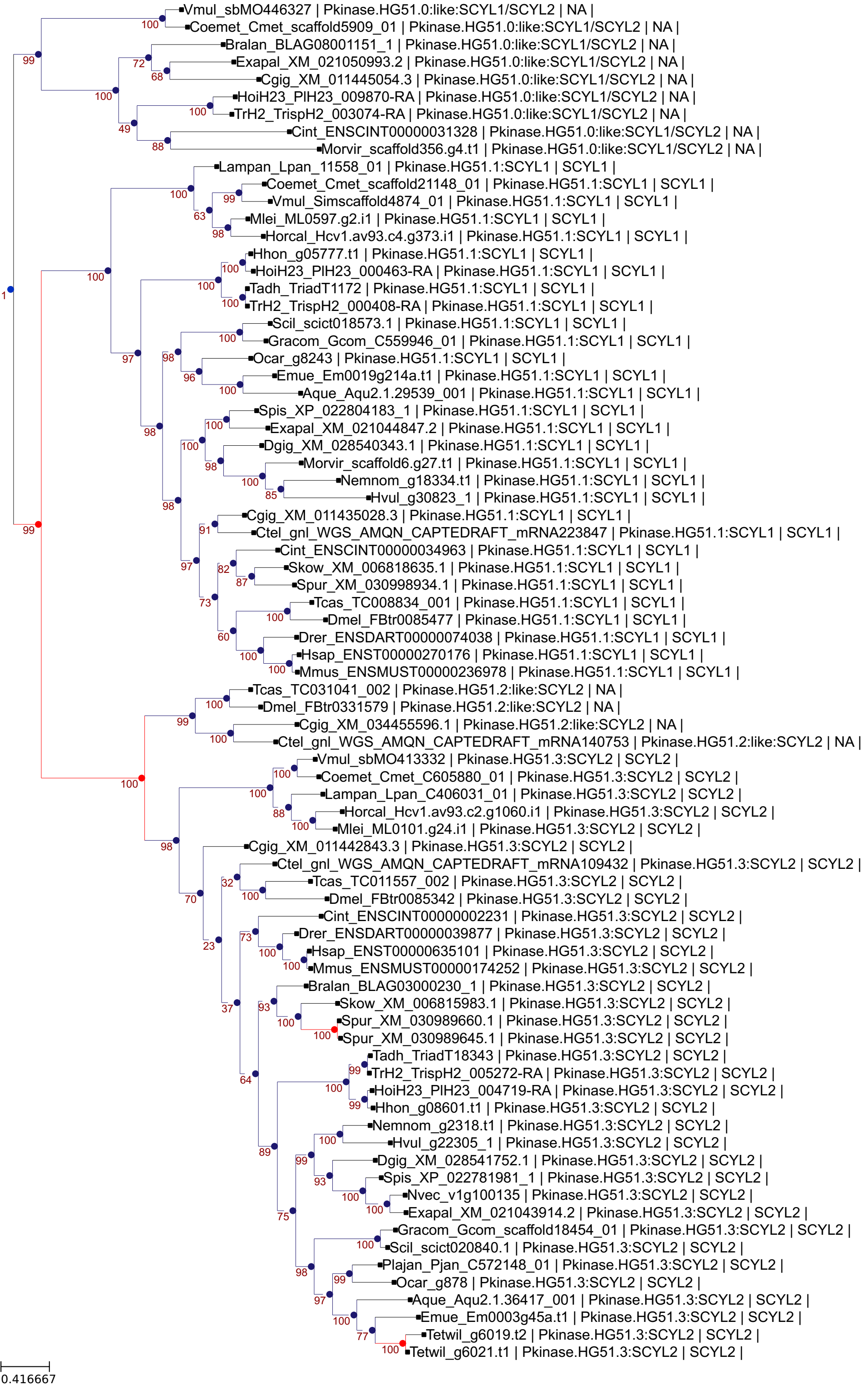
Vmul_sbMO446327 | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
Coemet_Cmet_scaffold5909_01 | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
Bralan_BLAG08001151_1 | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
Exapal_XM_021050993.2 | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
Cgig_XM_011445054.3 | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
HoiH23_PlH23_009870-RA | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
TrH2_TrispH2_003074-RA | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
Cint_ENSCINT00000031328 | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
Morvir_scaffold356.g4.t1 | Pkinase.HG51.0:like:SCYL1/SCYL2 | NA |
Lampan_Lpan_11558_01 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Coemet_Cmet_scaffold21148_01 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Vmul_Simscaffold4874_01 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Mlei_ML0597.g2.i1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Horcal_Hcv1.av93.c4.g373.i1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Hhon_g05777.t1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
HoiH23_PlH23_000463-RA | Pkinase.HG51.1:SCYL1 | SCYL1 |
Tadh_TriadT1172 | Pkinase.HG51.1:SCYL1 | SCYL1 |
TrH2_TrispH2_000408-RA | Pkinase.HG51.1:SCYL1 | SCYL1 |
Scil_scict018573.1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Gracom_Gcom_C559946_01 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Ocar_g8243 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Emue_Em0019g214a.t1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Aque_Aqu2.1.29539_001 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Spis_XP_022804183_1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Exapal_XM_021044847.2 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Dgig_XM_028540343.1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Morvir_scaffold6.g27.t1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Nemnom_g18334.t1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Hvul_g30823_1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Cgig_XM_011435028.3 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA223847 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Cint_ENSCINT00000034963 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Skow_XM_006818635.1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Spur_XM_030998934.1 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Tcas_TC008834_001 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Dmel_FBtr0085477 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Drer_ENSDART00000074038 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Hsap_ENST00000270176 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Mmus_ENSMUST00000236978 | Pkinase.HG51.1:SCYL1 | SCYL1 |
Tcas_TC031041_002 | Pkinase.HG51.2:like:SCYL2 | NA |
Dmel_FBtr0331579 | Pkinase.HG51.2:like:SCYL2 | NA |
Cgig_XM_034455596.1 | Pkinase.HG51.2:like:SCYL2 | NA |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA140753 | Pkinase.HG51.2:like:SCYL2 | NA |
Vmul_sbMO413332 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Coemet_Cmet_C605880_01 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Lampan_Lpan_C406031_01 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Horcal_Hcv1.av93.c2.g1060.i1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Mlei_ML0101.g24.i1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Cgig_XM_011442843.3 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA109432 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Tcas_TC011557_002 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Dmel_FBtr0085342 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Cint_ENSCINT00000002231 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Drer_ENSDART00000039877 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Hsap_ENST00000635101 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Mmus_ENSMUST00000174252 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Bralan_BLAG03000230_1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Skow_XM_006815983.1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Spur_XM_030989660.1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Spur_XM_030989645.1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Tadh_TriadT18343 | Pkinase.HG51.3:SCYL2 | SCYL2 |
TrH2_TrispH2_005272-RA | Pkinase.HG51.3:SCYL2 | SCYL2 |
HoiH23_PlH23_004719-RA | Pkinase.HG51.3:SCYL2 | SCYL2 |
Hhon_g08601.t1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Nemnom_g2318.t1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Hvul_g22305_1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Dgig_XM_028541752.1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Spis_XP_022781981_1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Nvec_v1g100135 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Exapal_XM_021043914.2 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Gracom_Gcom_scaffold18454_01 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Scil_scict020840.1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Plajan_Pjan_C572148_01 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Ocar_g878 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Aque_Aqu2.1.36417_001 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Emue_Em0003g45a.t1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Tetwil_g6019.t2 | Pkinase.HG51.3:SCYL2 | SCYL2 |
Tetwil_g6021.t1 | Pkinase.HG51.3:SCYL2 | SCYL2 |
0.416667