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TrH2 TrispH2 006828-RA | OG0 |
           <sup>100</sup><sup>1</sup>Tadh_wf_g8325.t1 | OG0 |
               •HoiH11_mRNA1014 | OG0 |
        100
                    •HoiH23_PIH23_010418-RA | OG0 |
                    •HoiH06 mRNA12249 | OG0 |
             100
                   Hhon g02110.t1 | OG0 |
                <sup>1001</sup>-HoiH04 _mRNA8936 | OG0 |
                              •Lcom_lcpid128128 | OG0 |
                          -Coemet_Cmet_C608188_01 | OG0 |
   88
                       <sup>1001</sup>•Coeast_Cast_sb798049 | OG0 |
                              Eupdun Edun_sb10690132 | OG0 |
                 100
                           -Lampan_Lpan_13335_01 | OG0 |
                             -Mertsp_Mesp_sb557019 | OG0 |
                      83
                                     -Eupdun_Edun_sb10615042 | OG0 |
                                  <sup>1001</sup>•Eupdun Edun sb10615043 | OG0 |
                        40
                                   Lampan Lpan 13336 01 | OG0 |
                          57
                                    Bolinf_Binf_sb1201577 | OG0 |
                                 <sup>76</sup> Mertsp_Mesp_sb557022 | OG0 |
                            67
                                        •Drygla_Dgla_sb307006 | OG0 |
                                           Beraby_Baby_sb336164 | OG0 |
                                        <sup>100</sup> Beraby Baby_sb336163 | OG0 |
                                            -Mertsp_Mesp_sb557020 | OG0 |
                                     85
                                             -Beraby_Baby_sb335095 | OG0 |
                                          <sup>100</sup>-Beraby_Baby_sb335094 | OG0 |
                                           Bolinf_Binf_sb1230795 | OG0 |
                                           •Bolinf Binf sb1227355 | OG0 |
                                            <sub>•</sub>Vmul_sbMO379032 | OG0 |
                                              Mertsp_Mesp_sb539539 | OG0 |
                                            66 Bolinf_Binf_sb1256219 | OG0 |
                                          <sub>[</sub>Mertsp_Mesp_sb535814 | OG0 |
                                              Eupdun Edun sb10623949 | OG0 |
                                                  •Pbac_sb2637958 | OG0 |
                                     35
                                                  Horcal Hcv1.av93.c9.g258.i1 | OG0 |
                                                <sup>99</sup><sup>1</sup>-Pbac sb2637957 | OG0 |
                  •Plajan_Pjan_C571626_01 | OG1 |
                    Popea EP00118 Oscarella_pearsei_P003100 | OG1 |
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                  <sup>100</sup>¹Ocar _g1770 | OG1 |
     76
                                •Clacor Ccor scaffold12273 01 | OG1 |
                    Clacor Ccor C521661 01 | OG1 |
                       Clacor Ccor scaffold8345 01 | OG1 |
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                       •Lcom_lcpid21076 | OG1 |
                           Gracom_Gcom_scaffold16854_01 | OG1 |
                   100
                            Scil_scict025208.1 | OG1 |
   86
                               •Gracom_Gcom_C528872_01 | OG1 |
                        69
                             •Leuniv_Lniv_scaffold8257_01 | OG1 |
                   Xestes_maker-XT_scaffold1772-snap-gene-0.5-mRNA-1 | OG1 |
                    Xestes_maker-XT_scaffold2662-augustus-gene-0.49-mRNA-1 | OG1 |
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                          •Aque_Aqu2.1.36075_001 | OG1 |
                         -Halamb_c48434_g1_i2mm.2659 | OG1 |
        93
                -Stycar maker-SC_scaffold10623-snap-gene-0.5-mRNA-1 | OG1 |
                    •Tetwil g17737.t1 | OG1 |
                          Stycar_maker-SC_scaffold2797-augustus-gene-0.4-mRNA-1 | OG1 |
            80
                       •Tetwil_g17735.t1 | OG1 |
                       |Stycar_maker-SC_scaffold3648-augustus-gene-0.3-mRNA-1 | OG1
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                             Tetwil g10870.t1 | OG1 |
                                     •Plespi Pspi scaffold20122 01 | OG1 |
                 76
                                 Stycar_maker-SC_scaffold3977-augustus-gene-0.4-mRNA-1 | OG1 |
                               •Spolac_c104157_g3_i1_m.38523c104157-g3 | OG1 |
                     86
                          <sup>100</sup><sup>1</sup>-Emue_Em0011g979a.t1 | OG1 |
                             •Emue Em0011g977a.t1 | OG1 |
                               <sub>!</sub>Spolac_c104157_g3_i1_m.38522c104157-g3 | OG1 |
                            •Spur_XM_030996301.1 | OG2 |
                  -Bralan_BLAG13000874_1 | OG2 |
                        Stycar maker-SC scaffold5564-augustus-gene-0.3-mRNA-1 | OG2 |
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                      Hsap ENST00000286398 | OG2 |
                         -Cscu_XM_023366047.1 | OG2 |
                                •Dpul mRNA10910 | OG2 |
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                                        •Tcas TC005757 001 | OG2 |
                       Ctel gnl WGS_AMQN_CAPTEDRAFT_mRNA95567 | OG2 |
                      Cgig XM 011425603.3 | OG2 |
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                         •Acagra_model.g7120.t1.1.5d3b8449 | OG2 |
                     <sup>1001</sup>Acagra_model.g7120.t1 | OG2 |
   83
                 rChem_TCONS_00005740 | OG2 |
                  -Chem_TCONS_00041380 | OG2 |
                   •Morvir scaffold33.g14.t1 | OG2 |
        98
                   ·Aaur scaffold32.g35.t1 | OG2 |
          100
                      •Aaur_scaffold32.g33.t1 | OG2 |
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                         Sanmal Sma 012071-T1 | OG2 |
                         Aaur scaffold32.g36.t1 | OG2 |
                 62
     95
                         •Nemnom g3744.t1 | OG2 |
                     <sup>1001</sup>-Rhoesc_Res_016301-T1 | OG2 |
                •Dgig_XM_028547831.1 | OG2 |
                  •Xesp_000986-T1 | OG2 |
                  •Nvec NVE8734 | OG2 |
       98
            100
                      •Exapal XM 021055800.2 | OG2 |
                         •Exapal_XM_021045947.2 | OG2 |
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                      •Spis_XP_022781443_1 | OG2 |
                       -Gasp_gasp1.m3.25128.m1 | OG2 |
            93
                    <sup>961</sup>•Fspp ffun1.m4.15886.m1 | OG2 |
              100
                       Gfas gfas1.m1.4902.m1 | OG2 |
                            -Amil Amillepora26173-RA | OG2 |
                       •Adig_XM_015923817.1 | OG2 |
                       Amil Amillepora26172-RA | OG2 |
0.416667
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