```
•Pbac sb2650947 | OG0 |
                           -Pbac_sb2655152 | OG0 |
                             Horcal_Hcv1.av93.c6.g74.i1 | OG0 |
                 100
                               •Pbac_sb2655151 | OG0 |
                              Beraby Baby sb340674 | OG0 |
                               •Mlei_ML0160.g4.i1 | OG0 |
                                •Lampan Lpan C418059 01 | OG0 |
                       84
 65
                                      •Coemet_Cmet_scaffold13606_01 | OG0 |
                                     •Vmul_sbMO407323 | OG0 |
                                     •Plespi Pspi scaffold16257 01 | OG0 |
                                  •Tetwil g4461.t1 | OG0 |
                           99
                                   Stycar_maker-SC_scaffold629-augustus-gene-0.3-mRNA-1 | OG0 |
              100
                                   •Emue Em0011g6a.t1 | OG0 |
                     100
                                    -Spolac c102294 g1 i4 m.28724c102294-g1 | OG0 |
                                 <sup>1001</sup>Spolac c102294 g1 i1 m.28718c102294-g1 | OG0 |
                 96
                                              Xestes maker-XT scaffold12993-augustus-gene-0.2-mRNA-1 | OG0 |
                                                   -Aque_Aqu2.1.03151_001 | OG0 |
                                     100
                                                <sup>1001</sup>-Aque_Aqu2.1.22804_001 | OG0 |
                                                   Halamb c54751 g2 i7mm.6128 | OG0 |
                                                <sup>1001</sup>-Halamb_c52248_g1_i4m.32549 | OG0 |
                                  Opea EP00118 Oscarella pearsei P011396 | OG0 |
               100
                                •Plajan Pjan scaffold21135 01 | OG0 |
                            Clacor_Ccor_scaffold5594_01 | OG0 |
    90
                                •Leuniv Lniv scaffold22822 01 | OG0 |
                  100
                                   -Scil scict017993.3 | OG0 |
                          100
                                      •Gracom_Gcom_scaffold8225_01 | OG0 |
                                      •Lcom_lcpid77986 | OG0 |
                    •Exapal XM 021044110.2 | OG0 |
                  Nvec_v1g100430 | OG0 |
           100
                <sup>100</sup>1•Nvec_NVE23246 | OG0 |
                    •Gfas gfas1.m1.396.m1 | OG0 |
        100
                      Adig_XM_015901733.1 | OG0 |
                   <sup>100</sup> -Amil_Amillepora25088-RA | OG0 |
            100
                       •Gasp gasp1.m3.4979.m1 | OG0 |
                      -Fspp ffun1.m4.11131.m1 | OG0 |
                     •Pocdam XM 027195379.1 | OG0 |
                  100 Spis_XP_022779759_1 | OG0 |
                       Hsap ENST00000354777 | OG0 |
                52
                                                     -Cscu XM 023371007.1 | OG0 |
             52
                    Bralan_BLAG13001040_1 | OG0 |
     84
                                Bralan BLAG13001087 1 | OG0 |
                       Skow XM 002735686.2 | OG0 |
               88
                           •Spur_XM_030996175.1 | OG0 |
                               Ctel gnl WGS AMQN CAPTEDRAFT mRNA179639 | OG0 |
                               Cgig XM 011417498.3 | OG0 |
                    99
                                •Acagra_model.g8154.t1 | OG0 |
                             Cscu XM 023371023.1 | OG0 |
                  89
                                    •Tcas_TC005345_001 | OG0 |
       50
                                 Dpul mRNA3386 | OG0 |
                              <sup>100</sup> Dpul_mRNA3387 | OG0 |
                          •Xesp_009341-T1 | OG0 |
                        •Dgig_XM_028538935.1 | OG0 |
                                 •HoiH23 PIH23 011225-RA | OG0 |
                                  Hhon g03522.t1 | OG0 |
                            100
                                <sup>100</sup><sup>1</sup>-HoiH04_mRNA6646 | OG0 |
         29
                         100
                                -•Tadh_TriadT57871 | OG0 |
                                 TrH2_TrispH2_001416-RA | OG0 |
                                <sup>601</sup>Tadh_wf_g6748.t1 | OG0 |
           31
                                        •Hvul g18842 1 | OG0 |
                             100
                                        •Chem_TCONS_00053970 | OG0 |
                                 •Morvir scaffold119.g27.t1 | OG0 |
                   100
                               Aaur scaffold6039.g1.t1 | OG0 |
                                       •Sanmal_Sma_011895-T1 | OG0 |
                                           •Aaur scaffold265.g19.t1 | OG0 |
                                   •Rhoesc_Res_016478-T1 | OG0 |
                               99<sup>1</sup>-Nemnom g6594.t1 | OG0 |
0.416667
```