

Yukihiro Kinjo

List of Publications by Year in descending order

Source: <https://exaly.com/author-pdf/3987710/publications.pdf>

Version: 2024-02-01

8
papers

179
citations

1478505

6
h-index

1588992

8
g-index

9
all docs

9
docs citations

9
times ranked

203
citing authors

#	ARTICLE	IF	CITATIONS
1	Vicariance and dispersal events inferred from mitochondrial genomes and nuclear genes (18S, 28S) shaped global <i>Cryptocercus</i> distributions. <i>Molecular Phylogenetics and Evolution</i> , 2022, 166, 107318.	2.7	4
2	The functional evolution of termite gut microbiota. <i>Microbiome</i> , 2022, 10, .	11.1	35
3	Digging deep: a revised phylogeny of Australian burrowing cockroaches (Blaberidae: Panesthiinae,) Tj ETQq1 1 0.784314 rgBT /Overlo evolution of burrowing. <i>Systematic Entomology</i> , 2021, 46, 767-783.	3.9	9
4	Enhanced Mutation Rate, Relaxed Selection, and the "Domino Effect" are associated with Gene Loss in <i>Blattabacterium</i> , A Cockroach Endosymbiont. <i>Molecular Biology and Evolution</i> , 2021, 38, 3820-3831.	8.9	13
5	Increased Mutation Rate Is Linked to Genome Reduction in Prokaryotes. <i>Current Biology</i> , 2020, 30, 3848-3855.e4.	3.9	44
6	Parallel and Gradual Genome Erosion in the <i>Blattabacterium</i> Endosymbionts of <i>Mastotermes darwiniensis</i> and <i>Cryptocercus</i> Wood Roaches. <i>Genome Biology and Evolution</i> , 2018, 10, 1622-1630.	2.5	14
7	An Efficient Strategy Developed for Next-Generation Sequencing of Endosymbiont Genomes Performed Using Crude DNA Isolated from Host Tissues: A Case Study of <i>Blattabacterium cuenoti</i> Inhabiting the Fat Bodies of Cockroaches. <i>Microbes and Environments</i> , 2015, 30, 208-220.	1.6	14
8	Maintenance of essential amino acid synthesis pathways in the <i>Blattabacterium cuenoti</i> symbiont of a wood-feeding cockroach. <i>Biology Letters</i> , 2013, 9, 20121153.	2.3	45