



G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Brucella abortus</i> 9-941 (Genbank: NC_006932)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum</i> sp. P-5-2 (Genbank: EF102833)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum guangzhouense</i> strain P2 (Genbank: EF125185)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum</i> sp. TSNRS4 (Genbank: EU249585)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Bacterium</i> IPN2 (Genbank: EU249926)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Bacterium</i> IPN9 (Genbank: EU249927)
A C T A G T T G T T G G G G A	T T C A T T	C T T A G T A	<i>Bacterium</i> IPN15 (Genbank: EU249928)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Bacterium</i> IPN20 (Genbank: EU249930)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> strain Y2-1 (Genbank: EU275247)
G T T A G C C G T T G G G G A	G T T T A C	N T T C G G T G	<i>Ochrobactrum</i> sp. WatG-BA (Genbank: AB272074)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum</i> sp. BMC4 (Genbank: AB272338)
G T T A G C C G T T C G G G C A	G T A T A C	G T T C G G T G	<i>Agrobacterium</i> sp. GW4 (Genbank: EF550174)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Pseudochrobactrum saccharolyticum</i> (Genbank: AM180484)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Pseudochrobactrum asaccharolyticum</i> (Genbank: AM180485)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum pseudogrignonense</i> (Genbank: AM422371)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Rhizobium</i> sp. TKW2 (Genbank: AY631061)
G T T A G C C G T T G G G T G	G T T T A C	G T C A G T G	<i>Bartonella clarridgeiae</i> strain Houston (Genbank: AB292603)
G T T A G C C G T T C G G G C G	G T T C A C	G C T C G G T G	<i>Bartonella</i> sp. Sm7696 (Genbank: EF370420)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> CCUG 12303 (Genbank: AM490615)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> CCUG 1838 (Genbank: AM490614)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> CCUG 43892 (Genbank: AM490611)
G T T A G C C G T T C G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> CCUG 50899 (AM490609)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum</i> sp. CGL-X (Genbank: DQ305290)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum</i> sp. 11a (Genbank: DQ342340)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum</i> sp. DM1 (Genbank: DQ437526)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> DSM 20150 (Genbank: AM490618)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> DSM 7216 (Genbank: AM490617)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G K -	Uncultured <i>Ochrobactrum</i> sp. (Genbank: EU000376)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> (Genbank: EU119263)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> LMG 35 (Genbank: AM490616)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> LMG 401 (Genbank: AM490612)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum anthropi</i> LMG 5446 (Genbank: AM490613)
G T T A G C C G T T C G G G C A	G T T T A C	G T C C G G T G	<i>Bartonella bacilliformis</i> NCTC12138 (Genbank: X60042)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum cytisi</i> 43(25)zhy (Genbank: AM411061)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum cytisi</i> ESC1 (Genbank: AY776289)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum cytisi</i> 252zhy (Genbank: AM411068)
G T T A G C C G T T C G G G G G	A T T T A T	C T T C G G T G	<i>Ochrobactrum gallinifaecis</i> Iso 196 (Genbank: AJ19939)
G T T A G C C G T T C G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum grignonense</i> Oga9a (Genbank: AM490620)
G T T A G C C G T T C G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum grignonense</i> Oga9c (Genbank: AM490619)
G T T A G C C G T T C G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum grignonense</i> SRC1 (Genbank: AB272351)
G T T A G C C G T T C G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum haemophilum</i> CCUG 38531 (Genbank: AM422370)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> CCM4352 (Genbank: AM490622)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> CCUG 43465 (Genbank: AM490610)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> E276 (Genbank: AM490624)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> LMG3301T (Genbank: AM490623)
G T T A G C C G T T C G G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> PR17 (Genbank: AM490632)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> RMA16449 (Genbank: AM490621)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> RR (Genbank: AM490625)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> TA 13 (Genbank: AM490628)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> TD 30 (Genbank: AM490625)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> TM73 (Genbank: AM490627)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> CCM7036 (Genbank: AM490631)
G T T A G C C G T T C G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum intermedium</i> TK14 (Genbank: AM490633)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum lupini</i> LUP23 (Genbank: AM490629)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum oryzae</i> MTCC4195 (Genbank: AM041247)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum oryzae</i> ws1846 (Genbank: AM490630)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Ochrobactrum pseudogrignonense</i> CCUG30717T (Genbank: AM422371)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum shiyanus</i> WSH-W04 (Genbank: AJ920029)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum tritici</i> LAIII-106 (Genbank: AM490634)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum tritici</i> TA 93 (Genbank: AM490635)
G T T A G C C G T T G G G G A	G T T T A C	C T T C G G T G	<i>Ochrobactrum tritici</i> WS1830 (Genbank: AM490636)
G T T A G C C G T T C G G G C A	G T T T A C	G T C C G G T G	<i>Sinorhizobium meliloti</i> LMG15285 (Genbank: AJ295071)
G C C A A C C G T T G G G C A	G C T T G C	T G C A G T G	<i>Corbulabacter subterraneus</i> (Genbank: AY078053)
G T T A G C C G T T C G G G G	G T T T A C	C T T C G G T G	<i>Pseudochrobactrum glaciale</i> (Genbank: AB369864)
G T T A G C C G T T C G G G C A	G T T T A C	G C T C G G T G	<i>Bartonella bacilliformis</i> KC584 (Genbank: AF442955)