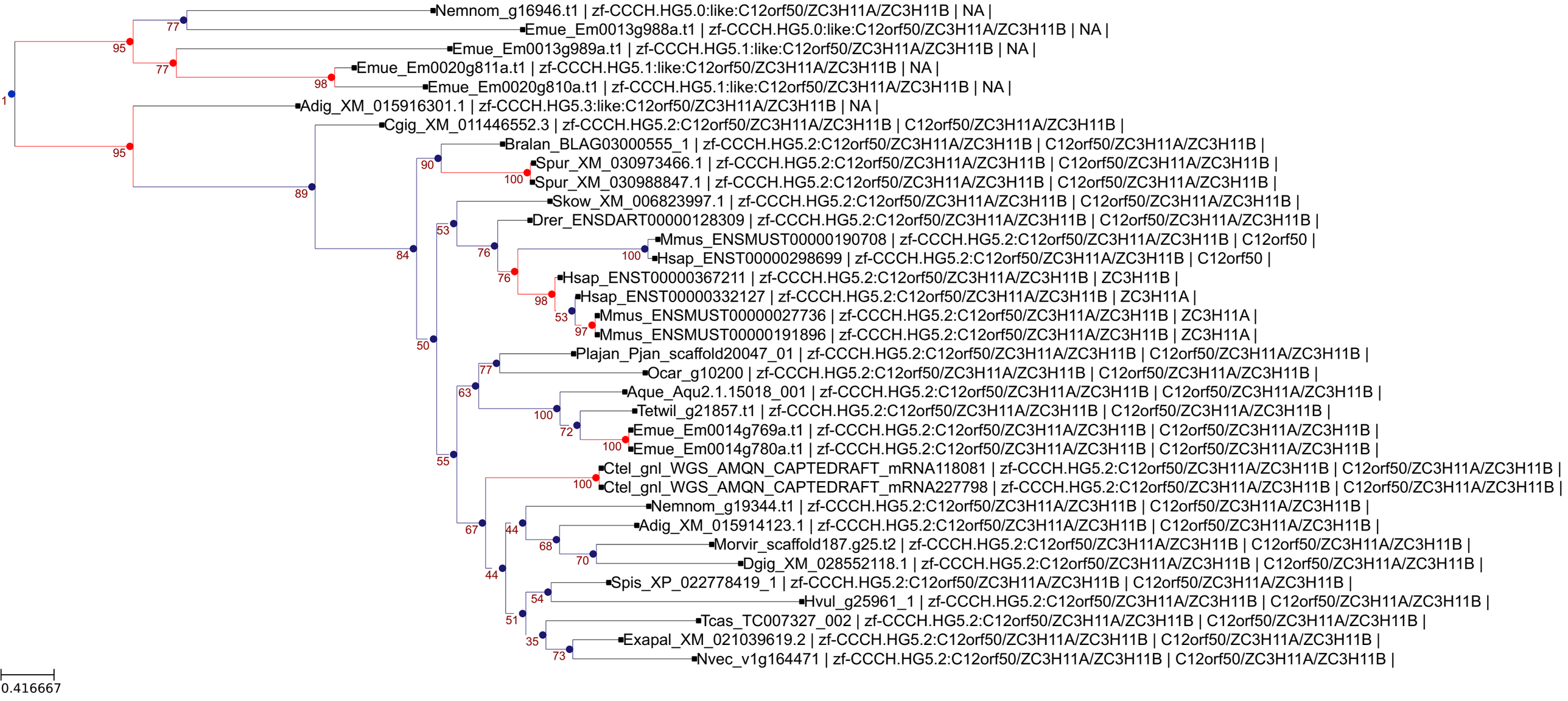
Nemnom_g16946.t1 | zf-CCCH.HG5.0:like:C12orf50/ZC3H11A/ZC3H11B | NA |
Emue_Em0013g988a.t1 | zf-CCCH.HG5.0:like:C12orf50/ZC3H11A/ZC3H11B | NA |
Emue_Em0013g989a.t1 | zf-CCCH.HG5.1:like:C12orf50/ZC3H11A/ZC3H11B | NA |
Emue_Em0020g811a.t1 | zf-CCCH.HG5.1:like:C12orf50/ZC3H11A/ZC3H11B | NA |
Emue_Em0020g810a.t1 | zf-CCCH.HG5.1:like:C12orf50/ZC3H11A/ZC3H11B | NA |
Adig_XM_015916301.1 | zf-CCCH.HG5.3:like:C12orf50/ZC3H11A/ZC3H11B | NA |
Cgig_XM_011446552.3 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Bralan_BLAG03000555_1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Spur_XM_030973466.1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Spur_XM_030988847.1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Skow_XM_006823997.1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Drer_ENSDART00000128309 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Mmus_ENSMUST00000190708 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50 |
Hsap_ENST00000298699 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50 |
Hsap_ENST00000367211 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | ZC3H11B |
Hsap_ENST00000332127 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | ZC3H11A |
Mmus_ENSMUST00000027736 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | ZC3H11A |
Mmus_ENSMUST00000191896 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | ZC3H11A |
Plajan_Pjan_scaffold20047_01 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Ocar_g10200 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Aque_Aqu2.1.15018_001 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Tetwil_g21857.t1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Emue_Em0014g769a.t1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Emue_Em0014g780a.t1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA118081 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA227798 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Nemnom_g19344.t1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Adig_XM_015914123.1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Morvir_scaffold187.g25.t2 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Dgig_XM_028552118.1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Spis_XP_022778419_1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Hvul_g25961_1 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Tcas_TC007327_002 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Exapal_XM_021039619.2 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
Nvec_v1g164471 | zf-CCCH.HG5.2:C12orf50/ZC3H11A/ZC3H11B | C12orf50/ZC3H11A/ZC3H11B |
0.416667