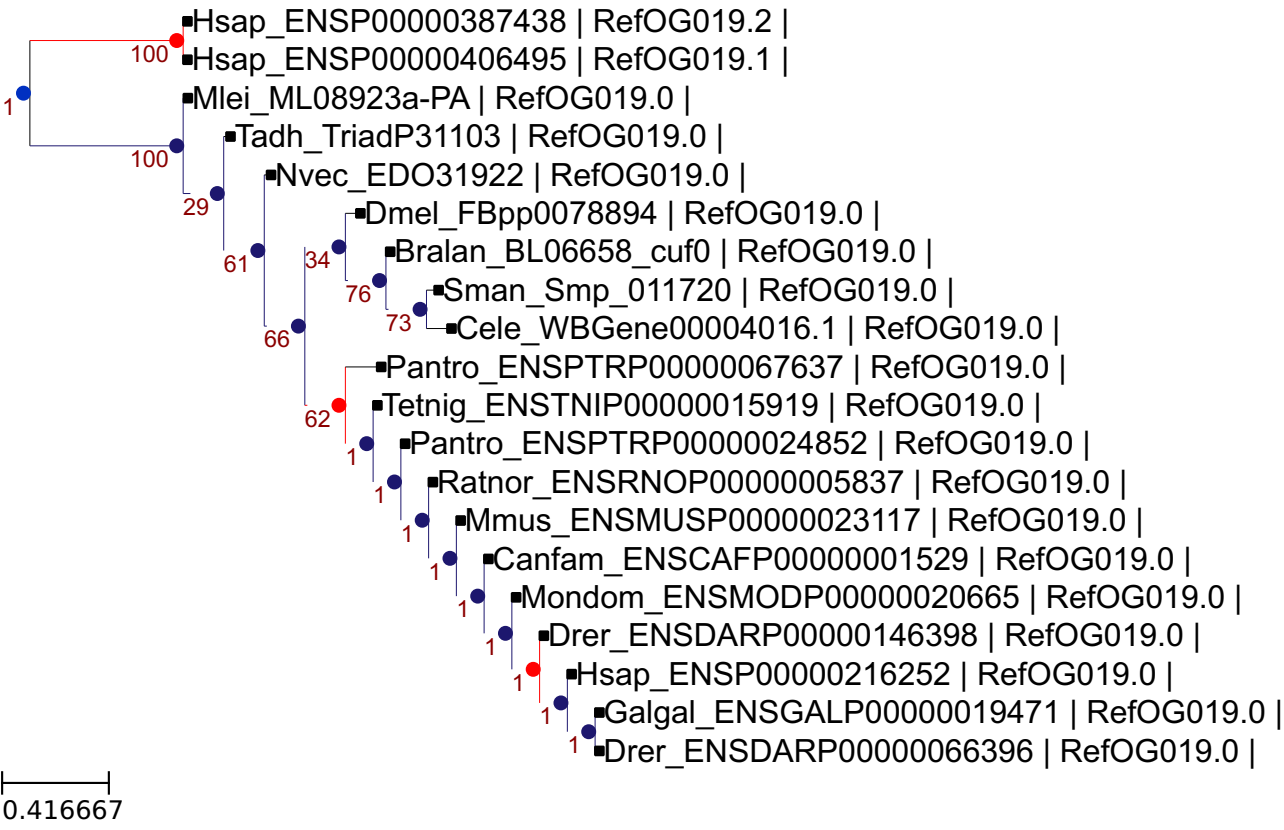
Hsap_ENSP00000387438 | RefOG019.2 |
Hsap_ENSP00000406495 | RefOG019.1 |
Mlei_ML08923a-PA | RefOG019.0 |
Tadh_TriadP31103 | RefOG019.0 |
Nvec_EDO31922 | RefOG019.0 |
Dmel_FBpp0078894 | RefOG019.0 |
Bralan_BL06658_cuf0 | RefOG019.0 |
Sman_Smp_011720 | RefOG019.0 |
Cele_WBGene00004016.1 | RefOG019.0 |
Pantro_ENSPTRP00000067637 | RefOG019.0 |
Tetnig_ENSTNIP00000015919 | RefOG019.0 |
Pantro_ENSPTRP00000024852 | RefOG019.0 |
Ratnor_ENSRNOP00000005837 | RefOG019.0 |
Mmus_ENSMUSP00000023117 | RefOG019.0 |
Canfam_ENSCAFP00000001529 | RefOG019.0 |
Mondom_ENSMODP00000020665 | RefOG019.0 |
Drer_ENSDARP00000146398 | RefOG019.0 |
Hsap_ENSP00000216252 | RefOG019.0 |
Galgal_ENSGALP00000019471 | RefOG019.0 |
Drer_ENSDARP00000066396 | RefOG019.0 |
100
1
100
29
61
34
76
73
66
62
1
1
1
1
1
1
1
1
0.416667