```
Auraur scaffold1371.g11.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
  Morvir_scaffold77.g75.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
      Hydvul_Sc4wPfr_804.g24111.t2 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
   100 Hydvul Sc4wPfr 804.g24111.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
Helrob HelroP128125 | SAM.HG4.0:PHC1/PHC2/PHC3 | islike:PHC1/PHC2/PHC3 |
              Emue 0008g976a | SAM.HG4.0:PHC1/PHC2/PHC3 | islike:PHC1/PHC2/PHC3 |
             Aque_Aqu2.1.36261_001 | SAM.HG4.0:PHC1/PHC2/PHC3 | islike:PHC1/PHC2/PHC3 |
     Helrob_HelroP108682 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
             Cint ENSCINP00000000976.3 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                    Nvec_NVE3274 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                   Exapal XP 020903333.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
               -Spis_XP_022788400_1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                  _Adig_XP_015762168.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3
                <sup>100</sup> •Adig_XP_015762166.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
  Ctel_CapteP138021 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
         Spur XP 030843112.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
            Apla XP 022110217.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
              Apla XP 022110214.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
            60 Apla_XP_022110216.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
           Ptyfla_pfl_40v0_9_20150316_1g11404.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
           -Skow XP 006819909.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                                 Aque_Aqu2.1.30297_001 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3
                              <sup>100</sup> Aque_Aqu2.1.04698_001 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3
                                    --Tetwil_Twilhelma_g18192.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                                100 Tetwil Twilhelma g21837.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                 Phoaus g14735.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3
                           -Dmel_FBpp0070416 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                        <sup>100</sup><sup>-</sup>•Dmel_FBpp0070417 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                               •Drer_ENSDARP00000100341.2 | SAM.HG4.0:PHC1/PHC2/PHC3 | islike:PHC1/PHC2/PHC3 |
                           •Drer ENSDARP00000058495.6 | SAM.HG4.0:PHC1/PHC2/PHC3 | islike:PHC1/PHC2/PHC3 |
                           Hsap ENSP00000420294.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC3 |
                            ---Hsap_ENSP00000437659.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1 |
                              Lgig_LotgiP116623 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1 |
                         •Hsap_ENSP00000257118.5 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC2
                             •Drer ENSDARP00000018055.9 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC2 |
                            •Drer ENSDARP00000073668.4 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC2 |
                         100 -Drer ENSDARP00000130027.2 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC2 |
               Bralan BL09892 cuf13 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                -Lpol_XP_022253629.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                 --Cscu_XP_023231831.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                 Lpol XP 013779753.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
               <sup>100</sup> Lpol_XP_022247581.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                    Lpol_XP_013771947.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                   Lpol_XP_022238631.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                    _PLpol_XP_022238633.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                  <sup>95</sup> Lpol XP 022238632.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                    Notgen_g1680.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                <sup>100</sup> Notgen g1682.t1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                      Cgig_EKC18951 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                   100 - Cgig EKC20704 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3
                       Dpul_EFX87557 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                       Tcas_XP_015833722.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                         Tcas_XP_015833720.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
                       70 Tcas_XP_015833719.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3
                           Tcas XP 015833721.1 | SAM.HG4.0:PHC1/PHC2/PHC3 | PHC1/PHC2/PHC3 |
      Diprot Gene.16336 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD1/SAMD13/SAMD7/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD1/SAMD13/SAMD7/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD1/SAMD13/SAMD7/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD1/SAMD13/SAMD7/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD13/SAMD7/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML1/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML1/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML1/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML1/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML1/SCML4/SFMBT1/SFMBT2 | L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML1/SCML4/SFMBT1/SFMBT2 | L3MBTL4/SAMD1/SAMD13/SAMD13/SCMH1/SCML1/SCML4/SFMBT2 | L3MBTL4/SAMD1/SAMD13/SAMD13/SCMH1/SCML1/SCML4/SFMBT2 | L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT2 | L3MBTL4/SAMD1/SAMD13/SAMD13/SCMH1/SCML4/SFMBT2 | L3MBTL4/SAMD1/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAMD13/SAM
             Ramvar_GAU97880.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 |
               Skow_XP_002737526.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                         -Apla XP 022104195.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                          _■Spur XP 003730022.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                        <sup>100</sup> ■Spur XP 011666256.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                        ana_g1080.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                                Lcom_lcpid71266 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                               ■Scil_scpid55889 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                        -Adig_XP_015759912.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                        -Spis_XP_022800980_1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                       ■Nvec_NVE95 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                          Exapal_XP_020916618.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                 ■Tadh_P17972 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                        -■Notgen_g34581.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                          ■Notgen_g34582.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                   Hsap_ENSP00000334668.3 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD7 |
                          Hsap_ENSP00000482138.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11 |
                            Drer_ENSDARP00000112558.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11 |
                        100<sup>1</sup>-Drer_ENSDARP00000145638.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11
                   ■Ctel CapteP226340 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                     ■Phoaus_g9698.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                           Ramvar_GAU98715.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                           -■Hduj_OQV17149.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                        -Dmel_FBpp0079750 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                   Tcas_XP_008197355.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                      Tcas_XP_015838503.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                         _Tcas_XP_008197354.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                       <sup>100</sup> ■Tcas_XP_015838502.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                       Cscu_XP_023226688.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                         Lpol_XP_022245021.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                        --Lpol_XP_013784168.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                         .■Lpol_XP_013779966.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7 |
                       100 - Lpol_XP_022247775.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD11/SAMD7
                 ■Tadh_P57390 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD13/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2
                   ■Morvir_scaffold128.g20.t2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2
                   Sman_Smp_175040.1_pep | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 |
                  Emul_CDS37817.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 |
                       ■Emul_CDS37816.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML4/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML4/SFMBT3/SCML4/SFMBT3/L3MBT2/SAMD13/SCMH1/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFMBT3/SCML4/SFM
                           -Adig_XP_015775087.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2
                           ■Spis_XP_022779947_1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 |
                -Aque_Aqu2.1.07575_001 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 |
                       Aque_Aqu2.1.07574_001 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 |
                   400 Aque_Aqu2.1.23364_001 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 |
                        ■Tetwil_Twilhelma_g19473.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 |
                              ■Emue_0023g421a | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 |
                          -■Emue_0023g423a | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4/SAMD1/SAMD13/SCMH1/SCML1/SCML4/SFMBT1/SFMBT2 |
                          Cgig_EKC24800 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                 ■Emul_CDS37271.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                               Sman_Smp_068230.1_pep | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                 -Morvir_scaffold182.g34.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                        •Ngru_EFC41874 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                 Exapal_XP_020899614.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                   -Nvec_NVE2604 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                  Spis XP 022777959 1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                  Adig_XP_015774759.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                    _•Adig_XP_015774751.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13 |
                                  97 Adig_XP_015774773.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SAMD1/SAMD13
                                Clyhem_TCONS_00000254-protein | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                             Lcom_lcpid44676 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                              -Scil scpid92276 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                               Pygbif_GCRY01003644.1.p1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                -■Phakir_PHAKIR_01310.mRNA.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                           Sman_Smp_137110.1_pep | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                    ■Emul_CDS35731.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                             ■Tadh_P58043 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                        Helrob_HelroP97993 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                    Dpul EFX84513 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                 Hydvul_Sc4wPfr_730.g15321.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                            ■Morvir_scaffold139.g44.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13
                                             -Auraur_scaffold32.g23.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                            -Adig_XP_015775377.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                     100 - Spis_XP_022797047_1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                            Nvec_NVE17345 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                         100 ♣Nvec_NVE17344 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                            .■Exapal_XP_020897472.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                         100 ■Exapal_XP_020897473.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                -Drer_ENSDARP00000086071.4 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD13 |
                                          100 Hsap_ENSP00000359705.3 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD13 |
                                                       -■Drer_ENSDARP00000131835.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1 |
                                                    Drer_ENSDARP00000114307.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1 |
                                                    Hsap_ENSP00000431971.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1 |
                                                 100 Hsap_ENSP00000500096.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1 |
                                            Cscu_XP_023235774.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                            Linana g17186.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                 •Cgig_EKC30194 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                <u>-</u>Linana_g26805.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                             Lgig_LotgiP120412 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                        -Lgig_LotgiP120410 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                       Lpol_XP_022251025.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                       ---Lpol_XP_022245554.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                      --Lpol_XP_013784204.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                       ■Cscu_XP_023227536.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                      Cscu_XP_023227518.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                   <sup>100</sup> ■Cscu_XP_023227519.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                  ■Phoaus_g15193.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                  Ctel_CapteP167868 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                          ■Cint ENSCINP00000029150.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                       Notgen g10399.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                   Skow_XP_002730583.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                       Bralan_BL11114_evm0 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                          . ■Apla_XP_022087897.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                                                      100 Apla_XP_022087899.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13
                                                          Spur_XP_030852226.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13
                                                       100™Spur_XP_030852562.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SAMD1/SAMD13 |
                              Ramvar_GAU98808.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 |
                          ■Tadh_P58096 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                         Cint_ENSCINP00000008413.3 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                            -■Morvir_scaffold288.g16.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                      -Hydvul_Sc4wPfr_347.g19301.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                              -Clyhem_TCONS_00048927-protein | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                           -Spis_XP_022804678_1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                            ■Spis_XP_022788944_1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                               -■Exapal_XP_020896089.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                -■Nvec_NVE714 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                            -Nvec_NVE716 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                _■Exapal_XP_020906324.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                <sup>1</sup>■Exapal_XP_020896074.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                         ■Tcas_XP_008199297.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                           Dpul_EFX86265 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                           Drer_ENSDARP00000123463.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3/L3MBTL4 |
                              Hsap_ENSP00000382976.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL4 |
                                 ■Drer_ENSDARP00000101801.3 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:L3MBTL3 |
                    100
                                --Hsap_ENSP00000354526.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3 |
                                -■Drer ENSDARP00000127490.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3 |
                            -■Apla_XP_022091091.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                       -■Skow_XP_006811109.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                          -Ptyfla pfl 40v0 9 20150316 1g21825.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                             -Lpol XP 022245005.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                 Cscu_XP_023234831.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                -Phoaus_g14882.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                              Linana_g17562.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                               -Notgen g4287.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                     •Cgig EKC39929 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4 |
                                     -Lgig_LotgiP216594 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4
                                     ■Ctel_CapteP161824 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | L3MBTL3/L3MBTL4
                                  ■Tetwil_Twilhelma_g26894.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                   ■Emue 0023g727a | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                               -Lcom_lcpid76155 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                -Scil_scpid23244 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                               Lcom_lcpid11952 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                                Scil scpid40697 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                             Scil scpid40059 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                                _Scil_scpid41679 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                              100 Scil_scpid40985 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                              Cint_ENSCINP00000025543.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                    ■Emul_CDS37224.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                   Sman_Smp_006250.2_pep | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                   ■Morvir_scaffold21.g96.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                    Auraur_scaffold310.g3.t2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                    -Clyhem_TCONS_00048363-protein | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                                        ■Hydvul_Sc4wPfr_582.g7498.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                              Spis XP 022788860 1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                  ■Exapal_XP_020912288.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                  Nvec_NVE23450 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                      ■Nvec_NVE23445 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                   100 Nvec_NVE23449 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                              Hsap_ENSP00000251900.4 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCML1/SCML2 |
                                                     Hsap ENSP00000369380.3 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCML1/SCML2 |
                                Drer_ENSDARP00000016700.7 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SCMH1/SCML4 |
                                    ■Drer_ENSDARP00000135816.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1 |
                              ■Drer_ENSDARP00000115766.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCML4 |
                                       —■Hsap_ENSP00000358016.3 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCML4 |
                              -Lpol_XP_013775899.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                              -■Lpol_XP_022247720.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                   Dpul_EFX76808 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                     .
-■Tcas_XP_966529.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                       -Dmel FBpp0081580 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                                    Spur_XP_030850317.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                   --Apia_XP_022081927.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                _■Skow_XP_006817556.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                  --Ptyfla_pfl_40v0_9_20150316_1g22287.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                                 Linana g11770.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                      -Cgig_EKC41694 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                      -Lgig LotgiP107440 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                        ■Notgen_g16582.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                         -■Ctel_CapteP120304 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                        -Phoaus_g3661.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4 |
                                           -Helrob_HelroP115837 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SCMH1/SCML1/SCML2/SCML4
                             Morvir_scaffold251.g10.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                 -Clyhem_TCONS_00063933-protein | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                          100 - Hydvul_Sc4wPfr_178.g10641.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                            -■Nvec_NVE15798 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                              Exapal_XP_020915016.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2
                            100 Exapal XP 020915017.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                            ■Spis_XP_022781210_1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                              Adig XP 015751169.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2
                            <sup>100</sup> Adig_XP_015751171.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                            -Spur XP 030836926.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                                 Apla XP 022084416.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                           Skow XP 006815581.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                                     -Ctel CapteP222398 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                  •Cgig EKC25175 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                  ■Linana g3576.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2
                                    -Phoaus g2441.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                    Tcas XP 967817.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                     -Dmel FBpp0288419 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                 -Lpol XP 013786572.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                               -■Cscu XP 023240089.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                             ■Gefoke MALSP 02774.mRNA.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                  ■Hydvul Sc4wPfr 719.g8329.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                                  -Clyhem TCONS 00034843-protein | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2
                                -Clyhem TCONS 00060732-protein | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                            .■Exapal_XP_020902826.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                         <sup>100</sup> ■Exapal_XP_028515490.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                            ─■Adig XP 015752853.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                            Spis XP 022780251 1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | islike:SFMBT1/SFMBT2 |
                               Morvir scaffold240.g29.t2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                   ■Clyhem_TCONS_00005280-protein | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2
                                  -■Hydvul_Sc4wPfr_264.g27404.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                               -Spis XP 022796962 1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                -Adig XP 015761803.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                 -Nvec NVE8986 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                  .■Exapal_XP_028513995.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                               <sup>100</sup> ■Exapal_XP_020897274.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                 Lpol XP 013794742.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                  ■Cgig EKC24670 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                         Helrob HelroP66618 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2
                                -Notgen_g5116.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                    Linana_g19034.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2
                                 Linana g19033.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2
                                        ■Ctel CapteP148410 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                         ■Phoaus g5105.t1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                               Skow XP 006811735.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                   Apla XP 022104927.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2
                                   Spur XP 030827835.1 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1/SFMBT2
                                -■Hsap ENSP00000355109.4 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT2 |
                                 •Drer ENSDARP00000143791.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT2 |
                                -- Hsap ENSP00000378235.2 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD1/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1 |
                                  PDrer ENSDARP00000041724.8 | SAM.HG4.1:L3MBTL3/L3MBTL4/SAMD1/SAMD7/SAMD11/SAMD13/SCMH1/SCML1/SCML2/SCML4/SFMBT1/SFMBT2 | SFMBT1 |
                               100 Indicate: 10
```