Craig M Hart

List of Publications by Year in descending order

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331670 454955 3,244 30 21 30 h-index citations g-index papers 31 31 31 3177 docs citations times ranked citing authors all docs

#	Article	IF	CITATIONS
1	Histone Methyltransferase Activity of a Drosophila Polycomb Group Repressor Complex. Cell, 2002, 111, 197-208.	28.9	1,416
2	Visualization of chromosomal domains with boundary element-associated factor BEAF-32. Cell, 1995, 81, 879-889.	28.9	303
3	A Drosophila ESC-E(Z) Protein Complex Is Distinct from Other Polycomb Group Complexes and Contains Covalently Modified ESC. Molecular and Cellular Biology, 2000, 20, 3069-3078.	2.3	147
4	Regulated inactivation of homologous gene expression in transgenic Nicotiana sylvestris plants containing a defense-related tobacco chitinase gene. Molecular Genetics and Genomics, 1992, 235, 179-188.	2.4	139
5	The scs′ Boundary Element: Characterization of Boundary Element-Associated Factors. Molecular and Cellular Biology, 1997, 17, 999-1009.	2.3	109
6	Facilitation of chromatin dynamics by SARs. Current Opinion in Genetics and Development, 1998, 8, 519-525.	3.3	105
7	A 61 bp enhancer element of the tobacco \hat{l}^2 -1,3-glucanase B gene interacts with one or more regulated nuclear proteins. Plant Molecular Biology, 1993, 21, 121-131.	3.9	95
8	Genome-Wide Mapping of Boundary Element-Associated Factor (BEAF) Binding Sites in <i>Drosophila melanogaster</i> Links BEAF to Transcription. Molecular and Cellular Biology, 2009, 29, 3556-3568.	2.3	95
9	Identification of a Class of Chromatin Boundary Elements. Molecular and Cellular Biology, 1998, 18, 7478-7486.	2.3	86
10	Evidence for an antagonistic relationship between the boundary element-associated factor BEAF and the transcription factor DREF. Chromosoma, 1999, 108, 375-383.	2.2	81
11	Evidence for a role of beta-1,3-glucanase in dicot seed germination. Plant Journal, 1994, 5, 273-278.	5.7	79
12	Developmental, hormonal, and pathogenesis-related regulation of the tobacco class I \hat{I}^2 -1,3-glucanase B promoter. Plant Molecular Biology, 1994, 25, 299-311.	3.9	73
13	Mapping geochemical singularity using multifractal analysis: Application to anomaly definition on stream sediments data from Funin Sheet, Yunnan, China. Journal of Geochemical Exploration, 2010, 104, 1-11.	3.2	69
14	BEAF Regulates Cell-Cycle Genes through the Controlled Deposition of H3K9 Methylation Marks into Its Conserved Dual-Core Binding Sites. PLoS Biology, 2008, 6, e327.	5.6	60
15	The Drosophila Boundary Element-Associated Factors BEAF-32A and BEAF-32B Affect Chromatin Structure. Genetics, 2006, 173, 1365-1375.	2.9	59
16	Characterization of BEAF Mutations Isolated by Homologous Recombination in Drosophila. Genetics, 2007, 176, 801-813.	2.9	49
17	Transcription antitermination by phage lambda gene Q protein requires a DNA segment spanning the RNA start site Genes and Development, 1987, 1, 217-226.	5.9	47
18	Identification of a multicopy chromatin boundary element at the borders of silenced chromosomal domains. Chromosoma, 2002, 110, 519-531.	2.2	47

#	Article	IF	CITATION
19	Genome-wide studies of the multi-zinc finger Drosophila Suppressor of Hairy-wing protein in the ovary. Nucleic Acids Research, 2012, 40, 5415-5431.	14.5	47
20	Studies of the Role of the Drosophila scs and scs′ Insulators in Defining Boundaries of a Chromosome Puff. Molecular and Cellular Biology, 2004, 24, 1470-1480.	2.3	36
21	Deletion Analysis of the Lambda tR1 Termination Region. Journal of Molecular Biology, 1994, 237, 255-265.	4.2	33
22	A genetic screen supports a broad role for the Drosophila insulator proteins BEAF-32A and BEAF-32B in maintaining patterns of gene expression. Molecular Genetics and Genomics, 2007, 277, 273-286.	2.1	21
23	4C-seq characterization of Drosophila BEAF binding regions provides evidence for highly variable long-distance interactions between active chromatin. PLoS ONE, 2018, 13, e0203843.	2.5	11
24	Promoter-Proximal Chromatin Domain Insulator Protein BEAF Mediates Local and Long-Range Communication with a Transcription Factor and Directly Activates a Housekeeping Promoter in <i>Drosophila</i> . Genetics, 2020, 215, 89-101.	2.9	10
25	Characterization of the Drosophila BEAF-32A and BEAF-32B Insulator Proteins. PLoS ONE, 2016, 11, e0162906.	2.5	10
26	Lack of the Drosophila BEAF insulator proteins alters regulation of genes in the Antennapedia complex. Molecular Genetics and Genomics, 2011, 285, 113-123.	2.1	4
27	Overlapping but Distinct Sequences Play Roles in the Insulator and Promoter Activities of the <i>Drosophila</i> BEAF-Dependent scs' Insulator. Genetics, 2020, 215, 1003-1012.	2.9	4
28	Targeted gene replacement by homologous recombination in Drosophila stimulates production of second-site mutations. Fly, 2010, 4, 12-17.	1.7	3
29	Do the BEAF insulator proteins regulate genes involved in cell polarity and neoplastic growth?. Developmental Biology, 2014, 389, 121-123.	2.0	3
30	Using a phiC31 "Disintegrase―to make new attP sites in the Drosophila genome at locations showing chromosomal position effects. PLoS ONE, 2018, 13, e0205538.	2.5	3