

Takuya Nakayama

List of Publications by Year in descending order

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41
papers

4,738
citations

257101

24
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41
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all docs

41
docs citations

41
times ranked

5258
citing authors

#	ARTICLE	IF	CITATIONS
1	Identification of Smad7, a TGF β ² -inducible antagonist of TGF β ² signalling. <i>Nature</i> , 1997, 389, 631-635.	13.7	1,684
2	Genome evolution in the allotetraploid frog <i>Xenopus laevis</i> . <i>Nature</i> , 2016, 538, 336-343.	13.7	849
3	Daughters against dpp modulates dpp organizing activity in <i>Drosophila</i> wing development. <i>Nature</i> , 1997, 389, 627-631.	13.7	402
4	Simple and efficient CRISPR/Cas9-mediated targeted mutagenesis in <i>Xenopus tropicalis</i> . <i>Genesis</i> , 2013, 51, 835-843.	0.8	251
5	A protein that binds to a cis-acting element of wheat histone genes has a leucine zipper motif. <i>Science</i> , 1989, 245, 965-967.	6.0	199
6	Physical and Functional Interaction of Murine and <i>Xenopus</i> Smad7 with Bone Morphogenetic Protein Receptors and Transforming Growth Factor β ² Receptors. <i>Journal of Biological Chemistry</i> , 1998, 273, 25364-25370.	1.6	143
7	A wheat histone H3 promoter confers cell division-dependent and -independent expression of the <i>gus</i> gene in transgenic rice plants. <i>Plant Journal</i> , 1993, 3, 241-252.	2.8	117
8	The activity and signaling range of mature BMP-4 is regulated by sequential cleavage at two sites within the prodomain of the precursor. <i>Genes and Development</i> , 2001, 15, 2797-2802.	2.7	115
9	Cas9-Based Genome Editing in <i>Xenopus tropicalis</i> . <i>Methods in Enzymology</i> , 2014, 546, 355-375.	0.4	96
10	Nuclear protein(s) binding to the conserved DNA hexameric sequence postulated to regulate transcription of wheat histone genes. <i>FEBS Letters</i> , 1987, 223, 273-278.	1.3	86
11	Smad6 functions as an intracellular antagonist of some TGF β ² family members during <i>Xenopus</i> embryogenesis. <i>Genes To Cells</i> , 1998, 3, 387-394.	0.5	73
12	Mutation of an upstream cleavage site in the BMP4 prodomain leads to tissue-specific loss of activity. <i>Development (Cambridge)</i> , 2006, 133, 1933-1942.	1.2	58
13	<i>Xenopus pax6</i> mutants affect eye development and other organ systems, and have phenotypic similarities to human aniridia patients. <i>Developmental Biology</i> , 2015, 408, 328-344.	0.9	58
14	Can't get no SMADisfaction: Smad proteins as positive and negative regulators of TGF β ² family signals. <i>BioEssays</i> , 1999, 21, 382-390.	1.2	47
15	Highly conserved hexamer, octamer and nonamer motifs are positive cis-regulatory elements of the wheat histone H3 gene. <i>FEBS Letters</i> , 1992, 300, 167-170.	1.3	40
16	Regulation of BMP/Dpp signaling during embryonic development. <i>Cellular and Molecular Life Sciences</i> , 2000, 57, 943-956.	2.4	37
17	Cisacting Sequences that Modulate Transcription of Wheat Histone H3 and 3' Processing of H3 Premature mRNA. <i>Plant and Cell Physiology</i> , 1989, 30, 825-832.	1.5	34
18	Molecular cloning and nucleotide sequences of cDNAs for histone H1 and H2B variants from wheat. <i>Nucleic Acids Research</i> , 1991, 19, 5077-5077.	6.5	31

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19	Dissection of inhibitory Smad proteins: both N- and C-terminal domains are necessary for full activities of <i>Xenopus</i> Smad6 and Smad7. <i>Mechanisms of Development</i> , 2001, 100, 251-262.	1.7	30
20	DNA-binding protein(s) interacts with a conserved nonameric sequence in the upstream regions of wheat histone genes. <i>FEBS Letters</i> , 1988, 239, 319-323.	1.3	29
21	A type I element composed of the hexamer (ACGTCA) and octamer (CGCGGATC) motifs plays a role(s) in meristematic expression of a wheat histone H3 gene in transgenic rice plants. <i>Plant Molecular Biology</i> , 1995, 27, 17-26.	2.0	29
22	<i>Xenopus</i> mutant reveals necessity of <i>rax</i> for specifying the eye field which otherwise forms tissue with telencephalic and diencephalic character. <i>Developmental Biology</i> , 2014, 395, 317-330.	0.9	28
23	no privacy, a <i>Xenopus tropicalis</i> mutant, is a model of human Hermansky-Pudlak Syndrome and allows visualization of internal organogenesis during tadpole development. <i>Developmental Biology</i> , 2017, 426, 472-486.	0.9	28
24	Specific Interaction of Nuclear Protein HBP-1 with the Conserved Hexameric Sequence ACGTCA in the Regulatory Region of Wheat Histone Genes. <i>Plant and Cell Physiology</i> , 1989, 30, 107-119.	1.5	25
25	Proximal promoter region of the wheat histone H3 gene confers S phase-specific gene expression in transformed rice cells. <i>Plant Molecular Biology</i> , 1993, 23, 553-565.	2.0	25
26	Structural and functional characterization of two wheat histone H2B promoters. <i>Plant Molecular Biology</i> , 1995, 28, 155-172.	2.0	24
27	Identification of three kinds of mutually related composite elements conferring S phase-specific transcriptional activation. <i>Plant Journal</i> , 1999, 18, 611-623.	2.8	21
28	Regulation of wheat histone gene expression. <i>Critical Reviews in Plant Sciences</i> , 1993, 12, 97-110.	2.7	19
29	Simple, fast, tissue-specific bacterial artificial chromosome transgenesis in <i>Xenopus</i> . <i>Genesis</i> , 2012, 50, 307-315.	0.8	19
30	Simple embryo injection of long single-stranded donor templates with the CRISPR/Cas9 system leads to homology-directed repair in <i>Xenopus tropicalis</i> and <i>Xenopus laevis</i> . <i>Genesis</i> , 2020, 58, e23366.	0.8	19
31	Structural characteristics of two wheat histone H2A genes encoding distinct types of variants and functional differences in their promoter activity. <i>Plant Molecular Biology</i> , 1997, 33, 791-802.	2.0	17
32	Cooperation of two distinct cis-acting elements is necessary for the S phase-specific activation of the wheat histone H3 promoter. <i>Plant Journal</i> , 1997, 11, 1219-1225.	2.8	17
33	High variability of expression profiles of homeologous genes for Wnt, Hh, Notch, and Hippo signaling pathways in <i>Xenopus laevis</i> . <i>Developmental Biology</i> , 2017, 426, 270-290.	0.9	16
34	A Zinc-Finger-Type Transcription Factor WZF-1 That Binds to a Novel cis-Acting Element Element of Histone Gene Promoters Represses Its Own Promoter. <i>Plant and Cell Physiology</i> , 1996, 37, 557-562.	1.5	15
35	Dissection of the wheat transcription factor HBP-1a(17) reveals a modular structure for the activation domain. <i>Molecular Genetics and Genomics</i> , 1997, 253, 553-561.	2.4	13
36	Roles of ADAM13-regulated Wnt activity in early <i>Xenopus</i> eye development. <i>Developmental Biology</i> , 2012, 363, 147-154.	0.9	12

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37	Chromosomal locations of the genes for histones and a histone gene-binding protein family HBP-1 in common wheat. <i>Plant Molecular Biology</i> , 1993, 22, 603-614.	2.0	10
38	Cell cycle-regulated gene expression in transgenic plant cells. <i>Genesis</i> , 1990, 11, 205-213.	3.1	9
39	Trans-activation of the wheat histone H3 promoter by Ga14 DNA-binding domain (1â€“94) in plant cells. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1995, 1263, 281-284.	2.4	8
40	Generation and Characterization of Developmental Mutations in <i>Xenopus tropicalis</i> . <i>Current Genomics</i> , 2003, 4, 673-685.	0.7	4
41	Organic small hairpin RNAs (OshR): A do-it-yourself platform for transgene-based gene silencing. <i>Methods</i> , 2013, 63, 101-109.	1.9	1