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Two Archaeal Metagenome-Assembled Genomes From El Tatio ...Provide New Insights Into The Crenarchaeota Phylum Andrés Santos 1,2, Pablo Bruna 1, Jaime Martinez-Urtaza 3, Francisco Solís 4, ... Temperature And PH Of The Sampling Point Were Measured In-situ. Afterwards, Samples Were Kept On Ice, Transported To The Laboratory And Frozen At 80 C Until DNA Extraction. Feb 5th, 2024Draft Genomes Of Cronobacter Sakazakii Strains Isolated ...C. Sakazakii Genomes Was 4393kb, With An Average Of 4055 Protein Coding Genes, And An Average Genome G+C Content Of 56.9%. The Genomes Contained Genes Related To Carbohydrate Transport And Metabolism, Amino Acid Transport And Metabolism, And Cell Wall/membrane Biogenesis. In Addition, We Identified Genes Encoding Proteins Feb 6th, 2024Positive Selection Within The Genomes Of SARS-CoV-2 And ...Sequence Alignment To Identify Branch Specific Positive Selection, It Is Necessary To Obtain A Query And A ... Reference Alignment. We Downloaded Six High Quality Reference Genomes From The ... To Build A Sequence Alignment. Next, We Refined The Alignment Using A Gene By Gene

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