Reza Salavati

List of Publications by Year in descending order

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933447 1058476 25 253 10 14 citations h-index g-index papers 25 25 25 371 all docs docs citations times ranked citing authors

#	Article	IF	CITATIONS
1	Global identification of conserved post-transcriptional regulatory programs in trypanosomatids. Nucleic Acids Research, 2013, 41, 8591-8600.	14.5	28
2	Insights into the insect salivary gland proteome: Diet-associated changes in caterpillar labial salivary proteins. Journal of Insect Physiology, 2013, 59, 351-366.	2.0	25
3	The DRBD13 RNA binding protein is involved in the insectâ€stage differentiation process of <i>Trypanosoma brucei</i> . FEBS Letters, 2015, 589, 1966-1974.	2.8	24
4	A Protein Complex Map of Trypanosoma brucei. PLoS Neglected Tropical Diseases, 2016, 10, e0004533.	3.0	19
5	Tail characteristics of Trypanosoma brucei mitochondrial transcripts are developmentally altered in a transcript-specific manner. International Journal for Parasitology, 2018, 48, 179-189.	3.1	19
6	Inhibitors of RNA editing as potential chemotherapeutics against trypanosomatid pathogens. International Journal for Parasitology: Drugs and Drug Resistance, 2012, 2, 36-46.	3.4	17
7	TrypsNetDB: An integrated framework for the functional characterization of trypanosomatid proteins. PLoS Neglected Tropical Diseases, 2017, 11, e0005368.	3.0	15
8	circTAIL-seq, a targeted method for deep analysis of RNA 3′ tails, reveals transcript-specific differences by multiple metrics. Rna, 2016, 22, 477-486.	3.5	14
9	The RNA binding activity of the first identified trypanosome protein with Z-DNA-binding domains. Scientific Reports, 2019, 9, 5904.	3.3	13
10	Morphometric Analysis of Larval Rostellar Hooks in Taenia multiceps of Sheep in Iran and Its Association with Mitochondrial Gene Variability. Iranian Journal of Parasitology, 2013, 8, 579-85.	0.6	12
11	The interaction of a Trypanosoma brucei KH-domain protein with a ribonuclease is implicated in ribosome processing. Molecular and Biochemical Parasitology, 2017, 211, 94-103.	1.1	11
12	Sequence-based functional annotation: what if most of the genes are unique to a genome?. Trends in Parasitology, 2010, 26, 225-229.	3.3	9
13	Pilot-Scale Compound Screening against RNA Editing Identifies Trypanocidal Agents. Journal of Biomolecular Screening, 2015, 20, 92-100.	2.6	9
14	Functional Genome Annotation by Combined Analysis across Microarray Studies of Trypanosoma brucei. PLoS Neglected Tropical Diseases, 2010, 4, e810.	3.0	7
15	The Oligonucleotide Binding (OB)-Fold Domain of KREPA4 Is Essential for Stable Incorporation into Editosomes. PLoS ONE, 2012, 7, e46864.	2.5	7
16	Deciphering RNA regulatory elements in trypanosomatids: one piece at a time or genome-wide?. Trends in Parasitology, 2014, 30, 234-240.	3.3	7
17	Sulfonated inhibitors of the RNA editing ligases validate the essential role of the MRP1/2 proteins in kinetoplastid RNA editing. Rna, 2020, 26, 827-835.	3.5	5
18	Mutational Analysis of Trypanosoma brucei RNA Editing Ligase Reveals Regions Critical for Interaction with KREPA2. PLoS ONE, 2015, 10, e0120844.	2.5	4

#	Article	IF	CITATION
19	Kinetoplastid RNA editing ligases 1 and 2 exhibit different electrostatic properties. Journal of Molecular Modeling, 2010, 16, 61-76.	1.8	3
20	Deciphering RNA Regulatory Elements Involved in the Developmental and Environmental Gene Regulation of Trypanosoma brucei. PLoS ONE, 2015, 10, e0142342.	2.5	2
21	Genetic and morphometric categorization of Taenia ovisfrom Sheep in Iran. Parasitology, 2019, 146, 563-568.	1.5	2
22	RNA Catalyst as a Reporter for Screening Drugs against RNA Editing in Trypanosomes. Journal of Visualized Experiments, 2014, , .	0.3	1
23	Structural Studies of <i>Trypanosoma brucei</i> RNA Editing Ligases and Their Binding Partner Proteins. Biochemistry, 2016, 55, 2319-2331.	2.5	O
24	Gene Function Discovery for Kinetoplastid Pathogens. Trends in Parasitology, 2019, 35, 8-12.	3.3	0
25	Characterization of putative mitochondrial processing peptidase αâ€subunit in Trypanosoma brucei. FASEB Journal, 2013, 27, 987.1.	0.5	O