```
•Spolac c90333 g1_i1_m.6253c90333-g1 | OG0 |
           •Emue Em0013g1124a.t1 | OG0 |
          •Tetwil_g1748.t1 | OG0 |
   93
              •Aque_Aqu2.1.36637_001 | OG0 |
                 •Halamb c20827 g1 i1m.3744 | OG0 |
                       -Xestes maker-XT scaffold11171-snap-gene-0.27-mRNA-1 | OG0 |
 88
            •Plajan Pjan C546285 01 | OG0 |
           Opea_EP00118_Oscarella_pearsei_P025396 | OG0 |
     100
        100 Ocar g9337 | OG0 |
   85
               Pbac sb2657239 | OG0 |
                 Pbac sb2657288 | OG0 |
                    •Coeast_Cast_sb749208 | OG0 |
                      Horcal Hcv1.av93.c5.g509.i1 | OG0 |
                              -Vmul sbMO418815 | OG0 |
                              Coemet Cmet scaffold21238 01 | OG0 |
                    56
                            <sup>100</sup>1-Coeast_Cast_sb779390 | OG0 |
                      80
                            Lampan_Lpan_C412115_01 | OG0 |
                                Eupdun_Edun_sb10636030 | OG0 |
                                 -Beraby_Baby_sb337287 | OG0 |
                                 •Mlei ML0010.g5.i1 | OG0 |
         Clacor_Ccor_C530537_01 | OG0 |
         Lcom | lcpid84386 | OG0 |
   99
              •Scil scict000705.1 | OG0 |
     100
              -Gracom_Gcom_C527510_01 | OG0 |
              •Leuniv_Lniv_scaffold13906_01 | OG0 |
              Bralan BLAG03001195_1 | OG0 |
              Hsap ENST00000263856 | OG0 |
            <sup>100</sup>L-Hsap_ENST00000604011 | OG0 |
 88
                •Acagra_model.g673.t1 | OG0 |
                 •Cgig XM 011445719.3 | OG0 |
                    •Tcas TC012251 001 | OG0 |
                        -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA216523 | OG0 |
                        -Cscu_XM_023354687.1 | OG0 |
                 •Spur_XM_011676406.2 | OG0 |
   93
              •Xesp 013903-T1 | OG0 |
          <sup>100</sup> Dgig_XM_028535359.1 | OG0 |
                 •Nvec v1g53582 | OG0 |
                  •Exapal_XM_021057996.1 | OG0 |
                •Gfas gfas1.m1.14474.m1 | OG0 |
          86
      48
                    Amil Amillepora01047-RA | OG0 |
            100
                 <sup>100</sup> Adig XM_015907506.1 | OG0 |
                      -Fspp ffun1.m4.8284.m1 | OG0 |
               98
                    <sup>99</sup> Gasp gasp1.m3.27996.m1 | OG0 |
                       Pocdam_XM_027185773.1 | OG0 |
        15
                    1001-Spis XP 022786013 1 | OG0 |
                Sanmal Sma 008689-T1 | OG0 |
                  Aaur scaffold329.g22.t1 | OG0 |
                   -Rhoesc_Res_003344-T1 | OG0 |
                 85 Nemnom_g9809.t1 | OG0 |
          62
                 •Morvir scaffold11.g23.t1 | OG0 |
                    •Chem TCONS 00004649 | OG0 |
                       <sub>•</sub>Hvul_g30858_1 | OG0 |
            50
                     <sup>1001</sup>Hvul _g30863_1 | OG0 |
                              -HoiH11 mRNA584 | OG0 |
              40
                                *Tadh TriadT50591 | OG0 |
                             TrH2_TrispH2_005146-RA | OG0 |
                             HoiH06 mRNA2486 | OG0 |
                               <sub>F</sub>HoiH23_PIH23_003269-RA | OG0 |
                                  <sub>r</sub>Hhon_g02262.t1 | OG0 |
                               <sup>100</sup> HoiH04_mRNA5809 | OG0 |
0.416667
```