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·Codhol_EP00042_Codosiga_hollandica_P040336 | OG0 |
             -Salkve_EP00050_Salpingoeca_kvevrii_P003420 | OG0 |
              Salurc_EP00051_Salpingoeca_urceolata_P004902 | OG0 |
                ·Salmac_EP00052_Salpingoeca_macrocollata_P027258 | OG0 |
100
               Salpun EP00053 Salpingoeca punica P000247 | OG0 |
                47
           -Saldol_EP00054_Salpingoeca_dolichothecata_P026548 | OG0 |
                  Didcos_EP00039_Didymoeca_costata_P033327 | OG0 |
                     Diagra_EP00040_Diaphanoeca_grandis_P013143 | OG0 |
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                    Stedip EP00041 Stephanoeca diplocostata P034260 | OG0 |
    38
                        •Acaspe_EP00035_Acanthoeca_spectabilis_P034587 | OG0 |
                        •Helnan_EP00037_Helgoeca_nana_P029536 | OG0 |
                  Salinf_EP00049_Salpingoeca_infusionum_P009578 | OG0 |
                    Microa_EP00043_Microstomoeca_roanoka_P020170 | OG0 |
                    •Sros EGD79932 | OG0 |
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                  Mbre EDQ84603 | OG0 |
                   -Chofle_EP00730_Choanoeca_flexa_P004571 | OG0 |
                    Choper_EP00045_Choanoeca_perplexa_P022700 | OG0 |
              •Lcom_lcpid1561 | OG0 |
               •Gracom_Gcom_scaffold17801_01 | OG0 |
                   Leuniv Lniv C801922 01 | OG0 |
             •Plajan_Pjan_scaffold15299_01 | OG0 |
  91
       100
             •Ocar_g8918 | OG0 |
          <sup>100</sup>l<sup>.</sup>Opea_EP00118_Oscarella_pearsei_P005341 | OG0 |
             Plespi_Pspi_scaffold20527_01 | OG0 |
    90
                 -Xestes maker-XT scaffold85870-snap-gene-0.3-mRNA-1 | OG0 |
                    •Halamb_c55952_g1_i1m.42139 | OG0 |
       100
                    •Aque_Aqu2.1.29852_001 | OG0 |
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                   •Tetwil_g6955.t2 | OG0 |
                     •Spolac_c99660_g1_i2_m.20265c39806-g1 | OG0 |
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                  <sup>100</sup>-Spolac c99660 g1_i3_m.20266c39806-g1 | OG0 |
                    <u>•</u>Emue_Em0015g744a.t1 | OG0 |
                     --Emue_Em0114g8a.t1 | OG0 |
               Eupdun_Edun_sb10665410 | OG0 |
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                  •Vmul SimC407394 | OG0 |
                    Coemet Cmet scaffold22583 01 | OG0 |
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                       Coeast_Cast_sb796027 | OG0 |
                     00 Coeast_Cast_sb796028 | OG0 |
            86
                   eHorcal_Hcv1.av93.c3.g1105.i1 | OG0 |
                <sup>100</sup> Pbac sb2639717 | OG0 |
                    Beraby Baby sb336145 | OG0 |
                     Bolinf_Binf_sb1267968 | OG0 |
                       Mertsp_Mesp_sb602649 | OG0 |
                     46 Bolinf_Binf_sb1267967 | OG0 |
                 •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA221643 | OG0 |
  93
                   Cgig XM 034443049.1 | OG0 |
          100
                  Acagra_model.g14549.t1.1.5d3b87c0 | OG0 |
                <sup>100</sup>-Acagra_model.g14549.t1 | OG0 |
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                Bralan_BLAG01002578_1 | OG0 |
                      •Hsap ENST00000673944 | OG0 |
                    Skow XM 006817118.1 | OG0 |
                     Spur XM 030992114.1 | OG0 |
                  <sup>100</sup>•Spur_XM_030975036.1 | OG0 |
                   Tadh_TriadT50643 | OG0 |
    98
                •HoiH11 mRNA6007 | OG0 |
              100
                         •HoiH06_mRNA7945 | OG0 |
                     <sup>100</sup> HoiH23_PIH23_000673-RA | OG0 |
                        Hhon g02655.t1 | OG0 |
                      <sup>1001</sup>HoiH04 mRNA16884 | OG0 |
                          TCONS 00024281 | OG0 |
       81
               100
                         -Hvul g23589 1 | OG0 |
                           scaffold32.g49.t1 | OG0 |
                      Sanmal_Sma_019300-T1 | OG0 |
                        Aaur scaffold436.g17.t1 | OG0 |
                          -Rhoesc Res 006593-T1 | OG0 |
         100
                       <sup>100</sup>-Nemnom g13644.t1 | OG0 |
                  -Dgig_XM_028540957.1 | OG0 |
               <sup>100</sup><sup>1</sup>⊸Xesp_004866-T1 | OG0 |
                   •Nvec v1g245829 | OG0 |
                     Spis XP 022781464 1 | OG0 |
                         •Gasp_gasp1.m3.21421.m1 | OG0 |
                        Fspp_ffun1.m4.6206.m1 | OG0 |
                          Amil_Amillepora22267-RA | OG0 |
                       <sup>100</sup><sup>1</sup>-Adig XM 015903213.1 | OG0 |
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