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Lcom lcpid236489 | MBT.HG1.0:like:PHF20L1:likeclu:1 | NA |
                         Ehux EOD31316 | MBT.HG1.0:like:PHF20L1:likeclu:1 | NA |
                           Gthe_EKX39041 | MBT.HG1.0:like:PHF20L1:likeclu:1 | NA |
                      Retfil ETO14459 | MBT.HG1.0:like:PHF20L1:likeclu:1 | NA |
                              Spun KNC99570 | MBT.HG1.0:like:PHF20L1:likeclu:1 | NA |
                      -Ptet CAK70058 | MBT.HG1.0:like:PHF20L1:likeclu:1 | NA |
                  100 Ptet CAK80324 | MBT.HG1.0:like:PHF20L1:likeclu:1 | NA |
                        Bnat_JGI_V11_86001 | MBT.HG1.3:like:PHF20L1:likeclu:1 | NA |
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                           Tcas XP 972506.1 | MBT.HG1.1:PHF20L1 | PHF20L1 |
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                               Lcom | lcpid13340 | MBT.HG1.1:PHF20L1 | PHF20L1 |
                           Helrob HelroP87257 | MBT.HG1.1:PHF20L1 | PHF20L1 |
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                                    100 Notgen_g39774.t1 | MBT.HG1.1:PHF20L1 | PHF20L1 |
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                    Gonavo_comp59428_c0_seq1_6_ORF8_234 | MBT.HG1.5:like:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2:likeclu:2 | NA |
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48
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                             Bralan_BL09568_cuf5 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                  Cgig_EKC39929 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                    ■Phoaus_g14882.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                   Linana_g17562.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                   -■Notgen_g4287.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                       Lgig LotgiP216594 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                       -Ctel_CapteP161824 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                 Cscu_XP_023234831.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                  Cscu_XP_023229945.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                   -■Cscu_XP_023229950.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                   -Lpol_XP_022237043.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
                                    -Lpol_XP_022245005.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | AL031681.3/L3MBTL1/L3MBTL3/L3MBTL4 |
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                                    Mlei_ML0473_g8_t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | islike:SCMH1/SCML2 |
                                        -Lcom_lcpid93286 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | islike:SCMH1/SCML2 |
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                                     Gonavo_comp61571_c0_seq2_6_ORF26_958 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | islike:SCMH1/SCML2 |
                                   Bnat_JGI_V11_72235 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | islike:SCMH1/SCML2 |
                             Lcom_lcpid76155 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                            Scil scpid23244 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
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                            Lcom | lcpid11952 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                             Scil_scpid40697 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                          Scil_scpid40059 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2
                             _ Scil_scpid41679 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
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                     •Tadh_P24969 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                          --Spis_XP_022788860_1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                             -Adig_XP_015756083.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2
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                                ■Morvir_scaffold21.g96.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                                 -Auraur_scaffold310.g3.t2 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                                -Clyhem_TCONS_00048363-protein | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
                                      Hydvul Sc4wPfr 582.g7498.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
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                               -Cscu_XP_023210195.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SCMH1/SCML2 |
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                                Hydvul_Sc4wPfr_719.g8329.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | islike:L3MBTL2/MBTD1/SFMBT1/SFMBT2 |
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                           Exapal_XP_020902826.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | islike:L3MBTL2/MBTD1/SFMBT1/SFMBT2
                           Exapal_XP_028515490.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | islike:L3MBTL2/MBTD1/SFMBT1/SFMBT2 |
                             ■Spis_XP_022796962_1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
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                             <sup>100</sup> Exapal_XP_020897274.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                             ■Morvir_scaffold240.g29.t2 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                               -Auraur_scaffold1294.g3.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
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                                       Sman Smp 074050.1 pep | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
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                                       ■Dpul EFX87706 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                               Skow XP 006811735.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
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                                   Spur_XP_030827835.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                               Bralan_BL04550_cuf21 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
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                                    --Drer_ENSDARP00000143791.2 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT2 |
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                                 100 Drer_ENSDARP00000154520.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1
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                                 Phoaus_g5105.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
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                                      -Ctel_CapteP148410 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
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                                 Linana_g19034.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                   ■Notgen_g5116.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2
                                       •Cgig_EKC24670 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                                  -Lgig_LotgiP123158 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                            99<sup>1</sup>-Lgig_LotgiP164102 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | SFMBT1/SFMBT2 |
                                 -Clyhem_TCONS_00063933-protein | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1 |
                           100 +Hydvul_Sc4wPfr_178.g10641.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1 |
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                                       Hydvul Sc4wPfr 178.g10645.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1 |
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                            100 Exapal_XP_020915016.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1 |
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                            100 Adig XP 015751169.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1 |
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                                     Apla XP 022084475.1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1 |
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                                      --Lgig LotgiP144696 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1 |
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                                    Linana g7133.t1 | MBT.HG1.2:AL031681.3/L3MBTL1/L3MBTL2/L3MBTL3/L3MBTL4/MBTD1/SCMH1/SCML2/SFMBT1/SFMBT2 | L3MBTL2/MBTD1
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