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List of Publications by Year in descending order

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#	Article	IF	CITATIONS
1	Comparative pharmacological characterization of D1-like dopamine receptors from Anopheles gambiae, Aedes aegypti and Culex quinquefasciatus suggests pleiotropic signaling in mosquito vector lineages. Parasites and Vectors, 2016, 9, 192.	2.5	15
2	Genomic insights into the Ixodes scapularis tick vector of Lyme disease. Nature Communications, 2016, 7, 10507.	12.8	450
3	Evaluation of <i>Aa</i> DOP2 Receptor Antagonists Reveals Antidepressants and Antipsychotics as Novel Lead Molecules for Control of the Yellow Fever Mosquito, <i>Aedes aegypti</i> . Journal of Pharmacology and Experimental Therapeutics, 2015, 352, 53-60.	2.5	12
4	Dopamine Receptor Antagonists as New Mode-of-Action Insecticide Leads for Control of Aedes and Culex Mosquito Vectors. PLoS Neglected Tropical Diseases, 2015, 9, e0003515.	3.0	34
5	Re-invigorating the insecticide discovery pipeline for vector control: GPCRs as targets for the identification of next gen insecticides. Pesticide Biochemistry and Physiology, 2013, 106, 141-148.	3.6	28
6	Transcriptome sequencing and annotation of the predatory mite Metaseiulus occidentalis (Acari:) Tj ETQq0 0 0 r Applied Acarology, 2013, 59, 283-296.	gBT /Over 1.6	lock 10 Tf 50 32
7	A "Genome-to-Lead―Approach for Insecticide Discovery: Pharmacological Characterization and Screening of Aedes aegypti D1-like Dopamine Receptors. PLoS Neglected Tropical Diseases, 2012, 6, e1478.	3.0	69
8	Observations on the entomopathogenic fungus Hirsutella citriformis attacking adult Diaphorina citri (Hemiptera: Psyllidae) in a managed citrus grove. BioControl, 2012, 57, 663-675.	2.0	40
9	Discovery of antagonists of tick dopamine receptors via chemical library screening and comparative pharmacological analyses. Insect Biochemistry and Molecular Biology, 2012, 42, 846-853.	2.7	20
10	Molecular and pharmacological characterization of two D1-like dopamine receptors in the Lyme disease vector, lxodes scapularis. Insect Biochemistry and Molecular Biology, 2011, 41, 563-571.	2.7	24
11	Genome organization of major tandem repeats in the hard tick, Ixodes scapularis. Chromosome Research, 2010, 18, 357-370.	2.2	26
12	Genome sequences of the human body louse and its primary endosymbiont provide insights into the permanent parasitic lifestyle. Proceedings of the National Academy of Sciences of the United States of America, 2010, 107, 12168-12173.	7.1	482
13	Isolation and characterization of an Isaria fumosorosea isolate infecting the Asian citrus psyllid in Florida. Journal of Invertebrate Pathology, 2008, 99, 96-102.	3.2	55
14	Morphological and molecular characterization of a Hirsutella species infecting the Asian citrus psyllid, Diaphorina citri Kuwayama (Hemiptera: Psyllidae), in Florida. Journal of Invertebrate Pathology, 2007, 95, 101-109.	3.2	66