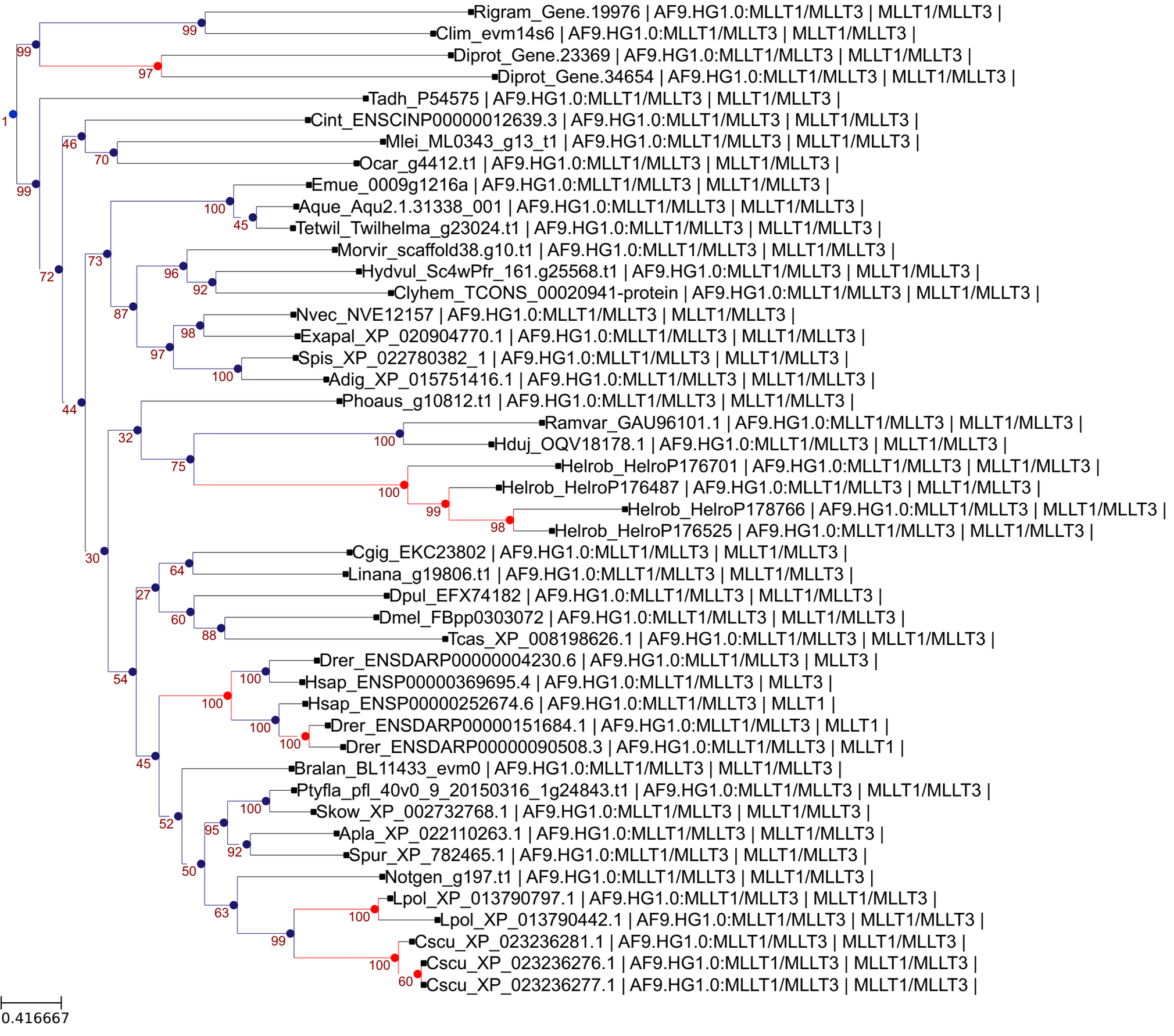
Rigram_Gene.19976 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Clim_evm14s6 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Diprot_Gene.23369 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Diprot_Gene.34654 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Tadh_P54575 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Cint_ENSCINP00000012639.3 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Mlei_ML0343_g13_t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Ocar_g4412.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Emue_0009g1216a | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Aque_Aqu2.1.31338_001 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Tetwil_Twilhelma_g23024.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Morvir_scaffold38.g10.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Hydvul_Sc4wPfr_161.g25568.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Clyhem_TCONS_00020941-protein | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Nvec_NVE12157 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Exapal_XP_020904770.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Spis_XP_022780382_1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Adig_XP_015751416.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Phoaus_g10812.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Ramvar_GAU96101.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Hduj_OQV18178.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Helrob_HelroP176701 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Helrob_HelroP176487 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Helrob_HelroP178766 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Helrob_HelroP176525 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Cgig_EKC23802 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Linana_g19806.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Dpul_EFX74182 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Dmel_FBpp0303072 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Tcas_XP_008198626.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Drer_ENSDARP00000004230.6 | AF9.HG1.0:MLLT1/MLLT3 | MLLT3 |
Hsap_ENSP00000369695.4 | AF9.HG1.0:MLLT1/MLLT3 | MLLT3 |
Hsap_ENSP00000252674.6 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1 |
Drer_ENSDARP00000151684.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1 |
Drer_ENSDARP00000090508.3 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1 |
Bralan_BL11433_evm0 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Ptyfla_pfl_40v0_9_20150316_1g24843.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Skow_XP_002732768.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Apla_XP_022110263.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Spur_XP_782465.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Notgen_g197.t1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Lpol_XP_013790797.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Lpol_XP_013790442.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Cscu_XP_023236281.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Cscu_XP_023236276.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
Cscu_XP_023236277.1 | AF9.HG1.0:MLLT1/MLLT3 | MLLT1/MLLT3 |
0.416667