

Table S1. 16S rRNA gene similarity between strain CJ426^T and related type strains

CJ426 ^T vs	Accession	Similarity (%)
<i>N. agricola</i> CC-SYL272 ^T	MZ452987	93.23
<i>N. aurantiaca</i> DSM 17617 ^T	DQ457019	94.31
<i>N. beijingensis</i> 3A5MI-3 ^T	MZ919349	93.16
<i>N. drilacis</i> DSM 25811 ^T	KC307772	94.14
<i>N. ginsenosidivorans</i> BS26 ^T	JQ349050	93.50
<i>N. hirudinis</i> DSM 25812 ^T	KC307771	94.66
<i>N. pedocola</i> JCM 31011 ^T	KT032154	94.13
<i>N. soli</i> DSM 19437 ^T	EF592608	95.52
<i>N. tibetebasis</i> 15-4 ^T	GU291295	92.34
<i>N. aquatica</i> RP-2 ^T	KT950742	95.39
<i>N. defluvii</i> I65 ^T	OM095447	97.77
<i>N. ginsengisoli</i> GR10-1 ^T	EU616816	96.58
<i>N. hibiscisoli</i> THG-DN5.5 ^T	KU505213	98.74
<i>N. yanshanensis</i> HAMBI 3031 ^T	FJ457040	98.11