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RASTtk: a modular and extensible implementation of the RAST algorithm for building custom annotation pipelines and annotating batches of genomes

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214	Identification of Five Serotypes of Enteropathogenic <i>Escherichia coli</i> from Diarrheic Calves and Healthy Cattle in Belgium and Comparative Genomics with Shigatoxigenic <i>E. coli</i> . 2022 , 9, 492	0
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212	Draft Genome Sequences of <i>Pseudomonas</i> sp. Strains MWU13-2100 and MWU13-2105 Isolated from Wild Cranberry Bog Soil in the Cape Cod National Seashore.	0
211	Emergence of Extensively Drug-Resistant ST170 <i>Citrobacter portucalensis</i> with Plasmids pK218-KPC, pK218-NDM, and pK218-SHV from a Tertiary Hospital, China.	0
210	Persistence of Rare <i>Salmonella</i> Typhi Genotypes Susceptible to First-Line Antibiotics in the Remote Islands of Samoa.	1
209	Genome analysis of a new sulphur disproportionating species <i>Thermosulfurimonas</i> strain F29 and comparative genomics of sulfur-disproportionating bacteria from marine hydrothermal vents. 2022 , 8,	0
208	Comparative analysis of <i>Listeria monocytogenes</i> plasmid transcriptomes reveals common and plasmid-specific gene expression patterns and high expression of noncoding RNAs. 2022 , 11,	0
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206	Bioinformatics study of expression from genomes of epidemiologically related MRSA CC398 isolates from human and wild animal samples. 2022 , 268, 104714	0
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203	Draft Genomic Sequences of <i>Pseudomonas moorei</i> Strains Isolated from Wild Cranberry Bogs in Massachusetts.	0

202	Novel species identification and deep functional annotation of electrogenic biofilms, selectively enriched in a microbial fuel cell array. 13,	0
201	Elevation of <i>Clavibacter michiganensis</i> subsp. <i>californiensis</i> to species level as <i>Clavibacter californiensis</i> sp. nov., merging and re-classification of <i>Clavibacter michiganensis</i> subsp. <i>chilensis</i> and <i>Clavibacter michiganensis</i> subsp. <i>phaseoli</i> as <i>Clavibacter phaseoli</i> sp. nov. based on complete genome <i>in silico</i> analyses. 2022, 72,	0
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196	Characterization of Tele, an LXG effector of <i>Streptococcus gallolyticus</i> , antagonized by a non-canonical immunity protein.	0
195	<i>Geoalkalibacter halelectricus</i> SAP -1 sp. nov. possessing extracellular electron transfer and mineral-reducing capabilities from a haloalkaline environment.	0
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