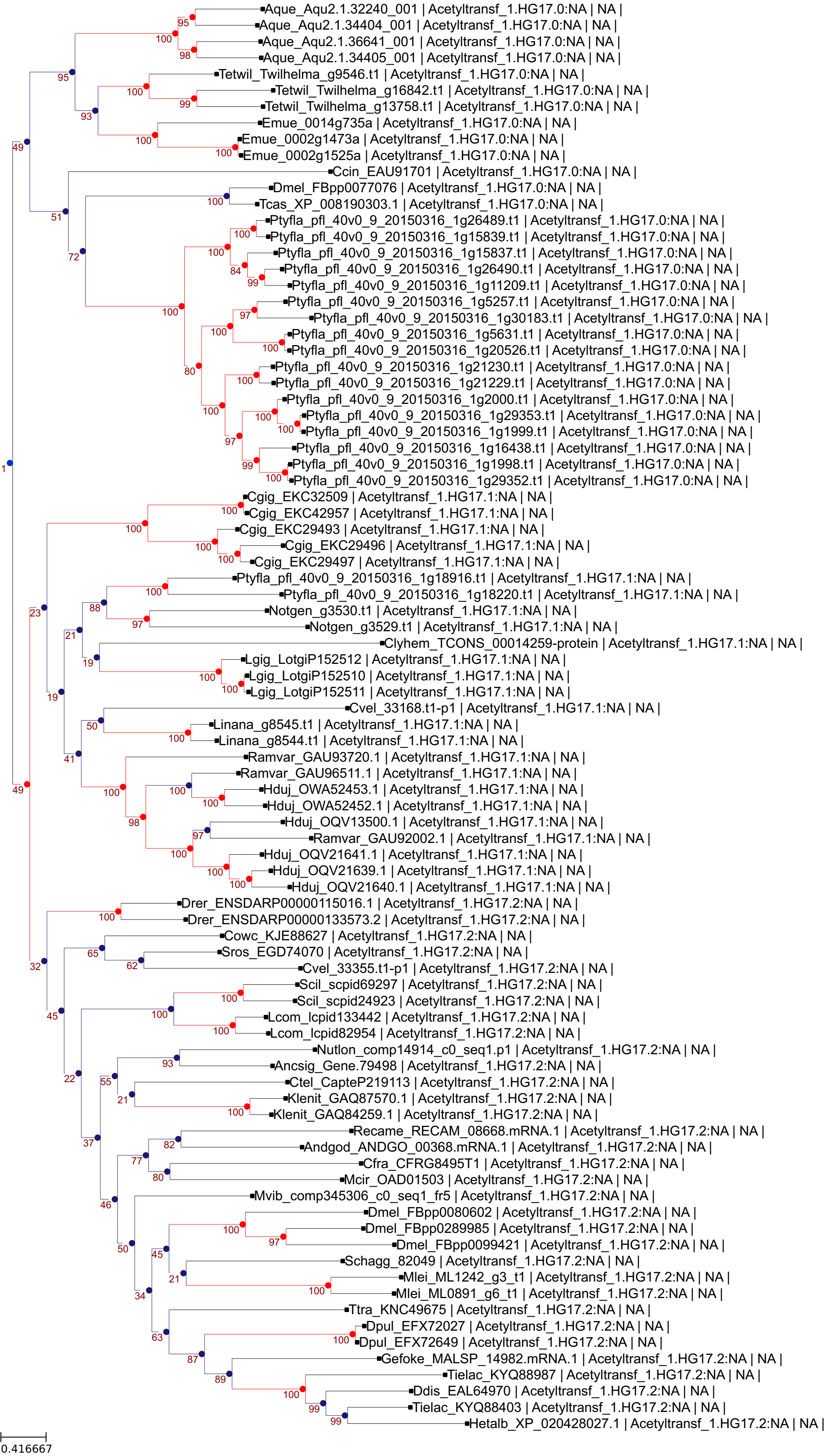
Aque_Aqu2.1.32240_001 | Acetyltransf_1.HG17.0:NA | NA |
Aque_Aqu2.1.34404_001 | Acetyltransf_1.HG17.0:NA | NA |
Aque_Aqu2.1.36641_001 | Acetyltransf_1.HG17.0:NA | NA |
Aque_Aqu2.1.34405_001 | Acetyltransf_1.HG17.0:NA | NA |
Tetwil_Twilhelma_g9546.t1 | Acetyltransf_1.HG17.0:NA | NA |
Tetwil_Twilhelma_g16842.t1 | Acetyltransf_1.HG17.0:NA | NA |
Tetwil_Twilhelma_g13758.t1 | Acetyltransf_1.HG17.0:NA | NA |
Emue_0014g735a | Acetyltransf_1.HG17.0:NA | NA |
Emue_0002g1473a | Acetyltransf_1.HG17.0:NA | NA |
Emue_0002g1525a | Acetyltransf_1.HG17.0:NA | NA |
Ccin_EAU91701 | Acetyltransf_1.HG17.0:NA | NA |
Dmel_FBpp0077076 | Acetyltransf_1.HG17.0:NA | NA |
Tcas_XP_008190303.1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g26489.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g15839.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g15837.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g26490.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g11209.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g5257.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g30183.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g5631.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g20526.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g21230.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g21229.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g2000.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g29353.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g1999.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g16438.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g1998.t1 | Acetyltransf_1.HG17.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g29352.t1 | Acetyltransf_1.HG17.0:NA | NA |
Cgig_EKC32509 | Acetyltransf_1.HG17.1:NA | NA |
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Ptyfla_pfl_40v0_9_20150316_1g18220.t1 | Acetyltransf_1.HG17.1:NA | NA |
Notgen_g3530.t1 | Acetyltransf_1.HG17.1:NA | NA |
Notgen_g3529.t1 | Acetyltransf_1.HG17.1:NA | NA |
Clyhem_TCONS_00014259-protein | Acetyltransf_1.HG17.1:NA | NA |
Lgig_LotgiP152512 | Acetyltransf_1.HG17.1:NA | NA |
Lgig_LotgiP152510 | Acetyltransf_1.HG17.1:NA | NA |
Lgig_LotgiP152511 | Acetyltransf_1.HG17.1:NA | NA |
Cvel_33168.t1-p1 | Acetyltransf_1.HG17.1:NA | NA |
Linana_g8545.t1 | Acetyltransf_1.HG17.1:NA | NA |
Linana_g8544.t1 | Acetyltransf_1.HG17.1:NA | NA |
Ramvar_GAU93720.1 | Acetyltransf_1.HG17.1:NA | NA |
Ramvar_GAU96511.1 | Acetyltransf_1.HG17.1:NA | NA |
Hduj_OWA52453.1 | Acetyltransf_1.HG17.1:NA | NA |
Hduj_OWA52452.1 | Acetyltransf_1.HG17.1:NA | NA |
Hduj_OQV13500.1 | Acetyltransf_1.HG17.1:NA | NA |
Ramvar_GAU92002.1 | Acetyltransf_1.HG17.1:NA | NA |
Hduj_OQV21641.1 | Acetyltransf_1.HG17.1:NA | NA |
Hduj_OQV21639.1 | Acetyltransf_1.HG17.1:NA | NA |
Hduj_OQV21640.1 | Acetyltransf_1.HG17.1:NA | NA |
Drer_ENSDARP00000115016.1 | Acetyltransf_1.HG17.2:NA | NA |
Drer_ENSDARP00000133573.2 | Acetyltransf_1.HG17.2:NA | NA |
Cowc_KJE88627 | Acetyltransf_1.HG17.2:NA | NA |
Sros_EGD74070 | Acetyltransf_1.HG17.2:NA | NA |
Cvel_33355.t1-p1 | Acetyltransf_1.HG17.2:NA | NA |
Scil_scpid69297 | Acetyltransf_1.HG17.2:NA | NA |
Scil_scpid24923 | Acetyltransf_1.HG17.2:NA | NA |
Lcom_lcpid133442 | Acetyltransf_1.HG17.2:NA | NA |
Lcom_lcpid82954 | Acetyltransf_1.HG17.2:NA | NA |
Nutlon_comp14914_c0_seq1.p1 | Acetyltransf_1.HG17.2:NA | NA |
Ancsig_Gene.79498 | Acetyltransf_1.HG17.2:NA | NA |
Ctel_CapteP219113 | Acetyltransf_1.HG17.2:NA | NA |
Klenit_GAQ87570.1 | Acetyltransf_1.HG17.2:NA | NA |
Klenit_GAQ84259.1 | Acetyltransf_1.HG17.2:NA | NA |
Recame_RECAM_08668.mRNA.1 | Acetyltransf_1.HG17.2:NA | NA |
Andgod_ANDGO_00368.mRNA.1 | Acetyltransf_1.HG17.2:NA | NA |
Cfra_CFRG8495T1 | Acetyltransf_1.HG17.2:NA | NA |
Mcir_OAD01503 | Acetyltransf_1.HG17.2:NA | NA |
Mvib_comp345306_c0_seq1_fr5 | Acetyltransf_1.HG17.2:NA | NA |
Dmel_FBpp0080602 | Acetyltransf_1.HG17.2:NA | NA |
Dmel_FBpp0289985 | Acetyltransf_1.HG17.2:NA | NA |
Dmel_FBpp0099421 | Acetyltransf_1.HG17.2:NA | NA |
Schagg_82049 | Acetyltransf_1.HG17.2:NA | NA |
Mlei_ML1242_g3_t1 | Acetyltransf_1.HG17.2:NA | NA |
Mlei_ML0891_g6_t1 | Acetyltransf_1.HG17.2:NA | NA |
Ttra_KNC49675 | Acetyltransf_1.HG17.2:NA | NA |
Dpul_EFX72027 | Acetyltransf_1.HG17.2:NA | NA |
Dpul_EFX72649 | Acetyltransf_1.HG17.2:NA | NA |
Gefoke_MALSP_14982.mRNA.1 | Acetyltransf_1.HG17.2:NA | NA |
Tielac_KYQ88987 | Acetyltransf_1.HG17.2:NA | NA |
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0.416667