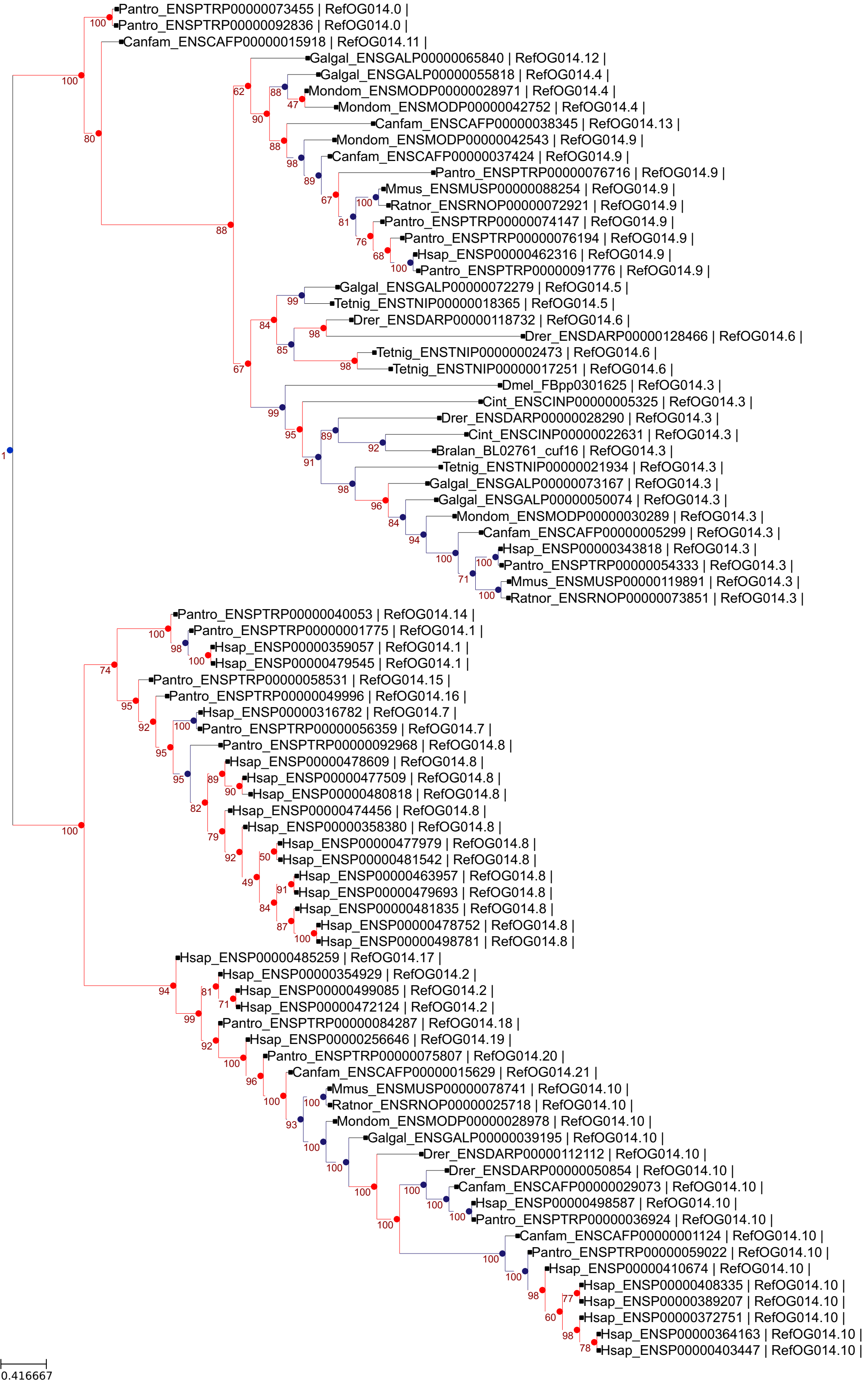
Pantro_ENSPTRP00000073455 | RefOG014.0 |
Pantro_ENSPTRP00000092836 | RefOG014.0 |
Canfam_ENSCAFP00000015918 | RefOG014.11 |
Galgal_ENSGALP00000065840 | RefOG014.12 |
Galgal_ENSGALP00000055818 | RefOG014.4 |
Mondom_ENSMODP00000028971 | RefOG014.4 |
Mondom_ENSMODP00000042752 | RefOG014.4 |
Canfam_ENSCAFP00000038345 | RefOG014.13 |
Mondom_ENSMODP00000042543 | RefOG014.9 |
Canfam_ENSCAFP00000037424 | RefOG014.9 |
Pantro_ENSPTRP00000076716 | RefOG014.9 |
Mmus_ENSMUSP00000088254 | RefOG014.9 |
Ratnor_ENSRNOP00000072921 | RefOG014.9 |
Pantro_ENSPTRP00000074147 | RefOG014.9 |
Pantro_ENSPTRP00000076194 | RefOG014.9 |
Hsap_ENSP00000462316 | RefOG014.9 |
Pantro_ENSPTRP00000091776 | RefOG014.9 |
Galgal_ENSGALP00000072279 | RefOG014.5 |
Tetnig_ENSTNIP00000018365 | RefOG014.5 |
Drer_ENSDARP00000118732 | RefOG014.6 |
Drer_ENSDARP00000128466 | RefOG014.6 |
Tetnig_ENSTNIP00000002473 | RefOG014.6 |
Tetnig_ENSTNIP00000017251 | RefOG014.6 |
Dmel_FBpp0301625 | RefOG014.3 |
Cint_ENSCINP00000005325 | RefOG014.3 |
Drer_ENSDARP00000028290 | RefOG014.3 |
Cint_ENSCINP00000022631 | RefOG014.3 |
Bralan_BL02761_cuf16 | RefOG014.3 |
Tetnig_ENSTNIP00000021934 | RefOG014.3 |
Galgal_ENSGALP00000073167 | RefOG014.3 |
Galgal_ENSGALP00000050074 | RefOG014.3 |
Mondom_ENSMODP00000030289 | RefOG014.3 |
Canfam_ENSCAFP00000005299 | RefOG014.3 |
Hsap_ENSP00000343818 | RefOG014.3 |
Pantro_ENSPTRP00000054333 | RefOG014.3 |
Mmus_ENSMUSP00000119891 | RefOG014.3 |
Ratnor_ENSRNOP00000073851 | RefOG014.3 |
Pantro_ENSPTRP00000040053 | RefOG014.14 |
Pantro_ENSPTRP00000001775 | RefOG014.1 |
Hsap_ENSP00000359057 | RefOG014.1 |
Hsap_ENSP00000479545 | RefOG014.1 |
Pantro_ENSPTRP00000058531 | RefOG014.15 |
Pantro_ENSPTRP00000049996 | RefOG014.16 |
Hsap_ENSP00000316782 | RefOG014.7 |
Pantro_ENSPTRP00000056359 | RefOG014.7 |
Pantro_ENSPTRP00000092968 | RefOG014.8 |
Hsap_ENSP00000478609 | RefOG014.8 |
Hsap_ENSP00000477509 | RefOG014.8 |
Hsap_ENSP00000480818 | RefOG014.8 |
Hsap_ENSP00000474456 | RefOG014.8 |
Hsap_ENSP00000358380 | RefOG014.8 |
Hsap_ENSP00000477979 | RefOG014.8 |
Hsap_ENSP00000481542 | RefOG014.8 |
Hsap_ENSP00000463957 | RefOG014.8 |
Hsap_ENSP00000479693 | RefOG014.8 |
Hsap_ENSP00000481835 | RefOG014.8 |
Hsap_ENSP00000478752 | RefOG014.8 |
Hsap_ENSP00000498781 | RefOG014.8 |
Hsap_ENSP00000485259 | RefOG014.17 |
Hsap_ENSP00000354929 | RefOG014.2 |
Hsap_ENSP00000499085 | RefOG014.2 |
Hsap_ENSP00000472124 | RefOG014.2 |
Pantro_ENSPTRP00000084287 | RefOG014.18 |
Hsap_ENSP00000256646 | RefOG014.19 |
Pantro_ENSPTRP00000075807 | RefOG014.20 |
Canfam_ENSCAFP00000015629 | RefOG014.21 |
Mmus_ENSMUSP00000078741 | RefOG014.10 |
Ratnor_ENSRNOP00000025718 | RefOG014.10 |
Mondom_ENSMODP00000028978 | RefOG014.10 |
Galgal_ENSGALP00000039195 | RefOG014.10 |
Drer_ENSDARP00000112112 | RefOG014.10 |
Drer_ENSDARP00000050854 | RefOG014.10 |
Canfam_ENSCAFP00000029073 | RefOG014.10 |
Hsap_ENSP00000498587 | RefOG014.10 |
Pantro_ENSPTRP00000036924 | RefOG014.10 |
Canfam_ENSCAFP00000001124 | RefOG014.10 |
Pantro_ENSPTRP00000059022 | RefOG014.10 |
Hsap_ENSP00000410674 | RefOG014.10 |
Hsap_ENSP00000408335 | RefOG014.10 |
Hsap_ENSP00000389207 | RefOG014.10 |
Hsap_ENSP00000372751 | RefOG014.10 |
Hsap_ENSP00000364163 | RefOG014.10 |
Hsap_ENSP00000403447 | RefOG014.10 |
0.416667