Molecular Cell Biology</i> , <b>2005</b> , 6, 127-38  | 38.7 | 330  |
| 22 | Separation of <i>Drosophila</i> RNA silencing complexes by native gel electrophoresis. <i>Methods in Molecular Biology</i> , <b>2005</b> , 309, 11-6  | 1.4  | 2    |
| 21 | ATP modulates siRNA interactions with an endogenous human Dicer complex. <i>Rna</i> , <b>2005</b> , 11, 1719-24   | 5.8  | 18   |
| 20 | "siRNAs and miRNAs": a meeting report on RNA silencing. <i>Rna</i> , <b>2004</b> , 10, 1165-73  | 5.8  | 10   |
| 19 | Molecular biology. Argonaute journeys into the heart of RISC. <i>Science</i> , <b>2004</b> , 305, 1409-10   | 33.3 | 46   |
| 18 | Thermodynamic and structural characterization of 2'-nitrogen-modified RNA duplexes. <i>Nucleic Acids Research</i> , <b>2004</b> , 32, 3446-55   | 20.1 | 14   |
| 17 | A Dicer-2-dependent 80s complex cleaves targeted mRNAs during RNAi in <i>Drosophila</i> . <i>Cell</i> , <b>2004</b> , 117, 83-94  | 56.2 | 348  |
| 16 | Distinct roles for <i>Drosophila</i> Dicer-1 and Dicer-2 in the siRNA/miRNA silencing pathways. <i>Cell</i> , <b>2004</b> , 117, 69-81  | 56.2 | 1016 |
| 15 | R2D2 leads the silencing trigger to mRNA's death star. <i>Cell</i> , <b>2003</b> , 115, 132-3   | 56.2 | 6    |
| 14 | 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> , <b>2000</b> , 6, 199-205 | 5.8  | 98   |

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|----|---|------|-----|
| 13 | 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> , <b>2000</b> , 39, 12939-52   | 3.2  | 72  |
| 12 | Bridging sulfur substitutions in the analysis of pre-mRNA splicing. <i>Methods</i> , <b>1999</b> , 18, 29-37  | 4.6  | 5   |
| 11 | Metal ion catalysis during splicing of premessenger RNA. <i>Nature</i> , <b>1997</b> , 388, 801-5   | 50.4 | 155 |
| 10 | Site-specific RNA crosslinking with 4-thiouridine. <i>Molecular Biology Reports</i> , <b>1994</b> , 20, 35-44   | 2.8  | 64  |
| 9  | 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> , <b>1992</b> , 89, 9739-43 | 11.5 | 126 |
| 8  | Heavily and Fully Modified RNAs Guide Efficient SpyCas9-Mediated Genome Editing   |      | 1   |
| 7  | One Anti-CRISPR to Rule Them All: Potent Inhibition of Cas9 Homologs Used for Genome Editing. <i>SSRN Electronic Journal</i> ,  | 1    | 1   |
| 6  | NmeCas9 is an intrinsically high-fidelity genome editing platform   |      | 4   |
| 5  | Orthogonal CRISPR-Cas genome editing and efficient inhibition with anti-CRISPRs in zebrafish embryos  |      | 1   |
| 4  | Potent Cas9 inhibition in bacterial and human cells by new anti-CRISPR protein families   |      | 1   |
| 3  | 5' Modifications Improve Potency and Efficacy of DNA Donors for Precision Genome Editing  |      | 8   |
| 2  | Tissue-specific Genome Editing in vivo by MicroRNA-repressible Anti-CRISPR Proteins   |      | 1   |
| 1  | Efficient Homology-directed Repair with Circular ssDNA Donors   |      | 7   |