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                      •Scil scict013276.2 | OG1 |
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1
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                      Vmul SimC403884 | OG1 |
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                      100
               83
                  87
                        100
                    85
                           100
                           100
                              100
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          99
        80
     34
                   •Dpul mRNA8898 | OG2 |
                  <sup>100</sup><sup>1</sup>Tadh_TriadT56467 | OG2 |
        58
                       •HoiH11 mRNA1777 | OG2 |
               100
                  100
   53
                     •Tcas_TC007521_001 | OG2 |
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                        58
          13
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                        100
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                 100
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0.416667
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Salmac_EP00052_Salpingoeca_macrocollata_P018636 | OG0 |
Salpun_EP00053_Salpingoeca_punica_P019972 | OG0 |
  •Salkve_EP00050_Salpingoeca_kvevrii_P007193 | OG0 |
     Codhol EP00042 Codosiga hollandica P052878 | OG0 |
        Codhol EP00042 Codosiga hollandica P052877 | OG0 |
       Didcos_EP00039_Didymoeca_costata_P019429 | OG0 |
            Diagra_EP00040_Diaphanoeca_grandis_P029369 | OG0 |
          Stedip_EP00041_Stephanoeca_diplocostata_P027444 | OG0 |
                Acaspe EP00035 Acanthoeca spectabilis P034440 | OG0 |
              Helnan EP00037 Helgoeca nana P032893 | OG0 |
 •Mylflu_EP00047_Mylnosiga_fluctuans_P010934 | OG0 |
      Saldol_EP00054_Salpingoeca_dolichothecata_P001539 | OG0 |
        ·Salurc_EP00051_Salpingoeca_urceolata_P014401 | OG0 |
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        Salinf EP00049 Salpingoeca infusionum P012549 | OG0 |
     Microa EP00043 Microstomoeca roanoka P008528 | OG0 |
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        Chofle EP00730 Choanoeca flexa P017643 | OG0 |
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      Leuniv_Lniv_scaffold18430_01 | OG1 |
   1001 Leuniv_Lniv_scaffold18431_01 | OG1 |
    -Coeast_Cast_sb819783 | OG1 |
 <sup>100</sup> Coemet_Cmet_C605288_01 | OG1 |
        Vmul sbMO442131 | OG1 |
    •Lampan Lpan C417521 01 | OG1 |
          Eupdun Edun sb10654075 | OG1 |
       <sup>1001</sup>•Eupdun_Edun_sb10654076 | OG1 |
        Mertsp Mesp sb593231 | OG1 |
          Mertsp Mesp sb593230 | OG1 |
            Mertsp_Mesp_sb593234 | OG1 |
              Mertsp_Mesp_sb593233 | OG1 |
                *Bolinf Binf sb1245591 | OG1 |
              971 Mertsp Mesp_sb593232 | OG1 |
           Horcal_Hcv1.av93.c7.g705.i1 | OG1 |
         <sup>991</sup>Horcal_Hcv1.av93.c7.g705.i2 | OG1 |
            •Pbac_sb2640957 | OG1 |
             Pbac sb2664062 | OG1 |
           66 Pbac sb2663265 | OG1 |
           Drygla_Dgla_sb306962 | OG1 |
         <sup>1001</sup>•Drygla_Dgla_sb306963 | OG1 |
            Drygla_Dgla_sb278994 | OG1 |
              Beraby_Baby_sb330343 | OG1 |
                Bolinf Binf sb1215772 | OG1 |
               <sup>22</sup> Mertsp_Mesp_sb520950 | OG1 |
               Mlei_ML2835.g10.i1 | OG1 |
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 <sup>1001</sup>-Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA225710 | OG2 |
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         Cscu_XM_023370690.1 | OG2 |
       <sup>79</sup> Cscu_XM_023375845.1 | OG2 |
    TrH2 TrispH2 001825-RA | OG2 |
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       <sup>100</sup>-HoiH06 mRNA13480 | OG2 |
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          •Lcom_lcpid124427 | OG2 |
   Spur XM 030980424.1 | OG2 |
<sup>100</sup> Spur _XM_030980491.1 | OG2 |
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     •Skow_XM_002735087.2 | OG2 |
     •Hsap_ENST00000231572 | OG2 |
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          •Plajan_Pjan_scaffold6839_01 | OG2 |
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       <sup>100</sup> Emue_Em0022g764a.t1 | OG2 |
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           <sup>1001</sup>Spis_XP_022795390_1 | OG2 |
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