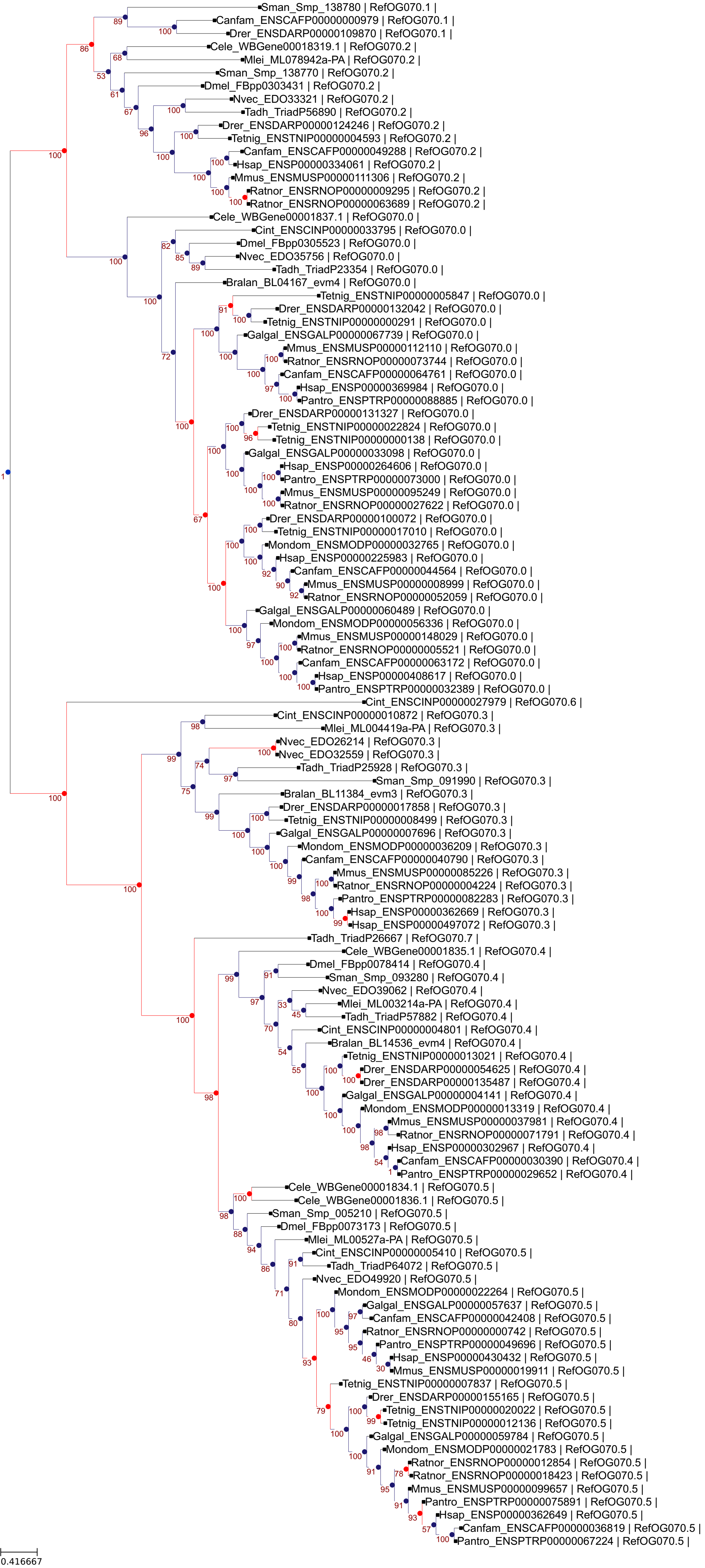
Sman_Smp_138780 | RefOG070.1 |
Canfam_ENSCAFP00000000979 | RefOG070.1 |
Drer_ENSDARP00000109870 | RefOG070.1 |
Cele_WBGene00018319.1 | RefOG070.2 |
Mlei_ML078942a-PA | RefOG070.2 |
Sman_Smp_138770 | RefOG070.2 |
Dmel_FBpp0303431 | RefOG070.2 |
Nvec_EDO33321 | RefOG070.2 |
Tadh_TriadP56890 | RefOG070.2 |
Drer_ENSDARP00000124246 | RefOG070.2 |
Tetnig_ENSTNIP00000004593 | RefOG070.2 |
Canfam_ENSCAFP00000049288 | RefOG070.2 |
Hsap_ENSP00000334061 | RefOG070.2 |
Mmus_ENSMUSP00000111306 | RefOG070.2 |
Ratnor_ENSRNOP00000009295 | RefOG070.2 |
Ratnor_ENSRNOP00000063689 | RefOG070.2 |
Cele_WBGene00001837.1 | RefOG070.0 |
Cint_ENSCINP00000033795 | RefOG070.0 |
Dmel_FBpp0305523 | RefOG070.0 |
Nvec_EDO35756 | RefOG070.0 |
Tadh_TriadP23354 | RefOG070.0 |
Bralan_BL04167_evm4 | RefOG070.0 |
Tetnig_ENSTNIP00000005847 | RefOG070.0 |
Drer_ENSDARP00000132042 | RefOG070.0 |
Tetnig_ENSTNIP00000000291 | RefOG070.0 |
Galgal_ENSGALP00000067739 | RefOG070.0 |
Mmus_ENSMUSP00000112110 | RefOG070.0 |
Ratnor_ENSRNOP00000073744 | RefOG070.0 |
Canfam_ENSCAFP00000064761 | RefOG070.0 |
Hsap_ENSP00000369984 | RefOG070.0 |
Pantro_ENSPTRP00000088885 | RefOG070.0 |
Drer_ENSDARP00000131327 | RefOG070.0 |
Tetnig_ENSTNIP00000022824 | RefOG070.0 |
Tetnig_ENSTNIP00000000138 | RefOG070.0 |
Galgal_ENSGALP00000033098 | RefOG070.0 |
Hsap_ENSP00000264606 | RefOG070.0 |
Pantro_ENSPTRP00000073000 | RefOG070.0 |
Mmus_ENSMUSP00000095249 | RefOG070.0 |
Ratnor_ENSRNOP00000027622 | RefOG070.0 |
Drer_ENSDARP00000100072 | RefOG070.0 |
Tetnig_ENSTNIP00000017010 | RefOG070.0 |
Mondom_ENSMODP00000032765 | RefOG070.0 |
Hsap_ENSP00000225983 | RefOG070.0 |
Canfam_ENSCAFP00000044564 | RefOG070.0 |
Mmus_ENSMUSP00000008999 | RefOG070.0 |
Ratnor_ENSRNOP00000052059 | RefOG070.0 |
Galgal_ENSGALP00000060489 | RefOG070.0 |
Mondom_ENSMODP00000056336 | RefOG070.0 |
Mmus_ENSMUSP00000148029 | RefOG070.0 |
Ratnor_ENSRNOP00000005521 | RefOG070.0 |
Canfam_ENSCAFP00000063172 | RefOG070.0 |
Hsap_ENSP00000408617 | RefOG070.0 |
Pantro_ENSPTRP00000032389 | RefOG070.0 |
Cint_ENSCINP00000027979 | RefOG070.6 |
Cint_ENSCINP00000010872 | RefOG070.3 |
Mlei_ML004419a-PA | RefOG070.3 |
Nvec_EDO26214 | RefOG070.3 |
Nvec_EDO32559 | RefOG070.3 |
Tadh_TriadP25928 | RefOG070.3 |
Sman_Smp_091990 | RefOG070.3 |
Bralan_BL11384_evm3 | RefOG070.3 |
Drer_ENSDARP00000017858 | RefOG070.3 |
Tetnig_ENSTNIP00000008499 | RefOG070.3 |
Galgal_ENSGALP00000007696 | RefOG070.3 |
Mondom_ENSMODP00000036209 | RefOG070.3 |
Canfam_ENSCAFP00000040790 | RefOG070.3 |
Mmus_ENSMUSP00000085226 | RefOG070.3 |
Ratnor_ENSRNOP00000004224 | RefOG070.3 |
Pantro_ENSPTRP00000082283 | RefOG070.3 |
Hsap_ENSP00000362669 | RefOG070.3 |
Hsap_ENSP00000497072 | RefOG070.3 |
Tadh_TriadP26667 | RefOG070.7 |
Cele_WBGene00001835.1 | RefOG070.4 |
Dmel_FBpp0078414 | RefOG070.4 |
Sman_Smp_093280 | RefOG070.4 |
Nvec_EDO39062 | RefOG070.4 |
Mlei_ML003214a-PA | RefOG070.4 |
Tadh_TriadP57882 | RefOG070.4 |
Cint_ENSCINP00000004801 | RefOG070.4 |
Bralan_BL14536_evm4 | RefOG070.4 |
Tetnig_ENSTNIP00000013021 | RefOG070.4 |
Drer_ENSDARP00000054625 | RefOG070.4 |
Drer_ENSDARP00000135487 | RefOG070.4 |
Galgal_ENSGALP00000004141 | RefOG070.4 |
Mondom_ENSMODP00000013319 | RefOG070.4 |
Mmus_ENSMUSP00000037981 | RefOG070.4 |
Ratnor_ENSRNOP00000071791 | RefOG070.4 |
Hsap_ENSP00000302967 | RefOG070.4 |
Canfam_ENSCAFP00000030390 | RefOG070.4 |
Pantro_ENSPTRP00000029652 | RefOG070.4 |
Cele_WBGene00001834.1 | RefOG070.5 |
Cele_WBGene00001836.1 | RefOG070.5 |
Sman_Smp_005210 | RefOG070.5 |
Dmel_FBpp0073173 | RefOG070.5 |
Mlei_ML00527a-PA | RefOG070.5 |
Cint_ENSCINP00000005410 | RefOG070.5 |
Tadh_TriadP64072 | RefOG070.5 |
Nvec_EDO49920 | RefOG070.5 |
Mondom_ENSMODP00000022264 | RefOG070.5 |
Galgal_ENSGALP00000057637 | RefOG070.5 |
Canfam_ENSCAFP00000042408 | RefOG070.5 |
Ratnor_ENSRNOP00000000742 | RefOG070.5 |
Pantro_ENSPTRP00000049696 | RefOG070.5 |
Hsap_ENSP00000430432 | RefOG070.5 |
Mmus_ENSMUSP00000019911 | RefOG070.5 |
Tetnig_ENSTNIP00000007837 | RefOG070.5 |
Drer_ENSDARP00000155165 | RefOG070.5 |
Tetnig_ENSTNIP00000020022 | RefOG070.5 |
Tetnig_ENSTNIP00000012136 | RefOG070.5 |
Galgal_ENSGALP00000059784 | RefOG070.5 |
Mondom_ENSMODP00000021783 | RefOG070.5 |
Ratnor_ENSRNOP00000012854 | RefOG070.5 |
Ratnor_ENSRNOP00000018423 | RefOG070.5 |
Mmus_ENSMUSP00000099657 | RefOG070.5 |
Pantro_ENSPTRP00000075891 | RefOG070.5 |
Hsap_ENSP00000362649 | RefOG070.5 |
Canfam_ENSCAFP00000036819 | RefOG070.5 |
Pantro_ENSPTRP00000067224 | RefOG070.5 |
0.416667