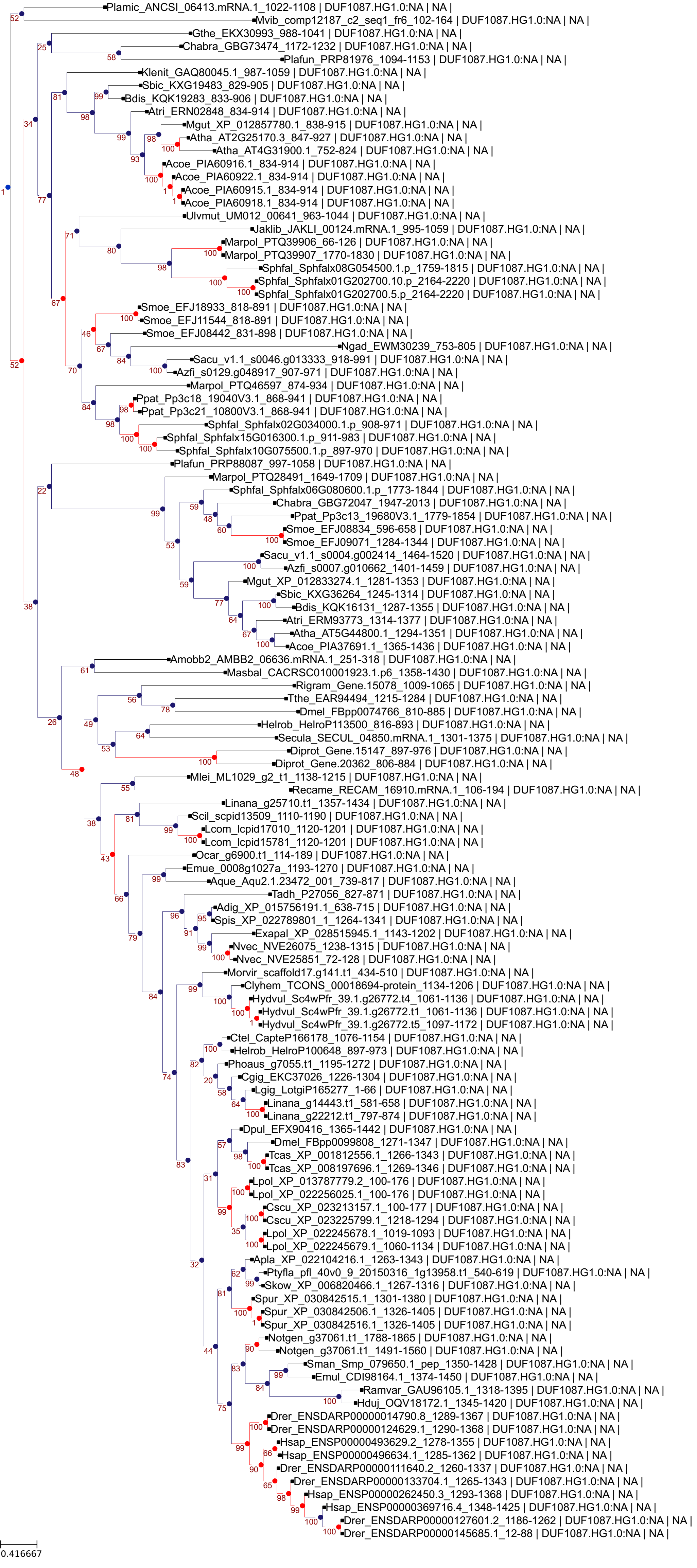
Plamic_ANCSI_06413.mRNA.1_1022-1108 | DUF1087.HG1.0:NA | NA |
Mvib_comp12187_c2_seq1_fr6_102-164 | DUF1087.HG1.0:NA | NA |
Gthe_EKX30993_988-1041 | DUF1087.HG1.0:NA | NA |
Chabra_GBG73474_1172-1232 | DUF1087.HG1.0:NA | NA |
Plafun_PRP81976_1094-1153 | DUF1087.HG1.0:NA | NA |
Klenit_GAQ80045.1_987-1059 | DUF1087.HG1.0:NA | NA |
Sbic_KXG19483_829-905 | DUF1087.HG1.0:NA | NA |
Bdis_KQK19283_833-906 | DUF1087.HG1.0:NA | NA |
Atri_ERN02848_834-914 | DUF1087.HG1.0:NA | NA |
Mgut_XP_012857780.1_838-915 | DUF1087.HG1.0:NA | NA |
Atha_AT2G25170.3_847-927 | DUF1087.HG1.0:NA | NA |
Atha_AT4G31900.1_752-824 | DUF1087.HG1.0:NA | NA |
Acoe_PIA60916.1_834-914 | DUF1087.HG1.0:NA | NA |
Acoe_PIA60922.1_834-914 | DUF1087.HG1.0:NA | NA |
Acoe_PIA60915.1_834-914 | DUF1087.HG1.0:NA | NA |
Acoe_PIA60918.1_834-914 | DUF1087.HG1.0:NA | NA |
Ulvmut_UM012_00641_963-1044 | DUF1087.HG1.0:NA | NA |
Jaklib_JAKLI_00124.mRNA.1_995-1059 | DUF1087.HG1.0:NA | NA |
Marpol_PTQ39906_66-126 | DUF1087.HG1.0:NA | NA |
Marpol_PTQ39907_1770-1830 | DUF1087.HG1.0:NA | NA |
Sphfal_Sphfalx08G054500.1.p_1759-1815 | DUF1087.HG1.0:NA | NA |
Sphfal_Sphfalx01G202700.10.p_2164-2220 | DUF1087.HG1.0:NA | NA |
Sphfal_Sphfalx01G202700.5.p_2164-2220 | DUF1087.HG1.0:NA | NA |
Smoe_EFJ18933_818-891 | DUF1087.HG1.0:NA | NA |
Smoe_EFJ11544_818-891 | DUF1087.HG1.0:NA | NA |
Smoe_EFJ08442_831-898 | DUF1087.HG1.0:NA | NA |
Ngad_EWM30239_753-805 | DUF1087.HG1.0:NA | NA |
Sacu_v1.1_s0046.g013333_918-991 | DUF1087.HG1.0:NA | NA |
Azfi_s0129.g048917_907-971 | DUF1087.HG1.0:NA | NA |
Marpol_PTQ46597_874-934 | DUF1087.HG1.0:NA | NA |
Ppat_Pp3c18_19040V3.1_868-941 | DUF1087.HG1.0:NA | NA |
Ppat_Pp3c21_10800V3.1_868-941 | DUF1087.HG1.0:NA | NA |
Sphfal_Sphfalx02G034000.1.p_908-971 | DUF1087.HG1.0:NA | NA |
Sphfal_Sphfalx15G016300.1.p_911-983 | DUF1087.HG1.0:NA | NA |
Sphfal_Sphfalx10G075500.1.p_897-970 | DUF1087.HG1.0:NA | NA |
Plafun_PRP88087_997-1058 | DUF1087.HG1.0:NA | NA |
Marpol_PTQ28491_1649-1709 | DUF1087.HG1.0:NA | NA |
Sphfal_Sphfalx06G080600.1.p_1773-1844 | DUF1087.HG1.0:NA | NA |
Chabra_GBG72047_1947-2013 | DUF1087.HG1.0:NA | NA |
Ppat_Pp3c13_19680V3.1_1779-1854 | DUF1087.HG1.0:NA | NA |
Smoe_EFJ08834_596-658 | DUF1087.HG1.0:NA | NA |
Smoe_EFJ09071_1284-1344 | DUF1087.HG1.0:NA | NA |
Sacu_v1.1_s0004.g002414_1464-1520 | DUF1087.HG1.0:NA | NA |
Azfi_s0007.g010662_1401-1459 | DUF1087.HG1.0:NA | NA |
Mgut_XP_012833274.1_1281-1353 | DUF1087.HG1.0:NA | NA |
Sbic_KXG36264_1245-1314 | DUF1087.HG1.0:NA | NA |
Bdis_KQK16131_1287-1355 | DUF1087.HG1.0:NA | NA |
Atri_ERM93773_1314-1377 | DUF1087.HG1.0:NA | NA |
Atha_AT5G44800.1_1294-1351 | DUF1087.HG1.0:NA | NA |
Acoe_PIA37691.1_1365-1436 | DUF1087.HG1.0:NA | NA |
Amobb2_AMBB2_06636.mRNA.1_251-318 | DUF1087.HG1.0:NA | NA |
Masbal_CACRSC010001923.1.p6_1358-1430 | DUF1087.HG1.0:NA | NA |
Rigram_Gene.15078_1009-1065 | DUF1087.HG1.0:NA | NA |
Tthe_EAR94494_1215-1284 | DUF1087.HG1.0:NA | NA |
Dmel_FBpp0074766_810-885 | DUF1087.HG1.0:NA | NA |
Helrob_HelroP113500_816-893 | DUF1087.HG1.0:NA | NA |
Secula_SECUL_04850.mRNA.1_1301-1375 | DUF1087.HG1.0:NA | NA |
Diprot_Gene.15147_897-976 | DUF1087.HG1.0:NA | NA |
Diprot_Gene.20362_806-884 | DUF1087.HG1.0:NA | NA |
Mlei_ML1029_g2_t1_1138-1215 | DUF1087.HG1.0:NA | NA |
Recame_RECAM_16910.mRNA.1_106-194 | DUF1087.HG1.0:NA | NA |
Linana_g25710.t1_1357-1434 | DUF1087.HG1.0:NA | NA |
Scil_scpid13509_1110-1190 | DUF1087.HG1.0:NA | NA |
Lcom_lcpid17010_1120-1201 | DUF1087.HG1.0:NA | NA |
Lcom_lcpid15781_1120-1201 | DUF1087.HG1.0:NA | NA |
Ocar_g6900.t1_114-189 | DUF1087.HG1.0:NA | NA |
Emue_0008g1027a_1193-1270 | DUF1087.HG1.0:NA | NA |
Aque_Aqu2.1.23472_001_739-817 | DUF1087.HG1.0:NA | NA |
Tadh_P27056_827-871 | DUF1087.HG1.0:NA | NA |
Adig_XP_015756191.1_638-715 | DUF1087.HG1.0:NA | NA |
Spis_XP_022789801_1_1264-1341 | DUF1087.HG1.0:NA | NA |
Exapal_XP_028515945.1_1143-1202 | DUF1087.HG1.0:NA | NA |
Nvec_NVE26075_1238-1315 | DUF1087.HG1.0:NA | NA |
Nvec_NVE25851_72-128 | DUF1087.HG1.0:NA | NA |
Morvir_scaffold17.g141.t1_434-510 | DUF1087.HG1.0:NA | NA |
Clyhem_TCONS_00018694-protein_1134-1206 | DUF1087.HG1.0:NA | NA |
Hydvul_Sc4wPfr_39.1.g26772.t4_1061-1136 | DUF1087.HG1.0:NA | NA |
Hydvul_Sc4wPfr_39.1.g26772.t1_1061-1136 | DUF1087.HG1.0:NA | NA |
Hydvul_Sc4wPfr_39.1.g26772.t5_1097-1172 | DUF1087.HG1.0:NA | NA |
Ctel_CapteP166178_1076-1154 | DUF1087.HG1.0:NA | NA |
Helrob_HelroP100648_897-973 | DUF1087.HG1.0:NA | NA |
Phoaus_g7055.t1_1195-1272 | DUF1087.HG1.0:NA | NA |
Cgig_EKC37026_1226-1304 | DUF1087.HG1.0:NA | NA |
Lgig_LotgiP165277_1-66 | DUF1087.HG1.0:NA | NA |
Linana_g14443.t1_581-658 | DUF1087.HG1.0:NA | NA |
Linana_g22212.t1_797-874 | DUF1087.HG1.0:NA | NA |
Dpul_EFX90416_1365-1442 | DUF1087.HG1.0:NA | NA |
Dmel_FBpp0099808_1271-1347 | DUF1087.HG1.0:NA | NA |
Tcas_XP_001812556.1_1266-1343 | DUF1087.HG1.0:NA | NA |
Tcas_XP_008197696.1_1269-1346 | DUF1087.HG1.0:NA | NA |
Lpol_XP_013787779.2_100-176 | DUF1087.HG1.0:NA | NA |
Lpol_XP_022256025.1_100-176 | DUF1087.HG1.0:NA | NA |
Cscu_XP_023213157.1_100-177 | DUF1087.HG1.0:NA | NA |
Cscu_XP_023225799.1_1218-1294 | DUF1087.HG1.0:NA | NA |
Lpol_XP_022245678.1_1019-1093 | DUF1087.HG1.0:NA | NA |
Lpol_XP_022245679.1_1060-1134 | DUF1087.HG1.0:NA | NA |
Apla_XP_022104216.1_1263-1343 | DUF1087.HG1.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g13958.t1_540-619 | DUF1087.HG1.0:NA | NA |
Skow_XP_006820466.1_1267-1316 | DUF1087.HG1.0:NA | NA |
Spur_XP_030842515.1_1301-1380 | DUF1087.HG1.0:NA | NA |
Spur_XP_030842506.1_1326-1405 | DUF1087.HG1.0:NA | NA |
Spur_XP_030842516.1_1326-1405 | DUF1087.HG1.0:NA | NA |
Notgen_g37061.t1_1788-1865 | DUF1087.HG1.0:NA | NA |
Notgen_g37061.t1_1491-1560 | DUF1087.HG1.0:NA | NA |
Sman_Smp_079650.1_pep_1350-1428 | DUF1087.HG1.0:NA | NA |
Emul_CDI98164.1_1374-1450 | DUF1087.HG1.0:NA | NA |
Ramvar_GAU96105.1_1318-1395 | DUF1087.HG1.0:NA | NA |
Hduj_OQV18172.1_1345-1420 | DUF1087.HG1.0:NA | NA |
Drer_ENSDARP00000014790.8_1289-1367 | DUF1087.HG1.0:NA | NA |
Drer_ENSDARP00000124629.1_1290-1368 | DUF1087.HG1.0:NA | NA |
Hsap_ENSP00000493629.2_1278-1355 | DUF1087.HG1.0:NA | NA |
Hsap_ENSP00000496634.1_1285-1362 | DUF1087.HG1.0:NA | NA |
Drer_ENSDARP00000111640.2_1260-1337 | DUF1087.HG1.0:NA | NA |
Drer_ENSDARP00000133704.1_1265-1343 | DUF1087.HG1.0:NA | NA |
Hsap_ENSP00000262450.3_1293-1368 | DUF1087.HG1.0:NA | NA |
Hsap_ENSP00000369716.4_1348-1425 | DUF1087.HG1.0:NA | NA |
Drer_ENSDARP00000127601.2_1186-1262 | DUF1087.HG1.0:NA | NA |
Drer_ENSDARP00000145685.1_12-88 | DUF1087.HG1.0:NA | NA |
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