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•Halamb_c39428_g1_i1m.11731 | OG0 |
           -Aque Aqu2.1.20295 001 | OG0 |
             •Aque Aqu2.1.06803 001 | OG0 |
              -Tetwil_g6625.t2 | OG0 |
74
          <sup>100</sup> Tetwil g14780.t1 | OG0 |
             |Spolac_c100638_g1_i1_m.23077c100638-g1 | OG0
  50
             Spolac_c100638_g1_i2_m.23078c100638-g1 | OG0 |
         100
                •Emue_Em0008g142a.t1 | OG0 |
                                   -Emue_Em0008g157a.t1 | OG0 |
                 •Aque_Aqu2.1.06804_001 | OG1 |
            TrH2 TrispH2_004551-RA | OG1 |
  87
         <sup>100</sup> Tadh_TriadT37861 | OG1 |
           -HoiH11_mRNA1858 | OG1 |
               <sub>"</sub>HoiH23_PIH23_005776-RA|OG1|
             <sup>99]</sup>HoiH06 mRNA7637 | OG1 |
                <sub>•</sub>Hhon g06925.t1 | OG1 |
             <sup>100</sup> HoiH04_mRNA14909 | OG1 |
                  ·Stycar_maker-SC_scaffold3234-snap-gene-0.6-mRNA-1 | OG1 |
                Opea_EP00118_Oscarella_pearsei_P025111 | OG1 |
    29
                •Plajan_Pjan_C539640_01 | OG1 |
        99
                        •Lcom_lcpid181040 | OG1 |
              95
                      Lcom_lcpid252644 | OG1 |
                    •Clacor_Ccor_scaffold10548_01 | OG1 |
       89
                        •Lcom_lcpid99323 | OG1 |
            98
                        Gracom_Gcom_scaffold10996_01 | OG1 |
  69
                           -Leuniv_Lniv_scaffold24633_01 | OG1 |
                                      •Scil_scict010901.2 | OG1 |
                 Skow_XM_002738499.2 | OG1 |
                     •Bralan_BLAG11000195_1 | OG1 |
                        •Hsap_ENST00000270586 | OG1 |
                                -Hsap_ENST00000374859 | OG1 |
                     •Spur_XM_030986824.1 | OG1 |
    69
                          •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA156053 | OG1 |
                         •Acagra_model.g30660.t1 | OG1 |
                         •Cgig_XM_011415038.3 | OG1 |
                              •Lcom_lcpid169951 | OG1 |
                        •Dpul mRNA22902 | OG1 |
                             •Plespi_Pspi_scaffold19264_01 | OG1 |
                             •Cscu_XM_023387572.1 | OG1 |
                          rtcas_TC008665_001 | OG1 |
                                     •Tcas_TC031599_001 | OG1 |
      60
                 •Nvec_v1g101124 | OG1 |
                 •Exapal_XM_021058524.2 | OG1 |
                  <sub>•</sub>Pocdam_XM_027192410.1 | OG1 |
          83
                   •Spis_XP_022793532_1 | OG1 |
                   •Fspp_ffun1.m4.7627.m1 | OG1 |
                      •Gasp_gasp1.m3.27646.m1 | OG1 |
                        •Gfas_gfas1.m1.19386.m1 | OG1 |
                         <sub>-</sub>-Amil_Amillepora24506-RA | OG1 |
                             •Adig_XM_015903718.1 | OG1 |
        76
                       •Lcom_lcpid151469 | OG1 |
                         Morvir scaffold17.g68.t1 | OG1 |
                          •Hvul_g7824_1 | OG1 |
                        •Aaur_scaffold697.g2.t1 | OG1 |
                          •Nemnom_g3882.t1 | OG1 |
                                -Sanmal_Sma_020703-T1 | OG1 |
          74
                         <sup>58</sup> •Rhoesc_Res_009814-T1 | OG1 |
                           -Lcom_lcpid133719 | OG1 |
                        •Dgig_XM_028538398.1 | OG1 |
                 95
                       •Xesp 012726-T1 | OG1 |
                            •Lcom | lcpid234220 | OG1 |
               94
                                         -Pbac_sb2644314 | OG1 |
                                    sb2644312 | OG1 |
                                     <sub>_</sub>-Bolinf_Binf_sb1252510 | OG1 |
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                                  <sup>1001</sup>Mertsp_Mesp_sb578272 | OG1 |
                                     •Drygla_Dgla_sb286945 | OG1 |
                            81
                                         Bolinf_Binf_sb1244827 | OG1 |
                                      99 Mertsp Mesp sb581493 | OG1 |
                                          Beraby Baby sb340933 | OG1 |
                                       <sup>100</sup> Drygla Dgla_sb281853 | OG1 |
                                     /mul_sbMO438473 | OG1 |
                                       Coemet Cmet C583474 01 | OG1 |
                                         Coeast Cast sb791171 | OG1 |
                                           Coeast_Cast_sb791173 | OG1 |
                                              Coeast Cast sb791174 | OG1 |
                                             <sup>28</sup> -Coeast Cast sb791172 | OG1 |
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