```
•Emue_Em0023g338a.t1 | OG0 |
                  -Spolac_c102139_g1_i2_m.28089c102139-g1 | OG0 |
                       Xestes maker-XT scaffold158192-augustus-gene-0.2-mRNA-1 | OG0 |
        97
                      •Halamb c52204_g2_i2mm.4398 | OG0 |
                  •Tetwil g11103.t1 | OG0 |
          88
                 •Stycar_snap_masked-SC_scaffold140251-processed-gene-0.9-mRNA-1 | OG0 |
               | Stycar maker-SC scaffold133857-augustus-gene-0.20-mRNA-1 | OG0
 76
                  •Plajan Pjan scaffold21118 01 | OG0 |
                Popea EP00118 Oscarella pearsei P003999 | OG0 |
        100
              100 Ocar q2068 | OG0 |
   43
                                                                         Drygla Dgla sb277458 | OG0 |
                            Eupdun Edun sb10653108 | OG0 |
                              Lampan Lpan C414005 01 | OG0 |
                 98
                                        -Pbac sb2644102 | OG0 |
                    64
                                    <sup>100</sup>1•Horcal _Hcv1.av93.c11.g1043.i1 | OG0 |
                                    Beraby Baby sb328912 | OG0 |
                                  •Mlei ML0144.g15.i1 | OG0 |
                            99
                                    Bolinf Binf sb1225888 | OG0 |
                                  Bolinf Binf sb1225885 | OG0 |
                                •Hvul_g28274_1 | OG1 |
               100
                               •Chem TCONS 00000990 | OG1 |
                   •Morvir scaffold369.g7.t2 | OG1 |
        100
                       Sanmal Sma 007612-T1 | OG1 |
                        -Aaur scaffold474.g13.t1 | OG1 |
              100
                       Rhoesc Res 011698-T1 | OG1 |
                 96
   65
                        •Nemnom_g13564.t1 | OG1 |
                        •Dgig XM 028550323.1 | OG1 |
                            •Xesp 000244-T1 | OG1 |
                    Exapal XM 021037948.2 | OG1 |
            98
      92
                  •Nvec v1g137654 | OG1 |
                    Gfas_gfas1.m1.23318.m1 | OG1 |
         94
                     Amil Amillepora23393-RA | OG1 |
                  <sup>100]</sup>•Adig_XM_015900371.1 | OG1 |
                      Gasp gasp1.m3.14454.m1 | OG1 |
             100
                  <sup>95</sup>-Fspp_ffun1.m4.685.m1 | OG1 |
                     -Spis XP 022790688 1 | OG1 |
                  <sup>100</sup> Pocdam XM 027203460.1 | OG1 |
                                                                         •Plajan Pjan scaffold23136 01 | OG1 |
                         TrH2 TrispH2 001488-RA | OG1 |
                       <sup>99</sup> Tadh TriadT57947 | OG1 |
                          •HoiH11 mRNA3718 | OG1 |
                    98
                             -HoiH23 PIH23 001111-RA | OG1 |
                           <sup>99</sup> HoiH06 mRNA7039 | OG1 |
                              Hhon g00264.t1 | OG1 |
                           100 HoiH04_mRNA6510 | OG1 |
                     Skow XM 002730318.2 | OG1 |
                       •Spur XM 030982811.1 | OG1 |
                   Bralan BLAG02000891 1 | OG1 |
                      •Hsap ENST00000360540 | OG1 |
                         •Hsap_ENST00000308824 | OG1 |
         95
                      Cscu_XM_023362418.1 | OG1 |
                             •Dpul mRNA10090 | OG1 |
                 88
                             -Tcas TC001944 001 | OG1 |
                       Cgig XM 020071961.2 | OG1 |
                      •Acagra_model.g24498.t1 | OG1 |
                         -Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA180377 | OG1 |
                                  •Ctel gnl WGS AMQN_CAPTEDRAFT_mRNA197057 | OG1 |
0.416667
```