```
■Pfal CAD52615 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA
                  ■lhof evm1s770 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
       ■Ihof evm1s582 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
         --Cowc KJE94520 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
            --Clim evm27s134 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
          -■Nk52_evm14s260 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
   ■Plafun PRP83573 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
      Plafun PRP87204 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA
                     ■Plafun PRP87667 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                         Plafun PRP77325 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                         --Plafun PRP81431 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                           --Tielac_KYQ90612 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA
                                ■Ddis_EAL71407 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                              Cavfas_EGG19762 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                        100 Hetalb_XP_020436604.1 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
          --Ddis_EAL68440 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                 --Tielac KYQ90942 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                            --Hetalb XP 020430277.1 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                           --Cavfas EGG19512 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                         ■Plafun_PRP89765 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                    ■Masbal CACRSC010000016.1.p195 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                        -Cavfas_EGG23797 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                    ---Hetalb_XP_020434345.1 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
              Ddis EAL70677 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
        99 Tielac_KYQ99908 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB3/EPHB4/EPHB6/... | NA |
                             Linana_g12477.t1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                   ■Phoaus_g17279.t1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                           ■Skow XP 006822734.1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                                    ■Meteor CRO19MET NODE 127.p2 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA
                                             100 Meteor_CRO19MET_NODE_124.p2 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                    ─■Hydvul_Sc4wPfr_740.1.g22464.t1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                  ■Plamic ANCSI 00442.mRNA.1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                                Helrob HelroP159151 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                                   --Cavfas EGG14275 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                                       -■Ddis_EAL61842 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                                            --Tielac KYQ90674 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                    Esil CBJ26852 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                        Sappar KDO24699 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA
                       ■Pinf PITG 19041T0 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                  ■Pinf PITG 22989T0 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                           ■Gonavo_comp60293_c0_seq1_6_ORF27_697 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                    Gonavo_comp54449_c0_seq1_6_ORF18_960 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                    --Gonavo comp48367 c1 seq1 6 ORF2 701 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                          --Aplker 52040 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                 Ocar_g6665.t1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                           _■Maljak_MALJA_17972.mRNA.1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                     100 ■Maljak_MALJA_13305.mRNA.1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                         Gonavo comp59570 c1 seq2 6 ORF37 1537 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                             ■Ttra KNC47820 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                 -Gonavo comp59309 c1 seq1 6 ORF7 1044 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                  ■Concor KXN68012 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                              Spun KND00975 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                   Spun KNC97082 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                               ■Coerev PIA13930.1 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                     -Cana ORZ40323 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                  _Amac KNE55500 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                             oo - Amac KNE60168 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                           ■Spom SPBC1D7.05.1 pep | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                            Scer YLR362W | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA
                                         •Umay KIS67763 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                                   -Cneo_AAW43189 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                            ■Tmel_CAZ81496 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                               -■Ncra EAA33758 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                           -■Rirr GBC12660 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                       ■Morelo OAQ26035 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                                                         -Concor KXN67506 | SAM.HG5.1:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
Acas ELR13102 | SAM.HG5.10:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6... | NA |
                    --Apla XP 022086323.1 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
              ■Notgen_g9027.t1 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
           Phoaus g1445.t1 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
      <sup>100</sup>┓ Phoaus g18318.t1 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                     ■Vitbra CEM16412.1 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                               ■Pfal CZT98734 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                 ■Vitbra_CEM08830.1 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/...
                     Tthe EAR94148 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                       --Ptet CAK71557 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                         ---Ptet_CAK62401 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                      ---Ptet_CAK79445 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                           _-Ptet CAK64142 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
                     100 -Ptet CAK79774 | SAM.HG5.2:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |
   •Gefoke MALSP 12413.mRNA.1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                         --Mlei_ML0901_g19_t1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
          Cowc KJE93839 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                    Sros EGD76196 | SAM.HG5.3:MAP3K20 | MAP3K20
                              ■Mbre EDQ85761 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                       Emue 0004g41a | SAM.HG5.3:MAP3K20 | MAP3K20 |
                     Tetwil_Twilhelma_g21156.t1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                <sup>100</sup> Tetwil_Twilhelma_g21156.t2 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                         Ocar_g1648.t1 | SAM.HG5.3:MAP3K20 | MAP3K20
                                         Scil scpid46783 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                Lcom | lcpid44634 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                           100 Lcom lcpid48819 | SAM.HG5.3:MAP3K20 | MAP3K20
                        Spis_XP_022782733_1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                           -Adia XP 015751006.1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                         ■Morvir scaffold255.g32.t1 | SAM.HG5.3:MAP3K20 | MAP3K20
                              Clyhem_TCONS_00034982-protein | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                ─■Hydvul_Sc4wPfr_1395.g14012.t1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                           --Hsap_ENSP00000364361.3 | SAM.HG5.3:MAP3K20 | MAP3K20 |
               100 -Drer ENSDARP00000093598.4 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                 -Cscu XP 023210465.1 | SAM.HG5.3:MAP3K20 | MAP3K20
                             -Lpol XP 013777524.1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                    -Cgig EKC38128 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                          ■Notgen_g3015.t1 | SAM.HG5.3:MAP3K20 | MAP3K20
                                    ____Phoaus g3739.t1 | SAM.HG5.3:MAP3K20 | MAP3K20
                                           -Linana g10844.t1 | SAM.HG5.3:MAP3K20 | MAP3K20
                                      Bralan BL16971 evm0 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                               --Apla_XP_022100778.1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                          --Skow_XP_006818517.1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
                                                                   ■Ptyfla pfl 40v0 9 20150316_1g9965.t1 | SAM.HG5.3:MAP3K20 | MAP3K20 |
          Cavfas EGG25012 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
              ---Hetalb XP 020426823.1 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                   ─■Ddis_EAL65677 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                     ■Tielac_KYR01693 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                     Acas ELR23232 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                     Acas ELR24935 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
                   ■Plafun PRP84750 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                   -Acas ELR20162 | SAM.HG5.4:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                            -Acas ELR22543 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
                 ■Maliak MALJA 02574.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:..
                       ■Gefoke MALSP 10725.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
                        ■Malcal MALCA 12227.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
                           Maljak MALJA 20283.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
                     100 Maljak MALJA 20952.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                      --Malcal_MALCA_10837.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                      -■Maljak MALJA 14766.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
               100 Maljak MALJA 14442.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                           ■Gefoke MALSP 11807.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
                         ■Gefoke MALSP 05601.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                       ■Gefoke MALSP 14812.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                          ■Malcal MALCA 08228.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                                  ■Maljak_MALJA_08530.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                                   ■Maĺcal_MALCA_01192.mRNA.1 | SAM.HG5.5:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                        -■Honfer_GBG33887 | SAM.HG5.6:like:ÉPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                  -Acas ELR19495 | SAM.HG5.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                          ■Bnat_JGI_V11_67859 | SAM.HG5.6:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                       ■Ttra KNC54854 | SAM.HG5.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                Acas ELR21887 | SAM.HG5.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                  -Acas ELR19817 | SAM.HG5.7:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
                  --Meteor CRO19MET NODE 885.p2 | SAM.HG5.11:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6... | NA |
                                                                    -Acas_ELR19386 | SAM.HG5.9:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA
                                                            --Ttra_KNC51622 | SAM.HG5.9:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |

    **Itia_kNC51022 | SAW::163:9:like:EPHA1/EPHA2/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHA10/EPHB1/EPHB5/EPHB6:... | NA |
    **Meteor_CRO19MET_NODE_7739.p1 | SAM.HG5.9:like:EPHA1/EPHA2/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
    **Rirr_GBC42727 | SAM.HG5.9:like:EPHA1/EPHA2/EPHA4/EPHA6/EPHA6/EPHA6/EPHA10/EPHB1/EPHB2/EPHB3/EPHB6:... | NA |
    **Ehux_EOD11627 | SAM.HG5.9:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |
    **Ehux_EOD30414 | SAM.HG5.9:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6:... | NA |

                                    ■Masbal CACRSC010000273.1.p48 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/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/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1
                             -Acas ELR15519 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA7/EPHB1/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                      -Acas ELR25726 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                      -Acas ELR19142 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                              Acas ELR13286 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHB1/EPHB1/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA5/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1
                                ■Bralan BL95005 evm1 | SAM.HG5.8: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/EP
                                    ---Mbre_EDQ88859 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA8/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                        -Sros EGD76978 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                    -•Mlei ML3591 g3 t1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA7/EPHB4/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA4/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA4/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6 | EPHA1/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
                                             -Ocar g5890.t1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                       Lcom | lcpid47939 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/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/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA1/EPHB1/EPHB4/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/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA
                                                       Scil scpid13953 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/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/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA5/EPHA6/EPHA1/EPHB1/EPHB4/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/EPHA1/EPHA1/EPHA1/EPHA
                                                 100 Scil scpid35129 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                                       Aque Aqu2.1.41463 001 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB3/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | 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/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/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/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/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/EP
                                                 100 Aque Aqu2.1.05134 001 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB4/EPHB4/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                 -Emue 0008g185a | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                 -Emue 0008g179a | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA4/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA6/EPHA6/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/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
                                                       Emue 0008g186a | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB1/EPHB3/EPHB4/EPHB6
                                                  100 = Emue 0008g180a | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB3/EPHB1/EPHB3/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6
                                                                   -■Tadh P12344 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                                   ■Morvir scaffold376.g15.t1 | SAM.HG5.8: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/EP
                                                    -Auraur scaffold453.α9.t1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB3/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                                                           -Clyhem TCONS 00000567-protein | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/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
                                                                         •Clyhem TCONS 00022872-protein | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB3/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                              --Clyhem TCONS 00024179-protein | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA1/EPHB4/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                             ■Hvdvul Sc4wPfr 697.1.g18496.t1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                                          ■Hydvul Sc4wPfr 740.1.g22478.t1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                                           Hydvul Sc4wPfr 740.1.g22478.t2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                    -•Nvec NVE23570 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 |
                                                    -Spis_XP_022784737_1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB6 | EPHA1/EPHA10/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB6/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB6/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA1/EPHB6/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA6/EPHA6/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/EP
                                                              Spis_XP_022796255_1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA7/EPHA8/EPHA6/EPHA6/EPHA6/EPHA7/EPHA8/EPHA6/EPHA6/EPHA6/EPHA6/EPHA7/EPHA8/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA7/EPHA8/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/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/EPHA
                                                         --Spis_XP_022806407_1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB4/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHB1/EPHB3/EPHB4/EPHB6
                                                             ---Adig_XP_015759865.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHB4/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/E
                                                           Nvec NVE5316 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/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/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1
                                                                   Nvec_NVE11033 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/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/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA1/EPHA5/EPHA6/EPHA1/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA10/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/EPHA1/EPHA1/EPHA1/EPHA1/EP
                                                                     100 Nvec NVE5319 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 |
                                                   ■Helrob HelroP178880 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/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/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/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
                                               Spur XP 011681770.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA10/EPHA2/EPHA3/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB3/EPHB4/EPHB6
                                                                  Linana g1515.t1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                                                 -Notgen_g12738.t1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                                                          -Ctel_CapteP198909 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHA6/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA6/EPHA6/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA3/EPHA5/EPHA6/EPHA6/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA3/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA4/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA6/EPHA6/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA6/EPHA6/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA6/EPHA6/EPHA8/EPHB1/EPHB2/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA6/EPHA6/EPHA8/EPHB8/EPHB3/EPHB4/EPHB6 | EPHA10/EPHA3/EPHA4/EPHA6/EPHA6/EPHA8/EPHB8/EPHB3/EPHB4/EPHB6 | EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA8/EPHB8/EPHB3/EPHB4/EPHB6 | EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA8/EPHB8/EPHB4/EPHB6 | EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA8/EPHB8/EPHB4/EPHB6 | EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA8/EPHB8/EPHB4/EPHB6 | EPHA10/EPHA10/EPHA3/EPHA6/EPHA6/EPHA6/EPHA8/EPHB8/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA4/EPHA6/EPHA6/EPHA6/EPHA8/EPHB4/EPHB6 | EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA6/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/EPHBA/E
                                                                      100 Ramvar_GAU98634.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB3/EPHB4/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA3/EPHA4/EPHA6/EPHA6/EPHA6/EPHB1/EPHB1/EPHB4/EPHB6 | 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/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA10/EPHA1
                                                                         ■Dmel FBpp0088153 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA7/EPHB1/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/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 |
                                                                                     -Lpol XP 022258078.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA7/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA2/EPHA3/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA3/EPHA4/EPHA6/EPHA6/EPHA7/EPHA8/EPHB1/EPHB4/EPHB6 | EPHA1/EPHA1/EPHA1/EPHA1/EPHA1/EPHA6/EPHA1/EPHA1/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/
                                                                                    T2 Lpol_XP_022258076.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/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/EPHB6 | EPHA1/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHA6/EPHB6/EPHB6/EPHB6 | EPHA1/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/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
                                                                                        100 Lpol_XP_022243645.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHB4/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB3/EPHB4/EPHB6 |
                                                                                     -Cscu XP 023220906.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/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
                                                                                         Cscu_XP_023213518.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/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/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/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
                                                                                          100 CSCU_XP_023226292.1 SAM.HG5.8: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/EP
                                                                                          -LDOI XP 022238397.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/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/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/EPHB1/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
                                                                                          -Lpol XP 022254188.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHB1/EPHB1/EPHB1/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                                         _PLpol XP 022244546.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHB1/EPHB1/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6
                                                         -Cint ENSCINP00000002487.3 | SAM.HG5.8: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 | EPHA1/EPHA1/EPHA3/EPHA4/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/EPHB1/EPHB3/EPHB4/EPHB6 |
                                                           -Cint ENSCINP00000025213.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB3/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                 -Cint ENSCINP00000014701.3 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA5/EPHA6/EPHA5/EPHB1/EPHB3/EPHB1/EPHB1/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA10/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                   -Cint ENSCINP00000022519.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA6/EPHA6/EPHB3/EPHB1/EPHB3/EPHB3/EPHB4/EPHB6 | EPHA1/EPHA2/EPHA3/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHB1/EPHB3/EPHB4/EPHB6
                                                                   -Drer ENSDARP00000113830.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB6 |
                                                                       Hsap ENSP00000404786.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB6 |
                                                                  100 Hsap ENSP00000481418.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB6
                                                                     --Hsap ENSP00000350896.3 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB4 |
                                                                    ____•Drer ENSDARP00000088414.3 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB4
                                                                       -Drer ENSDARP00000152897.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB4 |
                                                                  ■Drer ENSDARP00000140419.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB3 |
                                                                  -- Hsap ENSP00000332118.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB3
                                                                        -Drer ENSDARP00000111570.4 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB1
                                                                100  ■Hsap ENSP00000497153.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB1 |
                                                                       -Hsap ENSP00000383053.3 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB2
                                                                            -Drer ENSDARP00000043755.7 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB2
                                                                                  Drer ENSDARP00000112928.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB2 |
                                                                              100 - Drer ENSDARP00000151014.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHB2 |
                                                                            Hsap ENSP00000275815.3 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1 |
                                                                      100 Hsap ENSP00000494931.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA1 |
                                                                    ---Hsap ENSP00000351209.5 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2 |
                                                                      —■Drer ENSDARP00000044917.5 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2
                                                                             ■Drer ENSDARP00000011069.6 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2
                                                                        100 Drer ENSDARP00000148430.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA2
                                                                              ■Hsap ENSP00000397746.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA10 |
                                                                 --Hsap ENSP00000358309.4 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA7 |
                                                                       ■Drer ENSDARP00000093563.5 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA7
                                                                  100 Drer ENSDARP00000157495.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA7 |
                                                                         -- Hsap ENSP00000166244.3 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA8 |
                                                                        ■Drer ENSDARP00000121462.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA6 |
                                                                       -- Hsap ENSP00000374323.5 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA6 |
                                                                         -- Hsap ENSP00000337451.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA3
                                                                           -Drer ENSDARP00000075583.5 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA5 |
                                                                    100 Hsap ENSP00000480763.1 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA5
                                                                              --Drer ENSDARP00000096552.3 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | islike:EPHA4 |
                                                                            -Drer ENSDARP00000030606.8 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | islike:EPHA4 |
                                                                            -Hsap ENSP00000281821.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA4 |
                                                                                  ¬■Drer ENSDARP00000123962.2 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA4
                                                                                     -Drer_ENSDARP00000003161.9 | SAM.HG5.8:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6 | EPHA4 |
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-Cpar Cpar evm1s8670 | SAM.HG5.0:like:EPHA1/EPHA2/EPHA3/EPHA4/EPHA5/EPHA6/EPHA7/EPHA8/EPHA10/EPHB1/EPHB2/EPHB3/EPHB4/EPHB6/... | NA |