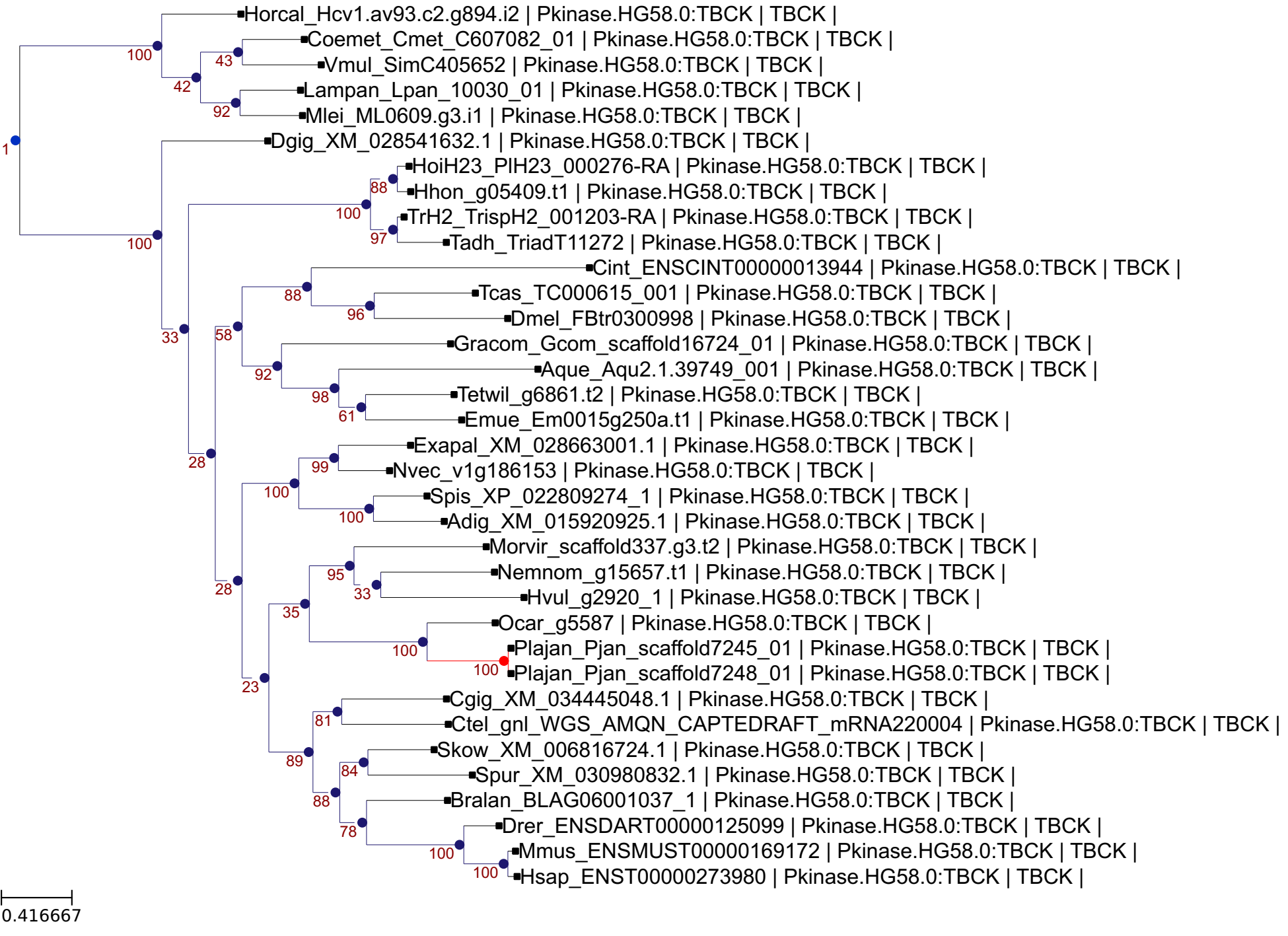
Horcal_Hcv1.av93.c2.g894.i2 | Pkinase.HG58.0:TBCK | TBCK |
Coemet_Cmet_C607082_01 | Pkinase.HG58.0:TBCK | TBCK |
Vmul_SimC405652 | Pkinase.HG58.0:TBCK | TBCK |
Lampan_Lpan_10030_01 | Pkinase.HG58.0:TBCK | TBCK |
Mlei_ML0609.g3.i1 | Pkinase.HG58.0:TBCK | TBCK |
Dgig_XM_028541632.1 | Pkinase.HG58.0:TBCK | TBCK |
HoiH23_PlH23_000276-RA | Pkinase.HG58.0:TBCK | TBCK |
Hhon_g05409.t1 | Pkinase.HG58.0:TBCK | TBCK |
TrH2_TrispH2_001203-RA | Pkinase.HG58.0:TBCK | TBCK |
Tadh_TriadT11272 | Pkinase.HG58.0:TBCK | TBCK |
Cint_ENSCINT00000013944 | Pkinase.HG58.0:TBCK | TBCK |
Tcas_TC000615_001 | Pkinase.HG58.0:TBCK | TBCK |
Dmel_FBtr0300998 | Pkinase.HG58.0:TBCK | TBCK |
Gracom_Gcom_scaffold16724_01 | Pkinase.HG58.0:TBCK | TBCK |
Aque_Aqu2.1.39749_001 | Pkinase.HG58.0:TBCK | TBCK |
Tetwil_g6861.t2 | Pkinase.HG58.0:TBCK | TBCK |
Emue_Em0015g250a.t1 | Pkinase.HG58.0:TBCK | TBCK |
Exapal_XM_028663001.1 | Pkinase.HG58.0:TBCK | TBCK |
Nvec_v1g186153 | Pkinase.HG58.0:TBCK | TBCK |
Spis_XP_022809274_1 | Pkinase.HG58.0:TBCK | TBCK |
Adig_XM_015920925.1 | Pkinase.HG58.0:TBCK | TBCK |
Morvir_scaffold337.g3.t2 | Pkinase.HG58.0:TBCK | TBCK |
Nemnom_g15657.t1 | Pkinase.HG58.0:TBCK | TBCK |
Hvul_g2920_1 | Pkinase.HG58.0:TBCK | TBCK |
Ocar_g5587 | Pkinase.HG58.0:TBCK | TBCK |
Plajan_Pjan_scaffold7245_01 | Pkinase.HG58.0:TBCK | TBCK |
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Cgig_XM_034445048.1 | Pkinase.HG58.0:TBCK | TBCK |
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Skow_XM_006816724.1 | Pkinase.HG58.0:TBCK | TBCK |
Spur_XM_030980832.1 | Pkinase.HG58.0:TBCK | TBCK |
Bralan_BLAG06001037_1 | Pkinase.HG58.0:TBCK | TBCK |
Drer_ENSDART00000125099 | Pkinase.HG58.0:TBCK | TBCK |
Mmus_ENSMUST00000169172 | Pkinase.HG58.0:TBCK | TBCK |
Hsap_ENST00000273980 | Pkinase.HG58.0:TBCK | TBCK |
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