Galgal_ENSGALP00000073898 | RefOG047.2 |
Tetnig_ENSTNIP00000016496 | RefOG047.0 |
Drer_ENSDARP00000147224 | RefOG047.0 |
Drer_ENSDARP00000087660 | RefOG047.0 |
Drer_ENSDARP00000152910 | RefOG047.0 |
Galgal_ENSGALP00000024750 | RefOG047.3 |
Bralan_BL18581_cuf6 | NA |
Bralan_BL08150_evm6 | NA |
Sman_Smp_002080 | NA |
Nvec_EDO45117 | NA |
Nvec_EDO37850 | NA |
Nvec_EDO37825 | NA |
Cint_ENSCINP00000016717 | RefOG047.1 |
Dmel_FBpp0088926 | RefOG047.1 |
Tadh_TriadP64181 | NA |
Nvec_EDO26870 | NA |
Mlei_ML018020a-PA | NA |
Cele_WBGene00008205.1 | RefOG047.1 |
Cele_WBGene00015540.1 | RefOG047.1 |
Cele_WBGene00015538.1 | RefOG047.1 |
Cele_WBGene00006416.1 | RefOG047.1 |
Tetnig_ENSTNIP00000014805 | RefOG047.1 |
Tetnig_ENSTNIP00000011512 | RefOG047.1 |
Drer_ENSDARP00000057868 | RefOG047.1 |
Drer_ENSDARP00000139220 | RefOG047.1 |
Galgal_ENSGALP00000003899 | RefOG047.1 |
Mondom_ENSMODP00000001856 | RefOG047.1 |
Canfam_ENSCAFP00000062817 | RefOG047.1 |
Hsap_ENSP00000361287 | RefOG047.1 |
Pantro_ENSPTRP00000004677 | RefOG047.1 |
Mmus_ENSMUSP00000153488 | RefOG047.1 |
Ratnor_ENSRNOP00000072105 | RefOG047.1 |
Tetnig_ENSTNIP00000011970 | RefOG047.1 |
Drer_ENSDARP00000105782 | RefOG047.1 |
Tetnig_ENSTNIP00000018925 | RefOG047.1 |
Drer_ENSDARP00000053962 | RefOG047.1 |
Galgal_ENSGALP00000069156 | RefOG047.1 |
Ratnor_ENSRNOP00000018170 | RefOG047.1 |
Mmus_ENSMUSP00000087118 | RefOG047.1 |
Mmus_ENSMUSP00000142136 | RefOG047.1 |
Hsap_ENSP00000303147 | RefOG047.1 |
Pantro_ENSPTRP00000020815 | RefOG047.1 |
Galgal_ENSGALP00000071065 | RefOG047.1 |
Canfam_ENSCAFP00000011509 | RefOG047.1 |
Mondom_ENSMODP00000019473 | RefOG047.1 |
0.416667