```
<sub>f</sub>Stycar_maker-SC_scaffold131343-augustus-gene-0.29-mRNA-1 | OG0 |
             <sup>1001</sup>-Stycar maker-SC_scaffold99123-augustus-gene-0.3-mRNA-1 | OG0 |
                    •Plespi_Pspi_scaffold20737_01 | OG0 |
 100
                               <sub>s</sub>Halamb_c49586_g1_i1mm.3148 | OG0 |
                            <sup>1001</sup>•Halamb_c50197_g2_i2mm.3396 | OG0 |
    90
                     Tetwil g13471.t2 | OG0 |
                     -Stycar_augustus_masked-SC_scaffold1458-processed-gene-0.1-mRNA-1 | OG0 |
      93
                         •Stycar_maker-SC_scaffold5880-snap-gene-0.3-mRNA-1 | OG0 |
                     Spolac_c104463_g2_i7_m.40690c104463-g2 | OG0 |
        97
                         Emue Em0022g432a.t1 | OG0 |
                 98
                          •Spolac_c104463_g3_i1_m.40687c104463-g3 | OG0 |
                            Emue_Em0022g433a.t1 | OG0 |
              100
                         •Spolac_c104463_g2_i4_m.40683c104463-g2 | OG0 |
                       Spolac_c104463_g2_i6_m.40688c104463-g2 | OG0 |
                          Spolac_c104463_g2_i1_m.40681c104463-g2 | OG0 |
                        <sup>1001</sup>Spolac c104463_g2_i5_m.40685c104463-g2 | OG0 |
                 •Plajan Pjan scaffold4748 01 | OG0 |
                     Plajan_Pjan_scaffold4749_01 | OG0 |
    63
                  <sup>100</sup>-Plajan_Pjan_scaffold4751_01 | OG0 |
                       Clacor_Ccor_scaffold11263_01 | OG0 |
      88
                    Leuniv_Lniv_C807180_01 | OG0 |
            100
                        •Lcom_lcpid25052 | OG0 |
                          Leuniv Lniv scaffold24778 01 | OG0 |
                            •Scil scict019919.1 | OG0 |
                     TrH2 TrispH2 005459-RA | OG0 |
                  <sup>1001</sup>•Tadh_TriadT25047 | OG0 |
 100
                       •HoiH11_mRNA1992 | OG0 |
               100
                           <sub>"</sub>HoiH06_mRNA15086 | OG0 |
                  99
                        <sup>1001</sup>•HoiH23_PIH23_004982-RA | OG0 |
                           <sub>[</sub>Hhon_g02853.t1 | OG0 |
                        <sup>100</sup> HoiH04 mRNA14603 | OG0 |
                      Bralan_BLAG06000556_1 | OG0 |
                         Skow_XM_006820450.1 | OG0 |
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                            •Spur_XM_030980300.1 | OG0 |
                           -Hsap_ENST00000381655 | OG0 |
    66
                         •Hsap_ENST00000381668 | OG0 |
        100
                       ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA176761 | OG0 |
                       -Cgig_XM_034444034.1 | OG0 |
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                   <sup>100</sup>•Acagra_model.g9151.t1.1.5d3b85d9 | OG0 |
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                           •Tcas_TC007810_001 | OG0 |
                             •Dpul_mRNA13378 | OG0 |
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                                  •Dpul_mRNA19691 | OG0 |
                             -Lcom_lcpid251742 | OG0 |
                          Hvul_g11267_1 | OG0 |
                        <sup>961</sup>-Hvul_g11268_1 | OG0 |
                           •Hvul_g6406_1 | OG0 |
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                              •Hvul g11266 1 | OG0 |
         80
                                Chem TCONS 00054822 | OG0 |
              93
                       •Morvir_scaffold197.g20.t1 | OG0 |
                        •Aaur_scaffold652.g3.t2 | OG0 |
                           -Rhoesc_Res_000394-T2 | OG0 |
           69
                        <sup>100</sup><sup>1</sup>•Nemnom_g15339.t1 | OG0 |
                      Dgig_XM_028545659.1 | OG0 |
                        Exapal XM 021059391.2 | OG0 |
                   100
                        •Nvec_v1g10129 | OG0 |
                          •Amil Amillepora08456-RA | OG0 |
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                          -Adig_XM_015914760.1 | OG0 |
                         •Gfas_gfas1.m1.1837.m1 | OG0 |
                            Gasp_gasp1.m3.26190.m1 | OG0 |
                               Fspp_ffun1.m4.9826.m1 | OG0 |
                        100
                                 Pocdam_XM_027204078.1 | OG0 |
                                   Pocdam_XM_027204091.1 | OG0 |
                                 <sup>99</sup> Spis_XP_022807594_1 | OG0 |
0.416667
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