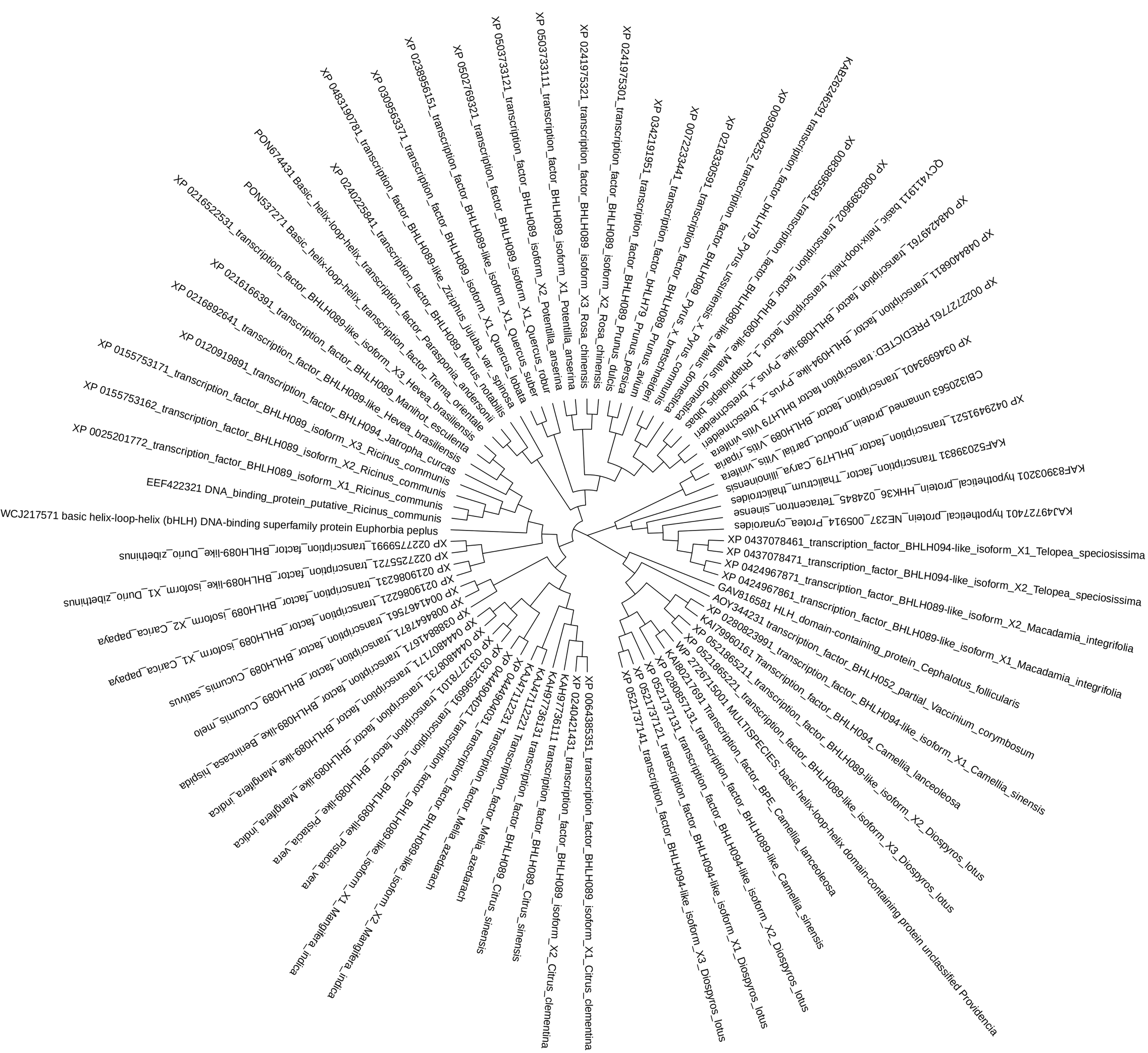
XP 0064385351_transcription_factor_BHLH089_isoform_X1_Citrus_clementina
XP 0240421431_transcription_factor_BHLH089_isoform_X2_Citrus_clementina
KAH97736111 transcription_factor_BHLH089_isoform_X2_Citrus_clementina
KAH97736131 transcription_factor_BHLH089_Citrus_sinensis
KAJ4711221 Transcription_factor_BHLH089_Citrus_sinensis
Xp 0444904031_transcription_factor_Melia_azedarach
XP 0312778101_transcription_factor_BHLH089-like_isotorm_X2_Mangifera_indica
XP 0312596691_transcription_factor_BHLH089-like_isotorm_X1_Mangifera_indica
XP 0444807171_transcription_factor_BHLH089-like_Pistacia_vera
XP 0388841671_transcription_factor_BHLH089-like_Pistacia_vera
XP 0084647871_transcription_factor_BHLH089-like_Mangifera_indica
XP 0044675611_transcription_factor_BHLH089-like_Mangifera_indica
XP 0219086221_transcription_factor_BHLH089_Cucumis_sativus
XP 0219086231_transcription_factor_BHLH089_isotorm_X1_Carica_papaya
XP 0227255721_transcription_factor_BHLH089_isotorm_X2_Carica_papaya
XP 0227759991_transcription_factor_BHLH089-like_Durio_zibethinus
WCJ217571 basic helix-loop-helix (bHLH) DNA-binding superfamily protein Euphorbia peplus
EEF422321 DNA_binding_protein_putative_Ricinus_communis
XP 0025201772_transcription_factor_BHLH089_isoform_X1_Ricinus_communis
XP 0155753162_transcription_factor_BHLH089_isoform_X2_Ricinus_communis
XP 0155753171_transcription_factor_BHLH089_isoform_X3_Ricinus_communis
XP 0120919891_transcription_factor_BHLH094_Jatropha_curcas
XP 0216892641_transcription_factor_BHLH089-like_Hevea_brasiliensis
XP 0216166391_transcription_factor_BHLH089-like_isoform_X3_Hevea_brasiliensis
PON537271 Basic_helix-loop-helix_transcription_factor_Trema_orientale
XP 0216522531_transcription_factor_BHLH089_Manihot_esculenta
PON674431 Basic_helix-loop-helix_transcription_factor_Parasponia_andersonii
XP 0240225841_transcription_factor_BHLH089_Morus_notabilis
XP 0483190781_transcription_factor_BHLH089-like_Ziziphus_jujuba_var._spinosa
XP 0309563371_transcription_factor_BHLH089_isoform_X1_Quercus_lobata
XP 0238956151_transcription_factor_BHLH089-like_isoform_X1_Quercus_suber
XP 0502769321_transcription_factor_BHLH089_isoform_X2_Potentilla_anserina
XP 0503733121_transcription_factor_BHLH089_isoform_X1_Quercus_robur
XP 0503733111_transcription_factor_BHLH089_isoform_X1_Potentilla_anserina
XP 0241975321_transcription_factor_BHLH089_isoform_X3_Rosa_chinensis
XP 0241975301_transcription_factor_BHLH089_isoform_X2_Rosa_chinensis
XP 0342191951_transcription_factor_BHLH79_Prunus_dulcis
XP 0072233441_transcription_factor_BHLH089_Prunus_persica
XP 0218330591_transcription_factor_BHLH089_Prunus_avium
XP 0093604252_transcription_factor_bHLH79_Pyrus_ussuriensis_x_Pyrus_x_bretschneideri
KAB2624629.1 transcription_factor_BHLH089_Malus_domestica
XP 0083895581_transcription_factor_BHLH089-like_Malus_domestica
XP 0083399602_transcription_factor_BHLH089_Rhaphiolepis_bibas
QCY41191.1 basic_helix-loop-helix_transcription_factor_BHLH089_x_bretschneideri
XP 0484249761_transcription_factor_BHLH089-like_Pyrus_x_bretschneideri
XP 0484406811_transcription_factor_BHLH094-like_Pyrus_x_bretschneideri
XP 0022727761 PREDICTED: transcription_factor_BHLH089_Vitis_vinifera
XP 0346993401_transcription factor bHLH79_Vitis_vinifera
CBI32056.3 unnamed_protein_product_partial_Vitis_vinifera
XP 0429491521_transcription_factor_bHLH79_Carya_illinoinensis
KAF5203983.1 Transcription_factor_Thalictrum_thalictroides
KAF8390320.1 hypothetical_protein_HHK36_024845_Tetracentron_sinense
KAJ4972740.1 hypothetical_protein_NE237_005914_Protea_cynaroides
XP 0437078461_transcription_factor_BHLH094-like_isoform_X1_Telopea_speciosissima
XP 0437078471_transcription_factor_BHLH094-like_isoform_X2_Telopea_speciosissima
XP 0424967871_transcription_factor_BHLH089-like_isoform_X2_Macadamia_integrifolia
GAV816581 HLH_domain-containing_protein_Cephalotus_follicularis
AOY344231 transcription_factor_BHLH052_partial_Vaccinium_corymbosum
XP 0280823991_transcription_factor_BHLH094-like_isoform_X1_Macadamia_integrifolia
KAI79960.1 Transcription_factor_BHLH094_Camellia_lanceoleosa
XP 0521865211_transcription_factor_BHLH089-like_isoform_X2_Diospyros_lotus
WP 2726715001 MULTISPECIES: basic helix-loop-helix domain-containing protein unclassified Providencia
KAI80217.1 Transcription_factor_BPE_Camellia_lanceoleosa
XP 0280857131_transcription_factor_BHLH089-like_Camellia_sinensis
XP 0521737131_transcription_factor_BHLH094-like_isoform_X2_Diospyros_lotus
XP 0521737121_transcription_factor_BHLH094-like_isoform_X1_Diospyros_lotus
XP 0521737141_transcription_factor_BHLH094-like_isoform_X3_Diospyros_lotus