```
Aque Aqu2.1.33739 001 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                       Emue_Em0005g129a.t1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                            Emue_Em0005g166a.t1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                         _=Emue_Em0005g141a.t1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                       86 Emue_Em0005g172a.t1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                    Exapal_XM_021050458.1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                       Spis_XP_022788197_1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                 100
                        •Adig_XM_015906425.1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
        93
                            Spur_XM_030978077.1 | zf-C2H2.HG11.0:KAT7 | KAT7
                                  Skow XM 006812112.1 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                                       Scil_scict005505.2 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                     Bralan_BLAG04000121_5 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                        Ctel gnl WGS AMQN CAPTEDRAFT mRNA228212 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                       Cgig_XM_034473766.1 | zf-C2H2.HG11.0:KAT7 | KAT7
                93
                                    Tcas_TC011945_001 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                         •Dmel_FBtr0079443 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                          Cint ENSCINT00000006517 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                  •Drer_ENSDART00000004896 | zf-C2H2.HG11.0:KAT7 | NA |
                                     •Drer ENSDART00000148239 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                             100
                                       Hsap_ENST00000259021 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                                   100 Mmus_ENSMUST00000092766 | zf-C2H2.HG11.0:KAT7 | KAT7 |
                      Nemnom g9993.t1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                      Cint_ENSCINT00000015965 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                            •Dgig_XM_028550421.1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18
     48
                            Adig_XM_015901157.1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                              •Nvec_v1g242817 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                            Exapal_XM_021038120.2 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18
                               Spis XP_022799853_1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                           100 -Adig_XM_015899404.1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                     Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA130298 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | NA |
                                           Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA89486 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | NA |
                                 Ctel gnl WGS AMQN CAPTEDRAFT mRNA179381 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                              •Cgig_XM_020071949.2 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                               •Tcas TC034259 005 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                                     •Dmel_FBtr0310495 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                               Spur_XM_030982807.1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                                Bralan_BLAG02000890_1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                                 Skow_NM_001168116.1 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1/MYT1L/ST18 |
                                    Mmus_ENSMUST00000049784 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1L |
                                 _<sup>100</sup> Hsap_ENST00000644820 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1L |
                                    •Drer_ENSDART00000153150 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1L |
                                       •Drer ENSDART00000187701 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1L |
                           100
                                    <sup>1</sup>■Hsap_ENST00000276480 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | ST18 |
                                       Mmus_ENSMUST00000081125 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1 |
                                   <sup>100</sup> Hsap_ENST00000536311 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1 |
                                       ■Drer ENSDART00000165427 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1
                                    <sup>100</sup> ■Drer_ENSDART00000135771 | zf-C2H2.HG11.1:MYT1/MYT1L/ST18 | MYT1 |
                                Dgig_XM_028557502.1 | zf-C2H2.HG11.2:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA
                             100 Dgig_XM_028563212.1 | zf-C2H2.HG11.2:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA
                                         Morvir_scaffold13.g148.t1 | zf-C2H2.HG11.2:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                                      Tetwil_g15266.t1 | zf-C2H2.HG11.2:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                         Tetwil_g103.t1 | zf-C2H2.HG11.2:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                           _■Tetwil_g21703.t2 | zf-C2H2.HG11.2:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                         60 Tetwil_g16838.t1 | zf-C2H2.HG11.2:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                    Gracom_Gcom_scaffold12139_01 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                         Skow_XM_006824895.1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                   Spis_XP_022786543_1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                          Dgig_XM_028541309.1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                      Exapal XM 021037692.2 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                                          Nemnom_g9992.t1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                                Hvul_g18961_1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                           Morvir_scaffold144.g16.t1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                              Nemnom g8449.t1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                                           Nemnom g8569.t1 | zf-C2H2.HG11.3:like:AL031681.2/KAT7/L3MBTL1/L3MBTL4/MYT1/MYT1L/ST18 | NA |
                           Skow_XM_006820496.1 | zf-C2H2.HG11.6:like:AL031681.2/L3MBTL1/L3MBTL4 | NA |
                          Mmus_ENSMUST00000035751 | zf-C2H2.HG11.4:AL031681.2/L3MBTL1 | AL031681.2/L3MBTL1 |
                                  •Drer ENSDART00000132604 | zf-C2H2.HG11.4:AL031681.2/L3MBTL1 | AL031681.2/L3MBTL1 |
                               Hsap ENST00000418998 | zf-C2H2.HG11.4:AL031681.2/L3MBTL1 | AL031681.2/L3MBTL1 |
         65
                            100 Hsap_ENST00000657241 | zf-C2H2.HG11.4:AL031681.2/L3MBTL1 | AL031681.2/L3MBTL1 |
                               Cint_ENSCINT00000008413 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                   Spur_XM_030982813.1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                          Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA161824 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                          Cgig XM 034458785.1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                              90
                                         Bralan_BLAG02000892_3 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                        Skow XM 006811046.1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                  100
                         81
                                     --Hsap_ENST00000400105 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                       -Mmus_ENSMUST00000093007 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                       •Drer_ENSDART00000155311 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                             •Drer_ENSDART00000110260 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                                  Tcas TC001922 001 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                      Hvul_g19303_1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                            100
                                     Morvir_scaffold288.g16.t1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                     Nemnom_g5800.t1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                          98
                                   •Dgig XM 028546543.1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                       Spis XP 022788944 1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                       <sub>-</sub>-Adig_XM_015912128.1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                    100 - Spis XP 022804678 1 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                      Exapal_XM_021040415.2 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                          ■Nvec_v1g187589 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
                                     99 -Exapal XM 021040430.2 | zf-C2H2.HG11.5:L3MBTL4 | L3MBTL4 |
0.416667
```