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Mbre EDQ92087 | OG0 |
              •Chofle_EP00730_Choanoeca_flexa_P004237 | OG0 |
    100
             •Chofle EP00730 Choanoeca flexa P004238 | OG0 |
       100
                •Choper_EP00045_Choanoeca_perplexa_P000492 | OG0 |
          •Salkve_EP00050_Salpingoeca_kvevrii_P006822 | OG0 |
 100
             Codhol EP00042 Codosiga hollandica P028376 | OG0 |
               •Salinf_EP00049_Salpingoeca_infusionum_P026154 | OG0 |
    95
                •Microa_EP00043_Microstomoeca_roanoka_P008827 | OG0 |
        73
           <sup>55</sup>-Sros EGD82757 | OG0 |
             Saldol EP00054_Salpingoeca_dolichothecata_P010183 | OG0 |
      70
                Mylflu EP00047 Mylnosiga fluctuans P001840 | OG0 |
        43
                    Salurc_EP00051_Salpingoeca_urceolata_P004827 | OG0 |
                       Salmac_EP00052_Salpingoeca_macrocollata_P030659 | OG0 |
           33
                       •Salpun_EP00053_Salpingoeca_punica_P000999 | OG0 |
             58
                    Didcos EP00039 Didymoeca costata P019091 | OG0 |
                         Acaspe EP00035 Acanthoeca spectabilis P021161 | OG0 |
                           -Helnan_EP00037_Helgoeca_nana_P011670 | OG0 |
                           -Stedip EP00041 Stephanoeca diplocostata P008243 | OG0 |
                     <sup>100</sup><sup>1</sup>⊸Diagra EP00040 Diaphanoeca grandis P017191 | OG0 |
         Aaur scaffold148.g13.t1 | OG0 |
            Nemnom g3054.t1 | OG0 |
          <sup>100</sup><sup>1</sup>•Rhoesc_Res_004312-T1 | OG0 |
    93
         •Morvir_scaffold250.g21.t1 | OG0 |
              Chem_TCONS_00017543 | OG0 |
      25
               --Hvul_g28_1 | OG0 |
              Clacor Ccor scaffold12097 01 | OG0 |
        80
                    Lcom_lcpid86962 | OG0 |
           100
                      -Scil_scict018977.1 | OG0 |
                  •Scil_scict010925.1 | OG0 |
                99 Lcom | lcpid114492 | OG0 |
          Exapal XM 021037553.2 | OG0 |
        <sup>99</sup> Nvec_v1g101531 | OG0 |
 100
             <sub>•</sub>Dgig_XM_028537688.1 | OG0 |
      91
          <sup>941</sup>•Xesp_021512-T1 | OG0 |
               -•Adig XM 015897301.1 | OG0 |
        70
             95
                     -Gasp gasp1.m3.12704.m1 | OG0 |
               Spis XP 022802457 1 | OG0 |
                  Fspp_ffun1.m4.25964.m1 | OG0 |
                 <sup>1</sup> Pocdam_XM_027189418.1 | OG0 |
                   Gfas_gfas1.m1.14529.m1 | OG0 |
                     Amil Amillepora25170-RA | OG0 |
                   <sup>33]</sup>Amil_Amillepora25174-RA | OG0 |
              Plespi Pspi scaffold15574 01 | OG0 |
           <sup>100</sup>Plespi_Pspi_scaffold15575_01 | OG0 |
              •Opea_EP00118_Oscarella_pearsei_P024948 | OG0 |
        79
                  •Tetwil g19867.t1 | OG0 |
                     Stycar_maker-SC_scaffold3995-augustus-gene-0.33-mRNA-1 | OG0 |
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                    Xestes_maker-XT_scaffold135537-augustus-gene-0.4-mRNA-1 | OG0 |
                     _*Aque_Aqu2.1.22337_001 | OG0 |
                   <sup>48]</sup>-Halamb_c38699_g1_i1m.11034 | OG0 |
               91
                    -Spolac_c95750_g1_i1_m.11976c95750-g1 | OG0 |
                       Emue Em0014g814a.t1 | OG0 |
      79
                    <sup>100</sup> Emue Em0014g785a.t1 | OG0 |
             Lcom_lcpid158400 | OG0 |
                     Lcom | lcpid190532 | OG0 |
          31
                  Spur XM 011680445.2 | OG0 |
               <sup>1001</sup>Spur_XM_030998703.1 | OG0 |
               •Cscu_XM_023375420.1 | OG0 |
                   <sub>"</sub>Tcas_TC033787_001 | OG0 |
        64
                 <sup>52</sup>-Dpul_mRNA13402 | OG0 |
                   Bralan BLAG04000351 1 | OG0 |
                       •Pbac_sb2644133 | OG0 |
                           /llei_ML1597.g4.i1 | OG0 |
                             Drygla_Dgla_sb287577 | OG0 |
           28
                       100
                          100<mark>1  PDrygla Dgla sb287576 | OG0 |</mark>
                             <sup>1</sup>•Drygla_Dgla_sb287578 | OG0 |
               Acagra_model.g17578.t1 | OG0 |
                  Cgig_XM_034468176.1 | OG0 |
                    •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA108511 | OG0 |
                       •Hsap ENST00000601498 | OG0 |
                         •Skow_XM_006824086.1 | OG0 |
                             HoiH04 mRNA1591 | OG0 |
                           80 HoiH06 mRNA3721 | OG0 |
                             •Hhon_g05177.t1 | OG0 |
                                  <sub>1</sub>+HoiH23_PIH23_001254-RA | OG0 |
                               <sup>36]</sup>HoiH11_mRNA3844 | OG0 |
                                  TrH2 TrispH2 006427-RA | OG0 |
                                <sup>99]</sup>Tadh TriadT52003 | OG0 |
0.416667
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