

W Augustine Dunn

List of Publications by Year in descending order

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12
papers

782
citations

933447

10
h-index

1199594

12
g-index

12
all docs

12
docs citations

12
times ranked

1181
citing authors

#	ARTICLE	IF	CITATIONS
1	Complex Modulation of the <i>Aedes aegypti</i> Transcriptome in Response to Dengue Virus Infection. <i>PLoS ONE</i> , 2012, 7, e50512.	2.5	138
2	RNA-seq analyses of blood-induced changes in gene expression in the mosquito vector species, <i>Aedes aegypti</i> . <i>BMC Genomics</i> , 2011, 12, 82.	2.8	133
3	Annotation and analysis of a large cuticular protein family with the R&R Consensus in <i>Anopheles gambiae</i> . <i>BMC Genomics</i> , 2008, 9, 22.	2.8	102
4	Developmental expression patterns of cuticular protein genes with the R&R Consensus from <i>Anopheles gambiae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2008, 38, 508-519.	2.7	101
5	CPF and CPFL, two related gene families encoding cuticular proteins of <i>Anopheles gambiae</i> and other insects. <i>Insect Biochemistry and Molecular Biology</i> , 2007, 37, 675-688.	2.7	84
6	Proteomic analysis of cast cuticles from <i>Anopheles gambiae</i> by tandem mass spectrometry. <i>Insect Biochemistry and Molecular Biology</i> , 2007, 37, 135-146.	2.7	81
7	Comparative genomics allows the discovery of cis-regulatory elements in mosquitoes. <i>Proceedings of the National Academy of Sciences of the United States of America</i> , 2009, 106, 3053-3058.	7.1	45
8	Strain Variation in the Transcriptome of the Dengue Fever Vector, <i>Aedes aegypti</i> . <i>G3: Genes, Genomes, Genetics</i> , 2012, 2, 103-114.	1.8	36
9	Changes in transcript abundance for cuticular proteins and other genes three hours after a blood meal in <i>Anopheles gambiae</i> . <i>Insect Biochemistry and Molecular Biology</i> , 2014, 44, 33-43.	2.7	28
10	An algorithm for the classification of mRNA patterns in eosinophilic esophagitis: Integration of machine learning. <i>Journal of Allergy and Clinical Immunology</i> , 2018, 141, 1354-1364.e9.	2.9	22
11	Mucosal Gene Expression in Pediatric and Adult Patients With Ulcerative Colitis Permits Modeling of Ideal Biopsy Collection Strategy for Transcriptomic Analysis. <i>Inflammatory Bowel Diseases</i> , 2018, 24, 2565-2578.	1.9	10
12	Genome annotation improvements from cross-phyla proteogenomics and time-of-day differences in malaria mosquito proteins using untargeted quantitative proteomics. <i>PLoS ONE</i> , 2019, 14, e0220225.	2.5	2