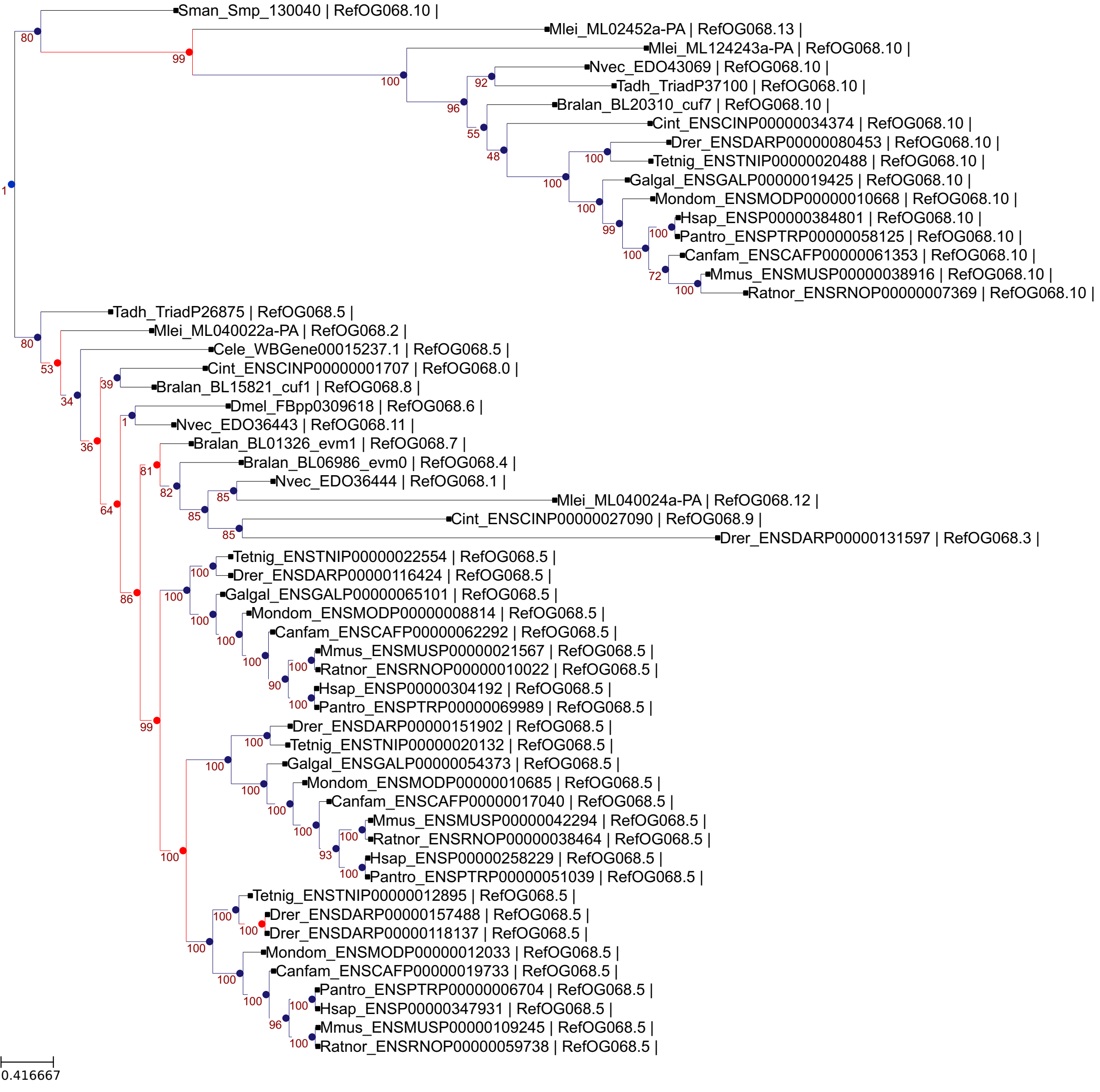
Sman_Smp_130040 | RefOG068.10 |
Mlei_ML02452a-PA | RefOG068.13 |
Mlei_ML124243a-PA | RefOG068.10 |
Nvec_EDO43069 | RefOG068.10 |
Tadh_TriadP37100 | RefOG068.10 |
Bralan_BL20310_cuf7 | RefOG068.10 |
Cint_ENSCINP00000034374 | RefOG068.10 |
Drer_ENSDARP00000080453 | RefOG068.10 |
Tetnig_ENSTNIP00000020488 | RefOG068.10 |
Galgal_ENSGALP00000019425 | RefOG068.10 |
Mondom_ENSMODP00000010668 | RefOG068.10 |
Hsap_ENSP00000384801 | RefOG068.10 |
Pantro_ENSPTRP00000058125 | RefOG068.10 |
Canfam_ENSCAFP00000061353 | RefOG068.10 |
Mmus_ENSMUSP00000038916 | RefOG068.10 |
Ratnor_ENSRNOP00000007369 | RefOG068.10 |
Tadh_TriadP26875 | RefOG068.5 |
Mlei_ML040022a-PA | RefOG068.2 |
Cele_WBGene00015237.1 | RefOG068.5 |
Cint_ENSCINP00000001707 | RefOG068.0 |
Bralan_BL15821_cuf1 | RefOG068.8 |
Dmel_FBpp0309618 | RefOG068.6 |
Nvec_EDO36443 | RefOG068.11 |
Bralan_BL01326_evm1 | RefOG068.7 |
Bralan_BL06986_evm0 | RefOG068.4 |
Nvec_EDO36444 | RefOG068.1 |
Mlei_ML040024a-PA | RefOG068.12 |
Cint_ENSCINP00000027090 | RefOG068.9 |
Drer_ENSDARP00000131597 | RefOG068.3 |
Tetnig_ENSTNIP00000022554 | RefOG068.5 |
Drer_ENSDARP00000116424 | RefOG068.5 |
Galgal_ENSGALP00000065101 | RefOG068.5 |
Mondom_ENSMODP00000008814 | RefOG068.5 |
Canfam_ENSCAFP00000062292 | RefOG068.5 |
Mmus_ENSMUSP00000021567 | RefOG068.5 |
Ratnor_ENSRNOP00000010022 | RefOG068.5 |
Hsap_ENSP00000304192 | RefOG068.5 |
Pantro_ENSPTRP00000069989 | RefOG068.5 |
Drer_ENSDARP00000151902 | RefOG068.5 |
Tetnig_ENSTNIP00000020132 | RefOG068.5 |
Galgal_ENSGALP00000054373 | RefOG068.5 |
Mondom_ENSMODP00000010685 | RefOG068.5 |
Canfam_ENSCAFP00000017040 | RefOG068.5 |
Mmus_ENSMUSP00000042294 | RefOG068.5 |
Ratnor_ENSRNOP00000038464 | RefOG068.5 |
Hsap_ENSP00000258229 | RefOG068.5 |
Pantro_ENSPTRP00000051039 | RefOG068.5 |
Tetnig_ENSTNIP00000012895 | RefOG068.5 |
Drer_ENSDARP00000157488 | RefOG068.5 |
Drer_ENSDARP00000118137 | RefOG068.5 |
Mondom_ENSMODP00000012033 | RefOG068.5 |
Canfam_ENSCAFP00000019733 | RefOG068.5 |
Pantro_ENSPTRP00000006704 | RefOG068.5 |
Hsap_ENSP00000347931 | RefOG068.5 |
Mmus_ENSMUSP00000109245 | RefOG068.5 |
Ratnor_ENSRNOP00000059738 | RefOG068.5 |
0.416667