

Company: AccuCompany

Sample: C028004 - 4722389-EM Sample.1

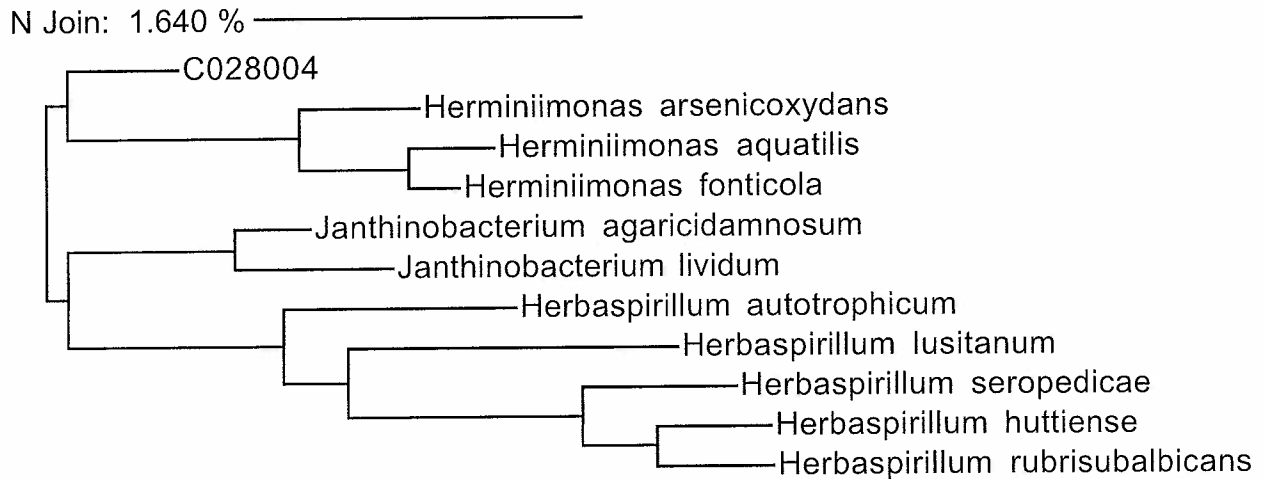
Date: Tuesday, April 8, 2008 9:43 AM

Sequence Alignment

Alignment: 524 C028004

- 1.92 % 522 *Janthinobacterium agaricidamnosum*
- 2.10 % 524 *Herminiimonas arsenicoxydans*
- 2.48 % 524 *Herbaspirillum autotrophicum*
- 2.68 % 522 *Janthinobacterium lividum*
- 2.77 % 524 *Herminiimonas aquatilis*
- 2.86 % 524 *Herminiimonas fonticola*
- 4.20 % 524 *Herbaspirillum lusitanum*
- 4.39 % 524 *Herbaspirillum seropedicae*
- 4.58 % 524 *Herbaspirillum huttiense*
- 4.77 % 524 *Herbaspirillum rubrisubalbicans*

Neighbor Joining Tree



Accugenix Database Search Result

Generated by: _____

Identification: Oxalobacteraceae

Reviewed by: _____

Confidence Level: Family

QA Review: _____

Batch #: 80404032

ACC1 - Due 4/11/08

NOT FOR USE IN INVITRO DIAGNOSTICS

SOP-GEN-049
Rev. 01Apr08 JM