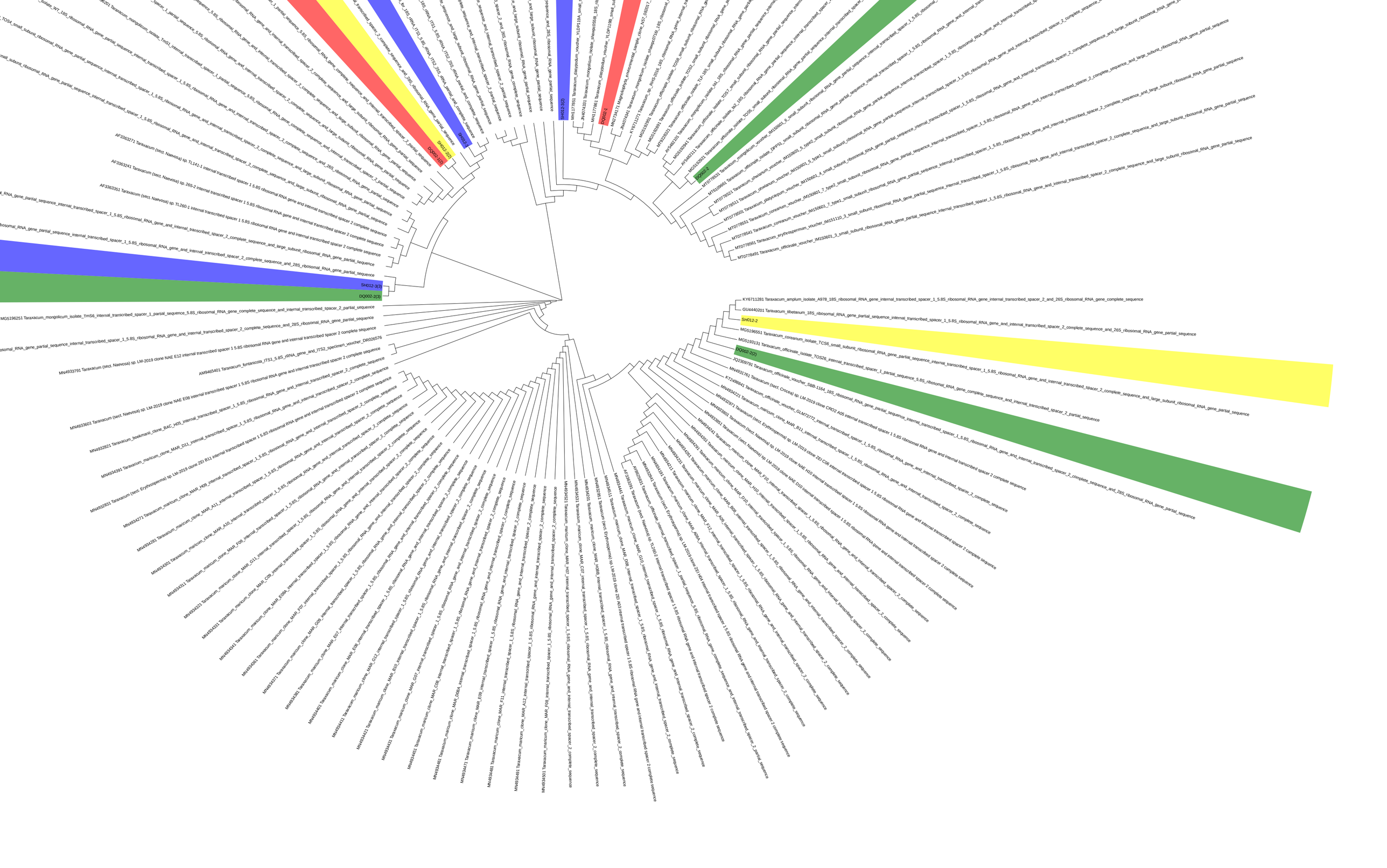
SH012-2(2)
DQ002-2(2)
SH012-3(3)
DQ002-2(3)
DQ002-2
SH012-2
KY6711281 Taraxacum_amplum_isolate_A978_18S_ribosomal_RNA_gene_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_internal_transcribed_spacer_2_and_26S_ribosomal_RNA_gene_complete_sequence
GU4440201 Taraxacum_tibetanum_18S_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_26S_ribosomal_RNA_gene_partial_sequence
MG5196551 Taraxacum_coreanum_isolate_TCS6_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MG5193131 Taraxacum_officinale_isolate_TOS26_internal_transcribed_spacer_1_partial_sequence_5.8S_ribosomal_RNA_gene_complete_sequence_and_internal_transcribed_spacer_2_partial_sequence
JQ2309791 Taraxacum_officinale_voucher_SBB-1164_18S_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4831761 Taraxacum (sect. Crocina) sp. LM-2019 clone CRO2 A05 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
KT2488441 Taraxacum_marsicum_clone_MAR_B11_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934221 Taraxacum (sect. Erythrospermia) sp. LM-2019 clone ZEI C06 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4932971 Taraxacum officinale_voucher_GLM73772_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4933601 Taraxacum (sect. Naevosa) sp. LM-2019 clone NAE D10 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934231 Taraxacum_marsicum_clone_MAR_F10_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934241 Taraxacum_marsicum_clone_MAR_D10_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934251 Taraxacum_marsicum_clone_MAR_E08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934261 Taraxacum_marsicum_clone_MAR_A05_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934271 Taraxacum_marsicum_clone_MAR_P12_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934281 Taraxacum_marsicum_clone_MAR_A08A_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934291 Taraxacum_marsicum_clone_MAR_E04 (sect. Erythrospermia) sp. LM-2019 clone ZEI H04 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934301 Taraxacum (sect. Erythrospermia) sp. LM-2019 clone ZEI A03 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934311 Taraxacum_marsicum_clone_MAR_C06_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934321 Taraxacum_officinale_internal_transcribed_spacer_1_partial_sequence_5.8S_ribosomal_RNA_gene_complete_sequence_and_internal_transcribed_spacer_2_partial_sequence
MN4934331 Taraxacum (sect. Naevosa) sp. LZ-2013 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934341 Taraxacum_marsicum_clone_MAR_G10_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934351 Taraxacum_marsicum_clone_MAR_F08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934361 Taraxacum (sect. Naevosa) sp. LM-2019 clone NAE E08 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934371 Taraxacum_marsicum_clone_MAR_H08B_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934381 Taraxacum_marsicum_clone_MAR_C07_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934391 Taraxacum_marsicum_clone_MAR_H07_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934401 Taraxacum_marsicum_clone_MAR_D08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934411 Taraxacum_marsicum_clone_MAR_E07_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934421 Taraxacum_marsicum_clone_MAR_E08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934431 Taraxacum_marsicum_clone_MAR_G12_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934441 Taraxacum_marsicum_clone_MAR_B10_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934451 Taraxacum_marsicum_clone_MAR_G07_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934461 Taraxacum_marsicum_clone_MAR_C08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934471 Taraxacum_marsicum_clone_MAR_D08A_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934481 Taraxacum_marsicum_clone_MAR_E09_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934491 Taraxacum_marsicum_clone_MAR_F11_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934501 Taraxacum_marsicum_clone_MAR_A12_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934511 Taraxacum_marsicum_clone_MAR_F08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934521 Taraxacum_marsicum_clone_MAR_H07 (sect. Erythrospermia) internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934531 Taraxacum_officinale_voucher_JM151110_3_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778491 Taraxacum_officinale_voucher_JM150601_3_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778501 Taraxacum_coreanum_voucher_JM150601_7_type1_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778511 Taraxacum_coreanum_voucher_JM150601_7_type2_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778521 Taraxacum_platycarpum_voucher_JM150601_4_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778531 Taraxacum_officinale_voucher_JM150601_5_type1_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778541 Taraxacum_officinale_voucher_JM150601_5_type2_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0199661 Taraxacum_officinale_isolate_DPPSL_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778551 Taraxacum_officinale_isolate_JC2_18S_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MT0778561 Taraxacum_officinale_isolate_TOS5_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MG5193201 Taraxacum_officinale_isolate_TOS7_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
AY5462111 Taraxacum_officinale_isolate_TIJ-18S_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MG5193241 Taraxacum_officinale_isolate_tu1_18S_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MG5193251 Taraxacum_officinale_isolate_TOS2_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MG5193261 Taraxacum_officinale_isolate_TOS8_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MG5193271 Taraxacum_sp._RHS-2015_18S_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
KY6711271 Taraxacum_sp._RHS-2015_18S_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MN2724341 Taraxacum_mongolicum_isolate_sharpo0716_18S_ribosomal_RNA_gene_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MN2724171 Magnoliophyta_environmental_sample_clone_H57_0920171_18S_ribosomal_RNA_gene_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MH1177851 Taraxacum_platypodum_voucher_YLDP1104_small_subunit_ribosomal_RNA_gene_partial_sequence
JN4074231 Taraxacum_mongolicum_isolate_shiwei0508_18S_ribosomal_RNA_gene_partial_sequence
MH1177861 Taraxacum_platypodum_voucher_YLDP1108_small_subunit_ribosomal_RNA_gene_partial_sequence
MT0778531 Taraxacum_mongolicum_voucher_JM150601_6_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence
MG5196251 Taraxacum_mongolicum_isolate_TmS6_internal_transcribed_spacer_1_partial_sequence_5.8S_ribosomal_RNA_gene_complete_sequence_and_internal_transcribed_spacer_2_partial_sequence
MN4933791 Taraxacum (sect. Naevosa) sp. LM-2019 clone NAE E12 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
AM9465401 Taraxacum_fontanicola_ITS1_5.8S_rRNA_gene_and_ITS2_specimen_voucher_DR026576
MN4933601 Taraxacum (sect. Naevosa) sp. LM-2019 clone NAE E08 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4932821 Taraxacum_beekmanii_clone_BAC_H05_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934391 Taraxacum_marsicum_clone_MAR_D11_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4932931 Taraxacum (sect. Erythrospermia) sp. LM-2019 clone ZEI B11 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934271 Taraxacum_marsicum_clone_MAR_H09_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934281 Taraxacum_marsicum_clone_MAR_A11_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934301 Taraxacum_marsicum_clone_MAR_A10_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934311 Taraxacum_marsicum_clone_MAR_F09_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934321 Taraxacum_marsicum_clone_MAR_G11_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934331 Taraxacum_marsicum_clone_MAR_C09_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934341 Taraxacum_marsicum_clone_MAR_E08A_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934351 Taraxacum_marsicum_clone_MAR_P07_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934361 Taraxacum_marsicum_clone_MAR_D09_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934371 Taraxacum_marsicum_clone_MAR_E07_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934381 Taraxacum_marsicum_clone_MAR_E08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934391 Taraxacum_marsicum_clone_MAR_G12_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934401 Taraxacum_marsicum_clone_MAR_B10_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934411 Taraxacum_marsicum_clone_MAR_G07_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934421 Taraxacum_marsicum_clone_MAR_C08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934431 Taraxacum_marsicum_clone_MAR_D08A_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934441 Taraxacum_marsicum_clone_MAR_E09_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934451 Taraxacum_marsicum_clone_MAR_F11_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934461 Taraxacum_marsicum_clone_MAR_A12_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934471 Taraxacum_marsicum_clone_MAR_F08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934481 Taraxacum_officinale_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_complete_sequence_and_internal_transcribed_spacer_2_partial_sequence
MN4934491 Taraxacum (sect. Naevosa) sp. LZ-2013 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934501 Taraxacum_marsicum_clone_MAR_G10_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934511 Taraxacum_marsicum_clone_MAR_F08_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence
MN4934521 Taraxacum (sect. Erythrospermia) sp. LM-2019 clone ZEI A03 internal transcribed spacer 1 5.8S ribosomal RNA gene and internal transcribed spacer 2 complete sequence
MN4934531 Taraxacum_officinale_voucher_JM151110_3_small_subunit_ribosomal_RNA_gene_partial_sequence_internal_transcribed_spacer_1_5.8S_ribosomal_RNA_gene_and_internal_transcribed_spacer_2_complete_sequence_and_large_subunit_ribosomal_RNA_gene_partial_sequence