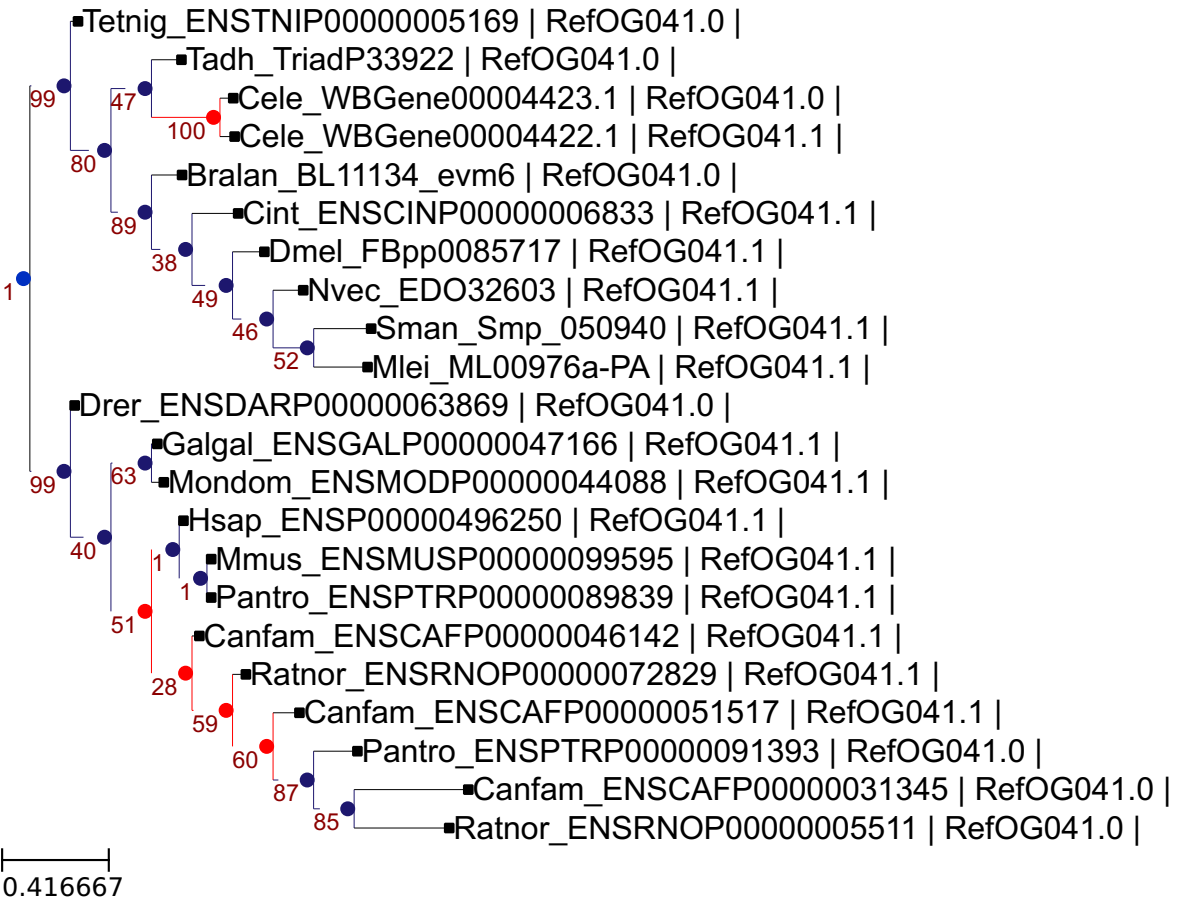
Tetnig_ENSTNIP00000005169 | RefOG041.0 |
Tadh_TriadP33922 | RefOG041.0 |
Cele_WBGene00004423.1 | RefOG041.0 |
Cele_WBGene00004422.1 | RefOG041.1 |
Bralan_BL11134_evm6 | RefOG041.0 |
Cint_ENSCINP00000006833 | RefOG041.1 |
Dmel_FBpp0085717 | RefOG041.1 |
Nvec_EDO32603 | RefOG041.1 |
Sman_Smp_050940 | RefOG041.1 |
Mlei_ML00976a-PA | RefOG041.1 |
Drer_ENSDARP00000063869 | RefOG041.0 |
Galgal_ENSGALP00000047166 | RefOG041.1 |
Mondom_ENSMODP00000044088 | RefOG041.1 |
Hsap_ENSP00000496250 | RefOG041.1 |
Mmus_ENSMUSP00000099595 | RefOG041.1 |
Pantro_ENSPTRP00000089839 | RefOG041.1 |
Canfam_ENSCAFP00000046142 | RefOG041.1 |
Ratnor_ENSRNOP00000072829 | RefOG041.1 |
Canfam_ENSCAFP00000051517 | RefOG041.1 |
Pantro_ENSPTRP00000091393 | RefOG041.0 |
Canfam_ENSCAFP00000031345 | RefOG041.0 |
Ratnor_ENSRNOP00000005511 | RefOG041.0 |
99
47
100
80
89
38
49
46
52
1
99
63
40
1
1
51
28
59
60
87
85
0.416667