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<sub>!</sub>Plajan_Pjan_scaffold10015_01 | OG0 |
        <sup>100</sup> Plajan_Pjan_scaffold10016_01 | OG0 |
                 Stycar_maker-SC_scaffold81092-augustus-gene-0.108-mRNA-1 | OG0 |
                 •Tetwil_g3822.t2 | OG0 |
        100
                      •Spolac c95524 g1 i1 m.11610c95524-g1 | OG0 |
                   <sup>100</sup><sup>1</sup>-Emue_Em0003g46a.t1 | OG0 |
              98
                      •Plespi_Pspi_C585688_01 | OG0 |
                           •Xestes_maker-XT_scaffold23098-snap-gene-0.22-mRNA-1 | OG0 |
                86
                            <sub>l</sub>-Halamb_c25132_g1_i1m.4679 | OG0 |
                         <sup>1001</sup>Aque_Aqu2.1.35139_001 | OG0 |
                             Drygla Dgla sb284458 | OG0 |
                   49
                                  Horcal Hcv1.av93.c4.g82.i1 | OG0 |
                               <sup>100</sup> Pbac_sb2654467 | OG0 |
                                 -Lampan_Lpan_C394599_01 | OG0 |
                                     •Vmul SimC387240 | OG0 |
                                   <sup>97</sup> Coemet Cmet_scaffold16138_01 | OG0 |
                                     <sub>|</sub>-Mlei_ML0464.g7.i1 | OG0 |
                                        -Beraby_Baby_sb338263 | OG0 |
                               ·Clacor_Ccor_C530953_01 | OG0 |
                                    -Scil scict018236.1 | OG0 |
                            100
                                  <sup>83</sup>-Lcom | lcpid80454 | OG0 |
                               100
                                    -Gracom_Gcom_scaffold14653_01 | OG0 |
                       49
                                  88 Leuniv_Lniv_scaffold23606_01 | OG0 |
                                  Lcom_lcpid146471 | OG0 |
                                      Codhol EP00042 Codosiga hollandica P018954 | OG0 |
                                   100
                                         •Codhol EP00042 Codosiga hollandica P018957 | OG0 |
                                          Saldol_EP00054_Salpingoeca_dolichothecata_P004910 | OG0 |
                              78
                                            Salurc EP00051 Salpingoeca urceolata P008938 | OG0 |
                                                •Salurc_EP00051_Salpingoeca_urceolata_P008939 | OG0 |
                                           -Stedip_EP00041_Stephanoeca_diplocostata_P019239 | OG0 |
                                    40
                                         <sup>691</sup>-Didcos EP00039_Didymoeca_costata_P028805 | OG0 |
                         81
                                      99
                                               Diagra_EP00040_Diaphanoeca_grandis_P001271 | OG0 |
                                              Acaspe_EP00035_Acanthoeca_spectabilis_P020706 | OG0 |
                                           1001-Helnan EP00037_Helgoeca_nana_P032337 | OG0 |
                                           Salkve_EP00050_Salpingoeca_kvevrii P011198 | OG0 |
                                  32
                                               Lcom | lcpid226399 | OG0 |
                                             •Mylflu_EP00047_Mylnosiga_fluctuans_P002515 | OG0 |
                                               Salpun EP00053 Salpingoeca punica P003809 | OG0 |
                                                <sub>!</sub>Salmac_EP00052_Salpingoeca_macrocollata_P024716 | OG0 |
                                             Galmac EP00052 Salpingoeca_macrocollata_P024715 | OG0 |
                                    78
                                             •Mbre EDQ88282 | OG0 |
                                               •Choper_EP00045_Choanoeca_perplexa_P020842 | OG0 |
                                            100<sup>1</sup>-Chofle_EP00730_Choanoeca_flexa_P019278 | OG0 |
                                             Salinf_EP00049_Salpingoeca_infusionum_P024690 | OG0 |
                                              -Sros EGD80568 | OG0 |
                                           100<sup>1</sup>-Microa EP00043_Microstomoeca_roanoka_P014784 | OG0 |
                                       -Skow_XM_002742135.2 | OG0 |
                                    63
                                         -Spur_XM_775701.5 | OG0 |
                                  66
                                         •Bralan BLAG03000791 1 | OG0 |
                                         •Hsap ENST00000551671 | OG0 |
                                        TrH2 TrispH2 010901-RA | OG0 |
                                     <sup>75</sup> Tadh_TriadT51633 | OG0 |
                                        <sub>e</sub>HoiH23_PIH23_003668-RA | OG0 |
                                         HoiH04_mRNA15520 | OG0 |
                                           *HoiH06 mRNA14482 | OG0 |
                                          <sup>1</sup> Hhon g01993.t1 | OG0 |
                                          •Tcas TC008443 001 | OG0 |
                                          -Cscu_XM_023384932.1 | OG0 |
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                                       86<sup>1</sup>-Dpul_mRNA2334 | OG0 |
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                                         Ctel gnl WGS AMQN CAPTEDRAFT mRNA228707 | OG0 |
                                          -Lcom | lcpid202002 | OG0 |
                                           —•Cgig_XM_011417915.3 | OG0 |
                                         <sup>⊌</sup>I ---•Acagra_model.g1227.t1 | OG0 |
                                        •Exapal_XM_021043840.1 | OG0 |
                                         Fspp ffun1.m4.27826.m1 | OG0 |
                                             •Nvec v1g223751 | OG0 |
                                                  Gasp_gasp1.m3.11828.m1 | OG0 |
                                             <sup>55</sup> Gfas_gfas1.m1.2889.m1 | OG0 |
                                               Amil_Amillepora13185-RA | OG0 |
                                                Pocdam XM 027185347.1 | OG0 |
                                  78
                                                <sup>981</sup>•Spis XP 022785674_1 | OG0 |
                                          <sub>-</sub>Xesp_024250-T1 | OG0 |
                                             Dgig_XM_028560947.1 | OG0 |
                                          <sup>100</sup> Dgig XM 028556815.1 | OG0 |
                                            Opea EP00118 Oscarella pearsei P025495 | OG0 |
                                              •Hvul_g2195 1 | OG0 |
                                               •Morvir_scaffold323.g12.t1 | OG0 |
                                                -Sanmal_Sma_024378-T1 | OG0 |
                                                 Aaur_scaffold500.g3.t1 | OG0 |
                                                    Nemnom_g13227.t1 | OG0 |
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0.416667
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