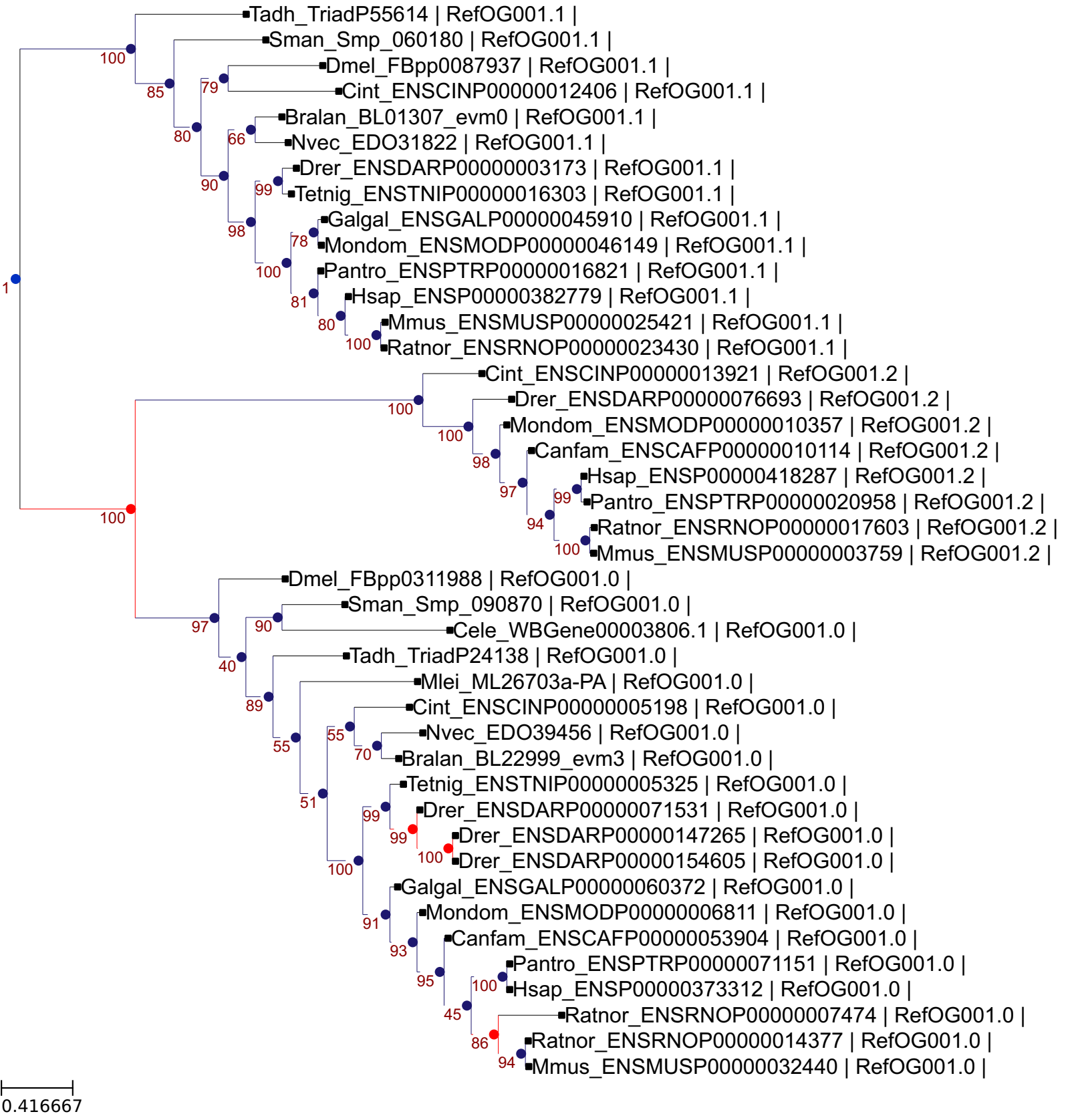
Tadh_TriadP55614 | RefOG001.1 |
Sman_Smp_060180 | RefOG001.1 |
Dmel_FBpp0087937 | RefOG001.1 |
Cint_ENSCINP00000012406 | RefOG001.1 |
Bralan_BL01307_evm0 | RefOG001.1 |
Nvec_EDO31822 | RefOG001.1 |
Drer_ENSDARP00000003173 | RefOG001.1 |
Tetnig_ENSTNIP00000016303 | RefOG001.1 |
Galgal_ENSGALP00000045910 | RefOG001.1 |
Mondom_ENSMODP00000046149 | RefOG001.1 |
Pantro_ENSPTRP00000016821 | RefOG001.1 |
Hsap_ENSP00000382779 | RefOG001.1 |
Mmus_ENSMUSP00000025421 | RefOG001.1 |
Ratnor_ENSRNOP00000023430 | RefOG001.1 |
Cint_ENSCINP00000013921 | RefOG001.2 |
Drer_ENSDARP00000076693 | RefOG001.2 |
Mondom_ENSMODP00000010357 | RefOG001.2 |
Canfam_ENSCAFP00000010114 | RefOG001.2 |
Hsap_ENSP00000418287 | RefOG001.2 |
Pantro_ENSPTRP00000020958 | RefOG001.2 |
Ratnor_ENSRNOP00000017603 | RefOG001.2 |
Mmus_ENSMUSP00000003759 | RefOG001.2 |
Dmel_FBpp0311988 | RefOG001.0 |
Sman_Smp_090870 | RefOG001.0 |
Cele_WBGene00003806.1 | RefOG001.0 |
Tadh_TriadP24138 | RefOG001.0 |
Mlei_ML26703a-PA | RefOG001.0 |
Cint_ENSCINP00000005198 | RefOG001.0 |
Nvec_EDO39456 | RefOG001.0 |
Bralan_BL22999_evm3 | RefOG001.0 |
Tetnig_ENSTNIP00000005325 | RefOG001.0 |
Drer_ENSDARP00000071531 | RefOG001.0 |
Drer_ENSDARP00000147265 | RefOG001.0 |
Drer_ENSDARP00000154605 | RefOG001.0 |
Galgal_ENSGALP00000060372 | RefOG001.0 |
Mondom_ENSMODP00000006811 | RefOG001.0 |
Canfam_ENSCAFP00000053904 | RefOG001.0 |
Pantro_ENSPTRP00000071151 | RefOG001.0 |
Hsap_ENSP00000373312 | RefOG001.0 |
Ratnor_ENSRNOP00000007474 | RefOG001.0 |
Ratnor_ENSRNOP00000014377 | RefOG001.0 |
Mmus_ENSMUSP00000032440 | RefOG001.0 |
0.416667