```
Tadh TriadT30823 | OG0 |
     <sup>100</sup> TrH2 TrispH2_009255-RA | OG0 |
                                    •HoiH23 PIH23 008756-RA | OG0 |
  88
            <sub>f</sub>Hhon_g10939.t1 | OG0 |
         <sup>100</sup> HoiH04 mRNA193 | OG0 |
           -HoiH23 PIH23 006006-RA | OG0 |
         -Lampan_Lpan_14280_01 | OG0 |
                            •Coemet_Cmet_scaffold21304_01 | OG0 |
              100
                            Horcal Hcv1.av93.c8.g749.i1 | OG0 |
                         <sup>100</sup><sup>1</sup>-Pbac sb2663952 | OG0 |
                 55
                                 Beraby_Baby_sb336641 | OG0 |
                   80
                             <sup>100</sup> Beraby_Baby_sb336640 | OG0 |
                     80
                              •Mlei_ML1844.g9.i1 | OG0 |
                              <sub>!</sub>Mertsp_Mesp_sb554887 | OG0 |
                           <sup>100</sup> Bolinf_Binf_sb1205320 | OG0 |
                  Morvir scaffold261.g13.t1 | OG0 |
                     Sanmal Sma 013536-T1 | OG0 |
  88
                  Nemnom_g3565.t1 | OG0 |
             100l
                  •Rhoesc_Res_009473-T1 | OG0 |
                      Cgig XM 034470048.1 | OG0 |
                           •Chem_TCONS_00067936 | OG0 |
                   100
                              •Hvul_g32513_1 | OG0 |
                         •Xesp_000979-T1 | OG0 |
                          •Dgig XM 028542943.1 | OG0 |
                  100
          59
                      <sup>99</sup> -Dgig XM 028542873.1 | OG0 |
     78
                     •Tetwil_g26778.t1 | OG0 |
                    •Nvec_v1g138093 | OG0 |
                      Exapal_XM_021048953.1 | OG0 |
              96
                        <sub>[</sub>•Adig_XM_015909715.1 | OG0 |
                62
                     <sup>100</sup><sup>1</sup>-Amil _Amillepora27321-RA | OG0 |
                          •Gfas_gfas1.m1.2695.m1 | OG0 |
                  100
                             •Pocdam_XM_027190950.1 | OG0 |
                     76
                          <sup>100</sup> Spis XP 022793883 1 | OG0 |
       69
                             Fspp ffun1.m4.27619.m1 | OG0 |
                          <sup>100</sup>Ū-Gasp_gasp1.m3.1455.m1 | OG0 |
                     •Dpul_mRNA4619 | OG0 |
                       •Cscu_XM_023386642.1 | OG0 |
                         •Cscu XM 023386645.1 | OG0 |
                100
                    <sup>100</sup> Cscu_XM_023386637.1 | OG0 |
                       •Skow XM 002736478.2 | OG0 |
                        Bralan_BLAG02001760_1 | OG0 |
                            •Hsap_ENST00000378842 | OG0 |
              85
                        Spur XM 030976914.1|OG0|
                           Cgig XM 034468731.1 | OG0 |
                            Acagra_model.g9425.t1.1.5d3b8600 | OG0 |
                                   -Tcas_TC032578_001 | OG0 |
                          •Spolac_c102408_g1_i2_m.29197c102408-g1 | OG0 |
                  84
                          Emue Em0009g986a.t1 | OG0 |
            66
                             •Stycar_maker-SC_scaffold2063-snap-gene-0.25-mRNA-1 | OG0 |
                             •Aque_Aqu2.1.22057_001 | OG0 |
                             <sub>!</sub>Halamb_c53894_g1_i9mm.5506 | OG0 |
                  73
                           <sup>1001</sup>-Halamb_c53894_g2_i7mm.5508 | OG0 |
                                Plespi_Pspi_scaffold4432_01 | OG0 |
                    60
                             <sup>100</sup>lPlespi Pspi scaffold4434_01 | OG0 |
                                        Ctel gnl WGS AMQN CAPTEDRAFT_mRNA147099 | OG0 |
                       60
                                                  -Spis XP 022795651 1 | OG0 |
                         Clacor_Ccor_C533139_01 | OG0 |
                          -Lcom_lcpid74627 | OG0 |
                  100
                              •Gracom Gcom scaffold14841 01 | OG0 |
                         <sup>100</sup>Ū-Leuniv_Lniv_C806652_01 | OG0 |
                99
                         lajan_Pjan_scaffold578_01 | OG0 |
                               <sub>[</sub>Opea_EP00118_Oscarella_pearsei_P013077 | OG0 |
                  88
                             <sup>57</sup>Opea EP00118 Oscarella pearsei P013079 | OG0 |
                                Ocar g9696 | OG0 |
                          100
                                    Opea_EP00118_Oscarella_pearsei_P013076 | OG0 |
                                  <sup>67</sup> Opea_EP00118_Oscarella_pearsei_P013078 | OG0 |
                                    Opea_EP00118_Oscarella_pearsei_P013082 | OG0 |
                                        Opea EP00118 Oscarella pearsei P013081 | OG0 |
                                    <sup>73</sup> Opea EP00118 Oscarella pearsei P013080 | OG0 |
0.416667
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