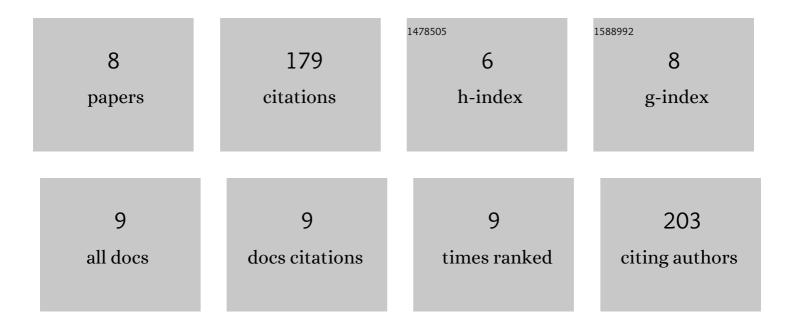
Yukihiro Kinjo

List of Publications by Year in descending order

Source: https://exaly.com/author-pdf/3987710/publications.pdf Version: 2024-02-01



#	Article	IF	CITATIONS
1	Vicariance and dispersal events inferred from mitochondrial genomes and nuclear genes (18S, 28S) shaped global Cryptocercus distributions. Molecular Phylogenetics and Evolution, 2022, 166, 107318.	2.7	4
2	The functional evolution of termite gut microbiota. Microbiome, 2022, 10, .	11.1	35
3	Digging deep: a revised phylogeny of Australian burrowing cockroaches (Blaberidae: Panesthiinae,) Tj ETQq1 1 0. evolution of burrowing. Systematic Entomology, 2021, 46, 767-783.	784314 rg 3.9	BT /Overloc 9
4	Enhanced Mutation Rate, Relaxed Selection, and the "Domino Effect―are associated with Gene Loss in <i>Blattabacterium</i> , A Cockroach Endosymbiont. Molecular Biology and Evolution, 2021, 38, 3820-3831.	8.9	13
5	Increased Mutation Rate Is Linked to Genome Reduction in Prokaryotes. Current Biology, 2020, 30, 3848-3855.e4.	3.9	44
6	Parallel and Gradual Genome Erosion in the Blattabacterium Endosymbionts of Mastotermes darwiniensis and Cryptocercus Wood Roaches. Genome Biology and Evolution, 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. Microbes and Environments, 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. Biology Letters, 2013, 9, 20121153.	2.3	45