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·Cgig_XM_011419609.3 | OG1 |
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                 •Sanmal Sma 015530-T1 | OG1 |
     92
               Aaur_scaffold1848.g1.t1 | OG1 |
       100
              •Nemnom_g8912.t1 | OG1 |
          <sup>1001</sup>--Rhoesc_Res_007818-T1 | OG1 |
            •Dgig_XM_028549234.1 | OG2 |
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       100
             •Xesp_002791-T1 | OG2 |
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              •Dgig_XM_028556196.1 | OG2 |
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               •Xesp_004363-T1 | OG2 |
                    •Lcom_lcpid127605 | OG2 |
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                           •Chem_TCONS_00006133 | OG2 |
   24
       91
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                       93
                      Aaur scaffold10.g59.t1 | OG2 |
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                          •Hvul_g19713_1 | OG2 |
                            -Chem_TCONS_00011186 | OG2 |
       65
                              •Hvul_g19718_1 | OG2 |
                        •Nvec_NVE18671 | OG2 |
                              •Nvec_v1g118161 | OG2 |
                             •Nvec_v1g131971 | OG2 |
                            Exapal_XM_021060028.2 | OG2 |
                      •Aaur_scaffold268.g17.t1 | OG2 |
              100
                         Aaur_scaffold1210.g1.t1 | OG2 |
                           •Nemnom_g3296.t1 | OG2 |
                         •Rhoesc_Res_008117-T1 | OG2 |
               •Acagra_model.g3636.t1 | OG0 |
                   Skow_XM_002739403.2 | OG0 |
                 -Bralan_BLAG01000732_1 | OG0 |
     69
              -Fspp_ffun1.m4.7924.m1 | OG0 |
          <sup>[100</sup>]-Fspp_ffun1.m4.7926.m1 | OG0 |
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                  -Spis XP 022782335 1 | OG0 |
               <sup>100</sup><sup>1</sup>-Pocdam_XM_027202097.1 | OG0 |
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                    Adig_XM_015907886.1 | OG0 |
                      JAmil_Amillepora14007-RA | OG0 |
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                         -•Adig XM 015907929.1 | OG0 |
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                  Popea_EP00118_Oscarella_pearsei_P025284 | OG0 |
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               <sup>100</sup> Ocar_g2111 | OG0 |
       59
                 •Spur_XM_003728089.3 | OG0 |
                 Spur XM 030987291.1 | OG0 |
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                Spur_XM_030995538.1 | OG0 |
              <sup>1001</sup>•Spur_XM_030995070.1 | OG0 |
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                                -Lcom | lcpid95957 | OG0 |
                            •Gracom_Gcom_C547136_01 | OG0 |
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         100
                           •Leuniv_Lniv_C805162_01 | OG0 |
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                       Lcom_lcpid46938 | OG0 |
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                    1001-Lcom | lcpid81510 | OG0 |
                       Lcom_lcpid41251 | OG0 |
                    <sup>100</sup>1-Lcom_lcpid71028 | OG0 |
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                          -Leuniv_Lniv_scaffold6525_01 | OG0 |
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                            •Gracom Gcom scaffold7316 01 | OG0 |
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              Plespi_Pspi_scaffold14720_01 | OG0 |
           <sup>100</sup>l-Plespi_Pspi_scaffold14722_01 | OG0 |
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                     Aque Aqu2.1.15854 001 | OG0 |
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                •Tetwil g11745.t1 | OG0 |
              <sup>1001</sup>•Tetwil g15405.t1 | OG0 |
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                       Stycar maker-SC scaffold4911-snap-gene-0.14-mRNA-1 | OG0 |
                      Stycar maker-SC scaffold7223-snap-gene-0.23-mRNA-1 | OG0 |
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                      •Spolac c91564 g1 i1 m.7017c91564-g1 | OG0 |
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                          -Emue_Em0021g866a.t1 | OG0 |
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                        <sup>100</sup> Emue Em0192g11a.t1 | OG0 |
                              -Tetwil g10867.t1 | OG0 |
                 TrH2 TrispH2 004269-RA | OG0 |
              <sup>100</sup> Tadh_TriadT24899 | OG0 |
                 •Tadh_TriadT25188 | OG0 |
         58
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                     -HoiH23 PIH23 010991-RA | OG0 |
                  <sup>100</sup> HoiH06 mRNA14451 | OG0 |
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                            <sup>1001</sup>Tadh_TriadT56969 | OG0 |
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                               TrH2 TrispH2 004271-RA | OG0 |
                            <sup>100</sup>Tadh TriadT56970 | OG0 |
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                           •HoiH06 mRNA2632 | OG0 |
                          HoiH23_PIH23_011726-RA | OG0 |
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                           •HoiH04 mRNA6013 | OG0 |
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                          90 Hhon_g06436.t1 | OG0 |
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                                                -HoiH23 PIH23 011725-RA | OG0 |
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                                  Hhon_g06437.t1 | OG0 |
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