Xylella fastidiosa subsp. fastidiosa strain XYL461 chromosome, complete genome

GenBank: CP109897.1

FASTA Graphics

Go to:

LOCUS CP109897 1194 bp DNA linear BCT 30-JAN-2023
DEFINITION Xylella fastidiosa subsp. fastidiosa strain XYL461 chromosome, complete genome.
ACCESSION CP109897 REGION: complement(1683309..1684502)
VERSION CP109897.1
DBLINK BioProject: PRJNA891297
BioSample: SAMN31318972
KEYWORDS .
SOURCE Xylella fastidiosa subsp. fastidiosa
ORGANISM Xylella fastidiosa subsp. fastidiosa
Bacteria; Pseudomonadati; Pseudomonadota; Gammaproteobacteria; Lysobacterales; Lysobacteraceae; Xylella.
REFERENCE 1 (bases 1 to 1194)
AUTHORS Velasco-Amo,M.D.P., Arias-Giraldo,L.F.F., Ecija,M.R., De La Fuente,L., Marco-Noales,E., Moralejo,E., Navas-Cort,J.A. and Landa,B.B.
TITLE Complete circularized genome resources of seven strains of Xylella fastidiosa subsp. fastidiosa using hybrid assembly reveals unknown plasmids
JOURNAL Phytopathology (2022) In press
PUBMED 36441872
REMARK Publication Status: Available-Online prior to print
REFERENCE 2 (bases 1 to 1194)
AUTHORS Landa,B., Arias-Giraldo,L.F., Roman-Ecija,M., Velasco-Amo,M.P., De
La Fuente,L., Marco-Noales,E. and Moralejo,E.
TITLE Direct Submission
JOURNAL Submitted (17-OCT-2022) Crop Protection, Institute for Sustainable
Agriculture - CSIC, Av. Menendez Pidal SN, Cordoba 14004, Spain
COMMENT The annotation was added by the NCBI Prokaryotic Genome Annotation
Pipeline (PGAP). Information about PGAP can be found here:
https://www.ncbi.nlm.nih.gov/genome/annotation_prok/

##Genome-Assembly-Data-START##
Assembly Date :: SEP-2022
Assembly Method :: Canu v. 2.2
Genome Representation :: Full

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