

# Eric U Selker

## List of Publications by Year in descending order

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72  
papers

10,404  
citations

76326

40  
h-index

91884

69  
g-index

111  
all docs

111  
docs citations

111  
times ranked

7400  
citing authors

#	ARTICLE	IF	CITATIONS
1	The genome sequence of the filamentous fungus <i>Neurospora crassa</i> . <i>Nature</i> , 2003, 422, 859-868.	27.8	1,528
2	Sequencing of <i>Aspergillus nidulans</i> and comparative analysis with <i>A. fumigatus</i> and <i>A. oryzae</i> . <i>Nature</i> , 2005, 438, 1105-1115.	27.8	1,250
3	A histone H3 methyltransferase controls DNA methylation in <i>Neurospora crassa</i> . <i>Nature</i> , 2001, 414, 277-283.	27.8	946
4	Premeiotic Instability of Repeated Sequences in <i>Neurospora crassa</i> . <i>Annual Review of Genetics</i> , 1990, 24, 579-613.	7.6	694
5	Lessons from the Genome Sequence of <i>Neurospora crassa</i> : Tracing the Path from Genomic Blueprint to Multicellular Organism. <i>Microbiology and Molecular Biology Reviews</i> , 2004, 68, 1-108.	6.6	572
6	Diverse Pathways Generate MicroRNA-like RNAs and Dicer-Independent Small Interfering RNAs in Fungi. <i>Molecular Cell</i> , 2010, 38, 803-814.	9.7	361
7	Trimethylated lysine 9 of histone H3 is a mark for DNA methylation in <i>Neurospora crassa</i> . <i>Nature Genetics</i> , 2003, 34, 75-79.	21.4	351
8	Structural Basis for the Product Specificity of Histone Lysine Methyltransferases. <i>Molecular Cell</i> , 2003, 12, 177-185.	9.7	307
9	GFP as a tool to analyze the organization, dynamics and function of nuclei and microtubules in <i>Neurospora crassa</i> . <i>Fungal Genetics and Biology</i> , 2004, 41, 897-910.	2.1	306
10	Structure of the <i>Neurospora</i> SET Domain Protein DIM-5, a Histone H3 Lysine Methyltransferase. <i>Cell</i> , 2002, 111, 117-127.	28.9	247
11	The methylated component of the <i>Neurospora crassa</i> genome. <i>Nature</i> , 2003, 422, 893-897.	27.8	214
12	HP1 Is Essential for DNA Methylation in <i>Neurospora</i> . <i>Molecular Cell</i> , 2004, 13, 427-434.	9.7	207
13	H3K27 methylation: a promiscuous repressive chromatin mark. <i>Current Opinion in Genetics and Development</i> , 2017, 43, 31-37.	3.3	207
14	Trichostatin A causes selective loss of DNA methylation in <i>Neurospora</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 1998, 95, 9430-9435.	7.1	192
15	In Vitro and in Vivo Analyses of a Phe/Tyr Switch Controlling Product Specificity of Histone Lysine Methyltransferases. <i>Journal of Biological Chemistry</i> , 2005, 280, 5563-5570.	3.4	166
16	Dispersed 5S RNA genes in <i>N. crassa</i> : Structure, expression and evolution. <i>Cell</i> , 1981, 24, 819-828.	28.9	150
17	Regional control of histone H3 lysine 27 methylation in <i>Neurospora</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2013, 110, 6027-6032.	7.1	147
18	Relics of repeat-induced point mutation direct heterochromatin formation in <i>Neurospora crassa</i> . <i>Genome Research</i> , 2009, 19, 427-437.	5.5	137

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19	<i>Neurospora crassa</i> , a Model System for Epigenetics Research. <i>Cold Spring Harbor Perspectives in Biology</i> , 2013, 5, a017921-a017921.	5.5	131
20	15 Repeat-induced gene silencing in fungi. <i>Advances in Genetics</i> , 2002, 46, 439-450.	1.8	127
21	DNA Methylation Is Independent of RNA Interference in <i>Neurospora</i> . <i>Science</i> , 2004, 304, 1939-1939.	12.6	116
22	Direct Interaction between DNA Methyltransferase DIM-2 and HP1 Is Required for DNA Methylation in <i>Neurospora crassa</i> . <i>Molecular and Cellular Biology</i> , 2008, 28, 6044-6055.	2.3	116
23	Tools for Fungal Proteomics: Multifunctional <i>Neurospora</i> Vectors for Gene Replacement, Protein Expression and Protein Purification. <i>Genetics</i> , 2009, 182, 11-23.	2.9	114
24	A Methylated <i>Neurospora</i> 5S rRNA Pseudogene Contains a Transposable Element Inactivated by Repeat-Induced Point Mutation. <i>Genetics</i> , 1998, 149, 1787-1797.	2.9	99
25	Loss of HP1 causes depletion of H3K27me3 from facultative heterochromatin and gain of H3K27me2 at constitutive heterochromatin. <i>Genome Research</i> , 2016, 26, 97-107.	5.5	96
26	Organization of ribosomal RNA genes in the fungus <i>Cochliobolus heterostrophus</i> . <i>Current Genetics</i> , 1988, 14, 573-582.	1.7	93
27	DNA Methylation and Normal Chromosome Behavior in <i>Neurospora</i> Depend on Five Components of a Histone Methyltransferase Complex, DCDC. <i>PLoS Genetics</i> , 2010, 6, e1001196.	3.5	93
28	Methylation of Histone H3 Lysine 36 Is Required for Normal Development in <i>Neurospora crassa</i> . <i>Eukaryotic Cell</i> , 2005, 4, 1455-1464.	3.4	88
29	Short TpA-rich segments of the $\hat{1}\hat{1}$ - $\hat{1}$ region induce DNA methylation in <i>Neurospora crassa</i> 1 Edited by K. Yamamoto. <i>Journal of Molecular Biology</i> , 2000, 300, 249-273.	4.2	72
30	The fungus <i>Neurospora crassa</i> displays telomeric silencing mediated by multiple sirtuins and by methylation of histone H3 lysine 9. <i>Epigenetics and Chromatin</i> , 2008, 1, 5.	3.9	72
31	The common ancestral core of vertebrate and fungal telomerase RNAs. <i>Nucleic Acids Research</i> , 2013, 41, 450-462.	14.5	70
32	Cytosine Methylation Associated With Repeat-Induced Point Mutation Causes Epigenetic Gene Silencing in <i>Neurospora crassa</i> . <i>Genetics</i> , 1997, 146, 509-523.	2.9	67
33	<i>Neurospora</i> chromosomes are organized by blocks of importin alpha-dependent heterochromatin that are largely independent of H3K9me3. <i>Genome Research</i> , 2016, 26, 1069-1080.	5.5	64
34	Synthesis of Signals for De Novo DNA Methylation in <i>Neurospora crassa</i> . <i>Molecular and Cellular Biology</i> , 2003, 23, 2379-2394.	2.3	63
35	Heterochromatin protein 1 forms distinct complexes to direct histone deacetylation and DNA methylation. <i>Nature Structural and Molecular Biology</i> , 2012, 19, 471-477.	8.2	63
36	Normal chromosome conformation depends on subtelomeric facultative heterochromatin in <i>Neurospora crassa</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, 15048-15053.	7.1	55

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37	Characterization of Chromosome Ends in the Filamentous Fungus <i>Neurospora crassa</i> . <i>Genetics</i> , 2009, 181, 1129-1145.	2.9	52
38	ASH1-catalyzed H3K36 methylation drives gene repression and marks H3K27me <sub>2/3</sub> -competent chromatin. <i>ELife</i> , 2018, 7, .	6.0	50
39	The DMM complex prevents spreading of DNA methylation from transposons to nearby genes in <i>Neurospora crassa</i> . <i>Genes and Development</i> , 2010, 24, 443-454.	5.9	49
40	Induction and maintenance of nonsymmetrical DNA methylation in <i>Neurospora</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2002, 99, 16485-16490.	7.1	46
41	REVERSAL OF A NEUROSPORA TRANSLOCATION BY CROSSING OVER INVOLVING DISPLACED rDNA, AND METHYLATION OF THE rDNA SEGMENTS THAT RESULT FROM RECOMBINATION. <i>Genetics</i> , 1986, 114, 791-817.	2.9	44
42	Identification of DIM-7, a protein required to target the DIM-5 H3 methyltransferase to chromatin. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2010, 107, 8310-8315.	7.1	41
43	Occurrence of Repeat Induced Point Mutation in Long Segmental Duplications of <i>Neurospora</i> . <i>Genetics</i> , 1997, 147, 125-136.	2.9	40
44	Mutations affecting the biosynthesis of S-adenosylmethionine cause reduction of DNA methylation in <i>Neurospora crassa</i> . <i>Nucleic Acids Research</i> , 1995, 23, 4818-4826.	14.5	38
45	H2B- and H3-Specific Histone Deacetylases Are Required for DNA Methylation in <i>Neurospora crassa</i> . <i>Genetics</i> , 2010, 186, 1207-1216.	2.9	38
46	Protein phosphatase PP1 is required for normal DNA methylation in <i>Neurospora</i> . <i>Genes and Development</i> , 2008, 22, 3391-3396.	5.9	32
47	Recurrent rewiring and emergence of RNA regulatory networks. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E2816-E2825.	7.1	32
48	Evolutionarily ancient BAH <sup>1</sup> PHD protein mediates Polycomb silencing. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2020, 117, 11614-11623.	7.1	30
49	Telomere repeats induce domains of H3K27 methylation in <i>Neurospora</i> . <i>ELife</i> , 2018, 7, .	6.0	30
50	Epigenetic Control of a Transposon-Inactivated Gene in <i>Neurospora</i> is Dependent on DNA Methylation. <i>Genetics</i> , 1996, 143, 137-146.	2.9	29
51	Dual chromatin recognition by the histone deacetylase complex HCHC is required for proper DNA methylation in <i>Neurospora crassa</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2016, 113, E6135-E6144.	7.1	28
52	Control of Development, Secondary Metabolism and Light-Dependent Carotenoid Biosynthesis by the Velvet Complex of <i>Neurospora crassa</i> . <i>Genetics</i> , 2019, 212, 691-710.	2.9	28
53	Induction of H3K9me <sub>3</sub> and DNA methylation by tethered heterochromatin factors in <i>Neurospora crassa</i> . <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2017, 114, E9598-E9607.	7.1	26
54	Gene silencing in filamentous fungi: RIP, MIP and quelling. <i>Journal of Genetics</i> , 1996, 75, 313-324.	0.7	25

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55	Neurospora Importin $\hat{\pm}$ Is Required for Normal Heterochromatic Formation and DNA Methylation. PLoS Genetics, 2015, 11, e1005083.	3.5	25
56	Substitutions in the Amino-Terminal Tail of Neurospora Histone H3 Have Varied Effects on DNA Methylation. PLoS Genetics, 2011, 7, e1002423.	3.5	22
57	Neurospora. Current Biology, 2011, 21, R139-R140.	3.9	15
58	Nucleosome Positioning by an Evolutionarily Conserved Chromatin Remodeler Prevents Aberrant DNA Methylation in <i>Neurospora</i> . Genetics, 2019, 211, 563-578.	2.9	13
59	Extensive and Varied Modifications in Histone H2B of Wild-Type and Histone Deacetylase 1 Mutant <i>Neurospora crassa</i> . Biochemistry, 2010, 49, 5244-5257.	2.5	12
60	Identification of a PRC2 Accessory Subunit Required for Subtelomeric H3K27 Methylation in <i>Neurospora crassa</i> . Molecular and Cellular Biology, 2020, 40, .	2.3	12
61	The Cullin-4 Complex DCDC Does Not Require E3 Ubiquitin Ligase Elements To Control Heterochromatin in <i>Neurospora crassa</i> . Eukaryotic Cell, 2015, 14, 25-28.	3.4	11
62	Marked <i>Neurospora crassa</i> Strains for Competition Experiments and Bayesian Methods for Fitness Estimates. G3: Genes, Genomes, Genetics, 2020, 10, 1261-1270.	1.8	11
63	The ACF chromatin-remodeling complex is essential for Polycomb repression. ELife, 2022, 11, .	6.0	10
64	Selection and Characterization of Mutants Defective in DNA Methylation in <i>Neurospora crassa</i> . Genetics, 2020, 216, 671-688.	2.9	7
65	Rapid and inexpensive preparation of genome-wide nucleosome footprints from model and non-model organisms. STAR Protocols, 2021, 2, 100486.	1.2	7
66	A Light-Inducible Strain for Genome-Wide Histone Turnover Profiling in <i>Neurospora crassa</i> . Genetics, 2020, 215, 569-578.	2.9	6
67	LSD1 prevents aberrant heterochromatin formation in <i>Neurospora crassa</i> . Nucleic Acids Research, 2020, 48, 10199-10210.	14.5	4
68	Robert L. Metzenberg, June 11, 1930–July 15, 2007: Geneticist Extraordinaire and “Model Human”. Genetics, 2008, 178, 611-619.	2.9	3
69	The 2005 Thomas Hunt Morgan Medal. Genetics, 2005, 169, 503-505.	2.9	2
70	Chromatin Structure and Modification. , 2014, , 113-123.		0
71	Transposable Elements and Repeat-Induced Point Mutation. , 0, , 124-131.		0
72	Reply to Hogan: Direct evidence of RNA–protein interactions and rewiring. Proceedings of the National Academy of Sciences of the United States of America, 2017, 114, E10854-E10855.	7.1	0