

# Adrian Francis Stewart

## List of Publications by Year in descending order

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76  
papers

9,998  
citations

81900

39  
h-index

74163

75  
g-index

86  
all docs

86  
docs citations

86  
times ranked

15600  
citing authors

#	ARTICLE	IF	CITATIONS
1	A conditional knockout resource for the genome-wide study of mouse gene function. <i>Nature</i> , 2011, 474, 337-342.	27.8	1,488
2	A new logic for DNA engineering using recombination in <i>Escherichia coli</i> . <i>Nature Genetics</i> , 1998, 20, 123-128.	21.4	1,123
3	High-efficiency deleter mice show that FLPe is an alternative to Cre-loxP. <i>Nature Genetics</i> , 2000, 25, 139-140.	21.4	1,073
4	Heterodimerization of the <i>Drosophila</i> ecdysone receptor with retinoid X receptor and ultraspiracle. <i>Nature</i> , 1993, 362, 471-475.	27.8	512
5	Rapid modification of bacterial artificial chromosomes by ET- recombination. <i>Nucleic Acids Research</i> , 1999, 27, 1555-1557.	14.5	475
6	DNA cloning by homologous recombination in <i>Escherichia coli</i> . <i>Nature Biotechnology</i> , 2000, 18, 1314-1317.	17.5	376
7	Full-length RecE enhances linear-linear homologous recombination and facilitates direct cloning for bioprospecting. <i>Nature Biotechnology</i> , 2012, 30, 440-446.	17.5	375
8	Dre recombinase, like Cre, is a highly efficient site-specific recombinase in <i>E. coli</i> , mammalian cells and mice. <i>DMM Disease Models and Mechanisms</i> , 2009, 2, 508-515.	2.4	254
9	Multiple epigenetic maintenance factors implicated by the loss of Mll2 in mouse development. <i>Development (Cambridge)</i> , 2006, 133, 1423-1432.	2.5	245
10	Mll2 is required for H3K4 trimethylation on bivalent promoters in embryonic stem cells, whereas Mll1 is redundant. <i>Development (Cambridge)</i> , 2014, 141, 526-537.	2.5	225
11	Mll2 Is Required in Oocytes for Bulk Histone 3 Lysine 4 Trimethylation and Transcriptional Silencing. <i>PLoS Biology</i> , 2010, 8, e1000453.	5.6	220
12	A reliable lacZ expression reporter cassette for multipurpose, knockout-first alleles. <i>Genesis</i> , 2004, 38, 151-158.	1.6	186
13	Temporally and spatially regulated somatic mutagenesis in mice. <i>Nucleic Acids Research</i> , 1998, 26, 1427-1432.	14.5	173
14	The H3K4 methyltransferase Setd1a is first required at the epiblast stage, whereas Setd1b becomes essential after gastrulation. <i>Development (Cambridge)</i> , 2014, 141, 1022-1035.	2.5	166
15	The histone 3 lysine 4 methyltransferase, Mll2, is only required briefly in development and spermatogenesis. <i>Epigenetics and Chromatin</i> , 2009, 2, 5.	3.9	154
16	Efficient FLP recombination in mouse ES cells and oocytes. <i>Genesis</i> , 2001, 31, 6-10.	1.6	151
17	Protein Interactions within the Set1 Complex and Their Roles in the Regulation of Histone 3 Lysine 4 Methylation. <i>Journal of Biological Chemistry</i> , 2006, 281, 35404-35412.	3.4	142
18	RecET direct cloning and Red $\beta$ recombineering of biosynthetic gene clusters, large operons or single genes for heterologous expression. <i>Nature Protocols</i> , 2016, 11, 1175-1190.	12.0	132

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19	An improved Flp deleter mouse in C57Bl/6 based on Flpo recombinase. <i>Genesis</i> , 2010, 48, 512-520.	1.6	128
20	MLL2 conveys transcription-independent H3K4 trimethylation in oocytes. <i>Nature Structural and Molecular Biology</i> , 2018, 25, 73-82.	8.2	127
21	Neuronal Kmt2a/Mll1 Histone Methyltransferase Is Essential for Prefrontal Synaptic Plasticity and Working Memory. <i>Journal of Neuroscience</i> , 2015, 35, 5097-5108.	3.6	126
22	Histone-Methyltransferase MLL2 (KMT2B) Is Required for Memory Formation in Mice. <i>Journal of Neuroscience</i> , 2013, 33, 3452-3464.	3.6	121
23	Mutation of cancer driver <i>MLL2</i> results in transcription stress and genome instability. <i>Genes and Development</i> , 2016, 30, 408-420.	5.9	112
24	ExoCET: exonuclease in vitro assembly combined with RecET recombination for highly efficient direct DNA cloning from complex genomes. <i>Nucleic Acids Research</i> , 2018, 46, e28-e28.	14.5	96
25	Synthetic CpG islands reveal DNA sequence determinants of chromatin structure. <i>ELife</i> , 2014, 3, e03397.	6.0	95
26	The histone demethylase UTX regulates stem cell migration and hematopoiesis. <i>Blood</i> , 2013, 121, 2462-2473.	1.4	93
27	High Conservation of the Set1/Rad6 Axis of Histone 3 Lysine 4 Methylation in Budding and Fission Yeasts. <i>Journal of Biological Chemistry</i> , 2003, 278, 8487-8493.	3.4	84
28	The Histone Methyltransferase Wbp7 Controls Macrophage Function through GPI Glycolipid Anchor Synthesis. <i>Immunity</i> , 2012, 36, 572-585.	14.3	79
29	KMT2A and KMT2B Mediate Memory Function by Affecting Distinct Genomic Regions. <i>Cell Reports</i> , 2017, 20, 538-548.	6.4	77
30	A Recombineering Pipeline to Make Conditional Targeting Constructs. <i>Methods in Enzymology</i> , 2010, 477, 125-144.	1.0	75
31	Conformational Adaptability of Red $\beta$ during DNA Annealing and Implications for Its Structural Relationship with Rad52. <i>Journal of Molecular Biology</i> , 2009, 391, 586-598.	4.2	73
32	MLL2, Not MLL1, Plays a Major Role in Sustaining MLL-Rearranged Acute Myeloid Leukemia. <i>Cancer Cell</i> , 2017, 31, 755-770.e6.	16.8	72
33	RAP1-mediated MEK/ERK pathway defects in Kabuki syndrome. <i>Journal of Clinical Investigation</i> , 2015, 125, 3585-3599.	8.2	69
34	A Practical Summary of Site-Specific Recombination, Conditional Mutagenesis, and Tamoxifen Induction of CreERT2. <i>Methods in Enzymology</i> , 2010, 477, 109-123.	1.0	53
35	Regular Nanoscale Protein Patterns via Directed Adsorption through Self-Assembled DNA Origami Masks. <i>ACS Applied Materials &amp; Interfaces</i> , 2016, 8, 31239-31247.	8.0	52
36	An Engineered Virus Library as a Resource for the Spectrum-wide Exploration of Virus and Vector Diversity. <i>Cell Reports</i> , 2017, 19, 1698-1709.	6.4	49

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37	Heterologous Production and Yield Improvement of Epothilones in Burkholderiales Strain DSM 7029. <i>ACS Chemical Biology</i> , 2017, 12, 1805-1812.	3.4	48
38	Expressing cytotoxic compounds in Escherichia coli Nissle 1917 for tumor-targeting therapy. <i>Research in Microbiology</i> , 2019, 170, 74-79.	2.1	48
39	SETD1A protects HSCs from activation-induced functional decline in vivo. <i>Blood</i> , 2018, 131, 1311-1324.	1.4	47
40	The histone 3 lysine 4 methyltransferase Setd1b is a maternal effect gene required for the oogenic gene expression program. <i>Development (Cambridge)</i> , 2017, 144, 2606-2617.	2.5	44
41	Single-Stranded DNA-Binding Protein and Exogenous RecBCD Inhibitors Enhance Phage-Derived Homologous Recombination in Pseudomonas. <i>IScience</i> , 2019, 14, 1-14.	4.1	43
42	Neuronal Deletion of Kmt2a/Mll1 Histone Methyltransferase in Ventral Striatum is Associated with Defective Spike-Timing-Dependent Striatal Synaptic Plasticity, Altered Response to Dopaminergic Drugs, and Increased Anxiety. <i>Neuropsychopharmacology</i> , 2016, 41, 3103-3113.	5.4	40
43	Enhanced Heterologous Spinosad Production from a 79-kb Synthetic Multioperon Assembly. <i>ACS Synthetic Biology</i> , 2019, 8, 137-147.	3.8	39
44	The epigenetic regulator Mll1 is required for Wnt-driven intestinal tumorigenesis and cancer stemness. <i>Nature Communications</i> , 2020, 11, 6422.	12.8	38
45	lncRNA Panct1 Maintains Mouse Embryonic Stem Cell Identity by Regulating TOBF1 Recruitment to Oct-Sox Sequences in Early G1. <i>Cell Reports</i> , 2017, 21, 3012-3021.	6.4	35
46	The H3K4 methyltransferase Setd1b is essential for hematopoietic stem and progenitor cell homeostasis in mice. <i>ELife</i> , 2018, 7, .	6.0	34
47	Recombineering, transfection, Western, IP and ChIP methods for protein tagging via gene targeting or BAC transgenesis. <i>Methods</i> , 2011, 53, 437-452.	3.8	33
48	The Histone Methyltransferase KMT2B Is Required for RNA Polymerase II Association and Protection from DNA Methylation at the <i>MagohB</i> CpG Island Promoter. <i>Molecular and Cellular Biology</i> , 2013, 33, 1383-1393.	2.3	33
49	MLL1 is required for PAX7 expression and satellite cell self-renewal in mice. <i>Nature Communications</i> , 2019, 10, 4256.	12.8	31
50	Cooperative genetic networks drive embryonic stem cell transition from naïve to formative pluripotency. <i>EMBO Journal</i> , 2021, 40, e105776.	7.8	31
51	KMT2B Is Selectively Required for Neuronal Transdifferentiation, and Its Loss Exposes Dystonia Candidate Genes. <i>Cell Reports</i> , 2018, 25, 988-1001.	6.4	28
52	Kmt2b conveys monovalent and bivalent H3K4me3 in mouse spermatogonial stem cells at germline and embryonic promoters. <i>Development (Cambridge)</i> , 2018, 145, .	2.5	26
53	The Set1 complex is dimeric and acts with Jhd2 demethylation to convey symmetrical H3K4 trimethylation. <i>Genes and Development</i> , 2019, 33, 550-564.	5.9	24
54	The contribution of homology arms to nuclease-assisted genome engineering. <i>Nucleic Acids Research</i> , 2017, 45, 8105-8115.	14.5	23

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55	RedEx: a method for seamless DNA insertion and deletion in large multimodular polyketide synthase gene clusters. <i>Nucleic Acids Research</i> , 2020, 48, e130-e130.	14.5	23
56	A Single-Strand Annealing Protein Clamps DNA to Detect and Secure Homology. <i>PLoS Biology</i> , 2015, 13, e1002213.	5.6	22
57	RAC-tagging: Recombineering And Cas9-assisted targeting for protein tagging and conditional analyses. <i>Scientific Reports</i> , 2016, 6, 25529.	3.3	22
58	Deficiency and haploinsufficiency of histone macroH2A1.1 in mice recapitulate hematopoietic defects of human myelodysplastic syndrome. <i>Clinical Epigenetics</i> , 2019, 11, 121.	4.1	21
59	MLL4 is required after implantation whereas MLL3 becomes essential during late gestation. <i>Development (Cambridge)</i> , 2020, 147, .	2.5	18
60	Twenty-Seven Tamoxifen-Inducible iCre-Driver Mouse Strains for Eye and Brain, Including Seventeen Carrying a New Inducible-First Constitutive-Ready Allele. <i>Genetics</i> , 2019, 211, 1155-1177.	2.9	17
61	msVolcano: A flexible web application for visualizing quantitative proteomics data. <i>Proteomics</i> , 2016, 16, 2491-2494.	2.2	16
62	Systems Analyses Reveal Shared and Diverse Attributes of Oct4 Regulation in Pluripotent Cells. <i>Cell Systems</i> , 2015, 1, 141-151.	6.2	15
63	DNA annealing by Red $\beta$ 2 is insufficient for homologous recombination and the additional requirements involve intra- and inter-molecular interactions. <i>Scientific Reports</i> , 2016, 6, 34525.	3.3	15
64	Distinct pathways affected by menin versus MLL1/MLL2 in MLL-rearranged acute myeloid leukemia. <i>Experimental Hematology</i> , 2019, 69, 37-42.	0.4	13
65	Loss of histone methyltransferase SETD1B in oogenesis results in the redistribution of genomic histone 3 lysine 4 trimethylation. <i>Nucleic Acids Research</i> , 2022, 50, 1993-2004.	14.5	13
66	Analysis of combinatorial chemokine receptor expression dynamics using multi-receptor reporter mice. <i>ELife</i> , 0, 11, .	6.0	12
67	The Kw Recombinase, an Integrase from <i>Kluyveromyces Waltii</i> . <i>FEBS Journal</i> , 1997, 248, 903-912.	0.2	11
68	Human adenovirus type 17 from species D transduces endothelial cells and human CD46 is involved in cell entry. <i>Scientific Reports</i> , 2018, 8, 13442.	3.3	10
69	Improved dsDNA recombineering enables versatile multiplex genome engineering of kilobase-scale sequences in diverse bacteria. <i>Nucleic Acids Research</i> , 2022, 50, e15-e15.	14.5	8
70	Postnatal expression of the lysine methyltransferase SETD1B is essential for learning and the regulation of neuron-enriched genes. <i>EMBO Journal</i> , 2022, 41, e106459.	7.8	7
71	Proteomic navigation using proximity-labeling. <i>Methods</i> , 2019, 164-165, 67-72.	3.8	6
72	Epigenetic modifier balances Mapk and Wnt signalling in differentiation of goblet and Paneth cells. <i>Life Science Alliance</i> , 2022, 5, e202101187.	2.8	6

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73	MLL1 is required for maintenance of intestinal stem cells. PLoS Genetics, 2021, 17, e1009250.	3.5	5
74	New methods for extracting function from the mammalian genome. Methods, 2019, 164-165, 1-2.	3.8	1
75	Protein-Assisted Room-Temperature Assembly of Rigid, Immobile Holliday Junctions and Hierarchical DNA Nanostructures. Molecules, 2020, 25, 5099.	3.8	1
76	The Role of MLL1 and MLL2 in MLL Fusion Oncoprotein-Initiated Leukemia. Blood, 2016, 128, 573-573.	1.4	0