Ryo Miyazaki

List of Publications by Citations

Source: https://exaly.com/author-pdf/9128826/ryo-miyazaki-publications-by-citations.pdf

Version: 2024-04-28

This document has been generated based on the publications and citations recorded by exaly.com. For the latest version of this publication list, visit the link given above.

The third column is the impact factor (IF) of the journal, and the fourth column is the number of citations of the article.

27 810 16 28 g-index

29 1,079 5.8 3.92 ext. papers ext. citations avg, IF L-index

#	Paper	IF	Citations
27	Community-wide plasmid gene mobilization and selection. ISME Journal, 2013, 7, 1173-86	11.9	96
26	The hidden life of integrative and conjugative elements. FEMS Microbiology Reviews, 2017, 41, 512-537	15.1	86
25	Genomic and functional analysis of the IncP-9 naphthalene-catabolic plasmid NAH7 and its transposon Tn4655 suggests catabolic gene spread by a tyrosine recombinase. <i>Journal of Bacteriology</i> , 2006 , 188, 4057-67	3.5	84
24	Isolation and characterization of naphthalene-catabolic genes and plasmids from oil-contaminated soil by using two cultivation-independent approaches. <i>Applied Microbiology and Biotechnology</i> , 2007 , 74, 501-10	5.7	70
23	Novel organization of aromatic degradation pathway genes in a microbial community as revealed by metagenomic analysis. <i>ISME Journal</i> , 2009 , 3, 1335-48	11.9	68
22	Complete nucleotide sequence of an exogenously isolated plasmid, pLB1, involved in gamma-hexachlorocyclohexane degradation. <i>Applied and Environmental Microbiology</i> , 2006 , 72, 6923-33	3 ^{4.8}	55
21	Intracellular excision and reintegration dynamics of the ICEclc genomic island of Pseudomonas knackmussii sp. strain B13. <i>Molecular Microbiology</i> , 2009 , 72, 1293-306	4.1	43
20	Distribution of gamma-hexachlorocyclohexane-degrading genes on three replicons in Sphingobium japonicum UT26. <i>FEMS Microbiology Letters</i> , 2006 , 256, 112-8	2.9	35
19	Cellular variability of RpoS expression underlies subpopulation activation of an integrative and conjugative element. <i>PLoS Genetics</i> , 2012 , 8, e1002818	6	32
18	Comparative genome analysis of Pseudomonas knackmussii B13, the first bacterium known to degrade chloroaromatic compounds. <i>Environmental Microbiology</i> , 2015 , 17, 91-104	5.2	30
17	A dual functional origin of transfer in the ICEclc genomic island of Pseudomonas knackmussii B13. <i>Molecular Microbiology</i> , 2011 , 79, 743-58	4.1	28
16	Vast Differences in Strain-Level Diversity in the Gut Microbiota of Two Closely Related Honey Bee Species. <i>Current Biology</i> , 2020 , 30, 2520-2531.e7	6.3	28
15	Cell differentiation to "mating bodies" induced by an integrating and conjugative element in free-living bacteria. <i>Current Biology</i> , 2013 , 23, 255-9	6.3	24
14	Sequencing and characterizing the genome of Estrella lausannensis as an undergraduate project: training students and biological insights. <i>Frontiers in Microbiology</i> , 2015 , 6, 101	5.7	21
13	Community analysis of gut microbiota in hornets, the largest eusocial wasps, Vespa mandarinia and V. simillima. <i>Scientific Reports</i> , 2019 , 9, 9830	4.9	17
12	A new large-DNA-fragment delivery system based on integrase activity from an integrative and conjugative element. <i>Applied and Environmental Microbiology</i> , 2013 , 79, 4440-7	4.8	17
11	An operon of three transcriptional regulators controls horizontal gene transfer of the integrative and conjugative element ICEclc in Pseudomonas knackmussii B13. <i>PLoS Genetics</i> , 2014 , 10, e1004441	6	14

LIST OF PUBLICATIONS

10	Insertion sequence-based cassette PCR: cultivation-independent isolation of gamma-hexachlorocyclohexane-degrading genes from soil DNA. <i>Applied Microbiology and Biotechnology</i> , 2008 , 79, 627-32	5.7	14
9	Characterization of the traD operon of naphthalene-catabolic plasmid NAH7: a host-range modifier in conjugative transfer. <i>Journal of Bacteriology</i> , 2008 , 190, 6281-9	3.5	11
8	Physiological and transcriptome changes induced by Pseudomonas putida acquisition of an integrative and conjugative element. <i>Scientific Reports</i> , 2018 , 8, 5550	4.9	9
7	The TetR-type MfsR protein of the integrative and conjugative element (ICE) ICEclc controls both a putative efflux system and initiation of ICE transfer. <i>Journal of Bacteriology</i> , 2014 , 196, 3971-9	3.5	7
6	Inhibitory effect of Pseudomonas putida nitrogen-related phosphotransferase system on conjugative transfer of IncP-9 plasmid from Escherichia coli. <i>FEMS Microbiology Letters</i> , 2013 , 345, 102-9	9 ^{2.9}	5
5	How can a dual oriT system contribute to efficient transfer of an integrative and conjugative element?. <i>Mobile Genetic Elements</i> , 2011 , 1, 82-84		4
4	Local Necrotic Cells Trigger Systemic Immune Activation via Gut Microbiome Dysbiosis in Drosophila. <i>Cell Reports</i> , 2020 , 32, 107938	10.6	4
3	Deep Sequencing Uncovers Caste-Associated Diversity of Symbionts in the Social Ant Camponotus japonicus. <i>MBio</i> , 2020 , 11,	7.8	3
2	A novel system of bacterial cell division arrest implicated in horizontal transmission of an integrative and conjugative element. <i>PLoS Genetics</i> , 2019 , 15, e1008445	6	2
1	Vast differences in strain-level diversity in the gut microbiota of two closely related honey bee species		2