

Takuya Nakayama

List of Publications by Year in descending order

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Version: 2024-02-01

41
papers

4,738
citations

257101

24
h-index

276539

41
g-index

41
all docs

41
docs citations

41
times ranked

5258
citing authors

| # | ARTICLE | IF | CITATIONS |
|----|---|------|-----------|
| 1 | 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 |
| 2 | 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 |
| 3 | 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 |
| 4 | Genome evolution in the allotetraploid frog <i>Xenopus laevis</i> . <i>Nature</i> , 2016, 538, 336-343. | 13.7 | 849 |
| 5 | <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 |
| 6 | Cas9-Based Genome Editing in <i>Xenopus tropicalis</i> . <i>Methods in Enzymology</i> , 2014, 546, 355-375. | 0.4 | 96 |
| 7 | <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 |
| 8 | 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 |
| 9 | Simple and efficient CRISPR/Cas9-mediated targeted mutagenesis in <i>Xenopus tropicalis</i> . <i>Genesis</i> , 2013, 51, 835-843. | 0.8 | 251 |
| 10 | Roles of ADAM13-regulated Wnt activity in early <i>Xenopus</i> eye development. <i>Developmental Biology</i> , 2012, 363, 147-154. | 0.9 | 12 |
| 11 | Simple, fast, tissue-specific bacterial artificial chromosome transgenesis in <i>Xenopus</i> . <i>Genesis</i> , 2012, 50, 307-315. | 0.8 | 19 |
| 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 | Generation and Characterization of Developmental Mutations in <i>Xenopus tropicalis</i> . <i>Current Genomics</i> , 2003, 4, 673-685. | 0.7 | 4 |
| 14 | 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 |
| 15 | 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 |
| 16 | Regulation of BMP/Dpp signaling during embryonic development. <i>Cellular and Molecular Life Sciences</i> , 2000, 57, 943-956. | 2.4 | 37 |
| 17 | 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 |
| 18 | Can't get no SMADisfaction: Smad proteins as positive and negative regulators of TGF- β family signals. <i>BioEssays</i> , 1999, 21, 382-390. | 1.2 | 47 |

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|----|--|------|-----------|
| 19 | Smad6 functions as an intracellular antagonist of some TGF- β family members during Xenopus embryogenesis. <i>Genes To Cells</i> , 1998, 3, 387-394. | 0.5 | 73 |
| 20 | Physical and Functional Interaction of Murine and Xenopus Smad7 with Bone Morphogenetic Protein Receptors and Transforming Growth Factor- β Receptors. <i>Journal of Biological Chemistry</i> , 1998, 273, 25364-25370. | 1.6 | 143 |
| 21 | Daughters against dpp modulates dpp organizing activity in Drosophila wing development. <i>Nature</i> , 1997, 389, 627-631. | 13.7 | 402 |
| 22 | Identification of Smad7, a TGF β -inducible antagonist of TGF- β signalling. <i>Nature</i> , 1997, 389, 631-635. | 13.7 | 1,684 |
| 23 | 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 |
| 24 | 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 |
| 25 | 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 |
| 26 | 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 |
| 27 | A type I element composed of the hexamer (ACGTCA) and octamer (CCCGGATC) 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 |
| 28 | Structural and functional characterization of two wheat histone H2B promoters. <i>Plant Molecular Biology</i> , 1995, 28, 155-172. | 2.0 | 24 |
| 29 | Trans-activation of the wheat histone H3 promoter by Ga14 DNA-binding domain (1-94) in plant cells. <i>Biochimica Et Biophysica Acta Gene Regulatory Mechanisms</i> , 1995, 1263, 281-284. | 2.4 | 8 |
| 30 | 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 |
| 31 | 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 |
| 32 | 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 |
| 33 | Regulation of wheat histone gene expression. <i>Critical Reviews in Plant Sciences</i> , 1993, 12, 97-110. | 2.7 | 19 |
| 34 | 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 |
| 35 | 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 |
| 36 | Cell cycle-regulated gene expression in transgenic plant cells. <i>Genesis</i> , 1990, 11, 205-213. | 3.1 | 9 |

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|----|---|-----|-----------|
| 37 | 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 |
| 38 | 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 |
| 39 | Cisacting Sequences that Modulate Transcription of Wheat Histone H3 and 3â€² Processing of H3 Premature mRNA. <i>Plant and Cell Physiology</i> , 1989, 30, 825-832. | 1.5 | 34 |
| 40 | 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 |
| 41 | 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 |