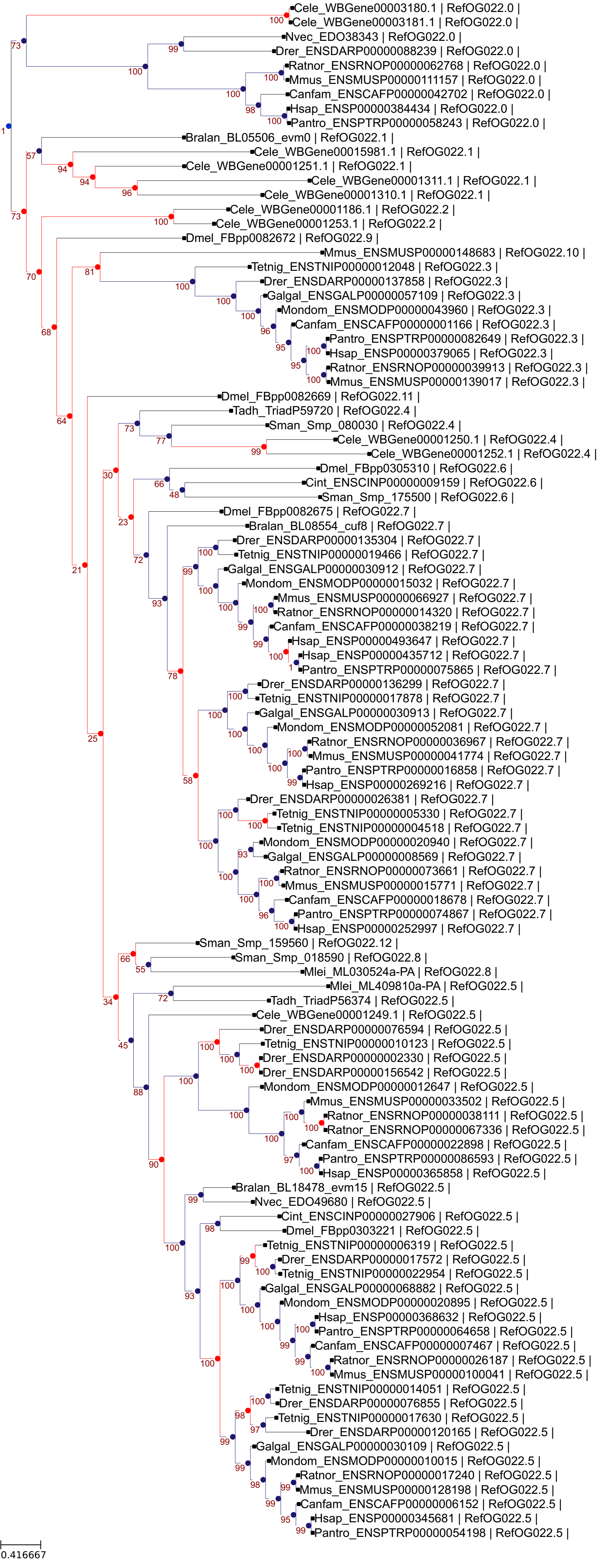
Cele_WBGene00003180.1 | RefOG022.0 |
Cele_WBGene00003181.1 | RefOG022.0 |
Nvec_EDO38343 | RefOG022.0 |
Drer_ENSDARP00000088239 | RefOG022.0 |
Ratnor_ENSRNOP00000062768 | RefOG022.0 |
Mmus_ENSMUSP00000111157 | RefOG022.0 |
Canfam_ENSCAFP00000042702 | RefOG022.0 |
Hsap_ENSP00000384434 | RefOG022.0 |
Pantro_ENSPTRP00000058243 | RefOG022.0 |
Bralan_BL05506_evm0 | RefOG022.1 |
Cele_WBGene00015981.1 | RefOG022.1 |
Cele_WBGene00001251.1 | RefOG022.1 |
Cele_WBGene00001311.1 | RefOG022.1 |
Cele_WBGene00001310.1 | RefOG022.1 |
Cele_WBGene00001186.1 | RefOG022.2 |
Cele_WBGene00001253.1 | RefOG022.2 |
Dmel_FBpp0082672 | RefOG022.9 |
Mmus_ENSMUSP00000148683 | RefOG022.10 |
Tetnig_ENSTNIP00000012048 | RefOG022.3 |
Drer_ENSDARP00000137858 | RefOG022.3 |
Galgal_ENSGALP00000057109 | RefOG022.3 |
Mondom_ENSMODP00000043960 | RefOG022.3 |
Canfam_ENSCAFP00000001166 | RefOG022.3 |
Pantro_ENSPTRP00000082649 | RefOG022.3 |
Hsap_ENSP00000379065 | RefOG022.3 |
Ratnor_ENSRNOP00000039913 | RefOG022.3 |
Mmus_ENSMUSP00000139017 | RefOG022.3 |
Dmel_FBpp0082669 | RefOG022.11 |
Tadh_TriadP59720 | RefOG022.4 |
Sman_Smp_080030 | RefOG022.4 |
Cele_WBGene00001250.1 | RefOG022.4 |
Cele_WBGene00001252.1 | RefOG022.4 |
Dmel_FBpp0305310 | RefOG022.6 |
Cint_ENSCINP00000009159 | RefOG022.6 |
Sman_Smp_175500 | RefOG022.6 |
Dmel_FBpp0082675 | RefOG022.7 |
Bralan_BL08554_cuf8 | RefOG022.7 |
Drer_ENSDARP00000135304 | RefOG022.7 |
Tetnig_ENSTNIP00000019466 | RefOG022.7 |
Galgal_ENSGALP00000030912 | RefOG022.7 |
Mondom_ENSMODP00000015032 | RefOG022.7 |
Mmus_ENSMUSP00000066927 | RefOG022.7 |
Ratnor_ENSRNOP00000014320 | RefOG022.7 |
Canfam_ENSCAFP00000038219 | RefOG022.7 |
Hsap_ENSP00000493647 | RefOG022.7 |
Hsap_ENSP00000435712 | RefOG022.7 |
Pantro_ENSPTRP00000075865 | RefOG022.7 |
Drer_ENSDARP00000136299 | RefOG022.7 |
Tetnig_ENSTNIP00000017878 | RefOG022.7 |
Galgal_ENSGALP00000030913 | RefOG022.7 |
Mondom_ENSMODP00000052081 | RefOG022.7 |
Ratnor_ENSRNOP00000036967 | RefOG022.7 |
Mmus_ENSMUSP00000041774 | RefOG022.7 |
Pantro_ENSPTRP00000016858 | RefOG022.7 |
Hsap_ENSP00000269216 | RefOG022.7 |
Drer_ENSDARP00000026381 | RefOG022.7 |
Tetnig_ENSTNIP00000005330 | RefOG022.7 |
Tetnig_ENSTNIP00000004518 | RefOG022.7 |
Mondom_ENSMODP00000020940 | RefOG022.7 |
Galgal_ENSGALP00000008569 | RefOG022.7 |
Ratnor_ENSRNOP00000073661 | RefOG022.7 |
Mmus_ENSMUSP00000015771 | RefOG022.7 |
Canfam_ENSCAFP00000018678 | RefOG022.7 |
Pantro_ENSPTRP00000074867 | RefOG022.7 |
Hsap_ENSP00000252997 | RefOG022.7 |
Sman_Smp_159560 | RefOG022.12 |
Sman_Smp_018590 | RefOG022.8 |
Mlei_ML030524a-PA | RefOG022.8 |
Mlei_ML409810a-PA | RefOG022.5 |
Tadh_TriadP56374 | RefOG022.5 |
Cele_WBGene00001249.1 | RefOG022.5 |
Drer_ENSDARP00000076594 | RefOG022.5 |
Tetnig_ENSTNIP00000010123 | RefOG022.5 |
Drer_ENSDARP00000002330 | RefOG022.5 |
Drer_ENSDARP00000156542 | RefOG022.5 |
Mondom_ENSMODP00000012647 | RefOG022.5 |
Mmus_ENSMUSP00000033502 | RefOG022.5 |
Ratnor_ENSRNOP00000038111 | RefOG022.5 |
Ratnor_ENSRNOP00000067336 | RefOG022.5 |
Canfam_ENSCAFP00000022898 | RefOG022.5 |
Pantro_ENSPTRP00000086593 | RefOG022.5 |
Hsap_ENSP00000365858 | RefOG022.5 |
Bralan_BL18478_evm15 | RefOG022.5 |
Nvec_EDO49680 | RefOG022.5 |
Cint_ENSCINP00000027906 | RefOG022.5 |
Dmel_FBpp0303221 | RefOG022.5 |
Tetnig_ENSTNIP00000006319 | RefOG022.5 |
Drer_ENSDARP00000017572 | RefOG022.5 |
Tetnig_ENSTNIP00000022954 | RefOG022.5 |
Galgal_ENSGALP00000068882 | RefOG022.5 |
Mondom_ENSMODP00000020895 | RefOG022.5 |
Hsap_ENSP00000368632 | RefOG022.5 |
Pantro_ENSPTRP00000064658 | RefOG022.5 |
Canfam_ENSCAFP00000007467 | RefOG022.5 |
Ratnor_ENSRNOP00000026187 | RefOG022.5 |
Mmus_ENSMUSP00000100041 | RefOG022.5 |
Tetnig_ENSTNIP00000014051 | RefOG022.5 |
Drer_ENSDARP00000076855 | RefOG022.5 |
Tetnig_ENSTNIP00000017630 | RefOG022.5 |
Drer_ENSDARP00000120165 | RefOG022.5 |
Galgal_ENSGALP00000030109 | RefOG022.5 |
Mondom_ENSMODP00000010015 | RefOG022.5 |
Ratnor_ENSRNOP00000017240 | RefOG022.5 |
Mmus_ENSMUSP00000128198 | RefOG022.5 |
Canfam_ENSCAFP00000006152 | RefOG022.5 |
Hsap_ENSP00000345681 | RefOG022.5 |
Pantro_ENSPTRP00000054198 | RefOG022.5 |
0.416667