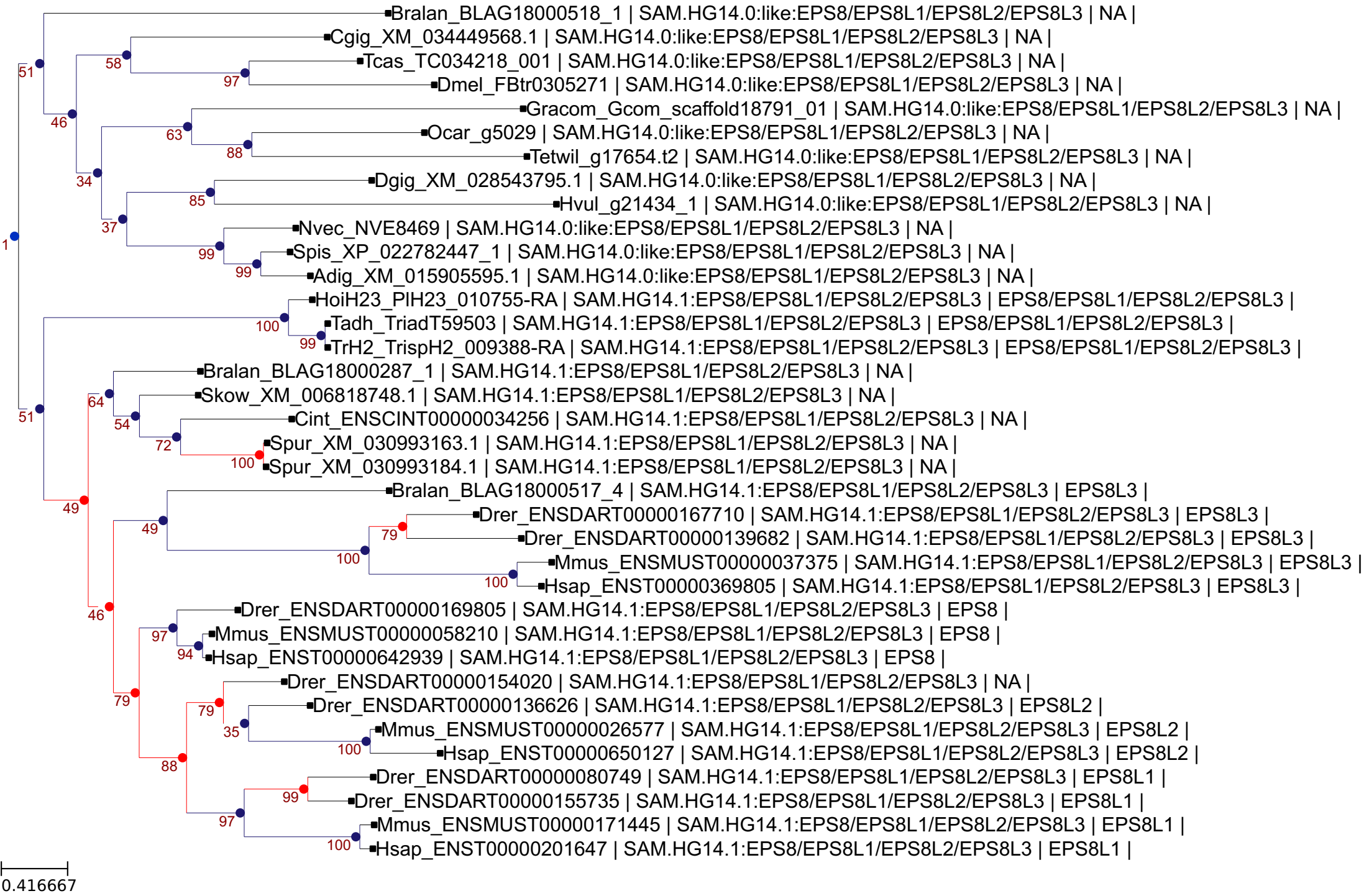
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Cgig_XM_034449568.1 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Tcas_TC034218_001 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Dmel_FBtr0305271 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Gracom_Gcom_scaffold18791_01 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Ocar_g5029 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Tetwil_g17654.t2 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Dgig_XM_028543795.1 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Hvul_g21434_1 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Nvec_NVE8469 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Spis_XP_022782447_1 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Adig_XM_015905595.1 | SAM.HG14.0:like:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
HoiH23_PlH23_010755-RA | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | EPS8/EPS8L1/EPS8L2/EPS8L3 |
Tadh_TriadT59503 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | EPS8/EPS8L1/EPS8L2/EPS8L3 |
TrH2_TrispH2_009388-RA | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | EPS8/EPS8L1/EPS8L2/EPS8L3 |
Bralan_BLAG18000287_1 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Skow_XM_006818748.1 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Cint_ENSCINT00000034256 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Spur_XM_030993163.1 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Spur_XM_030993184.1 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | NA |
Bralan_BLAG18000517_4 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | EPS8L3 |
Drer_ENSDART00000167710 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | EPS8L3 |
Drer_ENSDART00000139682 | SAM.HG14.1:EPS8/EPS8L1/EPS8L2/EPS8L3 | EPS8L3 |
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0.416667