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Tadh_TriadT24132 | OG0 |
             <sub>*</sub>TrH2_TrispH2_003833-RA | OG0 |
                •HoiH11_mRNA7826 | OG0 |
                    <sub>"</sub>HoiH23_PIH23_007030-RA | OG0 |
                 <sup>1001</sup>HoiH06_mRNA1844 | OG0 |
                    Hhon g02492.t1 | OG0 |
                 <sup>1001</sup>-HoiH04_mRNA17023 | OG0 |
               •Vmul_sbMO417406 | OG0 |
                 Coemet_Cmet_scaffold19477_01 | OG0 |
              <sup>1001</sup>•Coeast_Cast_sb813032 | OG0 |
                -Lampan_Lpan_12214_01 | OG0 |
                             -Pbac sb2644945 | OG0 |
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                     Pbac_sb2644946 | OG0 |
                   <sup>76]</sup>Horcal_Hcv1.av93.c1.g1511.i1 | OG0 |
                   <sub>"</sub>Bolinf_Binf_sb1187167 | OG0 |
                <sup>100</sup> Mertsp_Mesp_sb529414 | OG0 |
                     Mlei_ML0091.g3.i1 | OG0 |
                       <sub>-</sub>Bolinf_Binf_sb1215595 | OG0 |
                     <sup>991</sup>Vmul_sbMO400909 | OG0 |
                     Beraby_Baby_sb337603 | OG0 |
                       Drygla_Dgla_sb298983 | OG0 |
                         <u>-</u>Drygla_Dgla_sb298985 | OG0 |
                       <sup>84</sup>•Drygla_Dgla_sb298987 | OG0 |
             Clacor Ccor scaffold9903 01 | OG0 |
              Gracom Gcom scaffold15086 01 | OG0 |
                Lcom | lcpid35608 | OG0 |
                  -Leuniv_Lniv_scaffold19150_01 | OG0 |
              <sup>981</sup>-Scil_scict016828.1 | OG0 |
   87
              Plajan_Pjan_C549873_01 | OG0 |
              Opea_EP00118_Oscarella_pearsei_P009973 | OG0 |
                 •Opea_EP00118_Oscarella_pearsei_P009974 | OG0 |
                 •Tetwil g19110.t1 | OG0 |
                   •Plespi_Pspi_scaffold19718_01 | OG0 |
              Stycar_maker-SC_scaffold5809-snap-gene-0.11-mRNA-1 | OG0 |
                   Emue_Em0018g294a.t1 | OG0 |
                 -Xestes_augustus_masked-XT_scaffold51683-processed-gene-0.0-mRNA-1 | OG0 |
                      -Aque_Aqu2.1.38413_001 | OG0 |
                  <sup>100</sup> <mark>--</mark>Halamb_c48572_g1_i1m.24442 | OG0 |
           Hsap ENST00000219252 | OG0 |
               Skow XM 002741004.2 | OG0 |
                 •Bralan BLAG05001271 1 | OG0 |
     59
                    Spur XM 030975695.1 | OG0 |
                    Cscu_XM_023377356.1 | OG0 |
                     •Dpul_mRNA26178 | OG0 |
                     •Tcas_TC010438_001 | OG0 |
                     -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA155833 | OG0 |
                    Cgig_XM_011420332.3 | OG0 |
                   •Acagra_model.g28886.t1_model.g28888.t1 | OG0 |
   43
                 Chem_TCONS_00049502 | OG0 |
                 •Hvul_g7763_1 | OG0 |
             <sup>1001</sup>Hvul_g7766_1 | OG0 |
                 -Morvir_scaffold272.g7.t1 | OG0 |
                 •Sanmal_Sma_025138-T1 | OG0 |
                  -Rhoesc_Res_010612-T1 | OG0 |
                   -Nemnom_g18201.t1 | OG0 |
               •Dgig_XM_028536316.1 | OG0 |
                <sub>•</sub>Xesp_005699-T1 | OG0 |
                          •Xesp 001453-T1 | OG0 |
                             •Xesp_022511-T1 | OG0 |
                         100 Xesp_026762-T1 | OG0 |
       88
              Exapal_XM_021059959.1 | OG0 |
                •Nvec_v1g190101 | OG0 |
                   -Gasp_gasp1.m3.14867.m1 | OG0 |
                 <sup>77</sup><sup>L</sup>•Gfas_gfas1.m1.18908.m1 | OG0 |
                   Spis_XP_022791762_1 | OG0 |
99 Pocdam_XM_027183406.1 | OG0 |
              98
                        Fspp_ffun1.m4.3544.m1 | OG0 |
                        Amil_Amillepora10845-RA | OG0 |
                     <sup>100</sup>ً-Adig_XM_015899298.1 | OG0 |
0.416667
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