```
Ocar_g10302 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                -Adig_XM_015924122.1 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
    Horcal Hcv1.av93.c1.g1036.i2 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
    Horcal_Hcv1.av93.c1.g1036.i1 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
    1 Horcal Hcv1.av93.c1.g1036.i3 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
        Lampan Lpan C402619 01 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA
        ■Vmul Simscaffold5351 01 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
       Coemet_Cmet_scaffold10633_01 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
     Coemet_Cmet_scaffold10634_01 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
-Scil scict006816.1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
 ■Gracom Gcom C542520 01 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                              --Bralan_BLAG15101013_1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
   ■Tetwil g7485.t1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
       Emue Em0018g980a.t1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
    Emue Em0508g4a.t1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
             ■Horcal_Hcv1.av93.c11.g555.i1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA
          Horcal Hcv1.av93.c11.g555.i2 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA
                               Mlei_ML0866.g9.i1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                           100 MIEI_ML3427.g2.i1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                          Lampan Lpan 15754 01 RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... NA
                                       ■Vmul_Simscaffold16593_01 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                         Coemet_Cmet_scaffold380_01 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA
                                      Coemet_Cmet_scaffold381_01 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA
                                           Horcal_Hcv1.av93.c9.g92.i1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                           Bolinf_Binf_sb1258260 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                             Beraby Baby sb329256 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                          Bolinf Binf sb1230298 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                                Nmul_sbMO449155 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                             100 Mlei_ML0482.g5.i1 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                                Bolinf Binf sb1261489 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                                                  Bolinf_Binf_sb1261481 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA
                                                  Bolinf_Binf_sb1261490 | RRM.HG4.1:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
        Dmel_FBtr0085608 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
    ■Tcas TC009189 001 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
     -Cgig XM 020066571.2 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
     Skow_XM_002735552.2 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
  Bralan_BLAG04102199_2 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
      -Mmus ENSMUST00000172000 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
      -Cint ENSCINT00000017840 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
           Cint_ENSCINT00000032244 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
             ■Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA224043 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                     Plajan_Pjan_scaffold9435_01 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
            Cgig_XM_034467477.1 | RRM.HG4.2:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
            ■Emue Em0013g989a.t1 | RRM.HG4.13:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/S... | NA |
                     ■Gracom_Gcom_scaffold3987_01 | RRM.HG4.3:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                  100 Gracom_Gcom_scaffold3989_01 | RRM.HG4.3:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |
                     Tetwil_g10785.t2 | RRM.HG4.14:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                   Tetwil_g12196.t1 | RRM.HG4.6:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                          Aque Aqu2.1.40769 001 | RRM.HG4.6:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                  Emue Em0012g837a.t1 | RRM.HG4.6:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                       Dmel_FBtr0076835 | RRM.HG4.6:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                •Dmel_FBtr0070171 | RRM.HG4.6:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                     ■Dmel FBtr0070180 | RRM.HG4.6:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA
                                      -Dmel FBtr0112928 | RRM.HG4.6:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                             Scil scict014325.3 | RRM.HG4.9:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                          Gracom_Gcom_scaffold16569_01 | RRM.HG4.9:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA
                       Gracom_Gcom_scaffold16570_01 | RRM.HG4.9:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                               Ague Agu2.1.40898 001 | RRM.HG4.9:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                             Tetwil g6230.t1 | RRM.HG4.9:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                             Emue Em0018g1204a.t1 | RRM.HG4.9:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                             Emue Em0018g1217a.t1 | RRM.HG4.9:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                           Tcas TC000027 001 | RRM.HG4.10:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                          Gracom Gcom C557202 01 | RRM.HG4.10:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                         Scil scict007707.1 | RRM.HG4.10:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                     Cgig XM 034443213.1 | RRM.HG4.11:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                          .■Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA224539 | RRM.HG4.11:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                       Too Ictel gnl WGS AMQN CAPTEDRAFT_mRNA105051 | RRM.HG4.11:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                       Spur XM 030975297.1 | RRM.HG4.11:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                       Spur XM 011666085.2 | RRM.HG4.11:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                   Spur XM 030975180.1 | RRM.HG4.11:like:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                         -Aque Aqu2.1.40932 001 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                         ■Tetwil g6228.t1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                          ■Emue Em0018g1042a.t1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                       Emue Em0728g2a.t1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                         -Bolinf Binf sb1189926 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                         Horcal Hcv1.av93.c1.g1449.i2 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                       Horcal Hcv1.av93.c1.g1449.i3 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                             _Horcal Hcv1.av93.c1.g1449.i1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                             Horcal Hcv1.av93.c1.g1450.i1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                           I■MIei ML2143.g18.i1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                         Bolinf Binf sb1210583 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                             Beraby_Baby_sb335068 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                          Beraby Baby sb335069 RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                            -Lampan Lpan 14971 01 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL1/HNRNPCL2/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                              _-Coemet Cmet C595569 01 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                            99 -Wmul_Simscaffold6525_01 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                           Spur XM 030975158.1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                               Bralan BLAG07001238 1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                          ■Nemnom g9924.t1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                          ■Morvir scaffold17.g58.t1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                        Dgig_XM_028536723.1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                       Dgig_XM_028562114.1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                           ■Nvec v1g118934 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                       -■Exapal XM 021042032.2 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                                     HVUL g6745 1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                       -■Skow_XM_006814095.1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                         Skow XM 002732846.2 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | NA |
                                        Skow XM 006821134.1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                                  Bralan BLAG07001377 1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                                -Bralan BLAG15000792 1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                                 ■Bralan_BLAG15000331_1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                                      -Cgig XM 034480005.1 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                                    -Ctel gnl WGS AMQN CAPTEDRAFT mRNA223984 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                                       -Dmei FBtr0300248 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL
                                                   100 Tacas TC032696 001 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL |
                                               Hsap_ENST00000517638 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | RALYL |•
                                            | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 1000 | 
                                              -Drer ENSDART00000083999 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | RALY |
                                                _=Hsap ENST00000246194 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | RALY |
                                                4-Mmus ENSMUST00000029120 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | RALY |
                                                ■Drer_ENSDART00000075905 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4 |
                                           Too Indicate the standard of the control of the con
                                                 •Hsap ENST00000420743 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC |
                                               90 Mmus ENSMUST00000111610 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPC |
                                                     J■Hsap ENST00000621994 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4
                                                  100 Hsap_ENST00000317869 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4
                                                     Hsap ENST00000323770 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4
                                                   100 Hsap ENST00000617807 | RRM.HG4.12:HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/RALY/RALYL | HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4
                         Plaian Pian C569254 01 | RRM.HG4.4:MTHFSD | MTHFSD |
                         ■Emue Em0004g143a.t1 | RRM.HG4.4:MTHFSD | MTHFSD |
                             Tetwil g11157.t2 | RRM.HG4.4:MTHFSD | MTHFSD |
                            Aque Aqu2.1.42191 001 | RRM.HG4.4:MTHFSD | MTHFSD |
                                       Hsap ENST00000360900 | RRM.HG4.4:MTHFSD | MTHFSD |
                           Lampan Lpan 16051 01 | RRM.HG4.4:MTHFSD | MTHFSD |
                           Horcal_Hcv1.av93.c6.g443.i1 | RRM.HG4.4:MTHFSD | MTHFSD
                        Horcal Hcv1.av93.c6.g443.i2 | RRM.HG4.4:MTHFSD | MTHFSD |
                  Hvul g8438 1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                Morvir scaffold316.g10.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                 Nemnom g10668.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
              100 Nemnom_g19525.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
               Nvec_v1g202761 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                --Exapal_XM_021041989.2 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                 --Adig_XM_015913740.1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                 --Spis XP 022785795 1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                    ■Morvir_scaffold97.g2.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                         Nemnom g10666.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                <sup>100</sup> Nemnom_g19526.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                    Aque_Aqu2.1.39784_001 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                          Tetwil g2522.t2 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                          -■Tetwil_g2521.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                          Emue_Em0015g3a.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                       <sup>100</sup> Emue_Em0015g94a.t1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                     Gracom_Gcom_scaffold19114_01 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                <sup>100</sup> Scil scict025504.1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                        PDgig_XM_028544594.1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                     <sup>100</sup> Dgig_XM_028562692.1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                        •Dgig_XM_028562690.1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                          ■Vmul_Simscaffold5854_01 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                                Horcal Hcv1.av93.c3.g502.i1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                             100 Horcal_Hcv1.av93.c3.g502.i2 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                                  --Lampan_Lpan_14116_01 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                                -■Mlei_ML1430.g26.i1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                                  Beraby_Baby_sb345656 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                                   Beraby Baby sb345654 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                                   Beraby Baby sb345655 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                                   Beraby Baby sb345657 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9
                                 Beraby Baby sb345658 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                   <sup>™</sup>HoiH23 PIH23 007912-RA | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                   _*Tadh_TriadT49822 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                100 TrH2_TrispH2_004059-RA | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1/SRSF9 |
                Plajan_Pjan_C534281_01 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | SRSF9 |
                              Cint ENSCINT00000000458 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | SRSF9 |
                           -Cint_ENSCINT00000015147 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | SRSF9 |
                           •Drer_ENSDART00000002985 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | SRSF9
                             <sup>100</sup> Hsap_ENST00000229390 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | SRSF9 |
                       Bralan BLAG01001187 1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1
                            --Tcas_TC015810_001 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                             Dmel_FBtr0083272 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                                             Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA221687 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                              _=Cgig_XM_034443058.1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                               └-Cgig XM 011428506.3 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                       -■Skow_XM_006815747.1 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                              Spur_XM_011684379.2 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                              Drer_ENSDART00000080413 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                           96 Drer_ENSDART00000143566 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2/SRSF1 |
                             -Hsap ENST00000578794 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | AC015813.2 |
                                Hsap_ENST00000584773 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | SRSF1 |
                             69 Mmus_ENSMUST00000107920 | RRM.HG4.5:AC015813.2/SRSF1/SRSF9 | SRSF1 |
                      Nemnom_g11668.t1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                 Hvul g30512 1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                       Morvir scaffold187.g17.t2 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                       Nvec v1g196990 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                         Gracom_Gcom_C540236_01 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA
                                     Scil_scict009780.1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                    Ocar g414 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                ■Tetwil g10378.t1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                       Hhon_g03139.t1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                         TrH2_TrispH2_000901-RA | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                            ■Tadh_TriadT52131 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                   Cgig XM 011449723.3 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                         Ctel gnl WGS AMQN CAPTEDRAFT mRNA222188 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                      •Cgig XM 011417916.3 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                Bralan_BLAG03000790_1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                    Skow_XM_006825691.1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                         ■Spur XM 011671095.2 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA
                                                 Spur XM 030989255.1 | RRM.HG4.7:like:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
               --Scil scict001348.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
           100 Gracom Gcom scaffold8846_01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                 Gracom_Gcom_scaffold15917_01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
             100 Scil scict005000.1 RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 AL031681.2/SRSF4/SRSF5/SRSF6
                     -Adig XM 015895573.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
               -■Nvec_v1g26678 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                     -■Exapal_XM_021038922.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                              ■Mlei_ML1207.g45.i1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                          --Lampan Lpan 11315 01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                Coemet_Cmet_scaffold17172_01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                     -Wmul_Simscaffold15185_01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                       •Vmul Simscaffold4335 01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                          Lampan Lpan 4457 01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                              Coemet Cmet scaffold17601 01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA
                           100 Coemet Cmet scaffold17602 01 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                              ■Horcal_Hcv1.av93.c7.g962.i1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                  --Bolinf_Binf_sb1210245 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                   Mlei ML3233.g1.i1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                    <sup>1</sup>──■Vmul_sbMO386877 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA
                                    Bolinf_Binf_sb1226764 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                    Beraby_Baby_sb337796 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                 <sup>100</sup> Beraby_Baby_sb337797 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA
                   Hhon_g06258.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                      _=Tadh TriadT23659 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                    85<sup>7</sup>=HoiH23_PIH23_008789-RA | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                          Tadh_TriadT63986 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                       TrH2 TrispH2 002430-RA | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                        _=Hhon_g08244.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                       <sup>99 1</sup>•HoiH23 PIH23 003229-RA | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                         Spis XP 022780165 1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                         └-•Adig XM 015922827.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                Dgig XM 028538386.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                ■Dgig_XM_028538102.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                             __•Nvec_v1g85456 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                 --Exapal_XM_021038396.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                   -Hvul_g26616_1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                   -Nemnom_g17829.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                  -■Nemnom g4143.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                    -Morvir scaffold45.g65.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                        Dgig_XM_028555522.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                   -■Nvec_v1g136664 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                   --Exapal XM 028659598.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                   Morvir scaffold283.g8.t2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                       Nemnom_g11109.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                Dgig_XM_028556812.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                 --Adig XM 015917690.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                               Spis XP 022796108 1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                              Nvec v1g230980 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                 Exapal_XM_021048368.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA
                              100 Exapal XM 021048369.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                       -Aque Aqu2.1.28194 001 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                        --Tetwil g3838.t2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                            Emue Em0019g630a.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                            Emue_Em0666g5a.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                 Emue_Em0666g4a.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                               _Emue Em0019g629a.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                            100 Emue Em0666g6a.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                          ■Morvir scaffold546.g4.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                Nemnom_g5160.t1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                                  -■Hvul_g29527_1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                              Dgig XM 028544787.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                                ■Spis XP 022780528 1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                    --Adig XM 015915032.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                  Nvec v1g108976 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                    Exapal_XM_021038274.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                    Nvec v1g199162 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                             Ocar g3065 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                Cint ENSCINT00000033636 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                Cint ENSCINT00000010229 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                  Cint_ENSCINT00000032767 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                  -Cgig XM 020066693.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                                         Cint ENSCINT00000014783 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                  -Cgig_XM_011449346.3 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                        -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA175470 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                                        -Ctel gnl WGS AMQN CAPTEDRAFT mRNA220292 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                                     Tcas TC016051 001 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                         -Dmel FBtr0308195 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6
                                      100 Tcas TC012868 001 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF5/SRSF6 |
                                   Spur_XM_793653.5 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF5 |
                                <sup>100</sup> Spur_XM_030985554.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF5 |
                                     -Drer ENSDART00000181046 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | NA |
                                       Drer ENSDART00000134564 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF5 |
                                          Hsap ENST00000394366 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF5 |
                                       100 Mmus ENSMUST00000110351 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF5 |
                                     Skow XM 002732217.2 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF6 |
                                    └─-Skow XM 006813345.1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF6 |
                                       -Bralan BLAG19000716 1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF6
                                   100 - Bralan BLAG07000724 1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF6 | Bralan BLAG07000724 1 | RRM.HG4.8:AL031681.2/SRSF4/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF4/SRSF6 | AL031681.2/SRSF4/SRSF6 |
                                           Drer_ENSDART00000165033 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF4 |
                                            Hsap_ENST00000373795 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF4 |
                                          100 Mmus ENSMUST00000053819 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | SRSF4 |
                                             Drer ENSDART00000002318 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF6
                                         Drer ENSDART00000103747 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF6 |
                                             Mmus_ENSMUST00000130411 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF6 |
                                               Hsap_ENST00000244020 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF6 |
                                             99 Hsap ENST00000657241 | RRM.HG4.8:AL031681.2/SRSF4/SRSF5/SRSF6 | AL031681.2/SRSF6 |
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Hvul_g31738_1 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA

Exapal XM 028663324.1 | RRM.HG4.0:like:AC015813.2/AL031681.2/HNRNPC/HNRNPCL1/HNRNPCL2/HNRNPCL3/HNRNPCL4/MTHFSD/RALY/RALYL/SR... | NA |