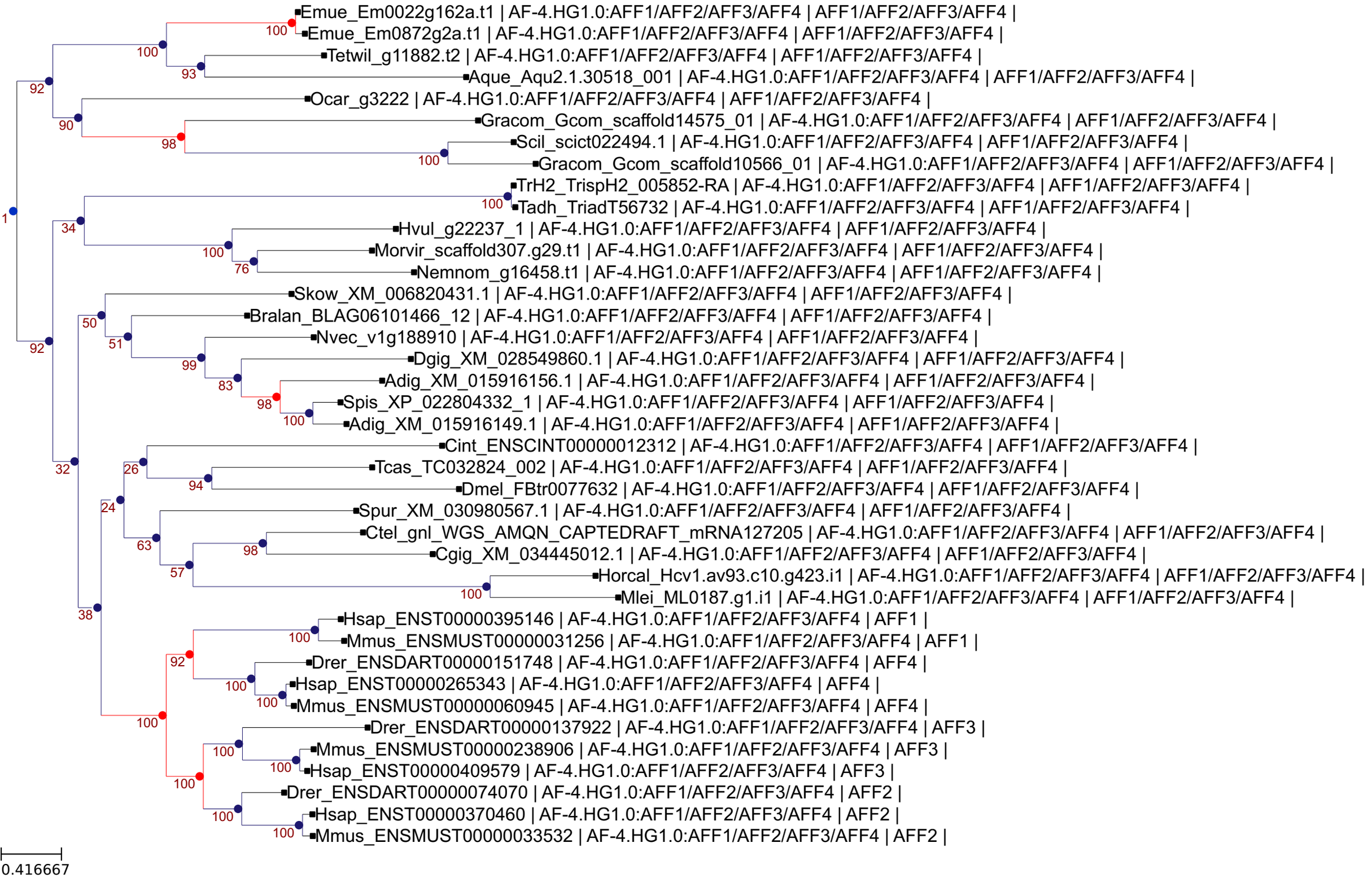
Emue_Em0022g162a.t1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Emue_Em0872g2a.t1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Tetwil_g11882.t2 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Aque_Aqu2.1.30518_001 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Ocar_g3222 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Gracom_Gcom_scaffold14575_01 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Scil_scict022494.1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Gracom_Gcom_scaffold10566_01 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
TrH2_TrispH2_005852-RA | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Tadh_TriadT56732 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Hvul_g22237_1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Morvir_scaffold307.g29.t1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Nemnom_g16458.t1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Skow_XM_006820431.1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Bralan_BLAG06101466_12 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Nvec_v1g188910 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Dgig_XM_028549860.1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Adig_XM_015916156.1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Spis_XP_022804332_1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Adig_XM_015916149.1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Cint_ENSCINT00000012312 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Tcas_TC032824_002 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Dmel_FBtr0077632 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Spur_XM_030980567.1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA127205 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Cgig_XM_034445012.1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Horcal_Hcv1.av93.c10.g423.i1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Mlei_ML0187.g1.i1 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1/AFF2/AFF3/AFF4 |
Hsap_ENST00000395146 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1 |
Mmus_ENSMUST00000031256 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF1 |
Drer_ENSDART00000151748 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF4 |
Hsap_ENST00000265343 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF4 |
Mmus_ENSMUST00000060945 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF4 |
Drer_ENSDART00000137922 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF3 |
Mmus_ENSMUST00000238906 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF3 |
Hsap_ENST00000409579 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF3 |
Drer_ENSDART00000074070 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF2 |
Hsap_ENST00000370460 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF2 |
Mmus_ENSMUST00000033532 | AF-4.HG1.0:AFF1/AFF2/AFF3/AFF4 | AFF2 |
0.416667