

Alejandro Manzano-MarÃ-n

List of Publications by Year in descending order

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Version: 2024-02-01

22
papers

1,150
citations

516710
16
h-index

677142
22
g-index

33
all docs

33
docs citations

33
times ranked

889
citing authors

#	ARTICLE	IF	CITATIONS
1	Evolutionarily recent dual obligatory symbiosis among adelgids indicates a transition between fungus- and insect-associated lifestyles. ISME Journal, 2022, 16, 247-256.	9.8	16
2	Human Follicular Mites: Ectoparasites Becoming Symbionts. Molecular Biology and Evolution, 2022, 39, .	8.9	6
3	Multiresistant Enterobacteriaceae in yellow-legged gull chicks in their first weeks of life. Ecology and Evolution, 2022, 12, .	1.9	4
4	Cytoplasmic incompatibility between Old and New World populations of a tramp ant. Evolution; International Journal of Organic Evolution, 2021, 75, 1775-1791.	2.3	13
5	Serial horizontal transfer of vitamin-biosynthetic genes enables the establishment of new nutritional symbionts in aphidsâ€™ di-symbiotic systems. ISME Journal, 2020, 14, 259-273.	9.8	79
6	Mitogenome of the blood feeding leech <i>Haementeria acuecueyetzin</i> (Hirudinida: Glossiphoniidae) from Tabasco, Mexico. Mitochondrial DNA Part B: Resources, 2020, 5, 3310-3312.	0.4	2
7	The Protector within: Comparative Genomics of APSE Phages across Aphids Reveals Rampant Recombination and Diverse Toxin Arsenals. Genome Biology and Evolution, 2020, 12, 878-889.	2.5	22
8	Draft genome of the European medicinal leech <i>Hirudo medicinalis</i> (Annelida, Clitellata,) Tj ETQq0 O O rgBT /Overlock 10 Tf 50_462 Td (Hi	3.3	27
9	No evidence for Wolbachia as a nutritional co-obligate endosymbiont in the aphid <i>Pentalonia nigronervosa</i> . Microbiome, 2020, 8, 72.	11.1	10
10	Cultivation-assisted genome of <i>Candidatus Fukatsuia symbiotica</i> ; the enigmatic â€˜X-typeâ€™ symbiont of aphids. Genome Biology and Evolution, 2019, 11, 3510-3522.	2.5	23
11	A Freeloader? The Highly Eroded Yet Large Genome of the <i>Serratia symbiotica</i> Symbiont of <i>Cinara strobi</i> . Genome Biology and Evolution, 2018, 10, 2178-2189.	2.5	29
12	Happens in the best of subfamilies: establishment and repeated replacements of coâ€¢oblige secondary endosymbionts within Lachninae aphids. Environmental Microbiology, 2017, 19, 393-408.	3.8	80
13	<i>Buchnera</i> has changed flatmate but the repeated replacement of coâ€¢oblige symbionts is not associated with the ecological expansions of their aphid hosts. Molecular Ecology, 2017, 26, 2363-2378.	3.9	103
14	Dissecting genome reduction and trait loss in insect endosymbionts. Annals of the New York Academy of Sciences, 2017, 1389, 52-75.	3.8	87
15	Comparative Mitogenomics of Leeches (Annelida: Clitellata): Genome Conservation and Placobdella-Specific trnD Gene Duplication. PLoS ONE, 2016, 11, e0155441.	2.5	18
16	Snapshots of a shrinking partner: Genome reduction in <i>Serratia symbiotica</i> . Scientific Reports, 2016, 6, 32590.	3.3	68
17	Reinventing the Wheel and Making It Round Again: Evolutionary Convergence in<i>Buchnera</i>â€“<i>Serratia</i> Symbiotic Consortia between the Distantly Related Lachninae Aphids<i>Tuberolachnus salignus</i>and<i>Cinara cedri</i>. Genome Biology and Evolution, 2016, 8, 1440-1458.	2.5	85
18	A novel intracellular mutualistic bacterium in the invasive ant <i>Cardiocondyla obscurior</i>. ISME Journal, 2016, 10, 376-388.	9.8	67

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19	Solving a Bloody Mess: B-Vitamin Independent Metabolic Convergence among Gammaproteobacterial Obligate Endosymbionts from Blood-Feeding Arthropods and the Leech <i>Haementeria officinalis</i> . <i>Genome Biology and Evolution</i> , 2015, 7, 2871-2884.	2.5	70
20	Settling Down: The Genome of <i>Serratia symbiotica</i> from the Aphid <i>Cinara tujafilina</i> Zooms in on the Process of Accommodation to a Cooperative Intracellular Life. <i>Genome Biology and Evolution</i> , 2014, 6, 1683-1698.	2.5	88
21	Comparative Genomics of <i>Serratia</i> spp.: Two Paths towards Endosymbiotic Life. <i>PLoS ONE</i> , 2012, 7, e47274.	2.5	29
22	<i>Serratia symbiotica</i> from the Aphid <i>Cinara cedri</i> : A Missing Link from Facultative to Obligate Insect Endosymbiont. <i>PLoS Genetics</i> , 2011, 7, e1002357.	3.5	208