Draft genome sequence of Chelonobacter oris strain 1662T, associated with respiratory disease in Hermann’s tortoises
Kudirkiene, Egle; Hansen, Mie Johanne; Bojesen, Anders Miki

Published in:
Genome Announcements

DOI:
10.1128/genomeA.01322-14

Publication date:
2014

Document Version
Publisher's PDF, also known as Version of record

Citation for published version (APA):
Kudirkiene, E., Hansen, M. J., & Bojesen, A. M. (2014). Draft genome sequence of Chelonobacter oris strain 1662T, associated with respiratory disease in Hermann’s tortoises. Genome Announcements, 2(6), [e01322-14]. https://doi.org/10.1128/genomeA.01322-14
Draft Genome Sequence of *Chelonobacter oris* Strain 1662<sup>T</sup>, Associated with Respiratory Disease in Hermann’s Tortoises

Egle Kudirkienė, Mie Johanne Hansen, Anders Miki Bojesen
Department of Veterinary Disease Biology, Faculty of Health and Medical Sciences, University of Copenhagen, Copenhagen, Denmark

*Chelonobacter oris* 1662<sup>T</sup> is a type strain of the recently described species of the *Pasteurellaceae* family. The strain was isolated from the choanae of a captive tortoise with signs of respiratory tract infection. The genome reported here is approximately 2.6 Mb in size and has a G+C content of 47.1%.

**References**


4. Hansen MJ, Bertelsen MF, Christensen H, Bojesen AM, Bisgaard M. 2012. *Otiodibacter oris* gen. nov., sp. nov., a member of the family *Pas-


