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Adig XP 015771483.1 1505-1601 | rve.HG17.22:NA | NA |
                 Auraur_scaffold921.g8.t1_186-311 | rve.HG17.4:NA | NA |
                   Morvir_scaffold58.g30.t1_1498-1621 | rve.HG17.4:NA | NA |
                    Auraur_scaffold3787.g1.t1_793-916 | rve.HG17.4:NA | NA |
                  Spis_XP_022796387_1_20-124 | rve.HG17.4:NA | NA |
                  . ■Adig XP 015751999.1 224-330 | rve.HG17.4:NA | NA |
                  ■Adig_XP_015757043.1_351-457 | rve.HG17.4:NA | NA |
                     Scil scpid32584 453-568 | rve.HG17.6:NA | NA |
          82
                  Auraur_scaffold1545.g3.t1_547-669 | rve.HG17.6:NA | NA |
                     Emue 0008g33a 17-114 | rve.HG17.10:NA | NA |
                               Scil scpid97941 192-309 | rve.HG17.10:NA | NA |
                      ■Emue_0007g1205a_1246-1341 | rve.HG17.11:NA | NA |
83
          94
                          Emue_1107g1a_533-648 | rve.HG17.11:NA | NA |
             92
                       .■Tetwil_Twilhelma_g749.t1_194-299 | rve.HG17.11:NA | NA |
                    <sup>100</sup> Tetwil_Twilhelma_g16834.t1_367-473 | rve.HG17.11:NA | NA |
                -Adig XP 015773647.1 93-189 | rve.HG17.0:NA | NA |
             Adig XP 015774073.1 52-156 | rve.HG17.0:NA | NA |
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                      Clyhem_TCONS_00060909-protein_398-500 | rve.HG17.1:NA | NA |
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                      Hydvul_Sc4wPfr_10.g18919.t1_1461-1562 | rve.HG17.1:NA | NA |
                          Exapal XP 020896007.1 43-168 | rve.HG17.1:NA | NA |
                            Adig_XP_015759415.1_35-158 | rve.HG17.1:NA | NA |
     51
                        <sup>100</sup> •Adig_XP_015771259.1_91-215 | rve.HG17.1:NA | NA |
                              Auraur_scaffold4180.g2.t1_131-254 | rve.HG17.1:NA | NA |
                              Auraur_scaffold2506.g1.t1_1-126 | rve.HG17.1:NA | NA |
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                             Spis XP 022779443 1 1401-1508 | rve.HG17.1:NA | NA |
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                               Spis XP 022798583 1 275-382 | rve.HG17.1:NA | NA |
                                   Adig_XP_015780681.1_52-168 | rve.HG17.1:NA | NA |
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                                     -Adig XP 015758779.1 46-148 | rve.HG17.1:NA | NA |
                                      Spis XP 022777809 1 86-207 | rve.HG17.1:NA | NA |
                                     Adig XP 015767465.1 94-200 | rve.HG17.1:NA | NA |
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                               Spis XP 022793401 1 44-168 | rve.HG17.1:NA | NA |
                                 Adig XP 015770723.1 46-168 | rve.HG17.1:NA | NA |
                              100 Adig XP 015752038.1 46-168 rve.HG17.1:NA NA
                                    -Adig XP 015777732.1 152-276 | rve.HG17.1:NA | NA |
                                100 Adig XP 015748702.1_295-419 | rve.HG17.1:NA | NA |
                 Skow_XP_006812920.1_19-143 | rve.HG17.2:NA | NA |
             Skow_XP_006817251.1_564-694 | rve.HG17.2:NA | NA |
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                Skow XP 006814449.1 31-162 | rve.HG17.2:NA | NA |
                ■Notgen g25623.t1 872-997 | rve.HG17.5:NA | NA |
                PNotgen_g21208.t1_1284-1394 | rve.HG17.5:NA | NA |
         99
                ┗Notgen_g21208.t1_1491-1601 | rve.HG17.5:NA | NA |
                       Skow_XP_002735894.1_741-870 | rve.HG17.7:NA | NA |
                      Spis XP 022791783 1 294-421 | rve.HG17.7:NA | NA |
      96
                       ■Nvec NVE15335 131-257 | rve.HG17.7:NA | NA |
                    <sup>100</sup> -Nvec_NVE9699_1335-1465 | rve.HG17.7:NA | NA |
                    Notgen_g37996.t1_197-325 | rve.HG17.23:NA | NA |
         62
                     Skow_XP_006815894.1_1-125 | rve.HG17.24:NA | NA |
                        Skow XP 006822499.1 21-148 | rve.HG17.25:NA | NA |
                98
                              Ptyfla pfl 40v0 9 20150316 1g21862.t1 785-908 | rve.HG17.19:NA | NA |
           82
                   88
                           <sup>100</sup> Ptyfla_pfl_40v0_9_20150316_1g25220.t1_872-996 | rve.HG17.19:NA | NA |
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                             Skow_XP_006820445.1_482-608 | rve.HG17.19:NA | NA |
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                             100 -Skow XP 006824596.1_716-835 | rve.HG17.19:NA | NA |
                        Notgen g33660.t1 10-134 | rve.HG17.18:NA | NA |
                     Notgen_g11120.t1_175-301 | rve.HG17.26:NA | NA |
                48
83
                           Skow_XP_006825937.1_72-196 | rve.HG17.20:NA | NA |
                               r=Skow_XP_006816667.1_600-721 | rve.HG17.20:NA | NA |
                   67
                                Skow XP 006817489.1 384-495 | rve.HG17.20:NA | NA |
                     87
                                 Skow_XP_006816472.1_239-361 | rve.HG17.21:NA | NA |
                                 ■Notgen_g33487.t1_1419-1549 | rve.HG17.21:NA | NA |
                                  -Notgen_g28984.t1_123-245 | rve.HG17.21:NA | NA |
                   Scil scpid43203 220-327 | rve.HG17.3:NA | NA |
                      Clyhem TCONS 00056363-protein 113-208 | rve.HG17.3:NA | NA |
     52
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                            Clyhem_TCONS_00003056-protein_156-261 | rve.HG17.3:NA | NA |
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                          Hydvul_Sc4wPfr_381.1.g32170.t1_1156-1270 | rve.HG17.3:NA | NA |
                            Hydvul_Sc4wPfr_1942.g11903.t1_1245-1345 | rve.HG17.3:NA | NA |
                         60 Hydvul Sc4wPfr 106.g23470.t1 72-172 | rve.HG17.3:NA | NA |
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                            47 -Cscu XP 023222405.1_694-831 | rve.HG17.9:NA | NA |
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                          Exapal_XP_020911222.1_466-605 | rve.HG17.9:NA | NA |
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                            Tetwil Twilhelma g7974.t1 4-118 | rve.HG17.9:NA | NA |
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                         <sup>100</sup> Tetwil_Twilhelma_g8094.t1_8-125 | rve.HG17.9:NA | NA |
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                               Scil_scpid3102_1488-1615 | rve.HG17.14:NA | NA |
                          Lcom lcpid5678 1966-2095 | rve.HG17.15:NA | NA |
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                         Spis XP 022778085 1 148-280 | rve.HG17.16:NA | NA |
                           Adig XP 015749350.1 241-373 | rve.HG17.16:NA | NA |
                        <sup>100</sup> Adig_XP_015765510.1_150-282 | rve.HG17.16:NA | NA |
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                                   Spur_XP_030834741.1_494-618 | rve.HG17.17:NA | NA |
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