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•Xesp 022641-T1 | OG0 |
                               •Xesp_022674-T1 | OG0 |
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                    Xesp_025060-T1 | OG0 |
                      •Xesp 028590-T1 | OG0 |
                         •Xesp 023975-T1|OG0|
                      •Xesp_026658-T1 | OG0 |
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     Xesp 028563-T1 | OG1
           Xesp_028560-T1 | OG1 |
         89 Xesp_022504-T1 | OG1 |
       *Xesp_009483-T1 | OG2 |
         *Xesp 025063-T1 | OG2 |
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            -Xesp 028559-T1 | OG2 |
         •Dgig XM 028543680.1 | OG2 |
                    Nvec v1g80474 | OG2 |
                     Exapal_XM_021049786.1 | OG2 |
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                     •Gfas_gfas1.m1.6285.m1 | OG2 |
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                       -Adig XM 015900543.1 | OG2 |
                     <sup>93</sup>L-Amil_Amillepora14733-RA | OG2 |
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                      •Gasp_gasp1.m3.23734.m1 | OG2 |
                        Fspp_ffun1.m4.1400.m1 | OG2 |
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                           <sub>[</sub>*Spis_XP_022788515_1 | OG2 |
                        <sup>1001</sup>•Pocdam XM_027184424.1 | OG2 |
                               •Lcom_lcpid183231 | OG2 |
                              •Hvul_g8490_1 | OG2 |
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                           ·Chem_TCONS_00000597 | OG2 |
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                                         •Hvul g8492 1 | OG2 |
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                                 rTrH2 TrispH2 002310-RA | OG2 |
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                                     •HoiH23_PIH23_001052-RA | OG2 |
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                                     <sub>•</sub>Hhon g01529.t1 | OG2 |
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                                      Cscu_XM_023371785.1 | OG2 |
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                                   Ctel gnl WGS AMQN CAPTEDRAFT mRNA159405 | OG2 |
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                           86 Opea_EP00118_Oscarella_pearsei_P007456 | OG2 |
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                                                   •Gracom Gcom scaffold10498 01 | OG2 |
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                                             -Scil scict020733.5 | OG2 |
                                            -Leuniv_Lniv_scaffold21052_01 | OG2 |
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                                         1001 Lcom_lcpid8974 | OG2 |
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                                     •Eupdun Edun sb10607897 | OG2 |
                                         •Lampan_Lpan_9213_01 | OG2 |
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                                            •Eupdun_Edun_sb10622604 | OG2 |
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                                              •Coemet_Cmet_scaffold22407_01 | OG2 |
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                                                  •Pbac_sb2639026 | OG2 |
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                                            88 Horcal_Hcv1.av93.c2.g1297.i1 | OG2 |
                                           Eupdun Edun sb10649300 | OG2 |
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                                        <sup>100</sup> Eupdun_Edun_sb10649304 | OG2 |
                                             Mertsp Mesp sb582490 | OG2 |
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                                             811 Mertsp_Mesp_sb582489 | OG2 |
                                        91
                                                Beraby_Baby_sb333614 | OG2 |
                                                   Bolinf_Binf_sb1263101 | OG2 |
                                                <sup>100</sup> Mertsp_Mesp_sb530338 | OG2 |
                                                   rVmul_sbMO379933 | OG2 |
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                                  •Plespi_Pspi_scaffold6024_01 | OG2 |
                                 Tetwil g4784.t1 | OG2 |
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                                      •Plespi_Pspi_scaffold984_01 | OG2 |
                                          Xestes_maker-XT_scaffold5404-snap-gene-0.7-mRNA-1 | OG2 |
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                                              <sub>r</sub>-Halamb_c56172_g3_i2m.42683 | OG2 |
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                                                rHalamb c54615_g1_i4m.38731 | OG2 |
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                                              <sup>98</sup> Halamb_c56172_g2_i2mm.7006 | OG2 |
                                              -Halamb_c56172_g1_i2mm.7005 | OG2 |
                                     96
                                           <sup>100</sup>•Halamb_c54520_g1_i4m.38508 | OG2 |
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                                                Aque_Aqu2.1.18093_001 | OG2 |
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                                   <sup>100</sup>Stycar_maker-SC_scaffold82321-augustus-gene-0.25-mRNA-1 | OG2 |
                                   •Spolac_c105082_g1_i1_m.44760c103425-g4 | OG2 |
                                      •Spolac_c95758_g1_i1_m.11994c103425-g4 | OG2 |
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                                        •Spolac_c97602_g1_i1_m.15367c103425-g4 | OG2 |
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                                            •Emue_Em0007g240a.t1 | OG2 |
                                                  •Emue_Em0007g136a.t1 | OG2 |
                                              <sup>98</sup> -Emue_Em0007g171a.t1 | OG2 |
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                                                                 -Emue_Em0007g235a.t1 | OG2 |
                                                           <u>r</u>Emue_Em0007g160a.t1 | OG2 |
                                                                -Emue_Em0007g150a.t1 | OG2 |
0.416667
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