

# Zilã; Luz Paulino Simões

## List of Publications by Year in descending order

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100  
papers

8,294  
citations

66343

42  
h-index

48315

88  
g-index

101  
all docs

101  
docs citations

101  
times ranked

6486  
citing authors

#	ARTICLE	IF	CITATIONS
1	Insights into social insects from the genome of the honeybee <i>Apis mellifera</i> . <i>Nature</i> , 2006, 443, 931-949.	27.8	1,648
2	Functional and Evolutionary Insights from the Genomes of Three Parasitoid <i>Nasonia</i> Species. <i>Science</i> , 2010, 327, 343-348.	12.6	808
3	Genomic signatures of evolutionary transitions from solitary to group living. <i>Science</i> , 2015, 348, 1139-1143.	12.6	357
4	The genomes of two key bumblebee species with primitive eusocial organization. <i>Genome Biology</i> , 2015, 16, 76.	8.8	330
5	Hormonal control of the yolk precursor vitellogenin regulates immune function and longevity in honeybees. <i>Experimental Gerontology</i> , 2004, 39, 767-773.	2.8	304
6	Vitellogenin regulates hormonal dynamics in the worker caste of a eusocial insect. <i>FEBS Letters</i> , 2005, 579, 4961-4965.	2.8	293
7	Validation of reference genes for gene expression studies in the honey bee, <i>Apis mellifera</i> , by quantitative real-time RT-PCR. <i>Apidologie</i> , 2008, 39, 372-385.	2.0	292
8	A worldwide survey of genome sequence variation provides insight into the evolutionary history of the honeybee <i>Apis mellifera</i> . <i>Nature Genetics</i> , 2014, 46, 1081-1088.	21.4	273
9	Disruption of vitellogenin gene function in adult honeybees by intra-abdominal injection of double-stranded RNA. <i>BMC Biotechnology</i> , 2003, 3, 1.	3.3	243
10	Molecular determinants of caste differentiation in the highly eusocial honeybee <i>Apis mellifera</i> . <i>BMC Developmental Biology</i> , 2007, 7, 70.	2.1	226
11	The vitellogenin of the honey bee, <i>Apis mellifera</i> : structural analysis of the cDNA and expression studies. <i>Insect Biochemistry and Molecular Biology</i> , 2003, 33, 459-465.	2.7	167
12	Inhibition of vitellogenin synthesis in <i>Apis mellifera</i> workers by a juvenile hormone analogue, pyriproxyfen. <i>Journal of Insect Physiology</i> , 2000, 46, 153-160.	2.0	160
13	Vitellogenin expression in queen ovaries and in larvae of both sexes of <i>Apis mellifera</i> . <i>Archives of Insect Biochemistry and Physiology</i> , 2005, 59, 211-218.	1.5	125
14	RNAi-mediated silencing of vitellogenin gene function turns honeybee ( <i>Apis mellifera</i> ) workers into extremely precocious foragers. <i>Die Naturwissenschaften</i> , 2008, 95, 953-961.	1.6	125
15	Effects of juvenile hormone and ecdysone on the timing of vitellogenin appearance in hemolymph of queen and worker pupae of <i>Apis mellifera</i> . <i>Journal of Insect Science</i> , 2002, 2, 1-8.	0.9	106
16	Identification of a juvenile hormone esterase-like gene in the honey bee, <i>Apis mellifera</i> L. Expression analysis and functional assays. <i>Comparative Biochemistry and Physiology - B Biochemistry and Molecular Biology</i> , 2008, 150, 33-44.	1.6	102
17	Developmental characterization, function and regulation of a Laccase2 encoding gene in the honey bee, <i>Apis mellifera</i> (Hymenoptera, Apinae). <i>Insect Biochemistry and Molecular Biology</i> , 2010, 40, 241-251.	2.7	95
18	The relationship between level of pollen in the diet, vitellogenin and juvenile hormone titres in Africanized <i>Apis mellifera</i> workers. <i>Journal of Apicultural Research</i> , 1996, 35, 27-36.	1.5	91

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19	The four hexamerin genes in the honey bee: structure, molecular evolution and function deduced from expression patterns in queens, workers and drones. <i>BMC Molecular Biology</i> , 2010, 11, 23.	3.0	89
20	Ecdysteroid titer and reproduction in queens and workers of the honey bee and of a stingless bee: loss of ecdysteroid function at increasing levels of sociality?. <i>Insect Biochemistry and Molecular Biology</i> , 2002, 32, 211-216.	2.7	88
21	Non-Target Effects of Green Fluorescent Protein (GFP)-Derived Double-Stranded RNA (dsRNA-GFP) Used in Honey Bee RNA Interference (RNAi) Assays. <i>Insects</i> , 2013, 4, 90-103.	2.2	85
22	Phenoloxidase activity in <i>Apis mellifera</i> honey bee pupae, and ecdysteroid-dependent expression of the prophenoloxidase mRNA. <i>Insect Biochemistry and Molecular Biology</i> , 2004, 34, 1257-1268.	2.7	80
23	Caste development and reproduction: a genome-wide analysis of hallmarks of insect eusociality. <i>Insect Molecular Biology</i> , 2006, 15, 703-714.	2.0	73
24	Expression analysis of putative vitellogenin and lipophorin receptors in honey bee ( <i>Apis mellifera</i> L.) queens and workers. <i>Journal of Insect Physiology</i> , 2008, 54, 1138-1147.	2.0	71
25	A non-invasive method for silencing gene transcription in honeybees maintained under natural conditions. <i>Insect Biochemistry and Molecular Biology</i> , 2009, 39, 157-160.	2.7	71
26	Ecdysteroid titers in pupae of highly social bees relate to distinct modes of caste development. <i>Journal of Insect Physiology</i> , 2002, 48, 783-790.	2.0	68
27	Molecular characterization of a cDNA encoding prophenoloxidase and its expression in <i>Apis mellifera</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2005, 35, 541-552.	2.7	68
28	Standard methods for physiology and biochemistry research in <i>Apis mellifera</i> . <i>Journal of Apicultural Research</i> , 2013, 52, 1-48.	1.5	65
29	Neuroigin-associated microRNA-932 targets actin and regulates memory in the honeybee. <i>Nature Communications</i> , 2014, 5, 5529.	12.8	65
30	Juvenile Hormone Biosynthesis Gene Expression in the corpora allata of Honey Bee ( <i>Apis mellifera</i> L.) Female Castes. <i>PLoS ONE</i> , 2014, 9, e86923.	2.5	64
31	Higher vitellogenin concentrations in honey bee workers may be an adaptation to life in temperate climates. <i>Insectes Sociaux</i> , 2005, 52, 316-319.	1.2	62
32	Dietary dopamine causes ovary activation in queenless <i>Apis mellifera</i> workers. <i>Apidologie</i> , 2003, 34, 281-289.	2.0	60
33	Bacterial infection activates the immune system response and dysregulates microRNA expression in honey bees. <i>Insect Biochemistry and Molecular Biology</i> , 2013, 43, 474-482.	2.7	55
34	The gene vitellogenin affects microRNA regulation in honey bee ( <i>Apis mellifera</i> ) fat body and brain. <i>Journal of Experimental Biology</i> , 2013, 216, 3724-32.	1.7	54
35	<i>Apis mellifera</i> ultraspiracle: cDNA sequence and rapid up-regulation by juvenile hormone. <i>Insect Molecular Biology</i> , 2004, 13, 459-467.	2.0	53
36	Hemocyte types and total and differential counts in unparasitized and parasitized <i>Anastrepha obliqua</i> (Diptera, Tephritidae) larvae. <i>Brazilian Journal of Biology</i> , 2002, 62, 689-699.	0.9	51

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37	Molecular cloning and expression of a hexamerin cDNA from the honey bee, <i>Apis mellifera</i> . <i>Journal of Insect Physiology</i> , 2005, 51, 1135-1147.	2.0	51
38	Developmental regulation of ecdysone receptor (EcR) and EcR-controlled gene expression during pharate-adult development of honeybees ( <i>Apis mellifera</i> ). <i>Frontiers in Genetics</i> , 2014, 5, 445.	2.3	49
39	Characterization and expression of theHex 110 gene encoding a glutamine-rich hexamerin in the honey bee, <i>Apis mellifera</i> . <i>Archives of Insect Biochemistry and Physiology</i> , 2006, 63, 57-72.	1.5	48
40	Trade-off between immune stimulation and expression of storage protein genes. <i>Archives of Insect Biochemistry and Physiology</i> , 2009, 71, 70-87.	1.5	47
41	A Honey Bee Hexamerin, HEX 70a, Is Likely to Play an Intranuclear Role in Developing and Mature Ovarioles and Testioles. <i>PLoS ONE</i> , 2011, 6, e29006.	2.5	47
42	The juvenile hormone (JH) epoxide hydrolase gene in the honey bee ( <i>Apis mellifera</i> ) genome encodes a protein which has negligible participation in JH degradation. <i>Journal of Insect Physiology</i> , 2010, 56, 1139-1146.	2.0	45
43	Non-coding transcription characterization and annotation. <i>RNA Biology</i> , 2012, 9, 274-282.	3.1	45
44	Morphometric and genetic changes in a population of <i>Apis mellifera</i> after 34 years of Africanization. <i>Genetics and Molecular Research</i> , 2009, 8, 709-717.	0.2	45
45	Genomewide analysis of admixture and adaptation in the Africanized honeybee. <i>Molecular Ecology</i> , 2017, 26, 3603-3617.	3.9	44
46	Transcriptome Analysis of Honeybee ( <i>Apis Mellifera</i> ) Haploid and Diploid Embryos Reveals Early Zygotic Transcription during Cleavage. <i>PLoS ONE</i> , 2016, 11, e0146447.	2.5	43
47	A honeybee storage protein gene, hex 70a, expressed in developing gonads and nutritionally regulated in adult fat body. <i>Journal of Insect Physiology</i> , 2008, 54, 867-877.	2.0	42
48	Honey bee ( <i>Apis mellifera</i> ) transferrin-gene structure and the role of ecdysteroids in the developmental regulation of its expression. <i>Insect Biochemistry and Molecular Biology</i> , 2004, 34, 415-424.	2.7	40
49	MicroRNA signatures characterizing caste-independent ovarian activity in queen and worker honeybees ( <i>Apis mellifera</i> L.). <i>Insect Molecular Biology</i> , 2016, 25, 216-226.	2.0	39
50	Evidence of at least two evolutionary lineages in <i>Melipona subnitida</i> (Apidae, Meliponini) suggested by mtDNA variability and geometric morphometrics of forewings. <i>Die Naturwissenschaften</i> , 2014, 101, 17-24.	1.6	38
51	Protein profiles of testes, seminal vesicles and accessory glands of honey bee pupae and their relation to the ecdysteroid titer. <i>Apidologie</i> , 2007, 38, 1-11.	2.0	36
52	Downregulation of ultraspiracle gene expression delays pupal development in honeybees. <i>Journal of Insect Physiology</i> , 2008, 54, 1035-1040.	2.0	34
53	Ecdysteroid-Dependent Expression of the Tweedle and Peroxidase Genes during Adult Cuticle Formation in the Honey Bee, <i>Apis mellifera</i> . <i>PLoS ONE</i> , 2011, 6, e20513.	2.5	33
54	A cuticle protein gene in the honeybee: Expression during development and in relation to the ecdysteroid titer. <i>Insect Biochemistry and Molecular Biology</i> , 2007, 37, 1272-1282.	2.7	32

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55	Silencing of <i>Apis mellifera</i> dorsal genes reveals their role in expression of the antimicrobial peptide defensin. <i>Insect Molecular Biology</i> , 2018, 27, 577-589.	2.0	31
56	A study of the constitutive heterochromatin and nucleolus organizer regions of <i>Isocopris inhiata</i> and <i>Diabroctis mimas</i> (Coleoptera: Scarabaeidae, Scarabaeinae) using C-banding, AgNO <sub>3</sub> staining and FISH techniques. <i>Genetics and Molecular Biology</i> , 2005, 28, 111-116.	1.3	30
57	Nutritionally Driven Differential Gene Expression Leads to Heterochronic Brain Development in Honeybee Castes. <i>PLoS ONE</i> , 2013, 8, e64815.	2.5	29
58	Evaluation of reference genes for gene expression analysis by real-time quantitative PCR (qPCR) in three stingless bee species (Hymenoptera: Apidae: Meliponini). <i>Scientific Reports</i> , 2019, 9, 17692.	3.3	26
59	Hormonal control and target genes of <i>ftz</i> expression in the honeybee <i>Apis mellifera</i> : a positive loop linking juvenile hormone, <i>ftz</i> , and <i>vitellogenin</i> . <i>Insect Molecular Biology</i> , 2019, 28, 145-159.	2.0	26
60	Vitellogenin gene expression in stingless bee workers differing in egg-laying behavior. <i>Insectes Sociaux</i> , 2007, 54, 70-76.	1.2	24
61	Farnesoic acid O-methyl transferase (FAMeT) isoforms: Conserved traits and gene expression patterns related to caste differentiation in the stingless bee, <i>Melipona scutellaris</i> . <i>Archives of Insect Biochemistry and Physiology</i> , 2008, 67, 97-106.	1.5	24
62	Methyl farnesoate epoxidase (mfe) gene expression and juvenile hormone titers in the life cycle of a highly eusocial stingless bee, <i>Melipona scutellaris</i> . <i>Journal of Insect Physiology</i> , 2017, 101, 185-194.	2.0	24
63	The use of Open Reading frame ESTs (ORESTES) for analysis of the honey bee transcriptome. <i>BMC Genomics</i> , 2004, 5, 84.	2.8	21
64	MicroRNA-34 directly targets pair-rule genes and cytoskeleton component in the honey bee. <i>Scientific Reports</i> , 2017, 7, 40884.	3.3	21
65	Organization, evolution and transcriptional profile of hexamerin genes of the parasitic wasp <i>Nasonia vitripennis</i> (Hymenoptera: Pteromalidae). <i>Insect Molecular Biology</i> , 2010, 19, 137-146.	2.0	20
66	Immunosenescence in honey bees ( <i>Apis mellifera</i> L.) is caused by intrinsic senescence and behavioral physiology. <i>Experimental Gerontology</i> , 2019, 119, 174-183.	2.8	20
67	Apoptosis Process in Mouse Leydig Cells during Postnatal Development. <i>Microscopy and Microanalysis</i> , 2003, 9, 68-73.	0.4	19
68	Exploring integument transcriptomes, cuticle ultrastructure, and cuticular hydrocarbons profiles in eusocial and solitary bee species displaying heterochronic adult cuticle maturation. <i>PLoS ONE</i> , 2019, 14, e0213796.	2.5	19
69	Hox Gene Expression Leads to Differential Hind Leg Development between Honeybee Castes. <i>PLoS ONE</i> , 2012, 7, e40111.	2.5	17
70	Comparative cytogenetics of three species of <i>Dichotomius</i> (Coleoptera, Scarabaeidae). <i>Genetics and Molecular Biology</i> , 2009, 32, 276-280.	1.3	15
71	Mars is close to venus – Female reproductive proteins are expressed in the fat body and reproductive tract of honey bee ( <i>Apis mellifera</i> L.) drones. <i>Journal of Insect Physiology</i> , 2010, 56, 1638-1644.	2.0	15
72	The nuclear and mitochondrial genomes of <i>Frieseomelitta varia</i> – a highly eusocial stingless bee (Meliponini) with a permanently sterile worker caste. <i>BMC Genomics</i> , 2020, 21, 386.	2.8	15

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73	The stomatogastric nervous system of the honey bee ( <i>Apis mellifera</i> ) in a critical phase of caste development. , 1998, 236, 139-149.		14
74	Active genic machinery for epigenetic <sc>RNA</sc> modifications in bees. <i>Insect Molecular Biology</i> , 2021, 30, 566-579.	2.0	14
75	Potential costs of bacterial infection on storage protein gene expression and reproduction in queenless <i>Apis mellifera</i> worker bees on distinct dietary regimes. <i>Journal of Insect Physiology</i> , 2012, 58, 1217-1225.	2.0	13
76	<sc>miRNA</sc>â€³4 and <sc>miRNA</sc>â€²10 target hexamerin genes enhancing their differential expression during early brain development of honeybee (<sc><i>Apis mellifera</i></sc>) castes. <i>Insect Molecular Biology</i> , 2021, 30, 594-604.	2.0	13
77	Effect of proctolin on the egg-laying activity of <i>Apis mellifera</i> queens. <i>Journal of Apicultural Research</i> , 2003, 42, 35-38.	1.5	10
78	Hormone-dependent protein patterns in integument and cuticular pigmentation in <i>Apis mellifera</i> during pharate adult development. <i>Journal of Insect Physiology</i> , 2001, 47, 1275-1282.	2.0	9
79	Immunity and physiological changes in adult honey bees ( <i>Apis mellifera</i> ) infected with <i>Nosema ceranae</i> : The natural colony environment. <i>Journal of Insect Physiology</i> , 2021, 131, 104237.	2.0	8
80	Occurrence of a prothoracicotropic hormone-like peptide in the developing nervous system of the honey bee ( <i>Apis mellifera</i> L). <i>Apidologie</i> , 1997, 28, 399-409.	2.0	8
81	Ovos produzidos por rainhas e operÃ¡rias de <i>Scaptotrigona depilis</i> (Hymenoptera, Apidae, Meliponina): morfometria e aspectos relacionados. <i>Iheringia - Serie Zoologia</i> , 2006, 96, 89-93.	0.5	7
82	Rapid method for DNA extraction from the honey bee <i>Apis mellifera</i> and the parasitic bee mite <i>Varroa destructor</i> using lysis buffer and proteinase K. <i>Genetics and Molecular Research</i> , 2013, 12, 4846-4854.	0.2	7
83	Casteâ€­specific gene expression underlying the differential adult brain development in the honeybee <sc><i>Apis mellifera</i></sc>. <i>Insect Molecular Biology</i> , 2021, 30, 42-56.	2.0	7
84	Transcriptome dynamics during metamorphosis of imaginal discs into wings and thoracic dorsum in <i>Apis mellifera</i> castes. <i>BMC Genomics</i> , 2021, 22, 756.	2.8	7
85	Mispatterning in the ommatidia of <i>Apis mellifera</i> pupae treated with a juvenile hormone analogue. <i>Journal of Morphology</i> , 2001, 249, 89-99.	1.2	6
86	Microsatellite loci for the carpenter bee <i>Xylocopa frontalis</i> (Apidae, Xylocopini). <i>Conservation Genetics Resources</i> , 2012, 4, 315-317.	0.8	6
87	Reproductive capacity and castes in eusocial stingless bees (Hymenoptera: Apidae). <i>Current Opinion in Insect Science</i> , 2019, 31, 20-28.	4.4	6
88	Vitellogenin content in fat body and ovary homogenates of workers and queens of <i>Melipona quadrifasciata anthidioides</i> during vitellogenesis. <i>Insect Science</i> , 2012, 19, 213-219.	3.0	5
89	Molecular underpinnings of the early brain developmental response to differential feeding in the honey bee <i>Apis mellifera</i> . <i>Biochimica Et Biophysica Acta - Gene Regulatory Mechanisms</i> , 2021, 1864, 194732.	1.9	5
90	<i>Ultraspiracle</i> of the stingless bees <i>Melipona scutellaris</i> and <i>Scaptotrigona depilis</i>: cDNA sequence and expression profiles during pupal development. <i>Apidologie</i> , 2007, 38, 462-471.	2.0	4

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91	SSD - a free software for designing multimeric mono-, bi- and trivalent shRNAs. <i>Genetics and Molecular Biology</i> , 2020, 43, e20190300.	1.3	4
92	Higher fertility of queenless workers in the Africanized honey bee. <i>Insectes Sociaux</i> , 1998, 45, 473-476.	1.2	3
93	Effects of treatment of the fat body trophocytes of <i>Melipona quadrifasciata anthidioides</i> nurse workers and virgin queens in culture by juvenile hormone III and ecdysterone (20â€œHE). <i>Microscopy Research and Technique</i> , 2013, 76, 20-27.	2.2	3
94	Circadian clock genes are differentially modulated during the daily cycles and chronological age in the social honeybee ( <i>Apis mellifera</i> ). <i>Apidologie</i> , 2018, 49, 71-83.	2.0	3
95	Vitellogenin of the solitary bees <i>Centris tarsata</i> and <i>Centris analis</i> (Hymenoptera: Apidae): cDNA structural analysis and gene expression. <i>Apidologie</i> , 2021, 52, 292-307.	2.0	3
96	In vitro secretion of ecdysteroid-dependent proteins and of a 70 kDa subunit reactive to anti-prophenoloxidase serum by <i>Apis mellifera</i> integument. <i>Apidologie</i> , 2003, 34, 377-388.	2.0	3
97	Worker bees ( <i>Apis mellifera</i> ) deprived of pollen in the first week of adulthood exhibit signs of premature aging. <i>Insect Biochemistry and Molecular Biology</i> , 2022, 146, 103774.	2.7	3
98	07-P001 Differential hind leg development in <i>Apis mellifera</i> castes. <i>Mechanisms of Development</i> , 2009, 126, S137.	1.7	1
99	Recombination mapping of the Brazilian stingless bee <i>Frieseomelitta varia</i> confirms high recombination rates in social hymenoptera. <i>BMC Genomics</i> , 2021, 22, 673.	2.8	1
100	07-P011 Differential expression of cuticle protein genes during metamorphosis of the honeybee, <i>Apis mellifera</i> . <i>Mechanisms of Development</i> , 2009, 126, S139-S140.	1.7	0