

Complete genome sequences of 17 clinical campylobacter jejuni strains from Chile

Bravo, V.
Porte, L.
Weitzel, T.
Varela, C.
Blondel, C.J.
Gonzalez-Escalona, N.

Abstract

Campylobacter jejuni is the leading cause of bacterial foodborne disease worldwide. Here, we report the complete annotated genomes and plasmid sequences of 17 Campylobacter jejuni strains isolated from patients with gastroenteritis in Santiago, Chile.

Indexed keywords

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