Kevin R Cook

List of Publications by Year
in descending order

25 papers	2,405 citations	567144 15 h-index	580701 25 g-index
27 all docs	27 docs citations	27 times ranked	3511 citing authors

#	Article	IF	Citations
1	Identification of novel split-GAL4 drivers for the characterization of enteroendocrine cells in the Drosophila melanogaster midgut. G3: Genes, Genomes, Genetics, 2022, 12, .	0.8	4
2	Identification of Split-GAL4 Drivers and Enhancers That Allow Regional Cell Type Manipulations of the Drosophila melanogaster Intestine. Genetics, 2020, 216, 891-903.	1.2	11
3	Identification and Characterization of Breakpoints and Mutations onDrosophila melanogasterBalancer Chromosomes. G3: Genes, Genomes, Genetics, 2020, 10, 4271-4285.	0.8	12
4	DrosophilaHeterochromatin Stabilization Requires the Zinc-Finger Protein Small Ovary. Genetics, 2019, 213, 877-895.	1.2	15
5	The joy of balancers. PLoS Genetics, 2019, 15, e1008421.	1.5	26
6	The Molecular and Genetic Characterization of Second Chromosome Balancers in Drosophila melanogaster. G3: Genes, Genomes, Genetics, 2018, 8, 1161-1171.	0.8	17
7	Mapping Second Chromosome Mutations to Defined Genomic Regions in Drosophila melanogaster. G3: Genes, Genomes, Genetics, 2018, 8, 9-16.	0.8	3
8	The U.S. Culture Collection Network Responding to the Requirements of the Nagoya Protocol on Access and Benefit Sharing. MBio, 2017, 8, .	1.8	30
9	Phenotypes Associated with Second ChromosomePElement Insertions inDrosophila melanogaster. G3: Genes, Genomes, Genetics, 2016, 6, 2665-2670.	0.8	5
10	A Common Suite of Coagulation Proteins Function in Drosophila Muscle Attachment. Genetics, 2016, 204, 1075-1087.	1.2	16
11	Third Chromosome Balancer Inversions Disrupt Protein-Coding Genes and Influence Distal Recombination Events inDrosophila melanogaster. G3: Genes, Genomes, Genetics, 2016, 6, 1959-1967.	0.8	32
12	Rare recombination events generate sequence diversity among balancer chromosomes in Drosophila melanogaster. Proceedings of the National Academy of Sciences of the United States of America, 2016, 113, E1352-61.	3.3	30
13	Effects of Gene Dose, Chromatin, and Network Topology on Expression in Drosophila melanogaster. PLoS Genetics, 2016, 12, e1006295.	1.5	38
14	Safeguarding gene drive experiments in the laboratory. Science, 2015, 349, 927-929.	6.0	254
15	The generation of chromosomal deletions to provide extensive coverage and subdivision of the Drosophila melanogaster genome. Genome Biology, 2012, 13, R21.	13.9	184
16	New research resources at the Bloomington Drosophila Stock Center. Fly, 2010, 4, 88-91.	0.9	52
17	A New Resource for Characterizing X-Linked Genes in Drosophila melanogaster: Systematic Coverage and Subdivision of the X Chromosome With Nested, Y-Linked Duplications. Genetics, 2010, 186, 1095-1109.	1.2	44
18	A spinosyn-sensitive Drosophila melanogaster nicotinic acetylcholine receptor identified through chemically induced target site resistance, resistance gene identification, and heterologous expression. Insect Biochemistry and Molecular Biology, 2010, 40, 376-384.	1.2	120

#	Article	IF	Citations
19	The ribosomal protein genes and Minute loci of Drosophila melanogaster. Genome Biology, 2007, 8, R216.	13.9	330
20	Systematic generation of high-resolution deletion coverage of the Drosophila melanogaster genome. Nature Genetics, 2004, 36, 288-292.	9.4	919
21	A high proportion of genes involved in position effect variegation also affect chromosome inheritance. Chromosoma, 2004, 112, 269-276.	1.0	13
22	The Drosophila Su(var)2-10 locus regulates chromosome structure and function and encodes a member of the PIAS protein family. Genes and Development, 2001, 15, 1334-1348.	2.7	175
23	Identification of Trans-Acting Genes Necessary for Centromere Function in Drosophila melanogaster Using Centromere-Defective Minichromosomes. Genetics, 1997, 145, 737-747.	1.2	17
24	Molecular characterization of ovarian tumors in drosophila. Mechanisms of Development, 1994, 47, 151-164.	1.7	26
25	A rosy future for heterochromatin.. Proceedings of the National Academy of Sciences of the United States of America, 1994, 91, 5219-5221.	3.3	32