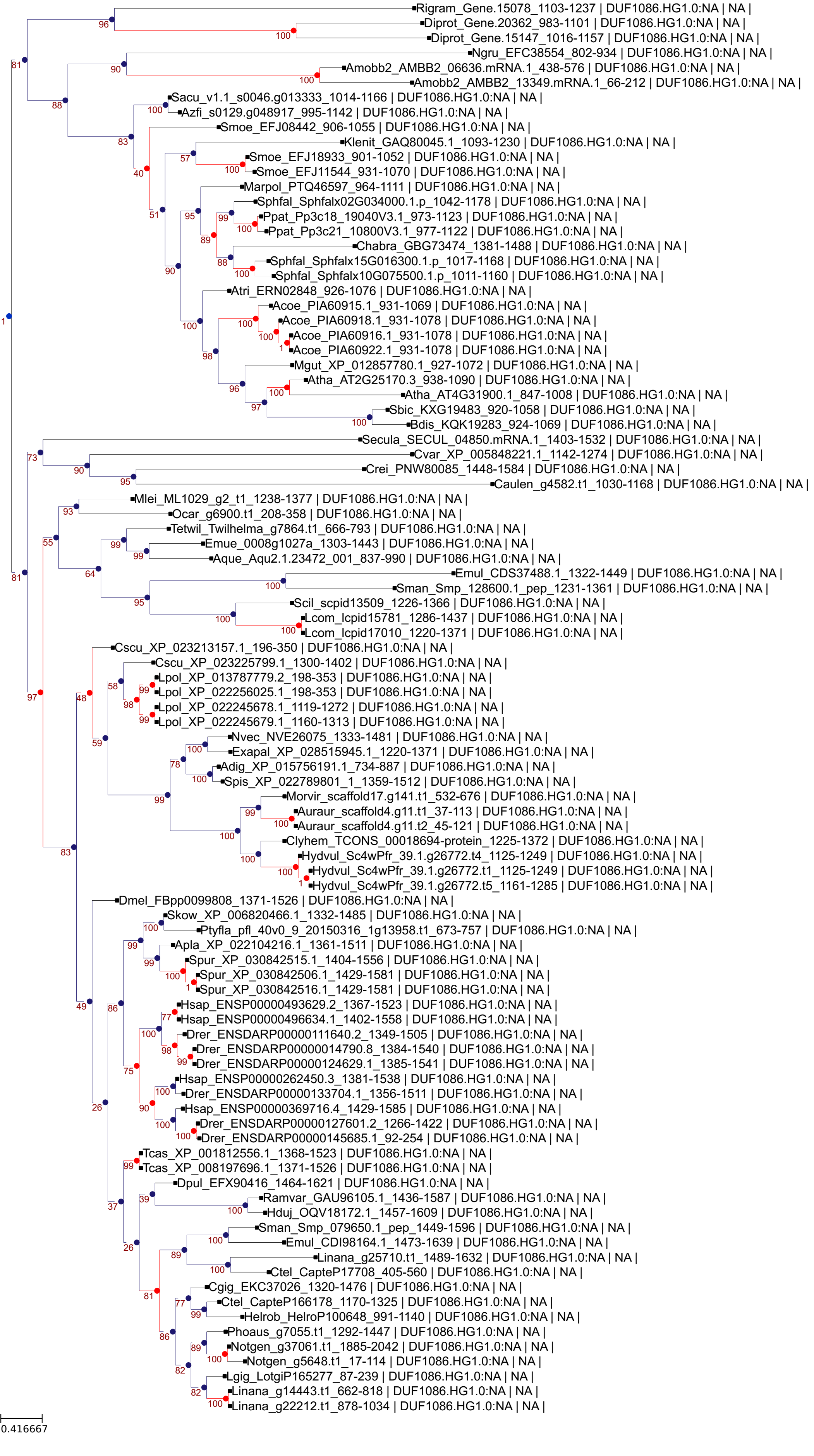
Rigram_Gene.15078_1103-1237 | DUF1086.HG1.0:NA | NA |
Diprot_Gene.20362_983-1101 | DUF1086.HG1.0:NA | NA |
Diprot_Gene.15147_1016-1157 | DUF1086.HG1.0:NA | NA |
Ngru_EFC38554_802-934 | DUF1086.HG1.0:NA | NA |
Amobb2_AMBB2_06636.mRNA.1_438-576 | DUF1086.HG1.0:NA | NA |
Amobb2_AMBB2_13349.mRNA.1_66-212 | DUF1086.HG1.0:NA | NA |
Sacu_v1.1_s0046.g013333_1014-1166 | DUF1086.HG1.0:NA | NA |
Azfi_s0129.g048917_995-1142 | DUF1086.HG1.0:NA | NA |
Smoe_EFJ08442_906-1055 | DUF1086.HG1.0:NA | NA |
Klenit_GAQ80045.1_1093-1230 | DUF1086.HG1.0:NA | NA |
Smoe_EFJ18933_901-1052 | DUF1086.HG1.0:NA | NA |
Smoe_EFJ11544_931-1070 | DUF1086.HG1.0:NA | NA |
Marpol_PTQ46597_964-1111 | DUF1086.HG1.0:NA | NA |
Sphfal_Sphfalx02G034000.1.p_1042-1178 | DUF1086.HG1.0:NA | NA |
Ppat_Pp3c18_19040V3.1_973-1123 | DUF1086.HG1.0:NA | NA |
Ppat_Pp3c21_10800V3.1_977-1122 | DUF1086.HG1.0:NA | NA |
Chabra_GBG73474_1381-1488 | DUF1086.HG1.0:NA | NA |
Sphfal_Sphfalx15G016300.1.p_1017-1168 | DUF1086.HG1.0:NA | NA |
Sphfal_Sphfalx10G075500.1.p_1011-1160 | DUF1086.HG1.0:NA | NA |
Atri_ERN02848_926-1076 | DUF1086.HG1.0:NA | NA |
Acoe_PIA60915.1_931-1069 | DUF1086.HG1.0:NA | NA |
Acoe_PIA60918.1_931-1078 | DUF1086.HG1.0:NA | NA |
Acoe_PIA60916.1_931-1078 | DUF1086.HG1.0:NA | NA |
Acoe_PIA60922.1_931-1078 | DUF1086.HG1.0:NA | NA |
Mgut_XP_012857780.1_927-1072 | DUF1086.HG1.0:NA | NA |
Atha_AT2G25170.3_938-1090 | DUF1086.HG1.0:NA | NA |
Atha_AT4G31900.1_847-1008 | DUF1086.HG1.0:NA | NA |
Sbic_KXG19483_920-1058 | DUF1086.HG1.0:NA | NA |
Bdis_KQK19283_924-1069 | DUF1086.HG1.0:NA | NA |
Secula_SECUL_04850.mRNA.1_1403-1532 | DUF1086.HG1.0:NA | NA |
Cvar_XP_005848221.1_1142-1274 | DUF1086.HG1.0:NA | NA |
Crei_PNW80085_1448-1584 | DUF1086.HG1.0:NA | NA |
Caulen_g4582.t1_1030-1168 | DUF1086.HG1.0:NA | NA |
Mlei_ML1029_g2_t1_1238-1377 | DUF1086.HG1.0:NA | NA |
Ocar_g6900.t1_208-358 | DUF1086.HG1.0:NA | NA |
Tetwil_Twilhelma_g7864.t1_666-793 | DUF1086.HG1.0:NA | NA |
Emue_0008g1027a_1303-1443 | DUF1086.HG1.0:NA | NA |
Aque_Aqu2.1.23472_001_837-990 | DUF1086.HG1.0:NA | NA |
Emul_CDS37488.1_1322-1449 | DUF1086.HG1.0:NA | NA |
Sman_Smp_128600.1_pep_1231-1361 | DUF1086.HG1.0:NA | NA |
Scil_scpid13509_1226-1366 | DUF1086.HG1.0:NA | NA |
Lcom_lcpid15781_1286-1437 | DUF1086.HG1.0:NA | NA |
Lcom_lcpid17010_1220-1371 | DUF1086.HG1.0:NA | NA |
Cscu_XP_023213157.1_196-350 | DUF1086.HG1.0:NA | NA |
Cscu_XP_023225799.1_1300-1402 | DUF1086.HG1.0:NA | NA |
Lpol_XP_013787779.2_198-353 | DUF1086.HG1.0:NA | NA |
Lpol_XP_022256025.1_198-353 | DUF1086.HG1.0:NA | NA |
Lpol_XP_022245678.1_1119-1272 | DUF1086.HG1.0:NA | NA |
Lpol_XP_022245679.1_1160-1313 | DUF1086.HG1.0:NA | NA |
Nvec_NVE26075_1333-1481 | DUF1086.HG1.0:NA | NA |
Exapal_XP_028515945.1_1220-1371 | DUF1086.HG1.0:NA | NA |
Adig_XP_015756191.1_734-887 | DUF1086.HG1.0:NA | NA |
Spis_XP_022789801_1_1359-1512 | DUF1086.HG1.0:NA | NA |
Morvir_scaffold17.g141.t1_532-676 | DUF1086.HG1.0:NA | NA |
Auraur_scaffold4.g11.t1_37-113 | DUF1086.HG1.0:NA | NA |
Auraur_scaffold4.g11.t2_45-121 | DUF1086.HG1.0:NA | NA |
Clyhem_TCONS_00018694-protein_1225-1372 | DUF1086.HG1.0:NA | NA |
Hydvul_Sc4wPfr_39.1.g26772.t4_1125-1249 | DUF1086.HG1.0:NA | NA |
Hydvul_Sc4wPfr_39.1.g26772.t1_1125-1249 | DUF1086.HG1.0:NA | NA |
Hydvul_Sc4wPfr_39.1.g26772.t5_1161-1285 | DUF1086.HG1.0:NA | NA |
Dmel_FBpp0099808_1371-1526 | DUF1086.HG1.0:NA | NA |
Skow_XP_006820466.1_1332-1485 | DUF1086.HG1.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g13958.t1_673-757 | DUF1086.HG1.0:NA | NA |
Apla_XP_022104216.1_1361-1511 | DUF1086.HG1.0:NA | NA |
Spur_XP_030842515.1_1404-1556 | DUF1086.HG1.0:NA | NA |
Spur_XP_030842506.1_1429-1581 | DUF1086.HG1.0:NA | NA |
Spur_XP_030842516.1_1429-1581 | DUF1086.HG1.0:NA | NA |
Hsap_ENSP00000493629.2_1367-1523 | DUF1086.HG1.0:NA | NA |
Hsap_ENSP00000496634.1_1402-1558 | DUF1086.HG1.0:NA | NA |
Drer_ENSDARP00000111640.2_1349-1505 | DUF1086.HG1.0:NA | NA |
Drer_ENSDARP00000014790.8_1384-1540 | DUF1086.HG1.0:NA | NA |
Drer_ENSDARP00000124629.1_1385-1541 | DUF1086.HG1.0:NA | NA |
Hsap_ENSP00000262450.3_1381-1538 | DUF1086.HG1.0:NA | NA |
Drer_ENSDARP00000133704.1_1356-1511 | DUF1086.HG1.0:NA | NA |
Hsap_ENSP00000369716.4_1429-1585 | DUF1086.HG1.0:NA | NA |
Drer_ENSDARP00000127601.2_1266-1422 | DUF1086.HG1.0:NA | NA |
Drer_ENSDARP00000145685.1_92-254 | DUF1086.HG1.0:NA | NA |
Tcas_XP_001812556.1_1368-1523 | DUF1086.HG1.0:NA | NA |
Tcas_XP_008197696.1_1371-1526 | DUF1086.HG1.0:NA | NA |
Dpul_EFX90416_1464-1621 | DUF1086.HG1.0:NA | NA |
Ramvar_GAU96105.1_1436-1587 | DUF1086.HG1.0:NA | NA |
Hduj_OQV18172.1_1457-1609 | DUF1086.HG1.0:NA | NA |
Sman_Smp_079650.1_pep_1449-1596 | DUF1086.HG1.0:NA | NA |
Emul_CDI98164.1_1473-1639 | DUF1086.HG1.0:NA | NA |
Linana_g25710.t1_1489-1632 | DUF1086.HG1.0:NA | NA |
Ctel_CapteP17708_405-560 | DUF1086.HG1.0:NA | NA |
Cgig_EKC37026_1320-1476 | DUF1086.HG1.0:NA | NA |
Ctel_CapteP166178_1170-1325 | DUF1086.HG1.0:NA | NA |
Helrob_HelroP100648_991-1140 | DUF1086.HG1.0:NA | NA |
Phoaus_g7055.t1_1292-1447 | DUF1086.HG1.0:NA | NA |
Notgen_g37061.t1_1885-2042 | DUF1086.HG1.0:NA | NA |
Notgen_g5648.t1_17-114 | DUF1086.HG1.0:NA | NA |
Lgig_LotgiP165277_87-239 | DUF1086.HG1.0:NA | NA |
Linana_g14443.t1_662-818 | DUF1086.HG1.0:NA | NA |
Linana_g22212.t1_878-1034 | DUF1086.HG1.0:NA | NA |
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