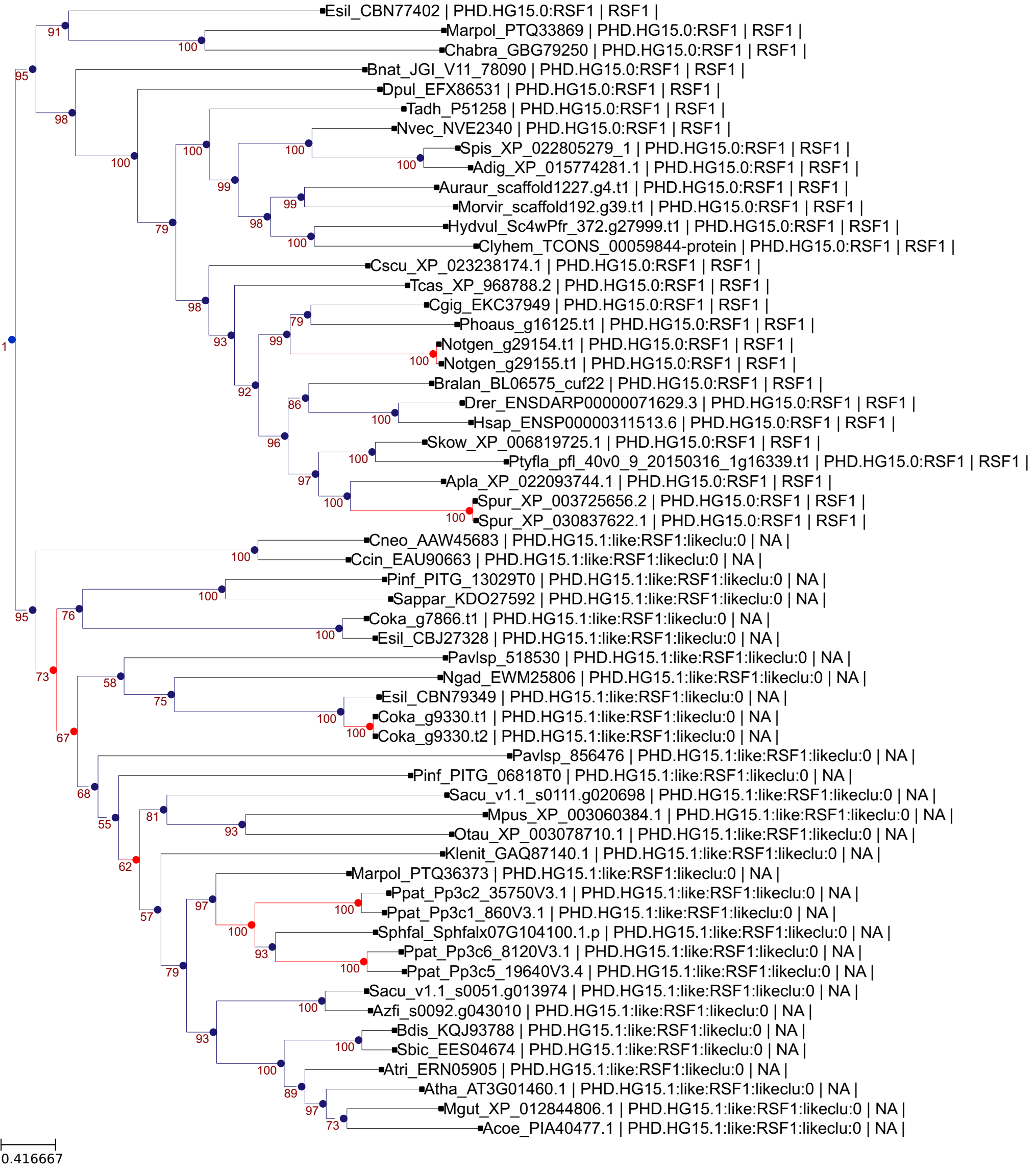
Esil_CBN77402 | PHD.HG15.0:RSF1 | RSF1 |
Marpol_PTQ33869 | PHD.HG15.0:RSF1 | RSF1 |
Chabra_GBG79250 | PHD.HG15.0:RSF1 | RSF1 |
Bnat_JGI_V11_78090 | PHD.HG15.0:RSF1 | RSF1 |
Dpul_EFX86531 | PHD.HG15.0:RSF1 | RSF1 |
Tadh_P51258 | PHD.HG15.0:RSF1 | RSF1 |
Nvec_NVE2340 | PHD.HG15.0:RSF1 | RSF1 |
Spis_XP_022805279_1 | PHD.HG15.0:RSF1 | RSF1 |
Adig_XP_015774281.1 | PHD.HG15.0:RSF1 | RSF1 |
Auraur_scaffold1227.g4.t1 | PHD.HG15.0:RSF1 | RSF1 |
Morvir_scaffold192.g39.t1 | PHD.HG15.0:RSF1 | RSF1 |
Hydvul_Sc4wPfr_372.g27999.t1 | PHD.HG15.0:RSF1 | RSF1 |
Clyhem_TCONS_00059844-protein | PHD.HG15.0:RSF1 | RSF1 |
Cscu_XP_023238174.1 | PHD.HG15.0:RSF1 | RSF1 |
Tcas_XP_968788.2 | PHD.HG15.0:RSF1 | RSF1 |
Cgig_EKC37949 | PHD.HG15.0:RSF1 | RSF1 |
Phoaus_g16125.t1 | PHD.HG15.0:RSF1 | RSF1 |
Notgen_g29154.t1 | PHD.HG15.0:RSF1 | RSF1 |
Notgen_g29155.t1 | PHD.HG15.0:RSF1 | RSF1 |
Bralan_BL06575_cuf22 | PHD.HG15.0:RSF1 | RSF1 |
Drer_ENSDARP00000071629.3 | PHD.HG15.0:RSF1 | RSF1 |
Hsap_ENSP00000311513.6 | PHD.HG15.0:RSF1 | RSF1 |
Skow_XP_006819725.1 | PHD.HG15.0:RSF1 | RSF1 |
Ptyfla_pfl_40v0_9_20150316_1g16339.t1 | PHD.HG15.0:RSF1 | RSF1 |
Apla_XP_022093744.1 | PHD.HG15.0:RSF1 | RSF1 |
Spur_XP_003725656.2 | PHD.HG15.0:RSF1 | RSF1 |
Spur_XP_030837622.1 | PHD.HG15.0:RSF1 | RSF1 |
Cneo_AAW45683 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Ccin_EAU90663 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Pinf_PITG_13029T0 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Sappar_KDO27592 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Coka_g7866.t1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Esil_CBJ27328 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Pavlsp_518530 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Ngad_EWM25806 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Esil_CBN79349 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Coka_g9330.t1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Coka_g9330.t2 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Pavlsp_856476 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Pinf_PITG_06818T0 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Sacu_v1.1_s0111.g020698 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Mpus_XP_003060384.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Otau_XP_003078710.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Klenit_GAQ87140.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Marpol_PTQ36373 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Ppat_Pp3c2_35750V3.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Ppat_Pp3c1_860V3.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Sphfal_Sphfalx07G104100.1.p | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Ppat_Pp3c6_8120V3.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Ppat_Pp3c5_19640V3.4 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Sacu_v1.1_s0051.g013974 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Azfi_s0092.g043010 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Bdis_KQJ93788 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Sbic_EES04674 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Atri_ERN05905 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
Atha_AT3G01460.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
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Acoe_PIA40477.1 | PHD.HG15.1:like:RSF1:likeclu:0 | NA |
0.416667