

Hitoshi Ueda

List of Publications by Year in descending order

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17
papers

592
citations

759233

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888059

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17
all docs

17
docs citations

17
times ranked

617
citing authors

#	ARTICLE	IF	CITATIONS
1	Proteasome activity determines pupation timing through the degradation speed of timer molecule Blimp-1. <i>Development Growth and Differentiation</i> , 2018, 60, 502-508.	1.5	7
2	Protease resistance of porcine acidic mammalian chitinase under gastrointestinal conditions implies that chitin-containing organisms can be sustainable dietary resources. <i>Scientific Reports</i> , 2017, 7, 12963.	3.3	29
3	A biological timer in the fat body comprised of Blimp-1, FTZ-F1 and Shade regulates pupation timing in <i>Drosophila melanogaster</i> . <i>Development (Cambridge)</i> , 2016, 143, 2410-6.	2.5	14
4	Autocrine regulation of ecdysone synthesis by FTZ-F1 octopamine receptor in the prothoracic gland is essential for <i>Drosophila</i> metamorphosis. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2015, 112, 1452-1457.	7.1	50
5	Function of the nuclear receptor FTZ-F1 during the pupal stage in <i>Drosophila melanogaster</i> . <i>Development Growth and Differentiation</i> , 2014, 56, 245-253.	1.5	42
6	The Binding of Multiple Nuclear Receptors to a Single Regulatory Region Is Important for the Proper Expression of EDG84A in <i>Drosophila melanogaster</i> . <i>Journal of Molecular Biology</i> , 2013, 425, 71-81.	4.2	7
7	Regulatory mechanisms of ecdysone-inducible Blimp-1 encoding a transcriptional repressor that is important for the prepupal development in <i>Drosophila</i> . <i>Development Growth and Differentiation</i> , 2011, 53, 697-703.	1.5	17
8	<i>Drosophila</i> Blimp-1 Is a Transient Transcriptional Repressor That Controls Timing of the Ecdysone-Induced Developmental Pathway. <i>Molecular and Cellular Biology</i> , 2007, 27, 8739-8747.	2.3	49
9	Anterior epidermis-specific expression of the cuticle gene EDG84A is controlled by many cis-regulatory elements in <i>Drosophila melanogaster</i> . <i>Development Genes and Evolution</i> , 2005, 215, 545-552.	0.9	9
10	A Simple and Quick Method to Isolate Nuclear Extracts from Pupae of <i>Drosophila melanogaster</i> . <i>Cytotechnology</i> , 2005, 49, 67-70.	1.6	4
11	Temporal regulation of the mid-prepupal gene FTZ-F1: DHR3 early late gene product is one of the plural positive regulators. <i>Genes To Cells</i> , 2003, 2, 559-569.	1.2	59
12	FTZ-F1 dependent and independent activation of Edg78E, a pupal cuticle gene, during the early metamorphic period in <i>Drosophila melanogaster</i> . <i>Development Growth and Differentiation</i> , 2002, 44, 419-425.	1.5	37
13	Identification of the core domain and the secondary structure of the transcriptional coactivator MBF1. <i>Genes To Cells</i> , 1999, 4, 415-424.	1.2	17
14	Transcriptional activation through interaction of MBF2 with TFIIA. <i>Genes To Cells</i> , 1997, 2, 143-153.	1.2	25
15	Intermittent Expression of BmFTZ-F1, a Member of the Nuclear Hormone Receptor Superfamily during Development of the Silkworm <i>Bombyx mori</i> . <i>Developmental Biology</i> , 1994, 162, 426-437.	2.0	96
16	Defining the sequence recognized with BmFTZ-F1, a sequence specific DNA binding factor in the silkworm, <i>Bombyx mori</i> , as revealed by direct sequencing of bound oligonucleotides and gel mobility shift competition analysis. <i>Nucleic Acids Research</i> , 1991, 19, 3689-3693.	14.5	68
17	Identification and purification of a <i>Bombyx mori</i> homologue of FTZ-F1. <i>Nucleic Acids Research</i> , 1990, 18, 7229-7234.	14.5	62