

Salkve_EP00050_Salpingoeca_kvevrii_P013577 | OG0 | 100 Salkve EP00050 Salpingoeca kvevrii P013578 | OG0 | Didcos_EP00039_Didymoeca_costata_P023432 | OG0 | Diagra EP00040 Diaphanoeca grandis P043313 | OG0 | Stedip_EP00041_Stephanoeca_diplocostata_P033039 | OG0 | Acaspe EP00035 Acanthoeca spectabilis P031721 | OG0 | •Helnan_EP00037_Helgoeca_nana_P010079 | OG0 | Salurc EP00051 Salpingoeca urceolata P003127 | OG0 | Mylflu EP00047 Mylnosiga fluctuans P015415 | OG0 | Saldol EP00054 Salpingoeca_dolichothecata_P038001 | OG0 | Salpun_EP00053_Salpingoeca_punica_P007492 | OG0 | •Salkve_EP00050_Salpingoeca_kvevrii_P009348 | OG0 | Salmac EP00052 Salpingoeca macrocollata P009983 | OG0 | Salmac_EP00052_Salpingoeca_macrocollata_P009984 | OG0 | Salpun EP00053 Salpingoeca punica P011567 | OG0 | Salpun EP00053 Salpingoeca punica P011569 | OG0 | Salpun EP00053 Salpingoeca punica P011571 | OG0 | Salinf_EP00049_Salpingoeca_infusionum_P019633 | OG0 | •Microa EP00043 Microstomoeca roanoka P002672 | OG0 | Sros EGD74592 | OG0 | •Mbre EDQ92988 | OG0 | Chofle_EP00730_Choanoeca_flexa_P013816 | OG0 | Choper EP00045 Choanoeca perplexa P019637 | OG0 | Lampan_Lpan_C404377_01 | OG0 | Coemet_Cmet_C591377_01 | OG0 | •Vmul SimC392124 | OG0 | •Coemet_Cmet_scaffold19737_01 | OG0 | •Vmul SimC390682 | OG0 | -Lampan_Lpan_6399_01 | OG0 | Beraby Baby sb339315 | OG0 | •Pbac sb2669113 | OG0 | •Pbac_sb2669112 | OG0 | •Coemet_Cmet_scaffold21529_01 | OG0 | •Vmul_SimC397570 | OG0 | •Lampan_Lpan_10303_01 | OG0 | Horcal Hcv1.av93.c2.g1578.i3 | OG0 | •Pbac_sb2637689 | OG0 | Mertsp Mesp sb556799 | OG0 | ¹⁰⁰ Mertsp_Mesp_sb556800 | OG0 | Mertsp Mesp sb556798 | OG0 | Bolinf Binf sb1295094 | OG0 | 65 Mertsp Mesp sb556797 | OG0 | Beraby_Baby_sb336794 | OG0 | ¹⁰⁰ Beraby_Baby_sb336795 | OG0 | •Mlei ML0438.g4.i1 | OG0 | •Mlei_ML0438.g5.i1 | OG0 | Beraby Baby sb352924 | OG0 | Drygla Dgla sb288234 | OG0 | ⁹⁴¹•Drygla_Dgla_sb288235 | OG0 | Drygla_Dgla sb288236 | OG0 | ¹⁰⁰¹Drygla_Dgla_sb288237 | OG0 | Ctel gnl WGS AMQN CAPTEDRAFT mRNA166467 | OG0 | •Cgig XM 011423379.3 | OG0 | ·Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA217713 | OG0 | 1001 Ctel gnl WGS AMQN_CAPTEDRAFT_mRNA98799 | OG0 | •Hsap ENST00000333503 | OG0 | Scil scict020507.2 | OG0 | Clacor Ccor C512035 01 | OG0 | Gracom Gcom scaffold225 01 | OG0 | Lcom | lcpid87788 | OG0 | •Leuniv_Lniv_scaffold20868_01 | OG0 | •Hvul g29624 1 | OG0 | Ivul_g29623_1 | OG0 | •Chem_TCONS_00000812 | OG0 | •Morvir_scaffold87.g25.t1 | OG0 | Sanmal_Sma_024401-T1 | OG0 | •Nemnom_g13383.t1 | OG0 | •Rhoesc_Res_017054-T1 | OG0 | •Aaur_scaffold218.g8.t1 | OG0 | •Aaur_scaffold218.g9.t1 | OG0 | Cscu XM 023363527.1 | OG0 | Tcas TC007560 001 | OG0 | -Dpul_mRNA17687 | OG0 | |Spur_XM_030988817.1 | OG0 •Skow_XM_006823871.1 | OG0 | Skow_XM_002740599.2 | OG0 | Skow XM 006815130.1 | OG0 | •Bralan_BLAG03000638_1 | OG0 | •Acagra_model.g1956.t1 | OG0 | •Acagra_model.g1959.t1 | OG0 | rCgig_XM_034476273.1 | OG0 | ¹⁰⁰ Cgig XM_011416663.3 | OG0 | •Aque_Aqu2.1.20362_001 | OG0 | •Tetwil_g4891.t2 | OG0 | -Stycar_maker-SC_scaffold8635-snap-gene-0.25-mRNA-1 | OG0 | -Stycar_snap_masked-SC_scaffold1985-processed-gene-0.1-mRNA-1 | OG0 | Emue Em0019g1050a.t1 | OG0 | Spolac_c101980_g1_i1_m.27476c101980-g1 | OG0 | _sSpolac_c101980_g1_i2_m.27477c101980-g1 | OG0 | ⁶¹Spolac_c101980_g1_i3_m.27478c101980-g1 | OG0 | Opea EP00118 Oscarella pearsei P015579 | OG0 | Opea_EP00118_Oscarella_pearsei_P015578 | OG0 | ⁵⁴Opea_EP00118_Oscarella_pearsei_P015580 | OG0 | _"HoiH23_PIH23_002299-RA | OG0 | ⁶⁶⁻HoiH06_mRNA13253 | OG0 | _•Hhon_g07619.t1 | OG0 | ¹⁰⁰ HoiH04_mRNA13687 | OG0 | •HoiH11_mRNA5451 | OG0 | TrH2_TrispH2_007525-RA | OG0 | ¹⁰⁰ Tadh_TriadT51940 | OG0 | _"Exapal_XM_021040865.2 | OG0 | ¹⁰⁰ Exapal_XM_021043570.2 | OG0 | •Dgig_XM_028549822.1 | OG0 | •Xesp_014081-T1 | OG0 | -Spis_XP_022792561_1 | OG0 | •Pocdam XM 027197966.1 | OG0 | Gasp_gasp1.m3.21780.m1 | OG0 | 99¹-Fspp_ffun1.m4.25666.m1 | OG0 | Gfas_gfas1.m1.9924.m1 | OG0 | Adig_XM_015904088.1 | OG0 | _"Amil_Amillepora24640-RA | OG0 | ¹⁰⁰ Adig XM_015904100.1 | OG0 |