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•Mylflu_EP00047_Mylnosiga_fluctuans_P002407 | OG0 |
               Salmac_EP00052_Salpingoeca_macrocollata_P002583 | OG0 |
         Salkve EP00050 Salpingoeca kvevrii P005878 | OG0 |
            ·Salpun_EP00053_Salpingoeca_punica_P007620 | OG0 |
                Didcos EP00039 Didymoeca costata P001844 | OG0 |
                      Saldol_EP00054_Salpingoeca_dolichothecata_P032112 | OG0 |
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                 Choper_EP00045_Choanoeca_perplexa_P006936 | OG0 |
                     Salinf_EP00049_Salpingoeca_infusionum_P017460 | OG0 |
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                80
                                 •Lcom_lcpid157972 | OG0 |
                           •Microa_EP00043_Microstomoeca_roanoka_P006669 | OG0 |
                       <sup>100</sup> Sros EGD80874 | OG0 |
     Codhol_EP00042_Codosiga_hollandica_P027455 | OG1 |
   <sup>991</sup>-Codhol_EP00042_Codosiga_hollandica_P027456 | OG1 |
              Salurc_EP00051_Salpingoeca_urceolata_P032104 | OG1 |
              Diagra_EP00040_Diaphanoeca_grandis_P030750 | OG1 |
                 Acaspe_EP00035_Acanthoeca_spectabilis_P020086 | OG1 |
               <sup>100</sup> Helnan EP00037_Helgoeca_nana_P029811 | OG1 |
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                •Dpul mRNA1553 | OG1 |
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            Cgig_XM_011450537.3 | OG1 |
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             •Skow XM 002741625.2 | OG1 |
                                -Lcom_lcpid253582 | OG1 |
                        Lcom_lcpid10273 | OG1 |
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                        Fspp ffun1.m4.3534.m1 | OG1 |
                        Gasp_gasp1.m3.14868.m1 | OG1 |
                        •Spis_XP_022803007_1 | OG1 |
                    <sup>100</sup> Pocdam XM 027191793.1 | OG1 |
                  <sub></sub>-Acagra_model.g28919.t1 | OG1 |
                <sup>981</sup>•Acagra_model.g28919.t1.1.5d3b8c72 | OG1 |
                        Dgig_XM_028536243.1 | OG1 |
                     <sup>98</sup> Xesp_028504-T1 | OG1 |
                         •Mlei_ML0117.g27.i1 | OG1 |
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                          Horcal Hcv1.av93.c8.g163.i1 | OG1 |
                        <sup>100</sup>lPbac_sb2663345 | OG1 |
                        •Hvul_g6104_1 | OG1 |
                                 •Coemet_Cmet_scaffold16020_01 | OG1 |
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                              •Mlei_ML2582.g47.i1 | OG1 |
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                               Horcal_Hcv1.av93.c1.g244.i1 | OG1 |
                            <sup>100</sup> Pbac_sb2638267 | OG1 |
                     46
                               •Morvir_scaffold243.g28.t1 | OG1 |
                               -Xestes_maker-XT_scaffold135647-augustus-gene-0.14-mRNA-1 | OG1 |
                                 <sub>-</sub>-Aque_Aqu2.1.38447_001 | OG1 |
                              <sup>100</sup>□-Halamb_c33449_g1_i1m.7661 | OG1 |
                                •Tetwil_g4965.t1 | OG1 |
                                     •Spolac_c9264_g1_i1_m.322c9264-g1 | OG1 |
                                  <sup>1001</sup>•Emue_Em0018g51a.t1 | OG1 |
                       •Bralan_BLAG07000869_1 | OG1 |
                             •Exapal_XM_021059962.2 | OG1 |
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                                     •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA219154 | OG1 |
                                      Lcom lcpid184470 | OG1 |
                           -Gfas_gfas1.m1.23595.m1 | OG1 |
                          <sub>-</sub>-Amil_Amillepora10840-RA | OG1 |
                             -Adig_XM_015899369.1 | OG1 |
                          •Nemnom_g18174.t1 | OG1 |
                        <sup>981</sup>•Rhoesc_Res_010867-T1 | OG1 |
                          Sanmal Sma 025577-T1 | OG1 |
                                   Opea_EP00118_Oscarella_pearsei_P025076 | OG1 |
                                ____Ocar_g3901 | OG1 |
                        58
                                   TrH2 TrispH2 005331-RA | OG1 |
                          84
                                     -Tadh wf g4578.t1 | OG1 |
                                     Hhon_g00981.t1 | OG1 |
                                   <sup>981</sup>HoiH04_mRNA13264 | OG1 |
                                     +HoiH11 mRNA7082 | OG1 |
                                   90 HoiH23 PIH23 001634-RA | OG1 |
                                     <sup>70]</sup>HoiH06 mRNA5267 | OG1 |
0.416667
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