```
Bolinf Binf sb1266424 | Pkinase.HG30.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                        --Dgig XM 028547863.1 | Pkinase.HG30.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... I NA I
                    -Morvir scaffold135.α22.t1 | Pkinase.HG30.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                   ■Skow XM 006817643.1 | Pkinase.HG30.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                   ■Dmel FBtr0073457 | Pkinase.HG30.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                     ■Cgig XM 034455542.1 | Pkinase.HG30.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                   -Cgig XM 034454968.1 | Pkinase.HG30.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA ∣
                                             -Coemet Cmet scaffold21140 01 | Pkinase.HG30.3:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                            sbMO400014 | Pkinase.HG30.3:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                       PHorcal Hcv1.av93.c3.g722.i2 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                        Horcal Hcv1.av93.c3.g722.i4 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                            ■Horcal Hcv1.av93.c3.g722.i1 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                            Horcal Hcv1.av93.c3.g722.i3 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                     •Vmul sbMO402403 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                           _■Bolinf Binf sb1225676 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                          -Mlei ML1309.g2.i1 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                          -Bolinf Binf sb1190876 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                           -■Bolinf Binf sb1190877 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                       Beraby Baby sb346884 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                 Beraby Baby sb346886 | Pkinase.HG30.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                   Scil scict008034.1 | Pkinase.HG30.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                                  ■Plajan Pjan scaffold22749 01 | Pkinase.HG30.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                              ■Ocar g2033 | Pkinase.HG30.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                       ■Plajan Pjan scaffold6423 01 | Pkinase.HG30.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                     -Scil scict004662.1 | Pkinase.HG30.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                      -■Scil_scict008193.1 | Pkinase.HG30.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                        Scil scict003470.4 | Pkinase.HG30.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                        ■Gracom Gcom scaffold13431 01 | Pkinase.HG30.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                       Gracom Gcom scaffold13432 01 | Pkinase.HG30.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                             Horcal Hcv1.av93.c1.g1556.i1 | Pkinase.HG30.9:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                          ■Plajan Pjan scaffold19921 01 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                                                                                        -Spis_XP_022778049_1 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                                 ■Coemet Cmet C596055 01 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                -Spis XP 022781446 1 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                                                              Adia XM 015899978.1 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                      Adia XM 015903536.1 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                             ■Horcal Hcv1.av93.c10.g609.i1 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                                                                                                        100 Horcal Hcv1.av93.c10.g609.i2 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                       -■Lampan Lpan 2354 01 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                                     -■Coemet_Cmet_scaffold21350_01 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                                                                                                                          ■Vmul_SimC402758 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                                    ■Beraby_Baby_sb328616 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                                 _■Mlei ML2577.g3.i1 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                                                                                                     --Bolinf Binf sb1238854 | Pkinase.HG30.8:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                     Bolinf Binf sb1317718 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                            ■Tetwil α9984.t2 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                -■Ague Agu2.1.44044 001 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                       PAgue Agu2.1.05490 001 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                  note in the following section in the section of the following in the first of the following in the following in the first of the following in the first of the fi
                                                     Aque Aqu2.1.15260 001 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                              -Aque Aqu2.1.08383 001 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA I
                                                               Aque Aqu2.1.15982 001 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                            ■Tetwil g10267.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                                          -Tetwil g17789.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                           Tetwil g20640.t2 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                             _■Tetwil_g24829.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                        <sup>100</sup> <sup>1</sup>-Tetwil_g24830.t2 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                -■Emue Em0012g256a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                               Emue Em0012g259a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                    ■Emue Em0003g1633a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                     ■Emue Em0003g1682a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA
                                                                     ■Emue_Em0003g1638a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                          -Emue Em0003g1680a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                        -■Emue Em0016g553a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                ■Emue Em0023g549a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                                   -■Tetwil_g1950.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                                 Emue_Em0023g529a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |

*Emue_Em0023g531a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |

*Emue_Em0023g550a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
                                              100 = Emue Em0023g551a.t1 | Pkinase.HG30.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA |
        ■Tetwil g15473.t2 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EP
                                 -Aque Aqu2.1.06131 001 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 |
                          Aque_Aqu2.1.41463_001 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPH
                      100 Aque Aqu2.1.05134 001 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHB6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1
       -Emue Em0008g185a.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB4/EPHB4/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA1/EPHB6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA1/EPHB6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA1/EPHB6/EPHA1/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA
100 - Emue_Em0008g179a.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA5/EPHA5/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
           =Emue_Em0008g186a.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB4/EPHB4/EPHB4/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA5/EPHA6/EPHA7/EPHB8/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA5/EPHA6/EPHA7/EPHB8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA5/EPHA6/EPHA7/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA1/EPHB6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA1/EPHA1/EPHA1/EPHA1/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA
                -Emue Em0008g180a.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB4/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                     Gracom Gcom scaffold1549 01 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA2/EPHA3/EPHA6/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA10/EPHA10/EPHA1/EPHA10/EPHA1/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPH
               100 Gracom Gcom scaffold1550 01 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHB3/EPHB4/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHB4/EPHB6 | EPHA1/EPHA4/EPHA5/EPHA4/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA4/EPHA6/EPHA4/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA4/EPHA6/EPHA4/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA4/EPHA6/EPHA4/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA4/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPH
                    ■Ocar_g5890 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA8/EPHB1/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 |
                   Plajan_Pjan_scaffold12315_01 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA1/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
              100 Plajan Pjan scaffold12318 01 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB4/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                 Dgig XM 028538530.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                         --Spis XP 022806407 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                               -Adig_XM_015904379.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                            -Nvec NVE9063 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                         ■Exapal XM 028659592.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                               --Exapal XM 021046568.2 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                        -■Exapal_XM_028659591.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                              <sub>-</sub>■Exapal_XM_028659595.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                        -Exapal_XM_028662969.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                    Nvec v1g17263 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                        Nvec v1g146109 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                             Nvec v1g12570 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                Nvec_v1g173481 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                     Nvec_v1g128719 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                                              -■Nvec v1g222388 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                     Spis XP 022796255 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                        ■Spis XP 022777731 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                             Spis_XP_022796256_1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                       __spis_XP_022796249_1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                   <sup>1</sup> ■Spis XP 022796259 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                              _■Adig XM 015923300.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                      -■Adig XM 015916546.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                               <sub>-</sub>-Adig_XM_015894207.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                           <sup>00</sup>¶•Adig_XM_015904776.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                -Adig XM 015920352.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                -Adig XM 015925403.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                       -Adig XM 015914302.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                            _-Adig_XM_015920355.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                        100<sup>™</sup>-■Adig XM 015920353.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                       ■Dgig XM 028538389.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                   ■Dgig XM 028538018.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                   ■Dgig XM 028538013.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                            ■Dgig_XM_028538533.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                  --Dgig_XM_028538525.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                ■Dgig XM 028538528.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                             -Nvec v1g234961 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                Exapal XM 021041496.2 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                           -Spis XP 022784737 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                   _Adig XM 015920675.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                               100 -Adia XM 015892326.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                       Morvir_scaffold376.g15.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |

_Hvul_g18122_1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                        -Hvul α10805 1 Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                      -■Hvul α18496 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                               -■Hvul_g18123_1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                                      -Hvul_g22471_1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                       ■Hvul g21527 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                            Hvul g22478 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                        100 - Hvul g21061 1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                    -■Morvir scaffold1262.g1.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                     Nemnom g1352.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                               100 Nemnom g19191.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA
                                                                                 Nemnom_g1362.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |

Nemnom_g1362.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |

Nemnom_g19198.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |

Nemnom_g1360.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |

Nemnom_g19197.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |

Nemnom_g4858.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                                           -Nemnom_g19194.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                                              -Nemnom g19195.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                                                Nemnom_g1356.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |

Nemnom_g19196.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                                                   61 Nemnom g1355.t1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                           Dgig XM 028538388.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1
                                            TrH2_TrispH2_006515-RA | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA6/EPHA5/EPHA6/EPHA6/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA6/EPHA6/EPHA6/EPHA6/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB6/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EPHB6/EP
                                       --Lampan Lpan_11294_01 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHB3/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6
                                                        ampan_Lpan_T1294_UTTPkinase.nG30.2:EPhA1/EPhA2/EPhA3/EPhA3/EPhA5/EPhA6/EPhA7/EPhA8/EPhA1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/EPhB1/
                                                                -Horcal_Hcv1.av93.c4.g440.i1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA6/EPHA6/EPHA6/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB6 | EPHA1/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA10/EPHA1/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EP
                                                                 Beraby_Baby_sb342831 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA6/EPHA5/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPH
                     Skow XM 006821124.1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1
                         -Spur_XM_011683468.2 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB
                               --Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA198909 | Pkinase.HG30.2:EPHA1/EPHA5/EPHA5/EPHA5/EPHA5/EPHB1/EPHB3/EPHB1/EPHB3/EPHB1/EPHB3/EPHB3/EPHB3/EPHB4/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA5/EPHA5/EPHA6/EPHA5/EPHA5/EPHA5/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA5/EPHA6/EPHA5/EPHA6/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA5/EPHA6/EPHA5/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA5/EPHA6/EPHA5/EPHB1/EPHB3/EPHB6 | EPHA1/EPHA5/EPHA6/EPHA5/EPHA6/EPHB3/EPHB3/EPHB6 | EPHA1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA5/EPHA6/EPHA6/EPHB1/EPHB3/EPHB6 | EPHA1/EPHB1/EPHB3/EPHB4/EPHB6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA5/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1
                               -Tcas_TC033957_001 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/
                               --Dmel_FBtr0089084 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA5/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA5/EPHA6/EPHA7/EPHA8/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                --Bralan_BLAG11101292_1 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 |
                                        ■Drer ENSDART00000037292 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                   100 - Drer ENSDART00000105774 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                            ■Drer_ENSDART00000149895 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA4 |
■Drer_ENSDART00000004338 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA4 |
■Hsap_ENST00000281821 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA4 |
                                         100 Mmus ENSMUST00000027451 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA4
                                              _•Mmus ENSMUST00000113403 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA5 |
                                                 -Drer ENSDART00000006787 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA3 |
                                                  □ Drer ENSDART00000102788 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA7 |
                                                      Hsap ENST00000369303 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA7
                                                   100 Mmus ENSMUST00000029964 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA7 |
                                                          -Hsap ENST00000166244 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA8 |
                                                    100 Mmus ENSMUST00000030420 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA8 |
                                                    --Drer ENSDART00000146530 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA6 |
                                                        _Hsap ENST00000389672 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA6 |
                                                    100 Mmus ENSMUST00000068860 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA6
                                                                           Cint ENSCINT00000013813 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2 |
                                                              -Cint ENSCINT00000014701 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2 |
                                                                      ■Cint ENSCINT00000022765 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2
                                                              ■Hsap ENST00000275815 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1
                                                         Mmus ENSMUST00000006614 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2 |
                                              Hsap ENST00000358432 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2
                                                     _ Drer ENSDART00000044918 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2 |
                                                         -Drer ENSDART00000025428 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2 |
                                                             ■Cint ENSCINT00000025460 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 |
                                                    -Cint ENSCINT00000002487 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 |
                                                    --Cint ENSCINT00000025459 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                               ■Drer ENSDART00000148120 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | NA |
                                                  -Drer ENSDART00000043756 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB2
                                                        _Hsap ENST00000400191 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB2 |
                                                    100 Mmus ENSMUST00000059287 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB2 |
                                                    -Drer ENSDART00000128457 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB1 |
                                                        Mmus ENSMUST00000035129 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB1 |
                                                     ■Drer ENSDART00000147483 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB6
                                                                                -Hsap ENST00000422643 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB6
                                                     100
                                                                        100 Mmus ENSMUST00000114732 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB6 |
                                                                 ■Hsap ENST00000330394 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB3
                                                                Mmus ENSMUST00000006112 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB3 |
                                                                  □-Drer ENSDART00000169242 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB3 |
                                                                   Drer ENSDART00000040209 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB3 |
                                                                  ■Hsap ENST00000358173 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB4 |
                                                                  mmus ENSMUST00000111055 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB4
                                                                   □ Drer ENSDART00000187955 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB4
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■Drer ENSDART00000097644 | Pkinase.HG30.2:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB4 |

■Tadh TriadT50842 | Pkinase.HG30.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/E... | NA

0.41666