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Clacor_Ccor_C540681_01 | OG0 |
                  -Leuniv_Lniv_scaffold20516_01 | OG0 |
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                    Lcom_lcpid95750 | OG0 |
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                      Gracom_Gcom_C554662_01 | OG0 |
                       Scil_scict022803.1 | OG0 |
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                                 •Lcom | lcpid161171 | OG1 |
                         •Vmul_sbMO447334 | OG1 |
                          •Coemet_Cmet_scaffold2634_01 | OG1 |
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                       <sup>100</sup> Coeast_Cast_sb771576 | OG1 |
                 92
                          Lampan Lpan C407573 01 | OG1 |
                                <sub>•</sub>Pbac_sb2667816 | OG1 |
                   88
                             <sup>100</sup>-Horcal_Hcv1.av93.c2.g619.i1 | OG1 |
                             Drygla_Dgla_sb301600 | OG1 |
                                Beraby_Baby_sb348165 | OG1 |
                                •Mlei ML0053.g11.i1 | OG1 |
                                Mlei ML0053.g10.i1 | OG1 |
                              <sup>96</sup>Vmul_sbMO392957 | OG1 |
                   •Plespi_Pspi_scaffold8261_01 | OG2 |
                       Xestes_maker-XT_scaffold4896-augustus-gene-0.38-mRNA-1 | OG2 |
                   Aque_Aqu2.1.32106_001 | OG2 |
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                   Halamb_c46612_g1_i1m.20914 | OG2 |
                                   Stycar_snap_masked-SC_scaffold7416-processed-gene-0.4-mRNA-1 | OG2 |
     81
                                      •Lcom_lcpid204213 | OG2 |
                    •Tetwil_g23948.t3 | OG2 |
               Emue_Em0021g321a.t1 | OG2 |
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                     •Emue Em0021g319a.t1 | OG2 |
                        •Lcom_lcpid210499 | OG3 |
                  •Plajan_Pjan_C548933_01 | OG3 |
                  Popea EP00118 Oscarella pearsei P027461 | OG3 |
                <sup>991</sup>⊸Ocar_g6887 | OG3 |
                -Dgig_XM_028552650.1 | OG3 |
                   •Xesp_008512-T1 | OG3 |
                  Exapal XM 021050918.2 | OG3 |
                   Nvec v1g13494 | OG3 |
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                 <sup>1001</sup>Nvec_v1g150238 | OG3 |
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          98
                     •Gfas_gfas1.m1.6855.m1 | OG3 |
                   •Spis_XP_022780810_1 | OG3 |
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                 <sup>99</sup>Ū-Pocdam_XM_027200316.1 | OG3 |
                      •Fspp_ffun1.m4.13003.m1 | OG3 |
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                      •Gasp gasp1.m3.19484.m1 | OG3 |
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                        r-Amil_Amillepora05059-RA | OG3 |
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                   Hsap ENST00000025399 | OG3 |
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                       •Bralan_BLAG18000685_1 | OG3 |
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                         Ctel gnl WGS AMQN CAPTEDRAFT mRNA229092 | OG3 |
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                          Cscu XM 023381899.1 | OG3 |
                              •Dpul_mRNA21540 | OG3 |
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                              •Tcas_TC006736_001 | OG3 |
                           •Hvul_g12789_1 | OG3 |
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                      •Morvir_scaffold379.g6.t1 | OG3 |
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                          •Sanmal_Sma_017429-T1 | OG3 |
                            •Nemnom_g8561.t1 | OG3 |
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                       <sup>100</sup><sup>1</sup>-Rhoesc_Res_002229-T1 | OG3 |
                         •HoiH06_mRNA13336 | OG3 |
                           •HoiH23 PIH23 002139-RA | OG3 |
                               <sub>e</sub>Hhon_g04697.t1 | OG3 |
                             <sup>36</sup> HoiH04_mRNA10469 | OG3 |
                                •HoiH11 mRNA5632 | OG3 |
                                  TrH2 TrispH2 002157-RA | OG3 |
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0.416667
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