Lista de estirpes tipo bacterianas disponíveis para distribuição no CRB-JD

Espécie	Código CRB-JD	Códigos em outras coleções	Referências
Azomonas macrocytogenes	BR 10692	LMG 1266 = WR 111 = ATCC 12335 = CIP 103928 = DSM 721 = NCAIM B.01790 = NCIMB 8700	NEW (P.B.) and TCHAN (Y.T.): Azomonas macrocytogenes (ex Baillie, Hodgkiss, and Norris 1962, 118) nom. rev. Int. J. Syst. Bacteriol., 1982, 32, 381-382.
Azorhizobium caulinodans	BR 5410	LMG 6465 = ORS 571 = ATCC 43989 = BCRC 15787 = DSM 5975	DREYFUS (B.), GARCIA (J.L.) and GILLIS (M.): Characterization of Azorhizobium caulinodans gen. nov., sp. nov., a stem-nodulating nitrogen-fixing bacterium isolated from Sesbania rostrata. Int. J. Syst. Bacteriol., 1988, 38, 89-98.
Azorhizobium doebereinerae	BR 5401	LMG 9993 = SEMIA 6401 = DSM 18977	MOREIRA (F.M.S.), CRUZ (L.), FARIA (S.M.), MARSH (T.), MARTÍNEZ-ROMERO (E.), PEDROSA (F.O.), PITARD (R.M.) and YOUNG (J.P.W.): Azorhizobium doebereinerae sp. Nov. Microsymbiont of Sesbania virgata (Caz.) Pers. Syst. Appl. Microbiol., 2006, 29, 197-206.
Azorhizophilus paspali (Azotobacter paspali)	BR 10345	LMG 3864 = ATCC 23833 = CCUG 53674 = CECT 4095 = DSM 2283 = NBRC 102228	THOMPSON (J.P.) and SKERMAN (V.B.D.): Azotobacteraceae: the taxonomy and ecology of the aerobic nitrogen-fixing bacteria. Academic Press, London, 1980, 405 pp.
Azospirillum brasilense	BR 11001	Sp 7 = ATCC 29145 = DSM 1690 = BCRC 12270 = LMG 13127	TARRAND (J.J.), KRIEG (N.R.) and DÖBEREINER (J.): A taxonomic study of the Spirillum lipoferum group, with descriptions of a new genus, Azospirillum gen. nov. and two species, Azospirillum lipoferum (Beijerinck) comb. nov. and Azospirillum brasilense sp. nov. Canadian Journal of Microbiology, 1978, 24, 967-980.
Azospirillum doebereinerae	BR 12281	GSF 71 = DSM 13131 = LMG 26176	ECKERT (B.), WEBER (O.B.), KIRCHHOF (G.), HALBRITTER (A.), STOFFELS (M.) and HARTMANN (A.): Azospirillum doebereinerae sp. nov., a nitrogen -fixing bacterium associated with the C4 -grass Miscanthus. Int. J. Syst. Evol. Microbiol., 2001, 51, 17-26.
Azospirillum fermentarium	BR 10693	LMG 27264 = CC-LY743 = BCRC 80505 = JCM 18688	LIN, S.-Y., LIU, Y.-C., HAMEED, A., HSU, Y.-H., LAI, W.-A., SHEN, F.-T. and YOUNG, C.-C. 2013. Azospirillum fermentarium sp. nov., a nitrogen-fixing species isolated from a fermenter. Int. J. Syst. Evol. Microbiol., 63, 3762-3768.

Azospirillum lipoferum	BR 11080	Sp 59 b = ATCC 29707 = DSM 1691= BCRC 12213 = LMG 13128	TARRAND (J.J.), KRIEG (N.R.) and DÖBEREINER (J.): A taxonomic study of the Spirillum lipoferum group, with descriptions of a new genus, Azospirillum gen. nov. and two species, Azospirillum lipoferum (Beijerinck) comb. nov. and Azospirillum brasilense sp. nov. Canadian Journal of Microbiology, 1978, 24, 967-980.
Azospirillum melinis	BR 12183	TMCY 0552 = DSM 17798 = LMG 24250	PENG (G.), WANG (H.), ZHANG (G.), HOU (W.), LIU (Y.), WANG (E.T.) and TAN (Z.): Azospirillum melinis sp. nov., a group of diazotrophs isolated from tropical molasses grass. Int. J. Syst. Evol. Microbiol., 2006, 56, 1263-1271.
Azospirillum oryzae	BR 10696	LMG 23844 = COC8 = IAM 15130 = JCM 21588 = NBRC 102291	XIE (C.H.) and YOKOTA (A.): Azospirillum oryzae sp. nov., a nitrogen-fixing bacterium isolated from the roots of the rice plant Oryza sativa. Int. J. Syst. Evol. Microbiol., 2005, 55, 1435-1438.
Azospirillum picis	BR 10330	DSM 19922 = CCUG 55431	LIN (S.Y.), YOUNG (C.C.), HUPFER (H.), SIERING (C.), ARUN (A.B.), CHEN (W.M.), LAI (W.A.), SHEN (F.T.), REKHA (P.D.) and YASSIN (A.F.): Azospirillum picis sp. nov., isolated from discarded tar. Int. J. Syst. Evol. Microbiol., 2009, 59, 761-765.
Azospirillum rugosum	BR 10695	DSM 19657 = IMMIB AFH-6 = CCUG 53966	YOUNG (C.C.), HUPFER (H.), SIERING (C.), HO (M.J.), ARUN (A.B.), LAI (W.A.), REKHA (P.D.), SHEN (F.T.), HUNG (M.H.), CHEN (W.M.) and YASSIN (A.F.): Azospirillum rugosum sp. nov., isolated from oil-contaminated soil. Int. J. Syst. Evol. Microbiol., 2008, 58, 959-963.
Azospirillum thiophilum	BR 10331	DSM 21654 = VKM B-2513	LAVRINENKO (K.), CHERNOUSOVA (E.), GRIDNEVA (E.), DUBININA (G.), AKIMOV (V.), KUEVER (J.), LYSENKO (A.) and GRABOVICH (M.): Azospirillum thiophilum sp. nov., a diazotrophic bacterium isolated from a sulfide spring. Int. J. Syst. Evol. Microbiol., 2010, 60, 2832-2837.
Azotobacter beijerinckii	BR 10657	LMG 1265 = ATCC 19360 = CIP 106282 = DSM 378 = JCM 20742 = NCAIM B.01800 = NRRL B-14367 = NRRL B-14640 = VKM B-161	LIPMAN (J.G.): Soil bacteriological studies. Report of the New Jersey Agricultural Experiment Station, 1904, 25, 237-289.
Azotobacter chroococcum	BR 10473	LMG 8756 = ATCC 9043 = DSM 2286 = JCM 20725 = JCM 21503 = NBRC 102613 = NCAIM B.01391 =	BEIJERINCK (M.W.): Über oligonitrophile Mikroben. Zentralblatt für Bakteriologie, Parasitenkunde, Infektionskrankheiten und Hygiene. Abteilung II, 1901, 7, 561-582.

		NRRL B-14346 = NRRL B-14637 = VKM B-1616	
Azotobacter nigricans subsp. nigricans	BR 10656	DSM 2288 = LMG 2816 = ATCC 35009 = UQM 1967	THOMPSON (J.P.) and SKERMAN (V.B.D.): Azotobacteraceae: the taxonomy and ecology of the aerobic nitrogen-fixing bacteria. Academic Press, London, 1980, 405 pp.
Azotobacter vinelandii	BR 10658	LMG 8758 = ATCC 478 = DSM 2289 = JCM 21475 = NBRC 102612 = NCCB 26007 = NRRL B-14641 = VKM B-1617	LIPMAN (J.G.): Experiments on the transformation and fixation of nitrogen by bacteria. Report of the New Jersey Agricultural Experiment Station, 1903, 24, 217-285.
Beijerinckia doebereinerae	BR 10344	LMG 2819 = CECT 7311 = DSM 19635	OGGERIN (M.), ARAHAL (D.R.), RUBIO (V.) and MARÍN (I.): Identification of Beijerinckia fluminensis strains CIP 106281T and UQM 1685T as Rhizobium radiobacter strains, and proposal of Beijerinckia doebereinerae sp. nov. to accommodate Beijerinckia fluminensis LMG 2819. Int. J. Syst. Evol. Microbiol., 2009, 59, 2323-2328.
Bradyrhizobium arachidis	BR 10337	LMG 26795 = CCBAU 051107 = CGMCC 1.12100 = HAMBI 3281	WANG (R.), CHANG (Y.L.), ZHENG (W.T.), ZHANG (D.), ZHANG (X.X.), SUI (X.H.), WANG (E.T.), HU (J.Q.), ZHANG (L.Y.), CHEN (W.X.). Bradyrhizobium arachidis sp. nov., isolated from effective nodules of Arachis hypogaea grown in China. Syst Appl Microbiol. 2013 Mar; 36(2): 101-5.
Bradyrhizobium betae	BR 10200	CNPSo 2079 = DSM 17288 = CECT 5829 = LMG 21987 = NBRC 103048	RIVAS (R.), WILLEMS (A.), PALOMO (J.L.), GARCÍA-BENAVIDES (P.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.), GILLIS (M.) and VELÁZQUEZ (E.): Bradyrhizobium betae sp. nov., isolated from roots of Beta vulgaris affected by tumour-like deformations. Int. J. Syst. Evol. Microbiol., 2004, 54, 1271-1275.

Bradyrhizobium canariense	BR 10180	DSM 16623 = ATCC BAA-1002 = LMG 22265	VINUESA (P.), LEÓN-BARRIOS (M.), SILVA (C.), WILLEMS (A.), JARABO-LORENZO (A.), PÉREZ-GALDONA (R.), WERNER (D.) and MARTÍNEZ-ROMERO (E.): Bradyrhizobium canariense sp. nov., an acid-tolerant endosymbiont that nodulates endemic genistoid legumes (Papilionoideae: Genisteae) from the Canary Islands, along with Bradyrhizobium japonicum bv. genistearum, Bradyrhizobium genospecies alpha and Bradyrhizobium genospecies beta. Int. J. Syst. Evol. Microbiol., 2005, 55, 569-575.
Bradyrhizobium cytisi	BR 10318	LMG 25866 = CECT 7749 = CTAW11	CHAHBOUNE (R.), CARRO (L.), PEIX (A.), BARRIJAL (S.), VELÁZQUEZ (E.) and BEDMAR (E.J.): Bradyrhizobium cytisi sp. nov., isolated from effective nodules of Cytisus villosus. Int. J. Syst. Evol. Microbiol., 2011, 61, 2922-2927.
Bradyrhizobium daqingense	BR 10474	HAMBI 3184 = CCBAU 15774 = CGMCC 1.10947 = LMG 26137	WANG (J.Y.), WANG (R.), ZHANG (Y.M.), LIU (H.C.), CHEN (W.F.), WANG (E.T.), SUI (X.H.) and W. X. CHEN (W.X.): Bradyrhizobium daqingense sp. nov., isolated from soybean nodules. Int. J. Syst. Evol. Microbiol., 2013, 63, 616-624.
Bradyrhizobium denitrificans (Blastobacter denitrificans)	BR 10327	LMG 8443 = ATCC 43295 = DSM 1113 = HAMBI 2266	VAN BERKUM (P.), LEIBOLD (J.M.) and EARDLY (B.D.): Proposal for combining Bradyrhizobium spp. (Aeschynomene indica) with Blastobacter denitrificans and to transfer Blastobacter denitrificans (Hirsh and Muller, 1985) to the genus Bradyrhizobium as Bradyrhizobium denitrificans (comb. nov.). Syst. Appl. Microbiol., 2006, 29, 207-215.
Bradyrhizobium diazoefficiens	BR 116	USDA 110 = BCRC 13528 = NBRC 14792 = SEMIA 5032 = CNPSo 46	DELAMUTA, J. R. M., RIBEIRO, R. A., ORMEÑO-ORRILLO, E., MELO, I. S., MARTÍNEZ-ROMERO, E. and HUNGRIA, M. 2013. Polyphasic evidence supporting the reclassification of Bradyrhizobium japonicum group Ia strains as Bradyrhizobium diazoefficiens sp. nov.. Int. J. Syst. Evol. Microbiol., 63, 3342-3351.
Bradyrhizobium elkanii	BR 10357	LMG 6134 = USDA 76 = ATCC 49852 = DSM 11554 = BCRC 15790	KUYKENDALL (L.D.), SAXENA (B.), DEVINE (T.E.) and UDELL (S.E.): Genetic diversity in Bradyrhizobium japonicum Jordan 1982 and a proposal for Bradyrhizobium elkanii sp. nov. Can. J. Microbiol., 1992, 38, 501-505.

Bradyrhizobium embrapense	BR 2212	CNPSo 2833 = CIAT 2372 = SEMIA 6208 = U674	DELAMUTA (J.R.), RIBEIRO (R.A.), ORMEÑO-ORRILLO (E.), PARMA (M.M.), MELO (I.S.), MARTÍNEZ-ROMERO (E.), HUNGRIA (M.). Bradyrhizobium tropiciagri sp. nov. and Bradyrhizobium embrapense sp. nov., nitrogen-fixing symbionts of tropical forage legumes. Int J Syst Evol Microbiol. 2015 Dec; 65(12): 4424-33.
Bradyrhizobium erythrophlei	BR 10486	HAMBI 3614 = CCBAU 53325 = CGMCC 1.13002 = LMG 28425	YAO, Y., SUI, X. H., ZHANG, X. X., WANG, E. T. and CHEN, W. X. 2015. Bradyrhizobium erythrophlei sp. nov. and Bradyrhizobium ferriligni sp. nov., isolated from effective nodules of Erythrophleum fordii. Int. J. Syst. Evol. Microbiol., 65, 1831-1837.
Bradyrhizobium ferriligni	BR 10482	HAMBI 3613 = CCBAU 51502 = CGMCC 1.13001	YAO, Y., SUI, X. H., ZHANG, X. X., WANG, E. T. and CHEN, W. X. 2015. Bradyrhizobium erythrophlei sp. nov. and Bradyrhizobium ferriligni sp. nov., isolated from effective nodules of Erythrophleum fordii. Int. J. Syst. Evol. Microbiol., 65, 1831-1837.
Bradyrhizobium huanghuaihaiense	BR 10336	LMG 26136 = CCBAU 23303 = HAMBI 3180	ZHANG (Y.M.), LI (Y.J.), CHEN (W.F.), WANG (E.T.), SUI (X.H.), LI (Q.Q.), ZHANG (Y.Z.), ZHOU (Y.G.) and CHEN (W.X.): Bradyrhizobium huanghuaihaiense sp. nov., an effective symbiotic bacterium isolated from soybean (Glycine max L.) nodules. Int. J. Syst. Evol. Microbiol., 2012, 62, 1951-1957.
Bradyrhizobium icense	BR 10399	CNPSo 2583 = LMTR 13 = HAMBI 3584	DURÁN, D., REY, L., MAYO, J., ZÚÑIGA-DÁVILA, D., IMPERIAL, J., RUIZ-ARGÄESO, T. S., MARTÍNEZ-ROMERO, E. and ORMEÑO-ORRILLO, E. 2014. Bradyrhizobium paxllaeri sp. nov. and Bradyrhizobium icense sp. nov., nitrogen-fixing rhizobial symbionts of Lima bean (Phaseolus lunatus L.) in Peru. Int. J. Syst. Evol. Microbiol., 64, 2072-2078.
Bradyrhizobium ingae	BR 10250	ERR 494 = HAMBI 3600	DA SILVA, K., DE MEYER, S. E., ROUWS, L. F. M., FARIAS, E. N. C., DOS SANTOS, M. A. O., O'HARA, G., ARDLEY, J. K., WILLEMS, A., PITARD, R. M. and ZILLI, J. E. 2014. Bradyrhizobium ingae sp. nov., isolated from effective nodules of Inga laurina grown in Cerrado soil. Int. J. Syst. Evol. Microbiol., 64, 3395-3401.

Bradyrhizobium iriomotense	BR 10354	LMG 24129 = EK05 = NBRC 102520	ISLAM (M.S.), KAWASAKI (H.), MURAMATSU (Y.), NAKAGAWA (Y.) and SEKI (T.): Bradyrhizobium iriomotense sp. nov., isolated from a tumor-like root of the legume Entada koshunensis from Iriomote Island in Japan. Biosci. Biotechnol. Biochem., 2008, 72, 1416-1429.
Bradyrhizobium japonicum	BR 114	USDA 06 = ATCC 10324 = BCRC 13518 = DSM 30131 = HAMBI 2314 = LMG 6138 = NBRC 14783 = USDA 505	JORDAN (D.C.) Transfer of Rhizobium japonicum Buchanan 1980 to Bradyrhizobium gen. nov., a genus of slow-growing, root nodule bacteria from leguminous plants. Int. J. Syst. Bacteriol., 1982, 32, 136-139.
Bradyrhizobium jicamae	BR 10585	LMG 24556 = PAC68 = CECT 7395	RAMÍREZ-BAHENA (M.H.), PEIX (A.), RIVAS (R.), CAMACHO (M.), RODRÍGUEZ-NAVARRO (D.N.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.), WILLEMS (A.) and VELÁZQUEZ (E.): Bradyrhizobium pachyrhizi sp. nov. and Bradyrhizobium jicamae sp. nov., isolated from effective nodules of Pachyrhizus erosus. Int. J. Syst. Evol. Microbiol., 2009, 59, 1929-1934.
Bradyrhizobium liaoningense	BR 10397	CNPSo 2483 = ATCC 700350 = CIP 104858 = NBRC 100396 = LMG 18230	XU (L.M.), GE (C.), CUI (Z.), LI (J.) and FAN (H.): Bradyrhizobium liaoningense sp. nov., isolated from the root nodules of soybeans. Int. J. Syst. Bacteriol., 1995, 45, 706-711.
Bradyrhizobium lupini	BR 10498	LMG 28514 = USDA 3051 = CECT 8630	PEIX, A., RAMÍREZ-BAHENA, M. H., FLORES-FÉLIX, J. D., ALONSO DE LA VEGA, P., RIVAS, R., MATEOS, P. F., IGUAL, J. M., MARTÍNEZ-MOLINA, E., TRUJILLO, M. E. and VELÁZQUEZ, E. 2015. Revision of the taxonomic status of the species Rhizobium lupini and reclassification as Bradyrhizobium lupini comb. nov. Int. J. Syst. Evol. Microbiol., 65, 1213-1219.
Bradyrhizobium manausense	BR 3351	CV1C4-31 = HAMBI 3596	SILVA, F. V., DE MEYER, S. E., SIMÕES-ARAÚJO, J. L., DA COSTA BARBÉ, T., XAVIER, G. R., O'HARA, G., ARDLEY, J. K., RUMJANEK, N. G., WILLEMS, A. and ZILLI, J. E. 2014. Bradyrhizobium manausense sp. nov., isolated from effective nodules of Vigna unguiculata grown in Brazilian Amazonian rainforest soils. Int. J. Syst. Evol. Microbiol., 64, 2358-2363.

Bradyrhizobium neotropicale	BR 10247	ERR 435 = HAMBI 3599	ZILLI, J. E., BARAÚNA, A. C., DA SILVA, K., DE MEYER, S. E., FARIAS, E. N. C., KAMINSKI, P. E., DA COSTA, I. B., ARDLEY, J. K., WILLEMS, A., CAMACHO, N. N., DOURADO, F. D. S. and O'HARA, G. 2014. Bradyrhizobium neotropicale sp. nov., isolated from effective nodules of Centrolobium paraense. Int. J. Syst. Evol. Microbiol., 64, 3950-3957.
Bradyrhizobium oligotrophicum (Agromonas oligotrophica)	BR 10343	LMG 10732 = ATCC 43045 = DSM 12412 = JCM 1494 = NCIMB 12151	RAMÍREZ-BAHENA (M.H.), CHAHBOUNE (R.), PEIX (A.) and VELÁZQUEZ (E.): Reclassification of Agromonas oligotrophica into the genus Bradyrhizobium as Bradyrhizobium oligotrophicum comb. nov. Int. J. Syst. Evol. Microbiol., 2013, 63, 1013-1016.
Bradyrhizobium ottawaense	BR 10475	HAMBI 3284 = OO99 = LMG 26739	YU, X., CLOUTIER, S., TAMBONG, J. T. and BROMFIELD, E. S. P. 2014. Bradyrhizobium ottawaense sp. nov., a symbiotic nitrogen fixing bacterium from root nodules of soybeans in Canada. Int. J. Syst. Evol. Microbiol., 64, 3202-3207.
Bradyrhizobium pachyrhizi	BR 10199	CNPSo 2077 = LMG 24246 = CECT 7396 = DSM 19631	RAMÍREZ-BAHENA (M.H.), PEIX (A.), RIVAS (R.), CAMACHO (M.), RODRÍGUEZ-NAVARRO (D.N.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.), WILLEMS (A.) and VELÁZQUEZ (E.): Bradyrhizobium pachyrhizi sp. nov. and Bradyrhizobium jicamae sp. nov., isolated from effective nodules of Pachyrhizus erosus. Int. J. Syst. Evol. Microbiol., 2009, 59, 1929-1934.
Bradyrhizobium paxllaeri	BR 10398	CNPSo 2582 = DSM 18454 = HAMBI 2911	DURÁN, D., REY, L., MAYO, J., ZÚÑIGA-DÁVILA, D., IMPERIAL, J., RUIZ-ARGÄESO, T. S., MARTÍNEZ-ROMERO, E. and ORMEÑO-ORRILLO, E. 2014. Bradyrhizobium paxllaeri sp. nov. and Bradyrhizobium icense sp. nov., nitrogen-fixing rhizobial symbionts of Lima bean (Phaseolus lunatus L.) in Peru. Int. J. Syst. Evol. Microbiol., 64, 2072-2078.
Bradyrhizobium rifense	10368	CNPSo 2468 = LMG 26781 = CECT 8066	CHAHBOUNE, R., CARRO, L., PEIX, A., RAMÍREZ-BAHENA, M. H., BARRIJAL, S., VELÁZQUEZ, E. and BEDMAR, E. J. 2012. Bradyrhizobium rifense sp. nov. isolated from effective nodules of Cytisus villosus grown in the Moroccan Rif. Syst. Appl. Microbiol., 35, 302-305.

Bradyrhizobium tropiciagri	BR 1009	CNPSo 1112 = SMS 303 = SEMIA 6148 = LMG 28867	DELAMUTA (J.R.), RIBEIRO (R.A.), ORMEÑO-ORRILLO (E.), PARMA (M.M.), MELO (I.S.), MARTÍNEZ-ROMERO (E.), HUNGRIA (M.). Bradyrhizobium tropiciagri sp. nov. and Bradyrhizobium embrapense sp. nov., nitrogen-fixing symbionts of tropical forage legumes. Int J Syst Evol Microbiol. 2015 Dec; 65(12): 4424-33.
Bradyrhizobium valentinum	BR 10400	CNPSo 2587 = LmjM3 = LMG 27619	DURÁN (D.), REY (L.), NAVARRO (A.), BUSQUETS (A.), IMPERIAL (J.), RUIZ-ARGÜESO (T.). Bradyrhizobium valentinum sp. nov., isolated from effective nodules of Lupinus mariae-josephae, a lupine endemic of basic-lime soils in Eastern Spain. Systematic and Applied Microbiology. V. 37, Issue 5, July 2014, Pages 336–341.
Bradyrhizobium viridifuturi	BR 1804	SEMIA 690 = CNPSo 991 = C100a = LMG 28866	HELENE (L.C.), DELAMUTA (J.R.), RIBEIRO (R.A.), ORMEÑO-ORRILLO (E.), ROGEL (M.A.), MERTÍNEZ-ROMERO (E.), HUNGRIA (M.). Bradyrhizobium viridifuturi sp. nov., encompassing nitrogen-fixing symbionts of legumes used for green manure and environmental services. Int J Syst Evol Microbiol. 2015 Dec; 65(12): 4441-8.
Bradyrhizobium yuanmingense	BR 10201	CNPSo 2080 = LMG 21827 = NBRC 100594	YAO (Z.Y.), KAN (F.L.), WANG (E.T.), WEI (G.H.) and CHEN (W.X.): Characterization of rhizobia that nodulate legume species of the genus Lespedeza and description of Bradyrhizobium yuanmingense sp. nov. Int. J. Syst. Evol. Microbiol., 2002, 52, 2219-2230.
Burkholderia gladioli	BR 12320	ATCC 10248 = DSM 4285	YABUUCHI (E.), KOSAKO (Y.), OYAIZU (H.), YANO (I.), HOTTA (H.), HASHIMOTO (Y.), EZAKI (T.) and ARAKAWA (M.): Proposal of Burkholderia gen. nov. and transfer of seven species of the genus Pseudomonas homology group II to the new genus, with the type species Burkholderia cepacia (Palleroni and Holmes 1981) comb. nov. Microbiol. Immunol., 1992, 36, 1251-1275.
Burkholderia glumae	BR 12322	ATCC 33617 = DSM 7169 = DSM 9512	URAKAMI (T.), ITO-YOSHIDA (C.), ARAKI (H.), KIJIMA (T.), SUZUKI (K.I.) and KOMAGATA (K.): Transfer of Pseudomonas plantarii and Pseudomonas glumae to Burkholderia as Burkholderia spp. and description of Burkholderia vandii sp. nov. Int. J. Syst. Bacteriol., 1994, 44, 235-245.

Burkholderia plantarii	BR 12324	ATCC 43733 = LMG 9035	URAKAMI (T.), ITO-YOSHIDA (C.), ARAKI (H.), KIJIMA (T.), SUZUKI (K.I.) and KOMAGATA (K.): Transfer of Pseudomonas plantarii and Pseudomonas glumae to Burkholderia as Burkholderia spp. and description of Burkholderia vandii sp. nov. Int. J. Syst. Bacteriol., 1994, 44, 235-245.
Burkholderia pyrrocinia	BR 12325	ATCC 15958 = DSM 10685	VANDAMME (P.), HOLMES (B.), VANCANNEYT (M.), COENYE (T.), HOSTE (B.), COOPMAN (R.), REVETS (H.), LAUWERS (S.), GILLIS (M.), KERSTERS (K.) and GOVAN (J.R.W.): Occurence of multiple genomovars of Burkholderia cepacia in cystic fibrosis patients and proposal of Burkholderia multivorans sp. nov. Int. J. Syst. Bacteriol., 1997, 47, 1188-1200.
Burkholderia vietnamiensis	BR 12311	TVV 75 = LMG 10929 = ATCC BAA-248	GILLIS (M.), VAN (T.V.), BARDIN (R.), GOOR (M.), HEBBAR (P.), WILLEMS (A.), SEGERS (P.), KERSTERS (K.), HEULIN (T.) and FERNANDEZ (M.P.): Polyphasic taxonomy in the genus Burkholderia leading to an emended description of the genus and proposition of Burkholderia vietnamiensis sp. nov. for N2-fixing isolates from rice in Vietnam. Int. J. Syst. Bacteriol., 1995, 45, 274-289.
Chelatococcus asaccharovorans	BR 10179	DSM 6462 = ATCC 51531 = LMG 25503	AULING (G.), BUSSE (H.J.), EGLI (T.), EL-BANNA (T.) and STACKEBRANDT (E.): Description of the gram-negative, obligately aerobic, nitrilotriacetate (NTA)-utilizing bacteria as Chelatobacter heintzii, gen. nov., sp. nov., and Chelatococcus asaccharovorans, gen. nov., sp. nov. Syst. Appl. Microbiol., 1993, 16, 104-112.
Cupriavidus taiwanensis	BR 3471	LMG 19424 = DSM 17343 = CIP 107171	VANDAMME (P.) and COENYE (T.): Taxonomy of the genus Cupriavidus: a tale of lost and found. Int. J. Syst. Evol. Microbiol., 2004, 54, 2285-2289.
Devosia neptuniae	BR 10334	LMG 21357 = CECT 5650 = CIP 108397	RIVAS (R.), WILLEMS (A.), SUBBA-RAO (N.S.), MATEOS (P.F.), DAZZO (F.B.), KROPPENSTEDT (R.M.), MARTÍNEZ-MOLINA (E.), GILLIS (M.) and VELÁZQUEZ (E.): Description of Devosia neptuniae sp. nov. that nodulates and fixes nitrogen in symbiosis with Neptunia natans, an aquatic legume from India. Syst. Appl. Microbiol., 2003, 26, 47-53.

Devosia yakushimensis	BR 10316	LMG 24299 = DSM 21277 = KCTC 22147 = NBRC 103855	BAUTISTA (V.V.), MONSALUD (R.G.) and YOKOTA (A.): Devosia yakushimensis sp. nov., isolated from root nodules of Pueraria lobata (Willd.) Ohwi. Int. J. Syst. Evol. Microbiol., 2010, 60, 627-632.
Ensifer americanum	BR 10401	CNPSo 2065 = LMG 22684 = ATCC BAA-532 = CIP 108390 = DSM 15007	WANG, Y. C., WANG, F., HOU, B. C., WANG, E. T., CHEN, W. F., SUI, X. H., CHEN, W. X., LI, Y. and ZHANG, Y. B. 2013. Proposal of Ensifer psoraleae sp. nov., Ensifer sesbaniae sp. nov., Ensifer morelense comb. nov. and Ensifer americanum comb. nov. Syst. Appl. Microbiol., 36, 467-473.
Ensifer fredii	BR 112	LMG 6217 = USDA 205 = ATCC 35423 = DSM 5851	YOUNG (J.M.): The genus name Ensifer Casida 1982 takes priority over Sinorhizobium Chen et al. 1988, and Sinorhizobium morelense Wang et al. 2002 is a later synonym of Ensifer adhaerens Casida 1982. Is the combination 'Sinorhizobium adhaerens' (Casida 1982) Willems et al. 2003 legitimate? Request for an Opinion. Int. J. Syst. Evol. Microbiol., 2003, 53, 2107-2110.
Ensifer medicae	BR 525	USDA 1037 = BCRC 15798 = HAMBI 2306 = LMG 19920 = NBRC 100384	YOUNG (J.M.): The genus name Ensifer Casida 1982 takes priority over Sinorhizobium Chen et al. 1988, and Sinorhizobium morelense Wang et al. 2002 is a later synonym of Ensifer adhaerens Casida 1982. Is the combination 'Sinorhizobium adhaerens' (Casida 1982) Willems et al. 2003 legitimate? Request for an Opinion. Int. J. Syst. Evol. Microbiol., 2003, 53, 2107-2110.
Ensifer meliloti	BR 7411	LMG 6133 = NZP 4027 = ATCC 9930 = BCRC 13516 = DSM 30135 = HAMBI 2148	YOUNG (J.M.): The genus name Ensifer Casida 1982 takes priority over Sinorhizobium Chen et al. 1988, and Sinorhizobium morelense Wang et al. 2002 is a later synonym of Ensifer adhaerens Casida 1982. Is the combination 'Sinorhizobium adhaerens' (Casida 1982) Willems et al. 2003 legitimate? Request for an Opinion. Int. J. Syst. Evol. Microbiol., 2003, 53, 2107-2110.

Ensifer mexicanus	BR 10215	CNPSo 2067 = ATCC BAA-1312 = DSM 18446 = HAMBI 2910	LLORET (L.), ORMEÑO-ORRILLO (E.), RINCÓN (R.), MARTÍNEZ-ROMERO (J.), ROGEL-HERNÁNDEZ (M.A.) and MARTÍNEZ-ROMERO (E.): Ensifer mexicanus sp. nov. a new species nodulating Acacia angustissima (Mill.) Kuntze in Mexico. Syst. Appl. Microbiol., 2007, 30, 280-290.
Ensifer saheli	BR 526	USDA 4893 = ATCC 51690 = BCRC 15799 = DSM 11273 = HAMBI 215 = LMG 7837 = NBRC 100386	YOUNG (J.M.): The genus name Ensifer Casida 1982 takes priority over Sinorhizobium Chen et al. 1988, and Sinorhizobium morelense Wang et al. 2002 is a later synonym of Ensifer adhaerens Casida 1982. Is the combination 'Sinorhizobium adhaerens' (Casida 1982) Willems et al. 2003 legitimate? Request for an Opinion. Int. J. Syst. Evol. Microbiol., 2003, 53, 2107-2110.
Ensifer sojae	BR 10312	LMG 25493 = CCBAU 05684 = HAMBI 3098	LI (Q.Q.), WANG (E.T.), CHANG (Y.L.), ZHANG (Y.Z.), ZHANG (Y.M.), SUI (X.H.), CHEN (W.F.) and CHEN (W.X.): Ensifer sojae sp. nov., isolated from root nodules of Glycine max grown in saline-alkaline soils. Int. J. Syst. Evol. Microbiol., 2011, 61, 1981-1988.
Ensifer terangae	BR 527	USDA 4894 = ATCC 51692 = BCRC 15800 = DSM 11282 = HAMBI 220 = LMG 7834 = NBRC 100385	YOUNG (J.M.): The genus name Ensifer Casida 1982 takes priority over Sinorhizobium Chen et al. 1988, and Sinorhizobium morelense Wang et al. 2002 is a later synonym of Ensifer adhaerens Casida 1982. Is the combination 'Sinorhizobium adhaerens' (Casida 1982) Willems et al. 2003 legitimate? Request for an Opinion. Int. J. Syst. Evol. Microbiol., 2003, 53, 2107-2110.
Ensifer xinjiangensis	BR 10214	CNPSo 2066 = ATCC 49357 = DSM 5852	YOUNG (J.M.): The genus name Ensifer Casida 1982 takes priority over Sinorhizobium Chen et al. 1988, and Sinorhizobium morelense Wang et al. 2002 is a later synonym of Ensifer adhaerens Casida 1982. Is the combination 'Sinorhizobium adhaerens' (Casida 1982) Willems et al. 2003 legitimate? Request for an Opinion. Int. J. Syst. Evol. Microbiol., 2003, 53, 2107-2110.

Gluconacetobacter azotocaptans	BR 10328	DSM 13594 = ATCC 700988 = BCC 36449 = LMG 21311	FUENTES-RAMÍREZ (L.E.), BUSTILLOS-CRISTALES (R.), TAPIA-HERNÁNDEZ (A.), JIMÉNEZ-SALGADO (T.), WANG (E.T.), MARTÍNEZ-ROMERO (E.) and CABALLERO-MELLADO (J.): Novel nitrogen-fixing acetic acid bacteria, Gluconacetobacter johannae sp. nov. and Gluconacetobacter azotocaptans sp. nov., associated with coffee plants. Int. J. Syst. Evol. Microbiol., 2001, 51, 1305-1314.
Gluconacetobacter diazotrophicus	BR 11281	Pal 5 = ATCC 49037 = LMG 7603 = DSM 5601 = BCRC 16088 = IBSBF 1863	YAMADA (Y.), HOSHINO (K.) and ISHIKAWA (T.): The phylogeny of acetic acid bacteria based on the partial sequences of 16S ribosomal RNA: the elevation of the subgenus Gluconoacetobacter to generic level. Biosci. Biotech. Biochem., 1997, 61, 1244-1251.
Gluconacetobacter johannae	BR 10697	DSM 13595 = CFN-Cf55 = ATCC 700987 = CIP 107160	FUENTES-RAMÍREZ (L.E.), BUSTILLOS-CRISTALES (R.), TAPIA-HERNÁNDEZ (A.), JIMÉNEZ-SALGADO (T.), WANG (E.T.), MARTÍNEZ-ROMERO (E.) and CABALLERO-MELLADO (J.): Novel nitrogen-fixing acetic acid bacteria, Gluconacetobacter johannae sp. nov. and Gluconacetobacter azotocaptans sp. nov., associated with coffee plants. Int. J. Syst. Evol. Microbiol., 2001, 51, 1305-1314.
Gluconacetobacter liquefaciens	BR 10543	LMG 1381 = ATCC 14835 = CCUG 18124 = CIP 103109 = DSM 5603 = IAM 1834 = NBRC 12388 = JCM 17840 = LMG 1382 = NCCB 76052	YAMADA (Y.), HOSHINO (K.) and ISHIKAWA (T.): The phylogeny of acetic acid bacteria based on the partial sequences of 16S ribosomal RNA: the elevation of the subgenus Gluconoacetobacter to generic level. Biosci. Biotech. Biochem., 1997, 61, 1244-1251.
Gluconacetobacter sacchari	BR 10698	DSM 12717 = SRI 1794 = CIP 106693	FRANKE (I.H.), FEGAN (M.), HAYWARD (C.), LEONARD (G.), STACKEBRANDT (E.) and SLY (L.I.): Description of Gluconacetobacter sacchari sp. nov., a new species of acetic acid bacterium isolated from the leaf sheath of sugar cane and from the pink sugar-cane mealy bug. Int. J. Syst. Bacteriol., 1999, 49, 1681-1693.

Herbaspirillum autotrophicum	BR 10684	LMG 4326 = ATCC 29984 = CCUG 12808 = DSM 732 = IAM 14942 = JCM 21424 = NBRC 15327 = VKM B-1394	DING (L.) and YOKOTA (A.): Proposals of Curvibacter gracilis gen. nov., sp. nov. and Herbaspirillum putei sp. nov. for bacterial strains isolated from well water and reclassification of [Pseudomonas] huttiensis, [Pseudomonas] lanceolata, [Aquaspirillum] delicatum and [Aquaspirillum] autotrophicum as Herbaspirillum huttiense comb. nov., Curvibacter lanceolatus comb. nov., Curvibacter delicatus comb. nov. and Herbaspirillum autotrophicum comb. nov. Int. J. Syst. Evol. Microbiol., 2004, 54, 2223-2230.
Herbaspirillum frisingense	BR 11790	GSF 30 = DSM 13128 = LMG 23164	KIRCHHOF (G.), ECKERT (B.), STOFFELS (M.), BALDANI (J.I.), REIS (V.M.) and HARTMANN (A.): Herbaspirillum frisingense sp. nov., a new nitrogen-fixing bacterial species that occurs in C4-fibre plants. Int. J. Syst. Evol. Microbiol., 2001, 51, 157-168.
Herbaspirillum hiltneri	BR 10685	LMG 23131 = N3 = DSM 17495	ROTHBALLER (M.), SCHMID (M.), KLEIN (I.), GATTINGER (A.), GRUNDMANN (S.) and HARTMANN (A.): Herbaspirillum hiltneri sp. nov., isolated from surface-sterilized wheat roots. Int. J. Syst. Evol. Microbiol., 2006, 56, 1341-1348.
Herbaspirillum huttiense	BR 10683	LMG 2199 = ATCC 14670 = CIP 103296 = DSM 10281 = IAM 14941 = JCM 21423 = NBRC 102521	DING (L.) and YOKOTA (A.): Proposals of Curvibacter gracilis gen. nov., sp. nov. and Herbaspirillum putei sp. nov. for bacterial strains isolated from well water and reclassification of [Pseudomonas] huttiensis, [Pseudomonas] lanceolata, [Aquaspirillum] delicatum and [Aquaspirillum] autotrophicum as Herbaspirillum huttiense comb. nov., Curvibacter lanceolatus comb. nov., Curvibacter delicatus comb. nov. and Herbaspirillum autotrophicum comb. nov. Int. J. Syst. Evol. Microbiol., 2004, 54, 2223-2230.
Herbaspirillum lusitanum	BR 10326	LMG 21710 = CCUG 48869 = CECT 5661 = CIP 108242 = DSM 17154	VALVERDE (A.), VELÁZQUEZ (E.), GUTIÉRREZ (C.), CERVANTES (E.), VENTOSA (A.) and IGUAL (J.M.): Herbaspirillum lusitanum sp. nov., a novel nitrogen-fixing bacterium associated with root nodules of Phaseolus vulgaris. Int. J. Syst. Evol. Microbiol., 2003, 53, 1979-1983.

Herbaspirillum rubrisubalbicans	BR 11192	M 4 = ATCC 19308 = DSM 9440 = DSM 11543 = BCRC 15833 = LMG 2286	BALDANI (J.I.), POT (B.), KIRCHHOF (G.), FALSEN (E.), BALDANI (V.L.D.), OLIVARES (F.L.), HOSTE (B.), KERSTERS (K.), HARTMANN (A.), GILLIS (M.) and DÖBEREINER (J.): Emended description of Herbaspirillum; inclusion of [Pseudomonas] rubrisubalbicans, a mild plant pathogen, as Herbaspirillum rubrisubalbicans comb. nov.; and classification of a group of clinical isolates (EF group 1) as Herbaspirillum species 3. Int. J. Syst. Bacteriol., 1996, 46, 802-810.
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Komagataeibacter hansenii	BR 10544	LMG 1527 = ATCC 35959 = BCC 6318 = CCUG 18123 = DSM 5602 = JCM 7643 = NBRC 14820 = NCIMB 8746	YAMADA (Y.), YUKPAN (P.), VU (H.T.L.), MURAMATSU (Y.), OCHAIKUL (D.), TANASAPUWAT (S.) and NAKAGAWA (Y.): Description of Komagataeibacter gen. nov., with proposals of new combinations (Acetobacteraceae). J. Gen. Appl. Microbiol., 2012, 58, 397-404.
Komagataeibacter intermedius	BR 10550	LMG 18909 = TF2 = BCC 36447 = CIP 105780 = DSM 11804 = JCM 16936	YAMADA (Y.), YUKPAN (P.), VU (H.T.L.), MURAMATSU (Y.), OCHAIKUL (D.), TANASAPUWAT (S.) and NAKAGAWA (Y.): Description of Komagataeibacter gen. nov., with proposals of new combinations (Acetobacteraceae). J. Gen. Appl. Microbiol., 2012, 58, 397-404.
Komagataeibacter kakiaceti	BR 10554	LMG 26206 = G5-1 = JCM 25156 = NRIC 0798	YAMADA, Y. 2014. Transfer of Gluconacetobacter kakiaceti, Gluconacetobacter medellinensis and Gluconacetobacter maltaceti to the genus Komagataeibacter as Komagataeibacter kakiaceti comb. nov., Komagataeibacter medellinensis comb. nov. and Komagataeibacter maltaceti comb. nov. Int. J. Syst. Evol. Microbiol., 64, 1670-1672.

Komagataeibacter kombuchae (Gluconacetobacter kombuchae)	BR 10553	LMG 23726 = RG3 = MTCC 6913	YAMADA (Y.), YUKPAN (P.), VU (H.T.L.), MURAMATSU (Y.) OCHAIKUL (D.) and NAKAGAWA (Y.): Subdivision of the genus Gluconacetobacter Yamada, Hoshino and Ishikawa 1998: the proposal of Komagatabacter gen. nov., for strains accomodated to the Gluconacetobacter xylinus group in the α-Proteobacteria. Ann. Microbiol., 2012, 62, 849-859.
Komagataeibacter maltaceti	BR 10545	LMG 1529 = NBRC 14815 = NCIMB 8752	YAMADA, Y. 2014. Transfer of Gluconacetobacter kakiaceti, Gluconacetobacter medellinensis and Gluconacetobacter maltaceti to the genus Komagataeibacter as Komagataeibacter kakiaceti comb. nov., Komagataeibacter medellinensis comb. nov. and Komagataeibacter maltaceti comb. nov. Int. J. Syst. Evol. Microbiol., 64, 1670-1672.
Komagataeibacter medellinensis	BR 10548	LMG 1693 = NBRC 3288 = Kondo 51	YAMADA, Y. 2014. Transfer of Gluconacetobacter kakiaceti, Gluconacetobacter medellinensis and Gluconacetobacter maltaceti to the genus Komagataeibacter as Komagataeibacter kakiaceti comb. nov., Komagataeibacter medellinensis comb. nov. and Komagataeibacter maltaceti comb. nov. Int. J. Syst. Evol. Microbiol., 64, 1670-1672.
Komagataeibacter nataicola	BR 10546	LMG 1536 = BCC 36443 = JCM 25120 = LMG 1536 = NRIC 0616	YAMADA (Y.), YUKPAN (P.), VU (H.T.L.), MURAMATSU (Y.), OCHAIKUL (D.), TANASAPUWAT (S.) and NAKAGAWA (Y.): Description of Komagataeibacter gen. nov., with proposals of new combinations (Acetobacteraceae). J. Gen. Appl. Microbiol., 2012, 58, 397-404.
Komagataeibacter rhaeticus	BR 10552	LMG 22126 = DST GL02 = BCC 36452 = DSM 16663 = JCM 17122	YAMADA (Y.), YUKPAN (P.), VU (H.T.L.), MURAMATSU (Y.), OCHAIKUL (D.), TANASAPUWAT (S.) and NAKAGAWA (Y.): Description of Komagataeibacter gen. nov., with proposals of new combinations (Acetobacteraceae). J. Gen. Appl. Microbiol., 2012, 58, 397-404.
Komagataeibacter saccharivorans	BR 10547	LMG 1582 = BCC 36444 = JCM 25121 = NRIC 0614	YAMADA (Y.), YUKPAN (P.), VU (H.T.L.), MURAMATSU (Y.), OCHAIKUL (D.), TANASAPUWAT (S.) and NAKAGAWA (Y.): Description of Komagataeibacter gen. nov., with proposals of new combinations (Acetobacteraceae). J. Gen. Appl. Microbiol., 2012, 58, 397-404.

Komagataeibacter swingsii	BR10551	LMG 22125 = DST GL01 = BCC 36451 = DSM 16373 = JCM 17123	YAMADA (Y.), YUKPAN (P.), VU (H.T.L.), MURAMATSU (Y.), OCHAIKUL (D.), TANASAPUWAT (S.) and NAKAGAWA (Y.): Description of Komagataeibacter gen. nov., with proposals of new combinations (Acetobacteraceae). J. Gen. Appl. Microbiol., 2012, 58, 397-404.
Mesorhizobium albiziae	BR 10306	LMG 23507 = CCBAU 61158 = USDA 4964	WANG (F.Q.), WANG (E.T.), LIU (J.), CHEN (Q.), SUI (X.H.), CHEN (W.F.) and CHEN (W.X.): Mesorhizobium albiziae sp. nov., a novel bacterium that nodulates Albizia kalkora in a subtropical region of China. Int. J. Syst. Evol. Microbiol., 2007, 57, 1192-1199.
Mesorhizobium amorphae	BR 10216	CNPSo 2068 = ACCC 19665 = LMG 18977	WANG (E.T.), VAN BERKUM (P.), SUI (X.H.), BEYENE (D.), CHEN (W.X.) and MARTÍNEZ-ROMERO (E.): Diversity of rhizobia associated with Amorpha fruticosa isolated from Chinese soils and description of Mesorhizobium amorphae sp. nov. Int. J. Syst. Bacteriol., 1999, 49, 51-65.
Mesorhizobium caraganae	BR 10308	LMG 24397 = CCBAU 11299 = HAMBI 2990	GUAN (S.H.), CHEN (W.F.), WANG (E.T.), LU (Y.L.), YAN (X.R.), ZHANG (X.X.) and CHEN (W.X.): Mesorhizobium caraganae sp. nov., a novel rhizobial species nodulated with Caragana spp. in China. Int. J. Syst. Evol. Microbiol., 2008, 58, 2646-2653.
Mesorhizobium chacoense	BR 10302	LMG 19008 = CECT 5336 = DSM 17287	VELÁZQUEZ (E.), IGUAL (J.M.), WILLEMS (A.), FERNÁNDEZ (M.P.), MUÑOZ (E.), MATEOS (P.F.), ABRIL (A.), TORO (N.), NORMAND (P.), CERVANTES (E.), GILLIS (M.) and MARTÍNEZ-MOLINA (E.): Mesorhizobium chacoense sp. nov., a novel species that nodulates Prosopis alba in the Chaco Arido region (Argentina). Int. J. Syst. Evol. Microbiol., 2001, 51, 1011-1021.
Mesorhizobium ciceri	BR 521	USDA 3383 = ATCC 51585 = BCRC 15795 = DSM 11540 = HAMBI 1750 = LMG 14989	JARVIS (B.D.W.), VAN BERKUM (P.), CHEN (W.X.), NOUR (S.M.), FERNANDEZ (M.P.), CLEYET-MAREL (J.C.) and GILLIS (M.): Transfer of Rhizobium loti, Rhizobium huakuii, Rhizobium ciceri, Rhizobium mediterraneum, and Rhizobium tianshanense to Mesorhizobium gen. nov. Int. J. Syst. Bacteriol., 1997, 47, 895-898.

Mesorhizobium huakuii	BR 524	USDA 4779 = ATCC 51122 = BCRC 15723 = DSM 6573 = HAMBI 1674 = LMG 14107	JARVIS (B.D.W.), VAN BERKUM (P.), CHEN (W.X.), NOUR (S.M.), FERNANDEZ (M.P.), CLEYET-MAREL (J.C.) and GILLIS (M.): Transfer of Rhizobium loti, Rhizobium huakuii, Rhizobium ciceri, Rhizobium mediterraneum, and Rhizobium tianshanense to Mesorhizobium gen. nov. Int. J. Syst. Bacteriol., 1997, 47, 895-898.
Mesorhizobium loti	BR 7801	LMG 6125 = ATCC 700743 = DSM 2626 = HAMBI 1129	JARVIS (B.D.W.), VAN BERKUM (P.), CHEN (W.X.), NOUR (S.M.), FERNANDEZ (M.P.), CLEYET-MAREL (J.C.) and GILLIS (M.): Transfer of Rhizobium loti, Rhizobium huakuii, Rhizobium ciceri, Rhizobium mediterraneum, and Rhizobium tianshanense to Mesorhizobium gen. nov. Int. J. Syst. Bacteriol., 1997, 47, 895-898.
Mesorhizobium mediterraneum	BR 522	USDA 3392 = ATCC 51670 = BCRC 15796 = DSM 11555 = HAMBI 2096 = LMG 14994 = NBRC 102497	JARVIS (B.D.W.), VAN BERKUM (P.), CHEN (W.X.), NOUR (S.M.), FERNANDEZ (M.P.), CLEYET-MAREL (J.C.) and GILLIS (M.): Transfer of Rhizobium loti, Rhizobium huakuii, Rhizobium ciceri, Rhizobium mediterraneum, and Rhizobium tianshanense to Mesorhizobium gen. nov. Int. J. Syst. Bacteriol., 1997, 47, 895-898.
Mesorhizobium plurifarium	BR 10305	LMG 11892 = HAMBI 208 = NBRC 102498	DE LAJUDIE (P.), WILLEMS (A.), NICK (G.), MOREIRA (F.), MOLOUBA (F.), HOSTE (B.), TORCK (U.), NEYRA (M.), COLLINS (M.D.), LINDSTRÖM (K.), DREYFUS (B.) and GILLIS (M.): Characterization of tropical tree rhizobia and description of Mesorhizobium plurifarium sp. nov. Int. J. Syst. Bacteriol., 1998, 48, 369-382.
Mesorhizobium silamurunense	BR 10309	LMG 24822 = CCBAU 01550 = HAMBI 3029 = DSM 29845	ZHAO (C.T.), WANG (E.T.), ZHANG (Y.M.), CHEN (W.F.), SUI (X.H.), CHEN (W.X.), LIU (H.C.) and ZHANG (X.X.): Mesorhizobium silamurunense sp. nov., isolated from root nodules of Astragalus species. Int. J. Syst. Evol. Microbiol., 2012, 62, 2180-2186.

Mesorhizobium tamadayense	BR 10320	LMG 26736 = CECT 8040	RAMÍREZ-BAHENA (M.H.), HERNÁNDEZ (M.), PEIX (A.), VELÁZQUEZ (E.) and LEÓN-BARRIOS (M.): Mesorhizobial strains nodulating Anagyris latifolia and Lotus berthelotii in Tamadaya ravine (Tenerife, Canary Islands) are two symbiovars of the same species, Mesorhizobium tamadayense sp. nov. Syst. Appl. Microbiol., 2012, 35, 334-341.
Mesorhizobium tarimense	BR 10533	LMG 24338 = CCBAU 83306 = HAMBI 2973	HAN (T.X.), HAN (L.L.), WU (L.J.), CHEN (W.F.), SUI (X.H.), GU (J.G.), WANG (E.T.) and CHEN (W.X.): Mesorhizobium gobiense sp. nov. and Mesorhizobium tarimense sp. nov., isolated from wild legumes growing in desert soils of Xinjiang, China. Int. J. Syst. Evol. Microbiol., 2008, 58, 2610-2618.
Mesorhizobium tianshanense	BR 523	USDA 3592 = BCRC 15797 = DSM 11417 = HAMBI 1870 = LMG 15767 = NBRC 102499	JARVIS (B.D.W.), VAN BERKUM (P.), CHEN (W.X.), NOUR (S.M.), FERNANDEZ (M.P.), CLEYET-MAREL (J.C.) and GILLIS (M.): Transfer of Rhizobium loti, Rhizobium huakuii, Rhizobium ciceri, Rhizobium mediterraneum, and Rhizobium tianshanense to Mesorhizobium gen. nov. Int. J. Syst. Bacteriol., 1997, 47, 895-898.
Methylobacterium nodulans	BR 10699	LMG 21967 = CNCM I 2342 = ORS 2060	JOURAND (P.), GIRAUD (E.), BÉNA (G.), SY (A.), WILLEMS (A.), GILLIS (M.), DREYFUS (B.) and DE LAJUDIE (P.): Methylobacterium nodulans sp. nov., for a group of aerobic, facultatively methylotrophic, legume root-nodule-forming and nitrogen-fixing bacteria. Int. J. Syst. Evol. Microbiol., 2004, 54, 2269-2273.
Microvirga lotononidis	BR 10204	WSM3557 = LMG 26455 = HAMBI 3237	ARDLEY (J.K.), PARKER (M.A.), DE MEYER (S.E.), TRENGOVE (R.D.), O'HARA (G.W.), REEVE (W.G.), YATES (R.J.), DILWORTH (M.J.), WILLEMS (A.) and HOWIESON (J.G.): Microvirga lupini sp. nov., Microvirga lotononidis sp. nov. and Microvirga zambiensis sp. nov. are alphaproteobacterial root-nodule bacteria that specifically nodulate and fix nitrogen with geographically and taxonomically separate legume hosts. Int. J. Syst. Evol. Microbiol., 2012, 62, 2579-2588.

Microvirga lupini	BR 10202	Lut6 = LMG 26460 = HAMBI 3236	ARDLEY (J.K.), PARKER (M.A.), DE MEYER (S.E.), TRENGOVE (R.D.), O'HARA (G.W.), REEVE (W.G.), YATES (R.J.), DILWORTH (M.J.), WILLEMS (A.) and HOWIESON (J.G.): Microvirga lupini sp. nov., Microvirga lotononidis sp. nov. and Microvirga zambiensis sp. nov. are alphaproteobacterial root-nodule bacteria that specifically nodulate and fix nitrogen with geographically and taxonomically separate legume hosts. Int. J. Syst. Evol. Microbiol., 2012, 62, 2579-2588.
Microvirga subterranea	BR 10191	DSM 14364 = ATCC BAA-295	KANSO (S.) and PATEL (B.K.C.): Microvirga subterranea gen. nov., sp. nov., a moderate thermophile from a deep subsurface Australian thermal aquifer. Int. J. Syst. Evol. Microbiol., 2003, 53, 401-406.
Microvirga vignae	BR 3299	7S60 = HAMBI 3457	RADL, V., SIMÕES-ARAÚJO, J. L., LEITE, J., PASSOS, S. R., MARTINS, L. M., XAVIER, G. R., RUMJANEK, N. G., BALDANI, J. I. and ZILLI, J. E. 2014. Microvirga vignae sp. nov., a root nodule symbiotic bacterium isolated from cowpea grown in semi-arid Brazil. Int. J. Syst. Evol. Microbiol., 64, 725-730.
Microvirga zambiensis	BR 10203	WSM3693 = LMG 26454 = HAMBI 3238	ARDLEY (J.K.), PARKER (M.A.), DE MEYER (S.E.), TRENGOVE (R.D.), O'HARA (G.W.), REEVE (W.G.), YATES (R.J.), DILWORTH (M.J.), WILLEMS (A.) and HOWIESON (J.G.): Microvirga lupini sp. nov., Microvirga lotononidis sp. nov. and Microvirga zambiensis sp. nov. are alphaproteobacterial root-nodule bacteria that specifically nodulate and fix nitrogen with geographically and taxonomically separate legume hosts. Int. J. Syst. Evol. Microbiol., 2012, 62, 2579-2588.
Nitrospirillum amazonense (Azospirillum amazonense)	BR 11142	Y 1 = ATCC 35119 = DSM 2787 = BCRC 14279 = LMG 22237	LIN, S.-Y., HAMEED, A., SHEN, F.-T., LIU, Y.-C., HSU, Y.-H., SHAHINA, M., LAI, W.-A. and YOUNG, C.-C. 2014. Description of Niveispirillum fermenti gen. nov., sp. nov., isolated from a fermentor in Taiwan, transfer of Azospirillum irakense (1989) as Niveispirillum irakense comb. nov., and reclassification of Azospirillum amazonense (1983) as Nitrospirillum amazonense gen. nov. Antonie van Leeuwenhoek, 105, 1149-1162.

Niveispirillum irakense	BR 10694	LMG 10653 = KBC1 = ATCC 51182 = BCRC 15764 = CIP 103311	LIN, S. Y., HAMEED, A., SHEN, F. T., LIU, Y. C., HSU, Y. H., SHAHINA, M., LAI, W. A. and YOUNG, C. C. 2014. Description of Niveispirillum fermenti gen. nov., sp. nov., isolated from a fermentor in Taiwan, transfer of Azospirillum irakense (1989) as Niveispirillum irakense comb. nov., and reclassification of Azospirillum amazonense (1983) as Nitrospirillum amazonense gen. nov. Antonie van Leeuwenhoek, 105, 1149-1162.
Noviherbaspirillum canariense (Herbaspirillum canariense)	BR 10687	LMG 26151 = SUEMI03 = CECT 7838	LIN, S.-Y., HAMEED, A., ARUN, A. B., LIU, Y.-C., HSU, Y.-H., LAI, W.-A., REKHA, P. D. and YOUNG, C.-C. 2013. Description of Noviherbaspirillum malthae gen. nov., sp. nov., isolated from an oil-contaminated soil, and proposal to reclassify Herbaspirillum soli, Herbaspirillum aurantiacum, Herbaspirillum canariense and Herbaspirillum psychrotolerans as Noviherbaspirillum soli comb. nov., Noviherbaspirillum aurantiacum comb. nov., Noviherbaspirillum canariense comb. nov. and Noviherbaspirillum psychrotolerans comb. nov. based on polyphasic analysis. Int. J. Syst. Evol. Microbiol., 63, 4100-4107.
Noviherbaspirillum soli (Herbaspirillum soli)	BR 10686	LMG 26149 = SUEMI10 = CECT 7840	LIN, S.-Y., HAMEED, A., ARUN, A. B., LIU, Y.-C., HSU, Y.-H., LAI, W.-A., REKHA, P. D. and YOUNG, C.-C. 2013. Description of Noviherbaspirillum malthae gen. nov., sp. nov., isolated from an oil-contaminated soil, and proposal to reclassify Herbaspirillum soli, Herbaspirillum aurantiacum, Herbaspirillum canariense and Herbaspirillum psychrotolerans as Noviherbaspirillum soli comb. nov., Noviherbaspirillum aurantiacum comb. nov., Noviherbaspirillum canariense comb. nov. and Noviherbaspirillum psychrotolerans comb. nov. based on polyphasic analysis. Int. J. Syst. Evol. Microbiol., 63, 4100-4107.
Ochrobactrum ciceri	BR 10332	DSM 22292 = CCUG 57879	IMRAN (A.), HAFEEZ (F.Y.), FRÜHLING (A.), SCHUMANN (P.), MALIK (K.A.) and STACKEBRANDT (E.): Ochrobactrum ciceri sp. nov., isolated from nodules of Cicer arietinum. Int. J. Syst. Evol. Microbiol., 2010, 60, 1548-1553.

Ochrobactrum cytisi	BR 10324	LMG 22713 = CCUG 54646 = CECT 7172 = CIP 109590 = DSM 19778	ZURDO-PIÑEIRO (J.L.), RIVAS (R.), TRUJILLO (M.E.), VIZCAÍNO (N.), CARRASCO (J.A.), CHAMBER (M.), PALOMARES (A.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.) and VELÁZQUEZ (E.): Ochrobactrum cytisi sp. nov., isolated from nodules of Cytisus scoparius in Spain. Int. J. Syst. Evol. Microbiol., 2007, 57, 784-788.
Ochrobactrum grignonense	BR 10322	LMG 18954 = BCRC 17249 = DSM 13338 = NBRC 102586	LEBUHN (M.), ACHOUAK (W.), SCHLOTER (M.), BERGE (O.), MEIER (H.), BARAKAT (M.), HARTMANN (A.) and HEULIN (T.): Taxonomic characterization of Ochrobactrum sp. isolates from soil samples and wheat roots, and description of Ochrobactrum tritici sp. nov. and Ochrobactrum grignonense sp. nov. Int. J. Syst. Evol. Microbiol. 2000, 50, 2207-2223.
Ochrobactrum lupini	BR 9078	DSM 16930 = LMG 22726 = NBRC 102587	TRUJILLO (M.E.), WILLEMS (A.), ABRIL (A.), PLANCHUELO (A.M.), RIVAS (R.), LUDEÑA (D.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.) and VELÁZQUEZ (E.): Nodulation of Lupinus albus by strains of Ochrobactrum lupini sp. nov. Appl. Environ. Microbiol., 2005, 71, 1318-1327.
Ochrobactrum oryzae	BR 10329	DSM 17471 = MTCC 4195 = NBRC 102588	TRIPATHI (A.K.), VERMA (S.C.), CHOWDHURY (S.P.), LEBUHN (M.), GATTINGER (A.) and SCHLOTER (M.): Ochrobactrum oryzae sp. nov., an endophytic bacterial species isolated from deep-water rice in India. Int. J. Syst. Evol. Microbiol., 2006, 56, 1677-1680.
Ochrobactrum tritici	BR 10323	LMG 18957 = BCRC 17250 = DSM 13340 = NBRC 102585	LEBUHN (M.), ACHOUAK (W.), SCHLOTER (M.), BERGE (O.), MEIER (H.), BARAKAT (M.), HARTMANN (A.) and HEULIN (T.): Taxonomic characterization of Ochrobactrum sp. isolates from soil samples and wheat roots, and description of Ochrobactrum tritici sp. nov. and Ochrobactrum grignonense sp. nov. Int. J. Syst. Evol. Microbiol. 2000, 50, 2207-2223.
Paraburkholderia caballeronis	BR 10519	LMG 26416 = TNe-841 = CIP 110324	MARTÍNEZ-AGUILAR, L., SALAZAR-SALAZAR, C., MÉNDEZ, R. D., CABALLERO-MELLADO, J., HIRSCH, A. M., VÁSQUEZ-MURRIETA, M. S. and ESTRADA-DE LOS SANTOS, P. 2014. Burkholderia caballeronis sp. nov., a nitrogen fixing species isolated from tomato (Lycopersicon esculentum) with the ability to effectively nodulate Phaseolus vulgaris. Antonie van Leeuwenhoek, 104, 1063-1071.

Paraburkholderia caledonica (Burkholderia caledonica)	BR 12329	ATCC BAA-462 = LMG 19076	COENYE (T.), LAEVENS (S.), WILLEMS (A.), OHLEN (M.), HANNANT (W.), GOVAN (J.R.W.), GILLIS (M.), FALSEN (E.) and VANDAMME (P.): Burkholderia fungorum sp. nov. and Burkholderia caledonica sp. nov., two new species isolated from the environment, animals and human clinical samples. Int. J. Syst. Evol. Microbiol., 2001, 51, 1099-1107. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia cariophylli (Burkholderia cariophylli)	BR 12316	ATCC 25418 = DSM 50341	YABUUCHI (E.), KOSAKO (Y.), OYAIZU (H.), YANO (I.), HOTTA (H.), HASHIMOTO (Y.), EZAKI (T.) and ARAKAWA (M.): Proposal of Burkholderia gen. nov. and transfer of seven species of the genus Pseudomonas homology group II to the new genus, with the type species Burkholderia cepacia (Palleroni and Holmes 1981) comb. nov. Microbiol. Immunol., 1992, 36, 1251-1275. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.

Paraburkholderia diazotrophica (Burkholderia diazotrophica)	BR 10319	LMG 26031 = BCRC 80259 = JPY 461 = KCTC 23308	SHEU (S.Y.), CHOU (J.H.), BONTEMPS (C.), ELLIOTT (G.N.), GROSS (E.), DOS REIS JUNIOR (F.B.), MELKONIAN (R.), MOULIN (L.), JAMES (E.K.), SPRENT (J.I.), YOUNG (J.P.W.) and CHEN (W.M.): Burkholderia diazotrophica sp. nov., isolated from root nodules of Mimosa spp. Int. J. Syst. Evol. Microbiol., 2013, 63, 435-441. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia dilworthii (Burkholderia dilworthii)	BR 10614	LMG 27173 = WSM3556 = HAMBI 3353	DE MEYER, S. E., CNOCKAERT, M., ARDLEY, J. K., VAN WYK, B.-E., VANDAMME, P. A. and HOWIESON, J. G. 2014. Burkholderia dilworthii sp. nov., isolated from Lebeckia ambigua root nodules. Int. J. Syst. Evol. Microbiol., 64, 1090-1095. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet. 5, 429.
Paraburkholderia glathei (Burkholderia glathei)	BR 12321	ATCC 29195 = DSM 50014	VIALLARD (V.), POIRIER (I.), COURNOYER (B.), HAURAT (J.), WIEBKIN (S.), OPHEL-KELLER (K.) and BALANDREAU (J.): Burkholderia graminis sp. nov., a rhizospheric Burkholderia species, and reassessment of [Pseudomonas] phenazinium, [Pseudomonas] pyrrocinia and [Pseudomonas] glathei as Burkholderia. Int. J. Syst. Bacteriol., 1998, 48, 549-563. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.

Paraburkholderia graminis (Burkholderia graminis)	BR 12323	C4D1M = ATCC 700544 = DSM 17151 = LMG 18924	VIALLARD (V.), POIRIER (I.), COURNOYER (B.), HAURAT (J.), WIEBKIN (S.), OPHEL-KELLER (K.) and BALANDREAU (J.): Burkholderia graminis sp. nov., a rhizospheric Burkholderia species, and reassessment of [Pseudomonas] phenazinium, [Pseudomonas] pyrrocinia and [Pseudomonas] glathei as Burkholderia. Int. J. Syst. Bacteriol., 1998, 48, 549-563. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia kururiensis (Burkholderia kururiensis)	BR 11897	KP 23 = ATCC 700977 = DSM 13646 = LMG 19447	ZHANG (H.), HANADA (S.), SHIGEMATSU (T.), SHIBUYA (K.), KAMAGATA (Y.), KANAGAWA (T.) and KURANE (R.): Burkholderia kururiensis sp. nov., a trichloroethylene (TCE)-degrading bacterium isolated from an aquifer polluted with TCE. Int. J. Syst. Evol. Microbiol., 2000, 50, 743-749. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia mimosarum (Burkholderia mimosarum)	BR 10610	LMG 23256 = PAS44 = CCUG 54296	CHEN (W.M.), JAMES (E.K.), COENYE (T.), CHOU (J.H.), BARRIOS (E.), DE FARIA (S.M.), ELLIOTT (G.N.), SHEU (S.Y.), SPRENT (J.I.) and VANDAMME (P.): Burkholderia mimosarum sp. nov., isolated from root nodules of Mimosa spp. from Taiwan and South America. Int. J. Syst. Evol. Microbiol., 2006, 56, 1847-1851. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.

Paraburkholderia nodosa (Burkholderia nodosa)	BR 3437	LMG 23741 = BCRC 17575	CHEN (W.M.), DE FARIA (S.M.), JAMES (E.K.), ELLIOTT (G.N.), LIN (K.Y.), CHOU (J.H.), SHEU (S.Y.), CNOCKAERT (M.), SPRENT (J.I.) and VANDAMME (P.): Burkholderia nodosa sp. nov., isolated from root nodules of the woody Brazilian legumes Mimosa bimucronata and Mimosa scabrella. Int. J. Syst. Evol. Microbiol., 2007, 57, 1055-1059. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia phenoliruptrix (Burkholderia phenoliruptrix)	BR 3459	AC 1100 = ATCC 39027 = ATCC 53867 = CIP 108591 = DSM 17773 = LMG 22037	COENYE (T.), HENRY (D.), SPEERT (D.P.) and VANDAMME (P.): Burkholderia phenoliruptrix sp. nov., to accommodate the 2,4,5-trichlorophenoxyacetic acid and halophenol-degrading strain AC1100. Syst. Appl. Microbiol., 2004, 27, 623-627. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia phymatum (Burkholderia phymatum)	BR 3486	STM 815 = DSM 17167 = LMG 21445	VANDAMME (P.), GORIS (J.), CHEN (W.M.), DE VOS (P.) and WILLEMS (A.): Burkholderia tuberum sp. nov. and Burkholderia phymatum sp. nov., nodulate the roots of tropical legumes. Syst. Appl. Microbiol., 2002, 25, 507-512. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.

Paraburkholderia rhynchosiae	BR 10615	LMG 27174 = WSM3937 = HAMBI 3354	DE MEYER, S. E., CNOCKAERT, M., ARDLEY, J. K., TRENGOVE, R. D., GARAU, G., HOWIESON, J. G. and VANDAMME, P. 2013. Burkholderia rhynchosiae sp. nov., isolated from Rhynchosia ferulifolia root nodules. Int. J. Syst. Evol. Microbiol., 63, 3944-3949.
Paraburkholderia sabiae (Burkholderia sabiae)	BR 3407	LMG 24235 = BCRC 17587 = SEMIA 6383	CHEN (W.M.), DE FARIA (S.M.), CHOU (J.H.), JAMES (E.K.), ELLIOTT (G.N.), SPRENT (J.I.), BONTEMPS (C.), YOUNG (J.P.W.) and VANDAMME (P.): Burkholderia sabiae sp. nov., isolated from root nodules of Mimosa caesalpiniifolia. Int. J. Syst. Evol. Microbiol., 2008, 58, 2174-2179. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia silvatlantica (Burkholderia silvatlantica)	BR 11907	SRMrh 20 = ATCC BAA-1244 = LMG 23149	PERIN (L.), MARTÍNEZ-AGUILAR (L.), PAREDES-VALDEZ (G.), BALDANI (J.I.), ESTRADA-DE LOS SANTOS (P.), REIS (V.M.) and CABALLERO-MELLADO (J.): Burkholderia silvatlantica sp. nov., a diazotrophic bacterium associated with sugar cane and maize. Int. J. Syst. Evol. Microbiol., 2006, 56, 1931-1937. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.

Paraburkholderia sprentiae (Burkholderia sprentiae)	BR 10616	LMG 27175 = WSM5005 = HAMBI 3357	DE MEYER, S. E., CNOCKAERT, M., ARDLEY, J. K., MAKER, G., YATES, R., HOWIESON, J. G. and VANDAMME, P. 2013. Burkholderia sprentiae sp. nov., isolated from Lebeckia ambigua root nodules. Int. J. Syst. Evol. Microbiol., 63, 3950-3957. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet. 5, 429.
Paraburkholderia symbiotica (Burkholderia symbiotica)	BR 10341	LMG 26032 = BCRC 80258 = JPY 345 = KCTC 23309	SHEU (S.Y.), CHOU (J.H.), BONTEMPS (C.), ELLIOTT (G.N.), GROSS (E.), JAMES (E.K.), SPRENT (J.I.), YOUNG (J.P.W.) and CHEN (W.M.): Burkholderia symbiotica sp. nov., isolated from root nodules of Mimosa spp. native to north-east Brazil. Int. J. Syst. Evol. Microbiol., 2012, 62, 2272-2278. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet. 5, 429.
Paraburkholderia tropica (Burkholderia tropica)	BR 11366	Ppe 8 = ATCC BAA-831 = DSM 15359 = LMG 22274	REIS (V.M.), ESTRADA-DE LOS SANTOS (P.), TENORIO-SALGADO (S.), VOGEL (J.), STOFFELS (M.), GUYON (S.), MAVINGUI (P.), BALDANI (V.L.D.), SCHMID (M.), BALDANI (J.I.), BALANDREAU (J.), HARTMANN (A.) and CABALLERO-MELLADO (J.): Burkholderia tropica sp. nov., a novel nitrogen-fixing, plant-associated bacterium. Int. J. Syst. Evol. Microbiol., 2004, 54, 2155-2162. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.

Paraburkholderia tuberum (Burkholderia tuberum)	BR 3487	STM 678 = DSM 18489 = LMG 21444	VANDAMME (P.), GORIS (J.), CHEN (W.M.), DE VOS (P.) and WILLEMS (A.): Burkholderia tuberum sp. nov. and Burkholderia phymatum sp. nov., nodulate the roots of tropical legumes. Syst. Appl. Microbiol., 2002, 25, 507-512. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Paraburkholderia unamae (Burkholderia unamae)	BR 12203	MTI 641 = ATCC BAA-744 = DSM 17197 = LMG 22722	CABALLERO-MELLADO (J.), MARTÍNEZ-AGUILAR (L.), PAREDES-VALDEZ (G.) and ESTRADA-DE LOS SANTOS (P.): Burkholderia unamae sp. nov., an N2-fixing rhizospheric and endophytic species. Int. J. Syst. Evol. Microbiol., 2004, 54, 1165-1172. Sawana, A., Adeolu, M. and Gupta, R.S. 2014. Molecular signatures and phylogenomic analysis of the genus Burkholderia: proposal for division of this genus into the emended genus Burkholderia containing pathogenic organisms and a new genus Paraburkholderia gen. nov. harboring environmental species. Front. Genet., 5, 429.
Phyllobacterium trifolii	BR 7608	DSM 17315 = LMG 22712 = CECT 7015	VALVERDE (A.), VELÁZQUEZ (E.), FERNÁNDEZ-SANTOS (F.), VIZCAÍNO (N.), RIVAS (R.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.), IGUAL (J.M.) and WILLEMS (A.): Phyllobacterium trifolii sp. nov., nodulating Trifolium and Lupinus in Spanish soils. Int. J. Syst. Evol. Microbiol., 2005, 55, 1985-1989.
Pseudomonas azotifigens	BR 10340	LMG 23662 = ATCC BAA-1049 = CCUG 53885 = CIP 108866 = DSM 17556	HATAYAMA (K.), KAWAI (S.), SHOUN (H.), UEDA (Y.) and NAKAMURA (A.): Pseudomonas azotifigens sp. nov., a novel nitrogen-fixing bacterium isolated from a compost pile. Int. J. Syst. Evol. Microbiol., 2005, 55, 1539-1544.
Pseudomonas stutzeri	BR 10342	LMG 10652 = A15 = R-4180	SIJDERIUS (R.): Heterotrophe bacterien, die thiosulfaat oxydeeren. Thesis, University Amsterdam, 1946, pp. 1-146. LEHMANN (K.B.) and NEUMANN (R.): Atlas und Grundriss der Bakteriologie und Lehrbuch der speziellen bakteriologischen Diagnostik, 1st ed., J.F. Lehmann, München, 1896.

Rhizobium aggregatum	BR 10672	DSM 1111 = Müller 161 = ATCC 43293 = IFAM 1003 = VKM B-2061	KAUR (J.), VERMA (M.) and LAL (R.): Rhizobium rosettiformans sp. nov., isolated from a hexachlorocyclohexane dump site, and reclassification of Blastobacter aggregatus Hirsch and Muller 1986 as Rhizobium aggregatum comb. nov. Int. J. Syst. Evol. Microbiol., 2011, 61, 1218-1225.
Rhizobium alamii	BR 10613	LMG 24466 = GBV016 = CFBP 7146	BERGE (O.), LODHI (A.), BRANDELET (G.), SANTAELLA (C.), RONCATO (M.A.), CHRISTEN (R.), HEULIN (T.) and ACHOUAK (W.): Rhizobium alamii sp. nov., an exopolysaccharide-producing species isolated from legume and non-legume rhizospheres. Int. J. Syst. Evol. Microbiol., 2009, 59, 367-372.
Rhizobium azibense	BR 10483	HAMBI 3541 = 23C2 = CCBAU 101087	MNASRI, B., LIU, T. Y., SAIDI, S., CHEN, W. F., CHEN, W. X., ZHANG, X. X. and MHAMDI, R. 2014. Rhizobium azibense sp. nov., a nitrogen fixing bacterium isolated from root-nodules of Phaseolus vulgaris. Int. J. Syst. Evol. Microbiol., 64, 1501-1506.
Rhizobium borbori	BR 10517	LMG 23925 = DN316 = CICC 10378	ZHANG (G.X.), REN (S.Z.), XU (M.Y.), ZENG (G.Q.), LUO (H.D.), CHEN (J.L.), TAN (Z.Y.) and SUN (G.P.): Rhizobium borbori sp. nov., aniline-degrading bacteria isolated from activated sludge. Int. J. Syst. Evol. Microbiol., 2011, 61, 816-822.
Rhizobium cellulosilyticum	BR 10611	LMG 23642 = ALA10B2 = CECT 7176 = DSM 18291	GARCÍA-FRAILE (P.), RIVAS (R.), WILLEMS (A.), PEIX (A.), MARTENS (M.), MARTÍNEZ-MOLINA (E.), MATEOS (P.F.) and VELÁZQUEZ (E.): Rhizobium cellulosilyticum sp. nov., isolated from sawdust of Populus alba. Int. J. Syst. Evol. Microbiol., 2007, 57, 844-848.
Rhizobium daejeonense	BR 10490	LMG 25353 = L61 = CCBAU 10050 = IAM 15042 = JCM 21505 = KCTC 12121 = NBRC 102495	QUAN (Z.X.), BAE (H.S.), BAEK (J.H.), CHEN (W.F.), IM (W.T.) and LEE (S.T.): Rhizobium daejeonense sp. nov., isolated from a cyanide treatment bioreactor. Int. J. Syst. Evol. Microbiol., 2005, 55, 2543-2549.
Rhizobium ecuadorense	BR 10406	CNPSo 671 = UMR 1450 = PIMAMPIRS I 5 = LMG 27578	RIBEIRO, R. A., MARTINS, T. B., ORMEÑO-ORRILLO, E., MARCON DELAMUTA, J. R., ROGEL, M. A., MARTÍNEZ-ROMERO, E. and HUNGRIA, M. 2015. Rhizobium ecuadorense sp. nov., an indigenous N2-fixing symbiont of the Ecuadorian common bean (Phaseolus vulgaris L.) genetic pool. Int. J. Syst. Evol. Microbiol., 65, 3162-3169.

Rhizobium endolithicum	BR 10485	HAMBI 3447 = JC140 = CCUG 64352 = KCTC 32077 = MTCC 11723	PARAG, B., SASIKALA, CH.. and RAMANA CH., V. 2014. Molecular and culture dependent characterization of endolithic bacteria in two beach sand samples and description of Rhizobium endolithicum sp. nov. Antonie van Leeuwenhoek, 104, 1235-1244.
Rhizobium endophyticum	BR 10314	LMG 26002 = ATCC BAA-2116 = HAMBI 3153 = CNPSo 2060	LÓPEZ-LÓPEZ (A.), ROGEL (M.A.), ORMEÑO-ORRILLO (E.), MARTÍNEZ-ROMERO (J.) and MARTÍNEZ-ROMERO (E.): Phaseolus vulgaris seed-borne endophytic community with novel bacterial species such as Rhizobium endophyticum sp. nov. Syst. Appl. Microbiol., 2010, 33, 322-327.
Rhizobium etli	BR 10026	CFN 42 = ATCC 51251 = BCRC 15792 = CECT 4651 = DSM 11541 = HAMBI 1727 = USDA 9032 = LMG 17827	SEGOVIA (L.), YOUNG (J.P.W.) and MARTÍNEZ-ROMERO (E.): Reclassification of American Rhizobium leguminosarum biovar Phaseoli type I strains as Rhizobium etli sp. nov. Int. J. Syst. Bacteriol., 1993, 43, 374-377.
Rhizobium fabae	BR 10210	CNPSo 2059 = LMG 23997	TIAN (C.F.), WANG (E.T.), WU (L.J.), HAN (T.X.), CHEN (W.F.), GU (C.T.), GU (J.G.) and CHEN (W.X.): Rhizobium fabae sp. nov., a bacterium that nodulates Vicia faba. Int. J. Syst. Evol. Microbiol., 2008, 58, 2871-2875.
Rhizobium freirei	BR 520	PRF 81 = CNPSo 122 = SEMIA 4080 = WDCM 440	DALL'AGNOL, R. F., RIBEIRO, R. A., ORMEÑO-ORRILLO, E., ROGEL, M. A., DELAMUTA, J. R. M., ANDRADE, D. S., MARTÍNEZ-ROMERO, E. and HUNGRIA, M. 2013. Rhizobium freirei sp. nov., a symbiont of Phaseolus vulgaris that is very effective at fixing nitrogen. Int. J. Syst. Evol. Microbiol., 63, 4167-4173.
Rhizobium galegae	BR 10055	LMG 6214 = HAMBI 540 = NZP 5563 = ATCC 43677	LINDSTRÖM (K.): Rhizobium galegae, a new species of legume root nodule bacteria. Int. J. Syst. Bacteriol., 1989, 39, 365-367.
Rhizobium grahamii	BR 10404	CNPSo 2485 = CCGE 502 = ATCC BAA-2124 = CFN 242 = HAMBI 3152	LÓPEZ-LÓPEZ (A.), ROGEL-HERNÁNDEZ (M.A.), BAROIS (I.), ORTIZ CEBALLOS (A.I.), MARTÍNEZ (J.), ORMEÑO-ORRILLO (E.) and MARTÍNEZ-ROMERO (E.): Rhizobium grahamii sp. nov., from nodules of Dalea leporina, Leucaena leucocephala and Clitoria ternatea, and Rhizobium mesoamericanum sp. nov., from nodules of Phaseolus vulgaris, siratro, cowpea and Mimosa pudica. Int. J. Syst. Evol. Microbiol., 2012, 62, 2264-2271.

Rhizobium halophytocola	BR 10680	DSM 21600 = YC6881 = KACC 13775	BIBI (F.), CHUNG (E.J.), KHAN (A.), JEON (C.O.) and CHUNG (Y.R.): Rhizobium halophytocola sp. nov., isolated from the root of a coastal dune plant. Int. J. Syst. Evol. Microbiol., 2012, 62, 1997-2003.
Rhizobium helanshanense	BR 10487	HAMBI 3083 = ACCC 16237	QIN (W.), DENG (Z.S.), XU (L.), WANG (N.N.), WEI (G.H.). Rhizobium helanshanense sp. nov., a bacterium that nodulates Sphaerophysa salsula (Pall.) DC. in China. Arch Microbiol. 2012 May; 194(5): 371-8.
Rhizobium herbae	BR 10489	HAMBI 3117 = CCBAU 83011 = LMG 25718	REN (D.W.), WANG (E.T.), CHEN (W.F.), SUI (X.H.), ZHANG (X.X.), LIU (H.C.) and CHEN (W.X.): Rhizobium herbae sp. nov. and Rhizobium giardinii-related bacteria, minor microsymbionts of various wild legumes in China. Int. J. Syst. Evol. Microbiol., 2011, 61, 1912-1920.
Rhizobium huautlense	BR 10396	CNPSo 2062 = S02 = ATCC BAA-115 = LMG 18254	WANG (E.T.), VAN BERKUM (P.), BEYENE (D.), SUI (X.H.), DORADO (O.), CHEN (W.X.) and MARTÍNEZ-ROMERO (E.): Rhizobium huautlense sp. nov., a symbiont of Sesbania herbacea that has a close phylogenetic relationship with Rhizobium galegae. Int. J. Syst. Bacteriol., 1998, 48, 687-699.
Rhizobium jaguaris	BR 10394	CNPSo 2465 = ATCC BAA-2445 = SJP1-2	RINCÓN-ROSALES, R., VILLALOBOS-ESCOBEDO, J. M., ROGEL, M. A., MARTINEZ, J., ORMEÑO-ORRILLO, E. and MARTÍNEZ-ROMERO, E. 2013. Rhizobium calliandrae sp. nov., Rhizobium mayense sp. nov. and Rhizobium jaguaris sp. nov., rhizobial species nodulating the medicinal legume Calliandra grandiflora. Int. J. Syst. Evol. Microbiol., 63, 3423-3429.
Rhizobium laguerreae	BR 10521	LMG 27434 = FB206 = CECT 8280	SAIDI, S., RAMÍREZ-BAHENA, M.-H., SANTILLANA, N., ZÚÑIGA, D., ÁLVAREZ-MARTÍNEZ, E., PEIX, A., MHAMDI, R. and VELÁZQUEZ, E. 2014. Rhizobium laguerreae sp. nov. nodulates Vicia faba on several continents. Int. J. Syst. Evol. Microbiol., 64, 242-247.
Rhizobium larrymoorei	BR 10515	LMG 21410 = ATCC 51759= CFBP 5473 = ICMP 14256 = NCPPB 4096	YOUNG (J.M.): Renaming of Agrobacterium larrymoorei Bouzar and Jones 2001 as Rhizobium larrymoorei (Bouzar and Jones 2001) comb. nov. Int. J. Syst. Evol. Microbiol., 2004, 54, 149.

Rhizobium leguminosarum (Rhizobium trifolii)	BR 7606	LMG 8820 = ATCC 14480 = Means 3D1K22a = BCRC 13519 = DSM 30141	RAMÍREZ-BAHENA (M.H.), GARCÍA-FRAILE (P.), PEIX (A.), VALVERDE (A.), RIVAS (R.), IGUAL (J.M.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.) and VELÁZQUEZ (E.): Revision of the taxonomic status of the species Rhizobium leguminosarum (Frank 1879) Frank 1889AL, Rhizobium phaseoli 1926AL and Rhizobium trifolii Dangeard 1926AL. R. trifolii is a later synonym of R. leguminosarum. Reclassification of the strain R. leguminosarum DSM 30132 (=NCIMB 11478) as Rhizobium pisi sp. nov. Int. J. Syst. Evol. Microbiol., 2008, 58, 2484-2490.
Rhizobium leucaenae	BR 10016	CFN 299 = CECT 4844 = CENA 183 = CNPSo 141 = JCM 21088 = LMG 9517 = ORS 651 = SEMIA 4083 = USDA 9039	RIBEIRO (R.A.), ROGEL (M.A.), LÓPEZ-LÓPEZ (A.), ORMEÑO-ORRILLO (E.), BARCELLOS (F.G.), MARTÍNEZ (J.), THOMPSON (F.L.), MARTÍNEZ-ROMERO (E.) and HUNGRIA (M.): Reclassification of Rhizobium tropici type A strains as Rhizobium leucaenae sp. nov. Int. J. Syst. Evol. Microbiol., 2012, 62, 1179-1184.
Rhizobium loessense	BR 10516	LMG 23187 = AS1.3401 = CCBAU 7190B = CIP 108030	WEI (G.H.), TAN (Z.Y.), ZHU (M.E.), WANG (E.T.), HAN (S.Z.) and CHEN (W.X.): Characterization of rhizobia isolated from legume species within the genera Astragalus and Lespedeza grown in the Loess Plateau of China and description of Rhizobium loessense sp. nov. Int. J. Syst. Evol. Microbiol., 2003, 53, 1575-1583.
Rhizobium lusitanum	BR 10207	CNPSo 2055 = LMG 22705 = CECT 7016	VALVERDE (A.), IGUAL (J.M.), PEIX (A.), CERVANTES (E.) and VELÁZQUEZ (E.): Rhizobium lusitanum sp. nov., a bacterium that nodulates Phaseolus vulgaris. Int. J. Syst. Evol. Microbiol., 2006, 56, 2631-2637.
Rhizobium mayense	BR 10367	CNPSo 2464 = ATCC BAA-2446 = NSJP1-1	RINCÓN-ROSALES, R., VILLALOBOS-ESCOBEDO, J. M., ROGEL, M. A., MARTINEZ, J., ORMEÑO-ORRILLO, E. and MARTÍNEZ-ROMERO, E. 2013. Rhizobium calliandrae sp. nov., Rhizobium mayense sp. nov. and Rhizobium jaguaris sp. nov., rhizobial species nodulating the medicinal legume Calliandra grandiflora. Int. J. Syst. Evol. Microbiol., 63, 3423-3429.

Rhizobium mesoamericanum	BR 10402	CNPSo 2482 = CCGE 501 = ATCC BAA-2123 = CIP 110148 = HAMBI 3151	LÓPEZ-LÓPEZ (A.), ROGEL-HERNÁNDEZ (M.A.), BAROIS (I.), ORTIZ CEBALLOS (A.I.), MARTÍNEZ (J.), ORMEÑO-ORRILLO (E.) and MARTÍNEZ-ROMERO (E.): Rhizobium grahamii sp. nov., from nodules of Dalea leporina, Leucaena leucocephala and Clitoria ternatea, and Rhizobium mesoamericanum sp. nov., from nodules of Phaseolus vulgaris, siratro, cowpea and Mimosa pudica. Int. J. Syst. Evol. Microbiol., 2012, 62, 2264-2271.
Rhizobium mesosinicum	BR 10212	CNPSo 2063 = LMG 24135 = CCBAU 25010	LIN (D.X.), CHEN (W.F.), WANG (F.Q.), HU (D.), WANG (E.T.), SUI (X.H.) and CHEN (W.X.): Rhizobium mesosinicum sp. nov., isolated from root nodules of three different legumes. Int. J. Syst. Evol. Microbiol., 2009, 59, 1919-1923.
Rhizobium miluonense	BR 10395	CNPSo 2056 = LMG 24208 = HAMBI 2971 = CCBAU 41251	GU (C.T.), WANG (E.T.), TIAN (C.F.), HAN (T.X.), CHEN (W.F.), SUI (X.H.) and CHEN (W.X.): Rhizobium miluonense sp. nov., a symbiotic bacterium isolated from Lespedeza root nodules. Int. J. Syst. Evol. Microbiol., 2008, 58, 1364-1368.
Rhizobium mongolense	BR 10513	LMG 19140 = ATCC BAA-116 = CIP 106112 = HAMBI 2349 = USDA 1844	VAN BERKUM (P.), BEYENE (D.), BAO (G.), CAMPBELL (T.A.) and EARDLY (B.D.): Rhizobium mongolense sp. nov. is one of three rhizobial genotypes identified which nodulate and form nitrogen-fixing symbioses with Medicago ruthenica [(L.) Ledebour]. Int. J. Syst. Bacteriol., 1998, 48, 13-22.
Rhizobium multihospitium	BR 10206	CNPSo 2054 = LMG 23946 = HAMBI 2975	HAN (T.X.), WANG (E.T.), WU (L.J.), CHEN (W.F.), GU (J.G.), GU (C.T.), TIAN (C.F.) and CHEN (W.X.): Rhizobium multihospitium sp. nov., isolated from multiple legume species native of Xinjiang, China. Int. J. Syst. Evol. Microbiol., 2008, 58, 1693-1699.
Rhizobium nepotum	BR 10520	LMG 26435 = 39/7 = CFBP 7436	PUŁAWSKA (J.), WILLEMS (A.), DE MEYER (S.E.) and SÜLE (S.): Rhizobium nepotum sp. nov. isolated from tumors on different plant species. Syst. Appl. Microbiol., 2012, 35, 215-220.

Rhizobium phaseoli	BR 10052	LMG 8819 = ATCC 14482 = BCRC 13520 = CECT 4115 = DSM 30137 = Means 3I6c15	RAMÍREZ-BAHENA (M.H.), GARCÍA-FRAILE (P.), PEIX (A.), VALVERDE (A.), RIVAS (R.), IGUAL (J.M.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.) and VELÁZQUEZ (E.): Revision of the taxonomic status of the species Rhizobium leguminosarum (Frank 1879) Frank 1889AL, Rhizobium phaseoli 1926AL and Rhizobium trifolii Dangeard 1926AL. R. trifolii is a later synonym of R. leguminosarum. Reclassification of the strain R. leguminosarum DSM 30132 (=NCIMB 11478) as Rhizobium pisi sp. nov. Int. J. Syst. Evol. Microbiol., 2008, 58, 2484-2490.
Rhizobium pisi (Rhizobium leguminosarum)	BR 10209	CNPSo 2058 = ATCC 10004 = LMG 14904 = DSM 30132	RAMÍREZ-BAHENA (M.H.), GARCÍA-FRAILE (P.), PEIX (A.), VALVERDE (A.), RIVAS (R.), IGUAL (J.M.), MATEOS (P.F.), MARTÍNEZ-MOLINA (E.) and VELÁZQUEZ (E.): Revision of the taxonomic status of the species Rhizobium leguminosarum (Frank 1879) Frank 1889AL, Rhizobium phaseoli 1926AL and Rhizobium trifolii Dangeard 1926AL. R. trifolii is a later synonym of R. leguminosarum. Reclassification of the strain R. leguminosarum DSM 30132 (=NCIMB 11478) as Rhizobium pisi sp. nov. Int. J. Syst. Evol. Microbiol., 2008, 58, 2484-2490.
Rhizobium pseudoryzae	BR 10333	J3-A127 = KCTC 23294 = DSM 26483	ZHANG (X.), SUN (L.), MA (X.), SUI (X.H.) and JIANG (R.): Rhizobium pseudoryzae sp. nov., isolated from the rhizosphere of rice. Int. J. Syst. Evol. Microbiol., 2011, 61, 2425-2429.
Rhizobium pusense	BR 10313	LMG 25623 = JCM 16209 = NCIMB 14639	PANDAY (D.), SCHUMANN (P.) and DAS (S.K.): Rhizobium pusense sp. nov., isolated from the rhizosphere of chickpea (Cicer arietinum L.). Int. J. Syst. Evol. Microbiol., 2011, 61, 2632-2639.
Rhizobium rhizogenes	BR 10477	HAMBI 1816 = ATCC 11325 = CIP 104328 = DSM 30148 = NBRC 13257 = CFBP 5520 = ICMP 5794 = JCM 20919 = LMG 150 = NCPPB 2991	YOUNG (J.M.), KUYKENDALL (L.D.), MARTÍNEZ -ROMERO (E.), KERR (A.) and SAWADA (H.): A revision of Rhizobium Frank 1889, with an emended description of the genus, and the inclusion of all species of Agrobacterium Conn 1942 and Allorhizobium undicola de Lajudie et al. 1998 as new combinations: Rhizobium radiobacter, R. rhizogenes, R. rubi, R. undicola and R. vitis. Int. J. Syst. Evol. Microbiol., 2001, 51, 89-103.

Rhizobium rosettiformans	BR 10682	DSM 26376 = W3 = CCM 7583 = MTCC 9454	KAUR (J.), VERMA (M.) and LAL (R.): Rhizobium rosettiformans sp. nov., isolated from a hexachlorocyclohexane dump site, and reclassification of Blastobacter aggregatus Hirsch and Muller 1986 as Rhizobium aggregatum comb. nov. Int. J. Syst. Evol. Microbiol., 2011, 61, 1218-1225.
Rhizobium rubi	BR 10476	HAMBI 1812 = ATCC 13335 = CFBP 5509 = CFBP 6448 = CIP 104332 = DSM 6772 = ICMP 6428 = JCM 20918 = LMG 156 = LMG 17935 = NBRC 13261 = NCPPB 1854	YOUNG (J.M.), KUYKENDALL (L.D.), MARTÍNEZ -ROMERO (E.), KERR (A.) and SAWADA (H.): A revision of Rhizobium Frank 1889, with an emended description of the genus, and the inclusion of all species of Agrobacterium Conn 1942 and Allorhizobium undicola de Lajudie et al. 1998 as new combinations: Rhizobium radiobacter, R. rhizogenes, R. rubi, R. undicola and R. vitis. Int. J. Syst. Evol. Microbiol., 2001, 51, 89-103.
Rhizobium selenitireducens	BR 10612	LMG 24075 = B1 = ATCC BAA-1503 = NRRL B-41997	HUNTER (W.J.), KUYKENDALL (L.D.) and MANTER (D.K.): Rhizobium selenireducens sp. nov.: a selenite-reducing α-Proteobacteria isolated from a bioreactor. Curr. Microbiol., 2007, 55, 455-460.
Rhizobium skierniewicense	BR 10518	LMG 26191 = Ch11 = CFBP 7420	PUŁAWSKA (J.), WILLEMS (A.) and SOBICZEWSKI (P.): Rhizobium skierniewicense sp. nov., isolated from tumours on chrysanthemum and cherry plum. Int. J. Syst. Evol. Microbiol., 2012, 62, 895-899.
Rhizobium smilacinae	BR 10508	LMG 27604 = PTYR-5 = CCTCC AB 2013016	ZHANG, L., SHI, X., SI, M., LI, C., ZHU, L., ZHAO, L., SHEN, X. and WANG, Y. 2014. Rhizobium smilacinae sp. nov., an endophytic bacterium isolated from the leaf of Smilacina japonica. Antonie van Leeuwenhoek, 106, 715-723.
Rhizobium sophorae	BR 10481	HAMBI 3615 = CCBAU 03386 = E5 = LMG 27901	JIAO, Y. S., YAN, H., JI, Z. J., LIU, Y. H., SUI, X. H., WANG, E. T., GUO, B. L., CHEN, W. X. and CHEN, W. F. 2015. Rhizobium sophorae sp. nov. and Rhizobium sophoriradicis sp. nov., nitrogen-fixing rhizobial symbionts of the medicinal legume Sophora flavescens. Int. J. Syst. Evol. Microbiol., 65, 497-503.
Rhizobium sophoriradicis	BR 10480	HAMBI 3510 = CCBAU 03470 = C-5-1 = LMG 27898	JIAO, Y. S., YAN, H., JI, Z. J., LIU, Y. H., SUI, X. H., WANG, E. T., GUO, B. L., CHEN, W. X. and CHEN, W. F. 2015. Rhizobium sophorae sp. nov. and Rhizobium sophoriradicis sp. nov., nitrogen-fixing rhizobial symbionts of the medicinal legume Sophora flavescens. Int. J. Syst. Evol. Microbiol., 65, 497-503.

Rhizobium sphaerophysae	BR 10479	HAMBI 3074 = CCNWGS0238 = ACCC17498	XU (L.), SHI (J.F.), ZHAO (P.), CHEN (W.M.), QIN (W.), TANG (M.) and WEI (G.H.): Rhizobium sphaerophysae sp. nov., a novel species isolated from root nodules of Sphaerophysa salsula in China. Antonie van Leeuwenhoek, 2011, 99, 845-854.
Rhizobium subbaraonis	BR 10681	DSM 24765 = JC85 = KCTC 23614	RAMANA (C.V.), PARAG (B.), GIRIJA (K.R.), RAM (B.R.), VENKATA RAMANA (V.) and SASIKALA (C.): Rhizobium subbaraonis sp. nov., an endolithic bacterium isolated from beach sand. Int. J. Syst. Evol. Microbiol., 2013, 63, 581-585.
Rhizobium taibaishanense	BR 10472	HAMBI 3214 = CCNWSX 0483 = ACCC 14971	YAO (L.J.), SHEN (Y.Y.), ZHAN (J.P.), XU (W.), CUI (G.L.) and WEI (G.H.): Rhizobium taibaishanense sp. nov., isolated from a root nodule of Kummerowia striata. Int. J. Syst. Evol. Microbiol., 2012, 62, 335-341.
Rhizobium tibeticum	BR 10213	CNPSo 2064 = DSM 21102 = LMG 24453	HOU (B.C.), WANG (E.T.), LI Jr (Y.), JIA (R.Z.), CHEN (W.F.), GAO (Y.), DONG (R.J.) and CHEN (W.X.): Rhizobium tibeticum sp. nov., a symbiotic bacterium isolated from Trigonella archiducis-nicolai (Širj.) Vassilcz. Int. J. Syst. Evol. Microbiol., 2009, 59, 3051-3057.
Rhizobium tropici	BR 322	CIAT 899 = SEMIA 4038 = ATCC 49672 = BCRC 15724 = DSM 11418 = HAMBI 1163 = LMG 9503	MARTÍNEZ-ROMERO (E.), SEGOVIA (L.), MERCANTE (F.M.), FRANCO (A.A.), GRAHAM (P.) and PARDO (M.A.): Rhizobium tropici, a novel species nodulating Phaseolus vulgaris L. beans and Leucaena sp. trees. Int. J. Syst. Bacteriol., 1991, 41, 417-426.
Rhizobium tubonense	BR 10484	HAMBI 3066 = CCBAU 85046 = LMG 25225	ZHANG (R.J.), HOU (B.C.), WANG (E.T.), LI Jr. (Y.), ZHANG (X.X.) and CHEN (W.X.): Rhizobium tubonense sp. nov., isolated from root nodules of Oxytropis glabra. Int. J. Syst. Evol. Microbiol., 2011, 61, 512-517.
Rhizobium undicola	BR 10205	CNPSo 207 = ATCC 700741 = LMG 11875 = ORS 992	YOUNG (J.M.), KUYKENDALL (L.D.), MARTÍNEZ -ROMERO (E.), KERR (A.) and SAWADA (H.): A revision of Rhizobium Frank 1889, with an emended description of the genus, and the inclusion of all species of Agrobacterium Conn 1942 and Allorhizobium undicola de Lajudie et al. 1998 as new combinations: Rhizobium radiobacter, R. rhizogenes, R. rubi, R. undicola and R. vitis. Int. J. Syst. Evol. Microbiol., 2001, 51, 89-103.

Rhizobium vallis	BR 10310	LMG 25295 = CCBAU 65647 = HAMBI 3073 = CNPSo 2484	WANG (F.), WANG (E.T.), WU (L.J.), SUI (X.H.), LI Jr. (Y.) and CHEN (W.X.): Rhizobium vallis sp. nov., isolated from nodules of three leguminous species. Int. J. Syst. Evol. Microbiol., 2011, 61, 2582-2588.
Rhizobium vignae	BR 10488	HAMBI 3039 = LMG 25447 = CCBAU 05176	REN (D.W.), CHEN (W.F.), SUI (X.H.), WANG (E.T.) and CHEN (W.X.): Rhizobium vignae sp. nov., a symbiotic bacterium isolated from multiple legume species. Int. J. Syst. Evol. Microbiol., 2011, 61, 580-586.
Rhizobium vitis	BR 10478	HAMBI 1817 = K309 = ATCC 49767 = CIP 105853 = ICMP 10752 = NBRC 15140 = JCM 21033 = LMG 8750 = NCPPB 3554	YOUNG (J.M.), KUYKENDALL (L.D.), MARTÍNEZ -ROMERO (E.), KERR (A.) and SAWADA (H.): A revision of Rhizobium Frank 1889, with an emended description of the genus, and the inclusion of all species of Agrobacterium Conn 1942 and Allorhizobium undicola de Lajudie et al. 1998 as new combinations: Rhizobium radiobacter, R. rhizogenes, R. rubi, R. undicola and R. vitis. Int. J. Syst. Evol. Microbiol., 2001, 51, 89-103.
Rhizobium yanglingense	BR 10514	LMG 19592 = SH 22623 = CCBAU 71623 = CIP 108028	TAN (Z.Y.), KAN (F.L.), PENG (G.X.), WANG (E.T.), REINHOLD-HUREK (B.) and CHEN (W.X.): Rhizobium yanglingense sp. nov., isolated from arid and semi-arid regions in China. Int. J. Syst. Evol. Microbiol., 2001, 51, 909-914.
Shinella kummerowiae	BR 10181	DSM 19334 = LMG 24136 = CCBAU 25048	LIN (D.X.), WANG (E.T.), TANG (H.), HAN (T.X.), HE (Y.R.), GUAN (S.H.) and CHEN (W.X.): Shinella kummerowiae sp. nov., a symbiotic bacterium isolated from root nodules of the herbal legume Kummerowia stipulacea. Int. J. Syst. Evol. Microbiol., 2008, 58, 1409-1413.
Sphingomonas oryziterrae	BR 10304	DSM 21455 = KCTC 22476	CHUNG (E.J.), JO (E.J.), YOON (H.S.), SONG (G.C.), JEON (C.O.) and CHUNG (Y.R.): Sphingomonas oryziterrae sp. nov. and Sphingomonas jinjuensis sp. nov. isolated from rhizosphere soil of rice (Oryza sativa L.). Int. J. Syst. Evol. Microbiol., 2011, 61, 2389-2394.
Stenotrophomonas maltophilia	BR 12332	ATCC 13637 = DSM 50170 = INCQS 103	PALLERONI (N.J.) and BRADBURY (J.F.): Stenotrophomonas, a new bacterial genus for Xanthomonas maltophilia (Hugh 1980) Swings et al. 1983. Int. J. Syst. Bacteriol., 1993, 43, 606-609.

Stenotrophomonas pavanii	BR 10317	LMG 25348 = CBMAI 564	RAMOS (P.L.), VAN TRAPPEN (S.), THOMPSON (F.L.), ROCHA (R.C.S.), BARBOSA (H.R.), DE VOS (P.) and MOREIRA-FILHO (C.A.): Screening for endophytic nitrogen-fixing bacteria in Brazilian sugar cane varieties used in organic farming and description of Stenotrophomonas pavanii sp. nov. Int. J. Syst. Evol. Microbiol., 2011, 61, 926-931.