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•Morvir_scaffold77.g63.t1 | OG0 |
           •Chem_TCONS_00055266 | OG0 |
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            •Hvul_g30984_1 | OG0 |
           •Sanmal_Sma_025368-T1 | OG0 |
             <sub>f</sub>Nemnom_g18249.t1 | OG0 |
          <sup>1001</sup>•Rhoesc_Res_017655-T1 | OG0 |
          •Plajan_Pjan_C533931_01 | OG0 |
            100 Ocar_g7184 | OG0 |
           Coemet_Cmet_scaffold21177 01 | OG0 |
   86
              Nmul sbMO400166 | OG0 |
           <sup>1001</sup>-Vmul_Simscaffold14837_01 | OG0 |
           <sub>-</sub>-Lampan_Lpan_C402229_01 | OG0 |
      100
                <sub>!</sub>Beraby_Baby_sb341198 | OG0 |
             <sup>1001</sup>•Drygla_Dgla_sb316289 | OG0 |
                  Drygla Dgla sb290211 | OG0 |
                        •Pbac_sb2648555 | OG0 |
           91
               80
                     <sup>72</sup> Pbac_sb2655622 | OG0 |
                       Horcal_Hcv1.av93.c5.g347.i1 | OG0 |
                     78 Coeast Cast sb773865 | OG0 |
             92
                     Bolinf Binf sb1230838 | OG0 |
                  <sup>100</sup>¹•Mertsp_Mesp_sb578454 | OG0 |
                     •Bolinf_Binf_sb1226099 | OG0 |
                       <sub>f</sub>Mlei_ML0844.g29.i1 | OG0 |
                         <sub>s</sub>Vmul_sbMO443215 | OG0 |
                       83<sup>1</sup>-Vmul_sbMO443212 | OG0 |
         •Spur_XM_030997941.1 | OG0 |
            •Lcom_lcpid153494 | OG0 |
               •Lcom_lcpid169287 | OG0 |
         Clacor_Ccor_scaffold10733_01 | OG0 |
            Gracom Gcom scaffold13914 01 | OG0 |
          <sup>90</sup><sup>-</sup>Scil _scict002425.1 | OG0 |
            -Gracom_Gcom_C537164_01 | OG0 |
              Lcom_lcpid116901 | OG0 |
            <sup>951</sup>-Lcom | lcpid115689 | OG0 |
           •Skow XM 002736012.2 | OG0 |
             •Bralan_BLAG05000642_1 | OG0 |
                eHsap_ENST00000433034 | OG0 |
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                    •Hsap ENST00000397256 | OG0 |
                •Cscu XM 023371841.1 | OG0 |
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                •Tcas TC015671 001 | OG0 |
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                   •Lcom_lcpid118477 | OG0 |
                   •Cgig_XM_011449511.3 | OG0 |
                <sub>[</sub>TrH2_TrispH2_000264-RA | OG0 |
             <sup>991</sup>Tadh TriadT38152 | OG0 |
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                  •HoiH04_mRNA9183 | OG0 |
                    <sup>e</sup>HoiH11 mRNA1582 | OG0 |
                       <sub>•</sub>HoiH23 PIH23 003095-RA | OG0 |
                    <sup>1001</sup>HoiH06_mRNA13594 | OG0 |
              •Acagra_model.g8552.t1 | OG0 |
            <sub>r</sub>-Nvec_v1g236559 | OG0 |
          <sup>97</sup>└-Exapal_XM_021038778.2 | OG0 |
             Gasp_gasp1.m3.24264.m1 | OG0 |
     53
               Spis_XP_022803938 1 | OG0 |
                 Gasp_gasp1.m3.8489.m1 | OG0 |
               <sup>41</sup> Pocdam_XM_027199884.1 | OG0 |
              Fspp_ffun1.m4.32984.m1 | OG0 |
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                    •Gfas_gfas1.m1.334.m1 | OG0 |
                     Amil Amillepora31902-RA | OG0 |
                   <sup>99</sup> Adig_XM_015914227.1 | OG0 |
                       -Dgig_XM_028536975.1 | OG0 |
                   <sup>[100]</sup>•Xesp_005516-T1 | OG0 |
                      •Tetwil g14048.t2 | OG0 |
                          Halamb c34809 g1 i1m.8333 | OG0 |
                             -Aque Aqu2.1.43637 001 | OG0 |
                          <sup>91]</sup>Xestes_maker-XT_scaffold1342-augustus-gene-0.37-mRNA-1 | OG0 |
                           •Plespi_Pspi_scaffold18919_01 | OG0 |
                             Emue_Em0014g372a.t1 | OG0 |
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0.416667
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