List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/5167681/publications.pdf Version: 2024-02-01



ΤΟΡΗ SHIMADA

#	Article	IF	CITATIONS
1	The genome sequence of Samia ricini , a new model species of lepidopteran insect. Molecular Ecology Resources, 2021, 21, 327-339.	4.8	12
2	Horizontal Gene Transfer and Gene Duplication of β-Fructofuranosidase Confer Lepidopteran Insects Metabolic Benefits. Molecular Biology and Evolution, 2021, 38, 2897-2914.	8.9	10
3	Duplication and diversification of trehalase confers evolutionary advantages on lepidopteran insects. Molecular Ecology, 2019, 28, 5282-5298.	3.9	28
4	High-quality genome assembly of the silkworm, Bombyx mori. Insect Biochemistry and Molecular Biology, 2019, 107, 53-62.	2.7	201
5	Two CCCH-type zinc finger domains in the Masc protein are dispensable for masculinization and dosage compensation in Bombyx mori. Insect Biochemistry and Molecular Biology, 2019, 104, 30-38.	2.7	21
6	Morphological and electrophysiological differences in tarsal chemosensilla between the wild silkmoth Bombyx mandarina and the domesticated species Bombyx mori. Arthropod Structure and Development, 2018, 47, 238-247.	1.4	5
7	Inhibitory role of the Bm8 protein in the propagation of Bombyx mori nucleopolyhedrovirus. Virus Research, 2018, 249, 124-131.	2.2	3
8	Bombyx ortholog of the Drosophila eye color gene brown controls riboflavin transport in Malpighian tubules. Insect Biochemistry and Molecular Biology, 2018, 92, 65-72.	2.7	15
9	Proteomic Analysis of Larval Integument in a Dominant Obese Translucent (Obs) Silkworm Mutant. Journal of Insect Science, 2018, 18, .	1.5	1
10	Accumulation of uric acid in the epidermis forms the white integument of Samia ricini larvae. PLoS ONE, 2018, 13, e0205758.	2.5	11
11	InÂvivo masculinizing function of the Ostrinia furnacalis Masculinizer gene. Biochemical and Biophysical Research Communications, 2018, 503, 1768-1772.	2.1	21
12	A single amino acid substitution in the Bombyx-specific mucin-like membrane protein causes resistance to Bombyx mori densovirus. Scientific Reports, 2018, 8, 7430.	3.3	12
13	A reexamination on the deficiency of riboflavin accumulation in Malpighian tubules in larval translucent mutants of the silkworm, Bombyx mori. Genetica, 2018, 146, 425-431.	1.1	4
14	Silkworms suppress the release of green leaf volatiles by mulberry leaves with an enzyme from their spinnerets. Scientific Reports, 2018, 8, 11942.	3.3	23
15	Bm-muted , orthologous to mouse muted and encoding a subunit of the BLOC-1 complex, is responsible for the otm translucent mutation of the silkworm Bombyx mori. Gene, 2017, 629, 92-100.	2.2	18
16	Artificial "ping-pong―cascade of PIWI-interacting RNA in silkworm cells. Rna, 2017, 23, 86-97.	3.5	10
17	The morphology of antennal lobe projection neurons is controlled by a POU-domain transcription factor Bmacj6 in the silkmoth Bombyx mori. Scientific Reports, 2017, 7, 14050.	3.3	0
18	Identification of a bipartite nuclear localization signal in the silkworm Masc protein. FEBS Letters, 2016, 590, 2256-2261.	2.8	11

#	Article	IF	CITATIONS
19	Bombyx mori nucleopolyhedrovirus BM5 protein regulates progeny virus production and viral gene expression. Virology, 2016, 498, 240-249.	2.4	15
20	Gene expression and localization analysis of Bombyx mori bidensovirus and its putative receptor in B. mori midgut. Journal of Invertebrate Pathology, 2016, 136, 50-56.	3.2	10
21	Identification of the silkworm quail gene reveals a crucial role of a receptor guanylyl cyclase in larval pigmentation. Insect Biochemistry and Molecular Biology, 2016, 68, 33-40.	2.7	16
22	Mapping and recombination analysis of two moth colour mutations, Black moth and Wild wing spot, in the silkworm Bombyx mori. Heredity, 2016, 116, 52-59.	2.6	18
23	Identification and functional analysis of a <i>Masculinizer</i> orthologue in <i>Trilocha varians</i> (Lepidoptera: Bombycidae). Insect Molecular Biology, 2015, 24, 561-569.	2.0	35
24	The Endosymbiotic Bacterium Wolbachia Selectively Kills Male Hosts by Targeting the Masculinizing Gene. PLoS Pathogens, 2015, 11, e1005048.	4.7	73
25	Functional analysis of antisense long non-coding RNAs transcribed from the Bombyx mori (Lepidoptera: Bombycidae) nucleopolyhedrovirus genome. Applied Entomology and Zoology, 2015, 50, 155-167.	1.2	1
26	Sexually biased transcripts at early embryonic stages of the silkworm depend on the sex chromosome constitution. Gene, 2015, 560, 50-56.	2.2	8
27	A novel sucrose hydrolase from the bombycoid silkworms Bombyx mori, Trilocha varians, and Samia cynthia ricini with a substrate specificity for sucrose. Insect Biochemistry and Molecular Biology, 2015, 61, 46-52.	2.7	13
28	The killing speed of egt-inactivated Bombyx mori nucleopolyhedrovirus depends on the developmental stage of B. mori larvae. Journal of Invertebrate Pathology, 2015, 126, 64-70.	3.2	13
29	Bombyx mori nucleopolyhedrovirus actin rearrangement-inducing factor 1 enhances systemic infection in B. mori larvae. Journal of General Virology, 2015, 96, 1938-1946.	2.9	12
30	Two Conserved Cysteine Residues Are Required for the Masculinizing Activity of the Silkworm Masc Protein. Journal of Biological Chemistry, 2015, 290, 26114-26124.	3.4	29
31	Positional cloning of the sexâ€linked giant egg (<i>Ge</i>) locus in the silkworm, <i>Bombyx mori</i> . Insect Molecular Biology, 2015, 24, 213-221.	2.0	8
32	Deletion analysis of a superoxide dismutase gene of Bombyx mori (Lepidoptera: Bombycidae) nucleopolyhedrovirus. Applied Entomology and Zoology, 2015, 50, 57-62.	1.2	5
33	Antennal lobe organization and pheromone usage in bombycid moths. Biology Letters, 2014, 10, 20140096.	2.3	21
34	Silkworm HP1a transcriptionally enhances highly expressed euchromatic genes via association with their transcription start sites. Nucleic Acids Research, 2014, 42, 11462-11471.	14.5	12
35	A single female-specific piRNA is the primary determiner of sex in the silkworm. Nature, 2014, 509, 633-636.	27.8	407
36	The BIR and BIR-like domains of Bombyx mori nucleopolyhedrovirus IAP2 protein are required for efficient viral propagation. Biochemical and Biophysical Research Communications, 2014, 454, 581-587.	2.1	14

#	Article	IF	CITATIONS
37	Anatomical and functional analysis of domestication effects on the olfactory system of the silkmoth <i>Bombyx mori</i> . Proceedings of the Royal Society B: Biological Sciences, 2014, 281, 20132582.	2.6	16
38	Dimerization and proper degradation of BmNPV IE2 are required for efficient virus growth in larvae of Bombyx mori (Lepidoptera: Bombycidae). Applied Entomology and Zoology, 2013, 48, 125-130.	1.2	1
39	Flavonoids from the cocoon of Rondotia menciana. Phytochemistry, 2013, 94, 108-112.	2.9	23
40	Reduced expression of the <i>dysbindin</i> -like gene in the <i>Bombyx mori ov</i> mutant exhibiting mottled translucency of the larval skin. Genome, 2013, 56, 101-108.	2.0	18
41	Mutation of a novel ABC transporter gene is responsible for the failure to incorporate uric acid in the epidermis of ok mutants of the silkworm, Bombyx mori. Insect Biochemistry and Molecular Biology, 2013, 43, 562-571.	2.7	51
42	Functional characterization of Bombyx mori nucleopolyhedrovirus CG30 protein. Virus Research, 2013, 174, 52-59.	2.2	16
43	Characterization of a novel chromodomain-containing gene from the silkworm, Bombyx mori. Gene, 2013, 527, 649-654.	2.2	8
44	Albino (al) is a tetrahydrobiopterin (BH4)-deficient mutant of the silkworm Bombyx mori. Insect Biochemistry and Molecular Biology, 2013, 43, 594-600.	2.7	31
45	Inhibition of Tumor Angiogenesis and Growth by a Small-Molecule Multi-FGF Receptor Blocker with Allosteric Properties. Cancer Cell, 2013, 23, 477-488.	16.8	138
46	Silkworm plasmatocytes are more resistant than other hemocyte morphotypes to Bombyx mori nucleopolyhedrovirus infection. Journal of Invertebrate Pathology, 2013, 112, 102-104.	3.2	10
47	Large Scale Full-Length cDNA Sequencing Reveals a Unique Genomic Landscape in a Lepidopteran Model Insect, <i>Bombyx mori</i> . G3: Genes, Genomes, Genetics, 2013, 3, 1481-1492.	1.8	87
48	Vitellogenin Receptor Mutation Leads to the Oogenesis Mutant Phenotype "scanty vitellin―of the Silkworm, Bombyx mori. Journal of Biological Chemistry, 2013, 288, 13345-13355.	3.4	76
49	The comprehensive epigenome map of piRNA clusters. Nucleic Acids Research, 2013, 41, 1581-1590.	14.5	29
50	Identification of Key Uric Acid Synthesis Pathway in a Unique Mutant Silkworm Bombyx mori Model of Parkinson's Disease. PLoS ONE, 2013, 8, e69130.	2.5	42
51	Precocious Metamorphosis in the Juvenile Hormone–Deficient Mutant of the Silkworm, Bombyx mori. PLoS Genetics, 2012, 8, e1002486.	3.5	135
52	The Baculovirus Uses a Captured Host Phosphatase to Induce Enhanced Locomotory Activity in Host Caterpillars. PLoS Pathogens, 2012, 8, e1002644.	4.7	78
53	Maximizing the amplitude of coherent phonons with shaped laser pulses. Journal of Applied Physics, 2012, 112, 113103.	2.5	9
54	Baculovirus-Encoded Protein BV/ODV-E26 Determines Tissue Tropism and Virulence in Lepidopteran Insects. Journal of Virology, 2012, 86, 2545-2555.	3.4	33

#	Article	IF	CITATIONS
55	A homolog of the human Hermansky–Pudluck syndrome-5 (HPS5) gene is responsible for the oa larval translucent mutants in the silkworm, Bombyx mori. Genetica, 2012, 140, 463-468.	1.1	28
56	Reinvestigation of the Sex Pheromone of the Wild Silkmoth Bombyx mandarina: The Effects of Bombykal and Bombykyl Acetate. Journal of Chemical Ecology, 2012, 38, 1031-1035.	1.8	26
57	Molecular Phylogeny, Laboratory Rearing, and Karyotype of the Bombycid Moth, <i>Trilocha varians</i> . Journal of Insect Science, 2012, 12, 1-17.	1.5	10
58	Positional cloning of a gene responsible for the <i>cts</i> mutation of the silkworm, <i>Bombyx mori</i> . Genome, 2012, 55, 493-504.	2.0	14
59	A role for transcription from a piRNA cluster in de novo piRNA production. Rna, 2012, 18, 265-273.	3.5	50
60	Altered expression of testis-specific genes, piRNAs, and transposons in the silkworm ovary masculinized by a W chromosome mutation. BMC Genomics, 2012, 13, 119.	2.8	10
61	Identification and characterization of host factors interacting with Bombyx mori nucleopolyhedrovirus ORF8. Journal of Microbiology, 2012, 50, 469-477.	2.8	6
62	Infection study of Bombyx mori macula-like virus (BmMLV) using a BmMLV-negative cell line and an infectious cDNA clone. Journal of Virological Methods, 2012, 179, 316-324.	2.1	23
63	Female sex pheromone and male behavioral responses of the bombycid moth Trilocha varians: comparison with those of the domesticated silkmoth Bombyx mori. Die Naturwissenschaften, 2012, 99, 207-215.	1.6	14
64	Identification and characterization of the fusion transcript, composed of the apterous homolog and a putative protein phosphatase gene, generated by 1.5-Mb interstitial deletion in the vestigial (Vg) mutant of Bombyx mori. Insect Biochemistry and Molecular Biology, 2011, 41, 306-312.	2.7	6
65	Mutations in an amino acid transporter gene are responsible for sex-linked translucent larval skin of the silkworm, Bombyx mori. Insect Biochemistry and Molecular Biology, 2011, 41, 680-687.	2.7	22
66	BmDJ-1 Is a Key Regulator of Oxidative Modification in the Development of the Silkworm, Bombyx mori. PLoS ONE, 2011, 6, e17683.	2.5	8
67	Positional cloning of silkworm white egg 2 (w-2) locus shows functional conservation and diversification of ABC transporters for pigmentation in insects. Genes To Cells, 2011, 16, 331-342.	1.2	62
68	Interspecies linkage analysis of mo, a Bombyx mori locus associated with mosaicism and gynandromorphism. Genetica, 2011, 139, 1323-1329.	1.1	2
69	Comparative analysis of budded virus infectivity of Bombyx mandarina and B. mori nucleopolyhedroviruses. Virus Genes, 2011, 43, 313-317.	1.6	8
70	The silkworm W chromosome is a source of female-enriched piRNAs. Rna, 2011, 17, 2144-2151.	3.5	50
71	Diversity in Copy Number and Structure of a Silkworm Morphogenetic Gene as a Result of Domestication. Genetics, 2011, 187, 965-976.	2.9	21
72	Zygotic amplification of secondary piRNAs during silkworm embryogenesis. Rna, 2011, 17, 1401-1407.	3.5	65

#	Article	IF	CITATIONS
73	Mass identification of transcriptional units expressed from the Bombyx mori nucleopolyhedrovirus genome. Journal of General Virology, 2011, 92, 200-203.	2.9	19
74	Sex-linked transcription factor involved in a shift of sex-pheromone preference in the silkmoth <i>Bombyx mori</i> . Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 18038-18043.	7.1	30
75	Sex pheromone desaturase functioning in a primitive <i>Ostrinia</i> moth is cryptically conserved in congeners' genomes. Proceedings of the National Academy of Sciences of the United States of America, 2011, 108, 7102-7106.	7.1	41
76	Role of the ubiquitin-proteasome system in Bombyx mori nucleopolyhedrovirus infection. Journal of General Virology, 2011, 92, 699-705.	2.9	43
77	The Silkworm-An Attractive BioResource Supplied by Japan. Experimental Animals, 2010, 59, 139-146.	1.1	42
78	Molecular analysis of sex chromosome-linked mutants in the silkworm Bombyx mori. Journal of Genetics, 2010, 89, 365-374.	0.7	11
79	Novel non-autonomous transposable elements on W chromosome of the silkworm, Bombyx mori. Journal of Genetics, 2010, 89, 375-387.	0.7	9
80	Characterization of a Bombyx mori nucleopolyhedrovirus mutant lacking both fp25K and p35. Virus Genes, 2010, 41, 144-148.	1.6	3
81	Molecular defect of isovalerylâ€CoA dehydrogenase in the <i>skunk</i> mutant of silkworm, <i>Bombyxâ€∫mori</i> . FEBS Journal, 2010, 277, 4452-4463.	4.7	3
82	Transgenic analysis of the <i>BmBLOS2</i> gene that governs the translucency of the larval integument of the silkworm, <i>Bombyx mori</i> . Insect Molecular Biology, 2010, 19, 659-667.	2.0	49
83	<i>Non-molting glossy</i> / <i>shroud</i> encodes a short-chain dehydrogenase/reductase that functions in the †Black Box' of the ecdysteroid biosynthesis pathway. Development (Cambridge), 2010, 137, 1991-1999.	2.5	163
84	Comparative Studies of Lepidopteran Baculovirus-Specific Protein FP25K: Development of a Novel <i>Bombyx mori</i> Nucleopolyhedrovirus-Based Vector with a Modified <i>fp25K</i> Gene. Journal of Virology, 2010, 84, 5191-5200.	3.4	27
85	The silkworm <i>Green b</i> locus encodes a quercetin 5- <i>O</i> -glucosyltransferase that produces green cocoons with UV-shielding properties. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 11471-11476.	7.1	100
86	Identification and molecular characterization of a sex chromosome rearrangement causing a soft and pliable (spli) larval body phenotype in the silkworm, Bombyx mori. Genome, 2010, 53, 45-54.	2.0	4
87	Yellow-e Determines the Color Pattern of Larval Head and Tail Spots of the Silkworm Bombyx mori. Journal of Biological Chemistry, 2010, 285, 5624-5629.	3.4	47
88	Molecular and functional characterization of an acetyl-CoA acetyltransferase from the adzuki bean borer moth Ostrinia scapulalis (Lepidoptera: Crambidae). Insect Biochemistry and Molecular Biology, 2010, 40, 74-78.	2.7	14
89	Recent transposition of <i>yabusame</i> , a novel <i>piggyBac</i> -like transposable element in the genome of the silkworm, <i>Bombyx mori</i> . Genome, 2010, 53, 585-593.	2.0	13
90	The Silkworm Mutant lemon (lemon lethal) Is a Potential Insect Model for Human Sepiapterin Reductase Deficiency. Journal of Biological Chemistry, 2009, 284, 11698-11705.	3.4	48

#	Article	IF	CITATIONS
91	The <i>Bombyx</i> ovary-derived cell line endogenously expresses PIWI/PIWI-interacting RNA complexes. Rna, 2009, 15, 1258-1264.	3.5	124
92	N-linked glycans located in the pro-region of Bombyx mori nucleopolyhedrovirus V-CATH are essential for the proper folding of V-CATH and V-CHIA. Journal of General Virology, 2009, 90, 170-176.	2.9	18
93	Bombyx mori nucleopolyhedrovirus ORF34 is required for efficient transcription of late and very late genes. Virology, 2009, 392, 230-237.	2.4	18
94	Abnormal red body coloration of the silkworm, <i>Bombyx mori</i> , is caused by a mutation in a novel kynureninase. Genes To Cells, 2009, 14, 129-140.	1.2	31
95	Identification and characterization of globin genes from two lepidopteran insects, Bombyx mori and Samia cynthia ricini. Gene, 2009, 431, 33-38.	2.2	9
96	Bombyx mori nucleopolyhedrovirus FP25K is essential for maintaining a steady-state level of v-cath expression throughout the infection. Virus Research, 2009, 140, 155-160.	2.2	10
97	A 25bp-long insertional mutation in the BmVarp gene causes the waxy translucent skin of the silkworm, Bombyx mori. Insect Biochemistry and Molecular Biology, 2009, 39, 287-293.	2.7	24
98	A Bombyx mandarina mutant exhibiting translucent larval skin is controlled by the molybdenum cofactor sulfurase gene. Genes and Genetic Systems, 2009, 84, 147-152.	0.7	8
99	Sex Chromosomes and Sex Determination in Bombyx mori. Contemporary Topics in Entomology Series, 2009, , .	0.3	2
100	Identification of the female-determining region of the W chromosome in Bombyx mori. Genetica, 2008, 133, 269-282.	1.1	41
101	Prominent downâ€regulation of storage protein genes after bacterial challenge in eriâ€silkworm, <i>Samia cynthia ricini</i> . Archives of Insect Biochemistry and Physiology, 2008, 67, 9-19.	1.5	10
102	Functional analysis of fourGloverin-like genes in the silkworm,Bombyx mori. Archives of Insect Biochemistry and Physiology, 2008, 67, 87-96.	1.5	55
103	<i>yellow</i> and <i>ebony</i> Are the Responsible Genes for the Larval Color Mutants of the Silkworm <i>Bombyx mori</i> . Genetics, 2008, 180, 1995-2005.	2.9	126
104	WildSilkbase: An EST database of wild silkmoths. BMC Genomics, 2008, 9, 338.	2.8	38
105	Mapping of sex-linked genes onto the genome sequence using various aberrations of the Z chromosome in Bombyx mori. Insect Biochemistry and Molecular Biology, 2008, 38, 1072-1079.	2.7	33
106	Bombyx small RNAs: Genomic defense system against transposons in the silkworm, Bombyx mori. Insect Biochemistry and Molecular Biology, 2008, 38, 1058-1065.	2.7	41
107	Genome-wide survey for baculoviral host homologs using the Bombyx genome sequence. Insect Biochemistry and Molecular Biology, 2008, 38, 1080-1086.	2.7	40
108	The genome of a lepidopteran model insect, the silkworm Bombyx mori. Insect Biochemistry and Molecular Biology, 2008, 38, 1036-1045.	2.7	592

TORU SHIMADA

#	Article	IF	CITATIONS
109	Identification of Bombyx mori 14-3-3 orthologs and the interactor Hsp60. Neuroscience Research, 2008, 61, 271-280.	1.9	22
110	The fibroblast growth factor homolog of Bombyx mori nucleopolyhedrovirus enhances systemic virus propagation in B. mori larvae. Virus Research, 2008, 137, 80-85.	2.2	15
111	Developmentally synchronized expression of two Bombyx mori Piwi subfamily genes, SIWI and BmAGO3 in germ-line cells. Biochemical and Biophysical Research Communications, 2008, 367, 755-760.	2.1	59
112	Bombyx mori nucleopolyhedrovirus SNF2 global transactivator homologue (Bm33) enhances viral pathogenicity in B. mori larvae. Journal of General Virology, 2008, 89, 3039-3046.	2.9	9
113	Deletion of a gene encoding an amino acid transporter in the midgut membrane causes resistance to a <i>Bombyx</i> parvo-like virus. Proceedings of the National Academy of Sciences of the United States of America, 2008, 105, 7523-7527.	7.1	77
114	β-Fructofuranosidase Genes of the Silkworm, Bombyx mori. Journal of Biological Chemistry, 2008, 283, 15271-15279.	3.4	104
115	Establishment of a Novel In Vivo Sex-Specific Splicing Assay System To Identify a <i>trans</i> -Acting Factor That Negatively Regulates Splicing of <i>Bombyx mori dsx</i> Female Exons. Molecular and Cellular Biology, 2008, 28, 333-343.	2.3	71
116	A silkworm–baculovirus model for assessing the therapeutic effects of antiviral compounds: characterization and application to the isolation of antivirals from traditional medicines. Journal of General Virology, 2008, 89, 188-194.	2.9	42
117	ERK- and JNK-Dependent Signaling Pathways Contribute to <i>Bombyx mori</i> Nucleopolyhedrovirus Infection. Journal of Virology, 2007, 81, 13700-13709.	3.4	109
118	Identification of differentially expressed host genes in Bombyx mori nucleopolyhedrovirus infected cells by using subtractive hybridization. Applied Entomology and Zoology, 2007, 42, 151-159.	1.2	26
119	Sex determination in the silkworm, Bombyx mori: A female determinant on the W chromosome and the sex-determining gene cascade. Seminars in Cell and Developmental Biology, 2007, 18, 379-388.	5.0	62
120	Mutational analysis of active site residues of chitinase from Bombyx mori nucleopolyhedrovirus. Virus Research, 2007, 124, 168-175.	2.2	30
121	SAGE analysis of early oogenesis in the silkworm, Bombyx mori. Insect Biochemistry and Molecular Biology, 2007, 37, 147-154.	2.7	15
122	Rescue of white egg 1 mutant by introduction of the wild-type Bombyx kynurenine 3?monooxygenase gene. Insect Science, 2007, 14, 85-92.	3.0	17
123	Isolation and characterization of sex chromosome rearrangements generating male muscle dystrophy and female abnormal oogenesis in the silkworm, Bombyx mori. Genetica, 2007, 130, 267-280.	1.1	8
124	Functional characterization of chitinase from Cydia pomonella granulovirus. Archives of Virology, 2007, 152, 1655-1664.	2.1	13
125	Comparative studies of Bombyx mori nucleopolyhedrovirus chitinase and its host ortholog, BmChi-h. Biochemical and Biophysical Research Communications, 2006, 345, 825-833.	2.1	27
126	N-linked glycans of Bombyx mori nucleopolyhedrovirus fibroblast growth factor are crucial for its secretion. Biochemical and Biophysical Research Communications, 2006, 350, 1069-1075.	2.1	20

#	ARTICLE	IF	CITATIONS
127	Glycine-rich protein genes, which encode a major component of the cuticle, have different developmental profiles from other cuticle protein genes in Bombyx mori. Insect Biochemistry and Molecular Biology, 2006, 36, 99-110.	2.7	46
128	The female-killing chromosome of the silkworm, Bombyx mori, was generated by translocation between the Z and W chromosomes. Genetica, 2006, 127, 253-265.	1.1	11
129	In vivo and in vitro analyses of a Bombyx mori nucleopolyhedrovirus mutant lacking functional vfgf. Virology, 2006, 355, 62-70.	2.4	50
130	Role of the silkworm argonaute2 homolog gene in double-strand break repair of extrachromosomal DNA. Nucleic Acids Research, 2006, 34, 1092-1101.	14.5	19
131	Lepidopteran Ortholog of Drosophila Breathless Is a Receptor for the Baculovirus Fibroblast Growth Factor. Journal of Virology, 2006, 80, 5474-5481.	3.4	48
132	THE GENETICS AND GENOMICS OF THE SILKWORM, <i>BOMBYX MORI</i> . Annual Review of Entomology, 2005, 50, 71-100.	11.8	432
133	Role of the male BmDSX protein in the sexual differentiation of Bombyx mori. Evolution & Development, 2005, 7, 58-68.	2.0	102
134	Partial deletions of the W chromosome due to reciprocal translocation in the silkworm Bombyx mori. Insect Molecular Biology, 2005, 14, 339-352.	2.0	29
135	Characteristics of two genes encoding proteins with an ADAM-type metalloprotease domain, which are induced during the molting periods inBombyx mori. Archives of Insect Biochemistry and Physiology, 2005, 59, 91-98.	1.5	6
136	The Bmdsx transgene including trimmed introns is sex-specifically spliced in tissues of the silkworm, Bombyx mori. Journal of Insect Science, 2005, 5, 17.	1.5	5
137	Novel Macula-Like Virus Identified in Bombyx mori Cultured Cells. Journal of Virology, 2005, 79, 5577-5584.	3.4	75
138	Retrotransposable elements on the W chromosome of the silkworm, <i>Bombyx mori</i> . Cytogenetic and Genome Research, 2005, 110, 144-151.	1.1	83
139	Simple sequence repeat-based consensus linkage map of Bombyx mori. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 16303-16308.	7.1	108
140	A baculovirus-encoded protein tyrosine phosphatase gene induces enhanced locomotory activity in a lepidopteran host. Proceedings of the National Academy of Sciences of the United States of America, 2005, 102, 2584-2589.	7.1	142
141	TheBmdsxtransgene including trimmed introns is sex-specifically spliced in tissues of the silkworm, Bombyx mori. Journal of Insect Science, 2005, 5, 1-6.	0.9	4
142	Accurate pre-mRNA splicing using a nuclear extract from Bombyx mori fat body. Insect Biochemistry and Molecular Biology, 2005, 35, 257-261.	2.7	0
143	The BmChi-h gene, a bacterial-type chitinase gene of Bombyx mori, encodes a functional exochitinase that plays a role in the chitin degradation during the molting process. Insect Biochemistry and Molecular Biology, 2005, 35, 1112-1123.	2.7	59
144	Identification of molting fluid carboxypeptidase A (MF-CPA) in Bombyx mori. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2005, 141, 314-322.	1.6	63

#	Article	IF	CITATIONS
145	SilkSatDb: a microsatellite database of the silkworm, Bombyx mori. Nucleic Acids Research, 2004, 33, D403-D406.	14.5	29
146	Characterization of the Baculovirus Bombyx Mori Nucleopolyhedrovirus Gene Homologous to the Mammalian FGF Gene Family. Virus Genes, 2004, 29, 211-217.	1.6	40
147	Reduced cysteine protease activity of the hemolymph of Bombyx mori larvae infected with fp25K-inactivated Bombyx mori nucleopolyhedrovirus results in the reduced postmortem host degradation. Archives of Virology, 2004, 149, 1773-82.	2.1	13
148	Daily Expression Patterns of <i>Cycle</i> and <i>Clock</i> Genes in the Head of the Silkworm, <i>Bombyx Mori</i> . Biotechnology and Biotechnological Equipment, 2004, 18, 77-81.	1.3	3
149	The Genome Sequence of Silkworm, Bombyx mori. DNA Research, 2004, 11, 27-35.	3.4	594
150	Change in the expressed gene patterns of the wing disc during the metamorphosis of Bombyx mori. Gene, 2004, 343, 133-142.	2.2	22
151	Expression profiling of baculovirus genes in permissive and nonpermissive cell lines. Biochemical and Biophysical Research Communications, 2004, 323, 599-614.	2.1	67
152	Microarray analysis of gene expression profiles in wing discs of Bombyx mori during pupal ecdysis. Insect Biochemistry and Molecular Biology, 2004, 34, 775-784.	2.7	34
153	W-derived BAC probes as a new tool for identification of the W chromosome and its aberrations in Bombyx mori. Chromosoma, 2003, 112, 48-55.	2.2	67
154	Analysis of the biological functions of a doublesex homologue in Bombyx mori. Development Genes and Evolution, 2003, 213, 345-354.	0.9	120
155	Genomic sequence of a 320-kb segment of the Z chromosome of Bombyx mori containing a kettin ortholog. Molecular Genetics and Genomics, 2003, 269, 137-149.	2.1	70
156	Detachment analysis of the translocated W chromosome shows that the female-specific randomly amplified polymorphic DNA (RAPD) marker, Female-218, is derived from the second chromosome fragment region of the translocated W chromosome of the sex-limited pB silkworm (Bombyx mori) strain. Hereditas, 2003, 138, 148-153.	1.4	2
157	The construction of an EST database for Bombyx mori and its application. Proceedings of the National Academy of Sciences of the United States of America, 2003, 100, 14121-14126.	7.1	245
158	Cloning of Cyc (Bmal1) homolog in Bombyx mori: structural analysis and tissue specific distributions. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2003, 134, 535-542.	1.6	25
159	Isolation and comparison of different ecdysone-responsive cuticle protein genes in wing discs of Bombyx mori. Insect Biochemistry and Molecular Biology, 2003, 33, 671-679.	2.7	50
160	A Bombyx mori gene, BmChi-h, encodes a protein homologous to bacterial and baculovirus chitinases. Insect Biochemistry and Molecular Biology, 2003, 33, 749-759.	2.7	83
161	Annotation pattern of ESTs from Spodoptera frugiperda Sf9 cells and analysis of the ribosomal protein genes reveal insect-specific features and unexpectedly low codon usage bias. Bioinformatics, 2003, 19, 2343-2350.	4.1	36
162	Characterization of the kynurenine 3-monooxygenase gene corresponding to the white egg 1 mutant in the silkworm Bombyx mori. Molecular Genetics and Genomics, 2002, 267, 1-9.	2.1	49

#	Article	IF	CITATIONS
163	Nested retrotransposons on the W chromosome of the wild silkworm <i>Bombyx mandarina</i> . Insect Molecular Biology, 2002, 11, 307-314.	2.0	23
164	Characterization of acyl-CoA-binding protein (ACBP) in the pheromone gland of the silkworm, Bombyx mori. Insect Biochemistry and Molecular Biology, 2001, 31, 603-609.	2.7	47
165	Isolation and expression of the ecdysteroid-inducible angiotensin-converting enzyme-related gene in wing discs of Bombyx mori. Insect Biochemistry and Molecular Biology, 2001, 31, 97-103.	2.7	38
166	Mass isolation of cuticle protein cDNAs from wing discs of Bombyx mori and their characterizations. Insect Biochemistry and Molecular Biology, 2001, 31, 1019-1028.	2.7	41
167	The mechanism of sex-specific splicing at the doublesex gene is different between Drosophila melanogaster and Bombyx mori. Insect Biochemistry and Molecular Biology, 2001, 31, 1201-1211.	2.7	113
168	Isolation and expression of an ecdysteroid-inducible neutral endopeptidase 24.11-like gene in wing discs of Bombyx mori. Insect Biochemistry and Molecular Biology, 2001, 31, 1213-1219.	2.7	22
169	A homologue of the Drosophila doublesex gene is transcribed into sex-specific mRNA isoforms in the silkworm, Bombyx mori. Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology, 2001, 128, 145-158.	1.6	129
170	Two novel Pao-like retrotransposons (Kamikaze and Yamato) from the silkworm species Bombyx mori and B. mandarina: common structural features of Pao-like elements. Molecular Genetics and Genomics, 2001, 265, 375-385.	2.1	38
171	Comparative Expressed-Sequence-Tag Analysis of Differential Gene Expression Profiles in BmNPV-Infected BmN Cells. Virology, 2001, 282, 348-356.	2.4	27
172	Identification and genetic mapping of RAPD markers linked to the densonucleosis refractoriness gene, nsd-2, in the silkworm, Bombyx mori Genes and Genetic Systems, 2000, 75, 93-96.	0.7	16
173	Molecular structure of a novel gypsy-Ty3-like retrotransposon (Kabuki) and nested retrotransposable elements on the W chromosome of the silkworm Bombyx mori. Molecular Genetics and Genomics, 2000, 263, 916-924.	2.4	46
174	Identification and characterisation of a silkworm ABC transporter gene homologous to Drosophila white. Molecular Genetics and Genomics, 2000, 264, 11-19.	2.4	36
175	The Chitinase Gene of the Silkworm, Bombyx mori, Contains a Novel Tc-like Transposable Element. Journal of Biological Chemistry, 2000, 275, 37725-37732.	3.4	26
176	cDNA cloning of acyl-CoA desaturase homologs in the silkworm, Bombyx mori. Gene, 2000, 246, 339-345.	2.2	53
177	The influence of triploidy on gene expression in the silkworm, Bombyx mori. Heredity, 1999, 82, 661-667.	2.6	13
178	Bm kettin, homologue of the Drosophila kettin gene, is located on the Z chromosome in Bombyx mori and is not dosage compensated. Heredity, 1999, 82, 170-179.	2.6	47
179	Molecular characterization of baculovirus Bombyx mori nucleopolyhedrovirus polyhedron mutants. Archives of Virology, 1999, 144, 1275-1285.	2.1	22
180	Diapause-associated transcription of BmEts, a gene encoding an ETS transcription factor homolog in Bombyx mori. Insect Biochemistry and Molecular Biology, 1999, 29, 339-347.	2.7	22

#	Article	IF	CITATIONS
181	Absence of dosage compensation at the transcription level of a sex-linked gene in a female heterogametic insect, Bombyx mori. Heredity, 1998, 81, 275-283.	2.6	43
182	Genetic- mapping of RAPD markers linked to the densonucleosis refractoriness gene, nsd-1, in the silkworm, Bombyx mori Genes and Genetic Systems, 1998, 73, 237-242.	0.7	14
183	A complete full-length non-LTR retrotransposon, BMC1, on the W chromosome of the silkworm, Bombyx mori Genes and Genetic Systems, 1998, 73, 353-358.	0.7	30
184	Identification of novel random amplified polymorphic DNAs (RAPDs) on the W chromosome of the domesticated silkworm, Bombyx mori, and the wild silkworm, B. mandarina, and their retrotransposable elementrelated nucleotide sequences Genes and Genetic Systems, 1998, 73, 243-254.	0.7	48
185	Estimation of the position effect and action mode of a semi-lethal factor locus on a DNA polymorphism linkage map in silkworm, Bombyx mori Genes and Genetic Systems, 1998, 73, 337-343.	0.7	0
186	Molecular structure of the copia-like retrotransposable element Yokozuna on the W chromosome of the silkworm, Bombyx mori Genes and Genetic Systems, 1998, 73, 345-352.	0.7	24
187	Absence of dosage compensation at the transcription level of a sex-linked gene in a female heterogametic insect, Bombyx mori. Heredity, 1998, 81, 275-283.	2.6	2
188	Evolution of the genome in silkmoths. Japanese Journal of Biometrics, 1998, 19, S15-S27.	0.0	0
189	Alanine Scanning Mutagenesis of the Switch I Region in the ATPase Site ofDictyostelium discoideumMyosin Ilâ€. Biochemistry, 1997, 36, 14037-14043.	2.5	90
190	Chromosomal localization of amplified esterase genes in insecticide resistant Culex mosquitoes. Insect Biochemistry and Molecular Biology, 1996, 26, 853-857.	2.7	10
191	Effects of Depletion of T Cell Subpopulations on the Course of Infection and Anti-Parasite Delayed Type Hypersensitivity Response in Mice Infected with Babesia microti and Babesia rodhaini Journal of Veterinary Medical Science, 1996, 58, 343-347.	0.9	18
192	Nucleotide Sequence of the Random Amplified Polymorphic DNA (RAPD) on the W Chromosome of the Silkworm, Bombyx mori (Lepidoptera: Bombycidae). Applied Entomology and Zoology, 1996, 31, 633-637.	1.2	11
193	Clinical Efficacy of Lansoprazole-Amoxicillin Treatment in Eradicating Helicobacter pylori. Journal of Clinical Gastroenterology, 1995, 20, S100-S103.	2.2	3
194	Linkage analysis of the gene encoding precursor protein of diapause hormone and pheromone biosynthesis-activating neuropeptide in the silkmoth, Bombyx mori. Genetical Research, 1995, 65, 105-111.	0.9	29
195	Linkage map of random amplified polymorphic DNAs (RAPDs) in the silkworm, <i>Bombyx mori</i> . Genetical Research, 1995, 66, 1-7.	0.9	78
196	PCR-based detection of Wolbachia, cytoplasmic incompatibility microorganisms, infected in natural populations of Laodelphax striatellus (Homoptera: Delphacidae) in central Japan: has the distribution of Wolbachia spread recently?. Insect Molecular Biology, 1995, 4, 237-243.	2.0	64
197	Phylogenetic Relationship of Silkmoths Inferred from Sequence Data of the Arylphorin Gene. Molecular Phylogenetics and Evolution, 1995, 4, 223-234.	2.7	42
198	Polymorphism and linkage analysis of the prothoracicotropic hormone gene in the silkmoth, <i>Bombyx mori</i> . Genetical Research, 1994, 63, 189-195.	0.9	9

#	Article	IF	CITATIONS
199	Comparison of Arylphorins among Species belonging to Saturniidae and Bombycidae(Lepidoptera) by Immunodiffusion Analyses Japanese Journal of Applied Entomology and Zoology, 1992, 36, 119-125.	0.1	2
200	Detection and Discrimination of <i>Mycoplasma pneumoniae</i> and <i>Mycoplasma genitalium</i> by the <i>in vitro</i> DNA Amplification. Microbiology and Immunology, 1992, 36, 21-27.	1.4	10
201	Distribution of split 5.8S ribosomal RNA in Diptera. Insect Molecular Biology, 1992, 1, 45-48.	2.0	11
202	β-Agonists modulate ACh-inhibition of a K current in intestinal smooth muscle cells. Biochemical and Biophysical Research Communications, 1991, 179, 327-332.	2.1	3
203	Hormonal control of vitellogenin mRNA levels in the male and female housefly, Musca domestica. Journal of Insect Physiology, 1991, 37, 383-390.	2.0	16
204	Sequence of 5.8S ribosomal RNA in the mosquito,Culex tritaeniorhynchus. Nucleic Acids Research, 1991, 19, 5435-5435.	14.5	9
205	Electrooptic Bistability of Ferroelectric Liquid Crystal Cell Prepared Using Obliquely Evaporated SiO Films. Molecular Crystals and Liquid Crystals, 1991, 201, 133-136.	0.7	4
206	Trimebutine maleate has inhibitory effects on the voltage-dependent Ca2+ inward current and other membrane currents in intestinal smooth muscle cells. Gastroenterologia Japonica, 1990, 25, 175-179.	0.3	7
207	Na+-dependent elevation of the acidic cell surface pH (microclimate pH) of rat jejunal villus cells induced by cyclic nucleotides and phorbol ester: possible mediators of the regulation of the Na+/H+ antiporter. Biochimica Et Biophysica Acta - Biomembranes, 1988, 937, 328-334.	2.6	34
208	The detection of mosaics and polyploids in a hereditary mosaic strain of the silk moth, Bombyx mori, using egg colour mutants. Genetical Research, 1988, 51, 223-229.	0.9	6
209	Diapause of the inter-specific F1 hybrids between Antheraea yamamai (Guerin-Meneville) and A. pernyi (GM.) (Lepidoptera: Saturniidae) Japanese Journal of Applied Entomology and Zoology, 1988, 32, 120-125.	0.1	4
210	Factors affecting the microclimate pH in rat jejunum Journal of Physiology, 1987, 392, 113-127.	2.9	47
211	Characterization of Arylphorin of the Eri-silkmoth, Samia cynthia ricini (DONOVAN) : Lepidoptera : Saturniidae. Applied Entomology and Zoology, 1987, 22, 543-552.	1.2	8
212	Expression of homeotic genes inBombyx mori estimated from asymmetry of dorsal closure in mutant/normal mosaics. The Journal of Experimental Zoology, 1986, 240, 335-342.	1.4	8