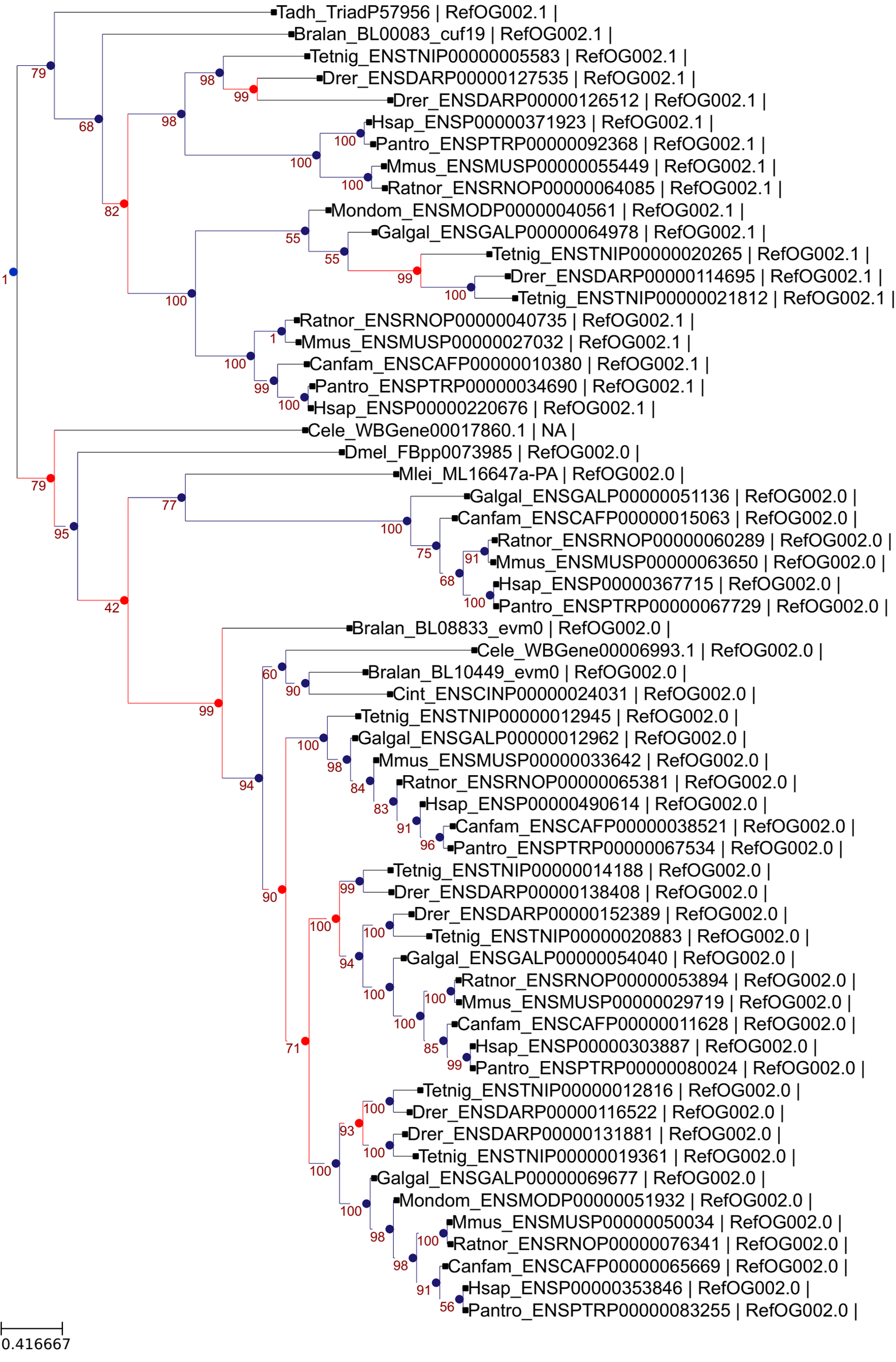
Tadh_TriadP57956 | RefOG002.1 |
Bralan_BL00083_cuf19 | RefOG002.1 |
Tetnig_ENSTNIP00000005583 | RefOG002.1 |
Drer_ENSDARP00000127535 | RefOG002.1 |
Drer_ENSDARP00000126512 | RefOG002.1 |
Hsap_ENSP00000371923 | RefOG002.1 |
Pantro_ENSPTRP00000092368 | RefOG002.1 |
Mmus_ENSMUSP00000055449 | RefOG002.1 |
Ratnor_ENSRNOP00000064085 | RefOG002.1 |
Mondom_ENSMODP00000040561 | RefOG002.1 |
Galgal_ENSGALP00000064978 | RefOG002.1 |
Tetnig_ENSTNIP00000020265 | RefOG002.1 |
Drer_ENSDARP00000114695 | RefOG002.1 |
Tetnig_ENSTNIP00000021812 | RefOG002.1 |
Ratnor_ENSRNOP00000040735 | RefOG002.1 |
Mmus_ENSMUSP00000027032 | RefOG002.1 |
Canfam_ENSCAFP00000010380 | RefOG002.1 |
Pantro_ENSPTRP00000034690 | RefOG002.1 |
Hsap_ENSP00000220676 | RefOG002.1 |
Cele_WBGene00017860.1 | NA |
Dmel_FBpp0073985 | RefOG002.0 |
Mlei_ML16647a-PA | RefOG002.0 |
Galgal_ENSGALP00000051136 | RefOG002.0 |
Canfam_ENSCAFP00000015063 | RefOG002.0 |
Ratnor_ENSRNOP00000060289 | RefOG002.0 |
Mmus_ENSMUSP00000063650 | RefOG002.0 |
Hsap_ENSP00000367715 | RefOG002.0 |
Pantro_ENSPTRP00000067729 | RefOG002.0 |
Bralan_BL08833_evm0 | RefOG002.0 |
Cele_WBGene00006993.1 | RefOG002.0 |
Bralan_BL10449_evm0 | RefOG002.0 |
Cint_ENSCINP00000024031 | RefOG002.0 |
Tetnig_ENSTNIP00000012945 | RefOG002.0 |
Galgal_ENSGALP00000012962 | RefOG002.0 |
Mmus_ENSMUSP00000033642 | RefOG002.0 |
Ratnor_ENSRNOP00000065381 | RefOG002.0 |
Hsap_ENSP00000490614 | RefOG002.0 |
Canfam_ENSCAFP00000038521 | RefOG002.0 |
Pantro_ENSPTRP00000067534 | RefOG002.0 |
Tetnig_ENSTNIP00000014188 | RefOG002.0 |
Drer_ENSDARP00000138408 | RefOG002.0 |
Drer_ENSDARP00000152389 | RefOG002.0 |
Tetnig_ENSTNIP00000020883 | RefOG002.0 |
Galgal_ENSGALP00000054040 | RefOG002.0 |
Ratnor_ENSRNOP00000053894 | RefOG002.0 |
Mmus_ENSMUSP00000029719 | RefOG002.0 |
Canfam_ENSCAFP00000011628 | RefOG002.0 |
Hsap_ENSP00000303887 | RefOG002.0 |
Pantro_ENSPTRP00000080024 | RefOG002.0 |
Tetnig_ENSTNIP00000012816 | RefOG002.0 |
Drer_ENSDARP00000116522 | RefOG002.0 |
Drer_ENSDARP00000131881 | RefOG002.0 |
Tetnig_ENSTNIP00000019361 | RefOG002.0 |
Galgal_ENSGALP00000069677 | RefOG002.0 |
Mondom_ENSMODP00000051932 | RefOG002.0 |
Mmus_ENSMUSP00000050034 | RefOG002.0 |
Ratnor_ENSRNOP00000076341 | RefOG002.0 |
Canfam_ENSCAFP00000065669 | RefOG002.0 |
Hsap_ENSP00000353846 | RefOG002.0 |
Pantro_ENSPTRP00000083255 | RefOG002.0 |
0.416667