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Bralan BLAG01000262 1 | MFS 1.HG17.0:like:MFSD4A/MFSD4B | NA |
                         Bralan BLAG14000670 1 | MFS 1.HG17.0:like:MFSD4A/MFSD4B | NA |
                            •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA186407 | MFS_1.HG17.0:like:MFSD4A/MFSD4B | NA |
                                           ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA185499 | MFS_1.HG17.0:like:MFSD4A/MFSD4B | NA |
  63
                                                      •Ctel gnl WGS AMQN CAPTEDRAFT mRNA206353 | MFS 1.HG17.0:like:MFSD4A/MFSD4B | NA |
                                            Ctel gnl WGS AMQN CAPTEDRAFT mRNA214193 | MFS 1.HG17.0:like:MFSD4A/MFSD4B | NA |
              100
                                              •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA216242 | MFS_1.HG17.0:like:MFSD4A/MFSD4B | NA |
                                   100
                                            •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA202354 | MFS_1.HG17.0:like:MFSD4A/MFSD4B | NA |
                                           -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA208843 | MFS_1.HG17.0:like:MFSD4A/MFSD4B | NA |
                                                -Ctel gnl WGS AMQN CAPTEDRAFT mRNA212130 | MFS 1.HG17.0:like:MFSD4A/MFSD4B | NA |
                                             Ctel gnl WGS AMQN CAPTEDRAFT mRNA215258 | MFS 1.HG17.0:like:MFSD4A/MFSD4B | NA |
                     •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA43971 | MFS_1.HG17.1:like:MFSD4A/MFSD4B | NA |
   88
                     •Cgig_XM_011429092.3 | MFS_1.HG17.1:like:MFSD4A/MFSD4B | NA |
                       ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA222889 | MFS_1.HG17.4:like:MFSD4A/MFSD4B | NA |
58
                                           *Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA214144 | MFS_1.HG17.5:like:MFSD4A/MFSD4B | NA |
                                  -Skow XM 006819307.1 | MFS 1.HG17.6:like:MFSD4A/MFSD4B | NA |
   91
                                Spur_XM_003727727.3 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                              <sup>100</sup> Spur XM 030988471.1 | MFS 1.HG17.2:MFSD4B | MFSD4B
                                      •Drer_ENSDART00000108955 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                 92
        99
                                    -Hsap_ENST00000672554 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                          100
                                       •Mmus_ENSMUST00000163705 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                                       Mmus_ENSMUST00000178045 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                                         -Mmus_ENSMUST00000170579 | MFS_1.HG17.2:MFSD4B | MFSD4B |
          50
                            •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA82130 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                  93
                                 •Cgig XM 011440618.3 | MFS 1.HG17.2:MFSD4B | MFSD4B |
                                                •Cgig_XM_011437408.3 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                                               •Cgig_XM_011429322.3 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                                                 •Cgig_XM_034454808.1 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                                               •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA205436 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                                                  Ctel gnl WGS AMQN CAPTEDRAFT mRNA215426 | MFS 1.HG17.2:MFSD4B | MFSD4B |
                             100
                                                        •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA199774 | MFS_1.HG17.2:MFSD4B | MFSD4B |
                             •Cint_ENSCINT00000028900 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                      •Gracom_Gcom_scaffold18611_01 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                       •Scil_scict014861.1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                   49
                                   •Plajan_Pjan_scaffold15642_01 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                Plajan_Pjan_scaffold15640_01 | MFS_1.HG17.3:MFSD4A | MFSD4A
                                                 -Emue Em0001g128a.t1 | MFS 1.HG17.3:MFSD4A | MFSD4A |
             40
                                              100 Emue Em0001g127a.t1 | MFS 1.HG17.3:MFSD4A | MFSD4A |
                                             •Aque_Aqu2.1.31952_001 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                         100
                                             -Tetwil_g8238.t1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                         •Tetwil_g8231.t1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                 100
                                                Tetwil_g8234.t1 | MFS_1.HG17.3:MFSD4A | MFSD4A
                                              1001 Tetwil g8233.t1 | MFS_1.HG17.3:MFSD4A | MFSD4A
               45
                                               •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA83377 | MFS_1.HG17.3:MFSD4A | NA |
                                                •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA211226 | MFS_1.HG17.3:MFSD4A | NA |
                                        •Dgig_XM_028544016.1 | MFS_1.HG17.3:MFSD4A | NA |
                                        •Spis_XP_022792506_1 | MFS_1.HG17.3:MFSD4A | NA |
                      96
                                           •Nvec_v1g33187 | MFS_1.HG17.3:MFSD4A | NA |
                           100
                                          Exapal_XM_021042974.2 | MFS_1.HG17.3:MFSD4A | NA |
                               100
                                        <sup>100</sup>-Exapal_XM_021048570.2 | MFS_1.HG17.3:MFSD4A | NA |
                                              Exapal_XM_021048580.2 | MFS_1.HG17.3:MFSD4A | NA |
                                     100
                                            <sup>1001</sup>Exapal XM 021042973.2 | MFS 1.HG17.3:MFSD4A | NA |
                                               Exapal XM 028660123.1 | MFS 1.HG17.3:MFSD4A | NA |
                                             <sup>100</sup> Exapal_XM_021048572.2 | MFS_1.HG17.3:MFSD4A | NA |
                                              •Bralan_BLAG05001352_1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                   73
                                                    •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA32514 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                  98
                                                Spur_XM_030972813.1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                 •Skow XM 006812389.1 | MFS 1.HG17.3:MFSD4A | MFSD4A |
                               97
                                                     -Hsap_ENST00000367147 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                 <sup>100</sup>└-Mmus_ENSMUST00000112370 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                         •Drer ENSDART00000010270 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                         •Drer_ENSDART00000029157 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                 •Exapal_XM_021045212.2 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                          100
                                                •Spis_XP_022791487_1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                      100
                                                         -Hvul_g30459_1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                           -Nemnom_g16926.t1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                  Spis XP 022791496 1 | MFS 1.HG17.3:MFSD4A | MFSD4A |
                               95
                                                   -Nvec_v1g201686 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                               <sup>100</sup> -Exapal_XM_021045203.2 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                  •Dgig_XM_028552141.1 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                       •Exapal_XM_021045192.2 | MFS_1.HG17.3:MFSD4A | MFSD4A |
                                                      •Nvec v1g92483 | MFS 1.HG17.3:MFSD4A | MFSD4A |
0.416667
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