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Lampan Lpan 12243 01 | OG0 |
                  •Vmul SimC383314 | OG0 |
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                  Coemet_Cmet_scaffold406_01 | OG0 |
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                <sup>100</sup> Coemet_Cmet_scaffold407_01 | OG0 |
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                Beraby Baby sb332002 | OG0 |
                  •Mlei ML2666.g3.i1 | OG0 |
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                     Drygla_Dgla_sb294727 | OG0 |
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                  <sup>100</sup>-Drygla_Dgla_sb294728 | OG0 |
                             -Pbac_sb2663124 | OG0 |
               43
                       Horcal Hcv1.av93.c4.g897.i3 | OG0 |
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                                  •Pbac sb2655345 | OG0 |
                           Horcal Hcv1.av93.c4.g897.i1 | OG0 |
                         31 Horcal_Hcv1.av93.c4.g897.i2 | OG0 |
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                Opea EP00118 Oscarella pearsei P024676 | OG0 |
             100 L
                  •Ocar_g1711 | OG0 |
                          •Plespi Pspi scaffold1441 01 | OG0 |
                             •Spolac_c97579_g1_i1_m.15334c97579-g1 | OG0 |
                                 Emue Em0016g1081a.t1 | OG0 |
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                         100
                             <sup>1001</sup>Emue Em0016g1099a.t1 | OG0 |
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                             Stycar maker-SC scaffold2898-augustus-gene-0.15-mRNA-1 | OG0 |
                                 •Halamb_c50630_g1_i1m.28770 | OG0 |
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                                     •Aque_Aqu2.1.22593_001 | OG0 |
                                    <sub>-</sub>-Halamb_c34288_g1_i1m.8050 | OG0 |
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                                      -Xestes maker-XT scaffold33705-augustus-gene-0.5-mRNA-1 | OG0 |
                                    Codhol EP00042 Codosiga hollandica P052354 | OG0 |
                              Salurc EP00051 Salpingoeca urceolata P019059 | OG0 |
                                  Salurc_EP00051_Salpingoeca_urceolata_P019064 | OG0 |
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                                  Didcos EP00039 Didymoeca costata P028303 | OG0 |
                                   Diagra EP00040 Diaphanoeca_grandis_P012464 | OG0 |
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                                      Stedip EP00041 Stephanoeca diplocostata P036465 | OG0 |
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                                         Acaspe EP00035 Acanthoeca spectabilis P023267 | OG0 |
                                          •Helnan_EP00037_Helgoeca_nana_P008055 | OG0 |
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                                   -Lcom_lcpid135685 | OG0 |
                                   •Salpun_EP00053_Salpingoeca_punica_P007094 | OG0 |
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                                   Salmac EP00052 Salpingoeca macrocollata P003227 | OG0 |
                                            •Mylflu_EP00047_Mylnosiga_fluctuans_P019733 | OG0 |
                                      <sup>100</sup> Mylflu_EP00047_Mylnosiga_fluctuans_P019734 | OG0 |
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                                    Salkve EP00050 Salpingoeca kvevrii P007565 | OG0 |
                                    Saldol EP00054 Salpingoeca dolichothecata_P004923 | OG0 |
                                       Salinf EP00049 Salpingoeca infusionum P026480 | OG0 |
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                                             Salinf_EP00049_Salpingoeca_infusionum_P026481 | OG0 |
                                     •Microa_EP00043_Microstomoeca_roanoka_P028212 | OG0 |
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                                    Sros EGD72394 | OG0 |
                                        •Mbre EDQ91699 | OG0 |
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                                      Choper EP00045 Choanoeca perplexa P003194 | OG0 |
                                         •Chofle EP00730 Choanoeca flexa P014274 | OG0 |
                                   <sup>100</sup> Chofle_EP00730_Choanoeca_flexa_P014275 | OG0 |
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                               •Clacor Ccor C508133 01 | OG0 |
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                           -Scil scict013994.1 | OG0 |
                   62
                                •Lcom | lcpid77861 | OG0 |
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                          <sup>100</sup> Gracom_Gcom_scaffold8418_01 | OG0 |
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                                Leuniv_Lniv_scaffold22055_01 | OG0 |
                                   Lcom | lcpid82924 | OG0 |
                                   Gracom Gcom scaffold15200 01 | OG0 |
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                          Bralan BLAG08000582 1 | OG0 |
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                            <sup>100</sup> Spur XM_030997591.1 | OG0 |
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                                 •Acagra_model.g20998.t1 | OG0 |
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                                  •Cgig_XM_034447088.1 | OG0 |
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                                          <sup>100</sup>-Dpul_mRNA6638 | OG0 |
                              -TrH2 TrispH2 003059-RA | OG0 |
                           <sup>[100]</sup>Tadh_TriadT63630 | OG0 |
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                                  <sub>[</sub>HoiH23_PIH23_000014-RA | OG0 |
                               100 - HoiH06 mRNA7927 | OG0 |
                                 <sup>e</sup>Hhon g01241.t1 | OG0 |
                               65 HoiH04 mRNA7939 | OG0 |
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                 46
                          <sup>100</sup><sup>1</sup> Dgig_XM_028552010.1 | OG0 |
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                                 Morvir scaffold152.g51.t1 | OG0 |
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                                      _-Nemnom_g10550.t1 | OG0 |
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                                <sub>.</sub>•Adig_XM_015910318.1 | OG0 |
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                                *Spis XP 022789692 1 | OG0 |
                             <sup>1001</sup>•Pocdam XM_027184086.1 | OG0 |
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