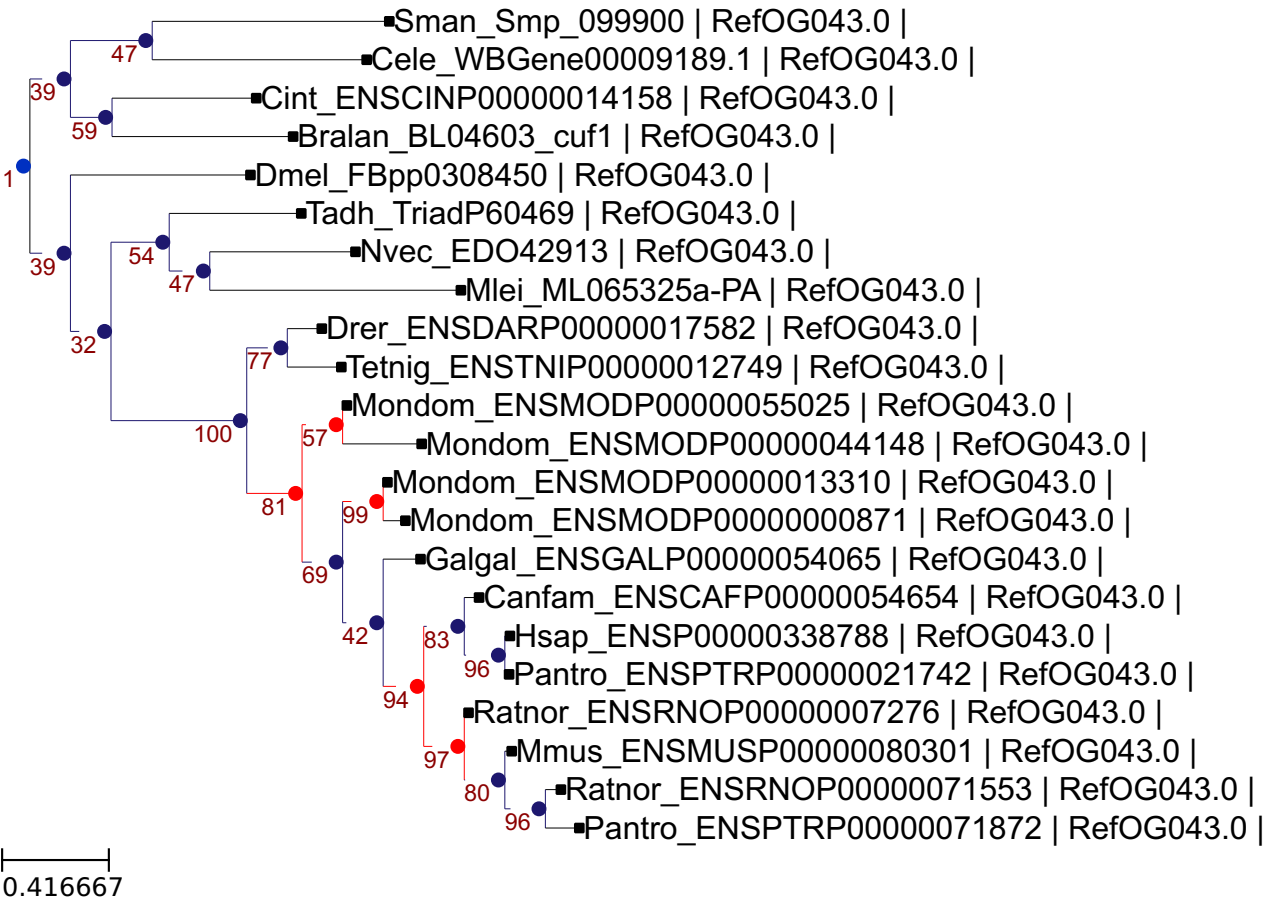
Sman_Smp_099900 | RefOG043.0 |
Cele_WBGene00009189.1 | RefOG043.0 |
Cint_ENSCINP00000014158 | RefOG043.0 |
Bralan_BL04603_cuf1 | RefOG043.0 |
Dmel_FBpp0308450 | RefOG043.0 |
Tadh_TriadP60469 | RefOG043.0 |
Nvec_EDO42913 | RefOG043.0 |
Mlei_ML065325a-PA | RefOG043.0 |
Drer_ENSDARP00000017582 | RefOG043.0 |
Tetnig_ENSTNIP00000012749 | RefOG043.0 |
Mondom_ENSMODP00000055025 | RefOG043.0 |
Mondom_ENSMODP00000044148 | RefOG043.0 |
Mondom_ENSMODP00000013310 | RefOG043.0 |
Mondom_ENSMODP00000000871 | RefOG043.0 |
Galgal_ENSGALP00000054065 | RefOG043.0 |
Canfam_ENSCAFP00000054654 | RefOG043.0 |
Hsap_ENSP00000338788 | RefOG043.0 |
Pantro_ENSPTRP00000021742 | RefOG043.0 |
Ratnor_ENSRNOP00000007276 | RefOG043.0 |
Mmus_ENSMUSP00000080301 | RefOG043.0 |
Ratnor_ENSRNOP00000071553 | RefOG043.0 |
Pantro_ENSPTRP00000071872 | RefOG043.0 |
47
39
59
1
39
54
47
32
77
100
57
81
99
69
42
83
96
94
97
80
96
0.416667