

•Salkve_EP00050_Salpingoeca_kvevrii_P001704 | OG0 | Salurc_EP00051_Salpingoeca_urceolata_P011233 | OG0 | Salurc_EP00051_Salpingoeca_urceolata_P011234 | OG0 | ·Codhol_EP00042_Codosiga_hollandica_P049340 | OG0 | Didcos EP00039 Didymoeca costata P030781 | OG0 | Diagra EP00040 Diaphanoeca grandis P004856 | OG0 | Stedip_EP00041_Stephanoeca_diplocostata_P029404 | OG0 | -Acaspe_EP00035_Acanthoeca_spectabilis_P003188 | OG0 | •Helnan_EP00037_Helgoeca_nana_P007264 | OG0 | •Mylflu_EP00047_Mylnosiga_fluctuans P003541 | OG0 | Saldol EP00054 Salpingoeca dolichothecata P009417 | OG0 | Salpun_EP00053_Salpingoeca_punica_P010836 | OG0 | Salmac_EP00052_Salpingoeca_macrocollata_P016269 | OG0 | 1001-Salmac EP00052_Salpingoeca_macrocollata_P016270 | OG0 | Microa EP00043 Microstomoeca roanoka P007565 | OG0 | •Sros EGD73079 | OG0 | rSalinf_EP00049_Salpingoeca_infusionum_P026350 | OG0 | Salinf_EP00049_Salpingoeca_infusionum_P026351 | OG0 | Choper EP00045 Choanoeca perplexa P014749 | OG0 | •Chofle EP00730 Choanoeca flexa P016448 | OG0 | Eupdun Edun sb10661179 | OG0 | 88 Eupdun_Edun_sb10661176 | OG0 | Eupdun_Edun_sb10661180 | OG0 | 65 Eupdun_Edun_sb10661181 | OG0 | Eupdun Edun sb10661177 | OG0 | 41 Eupdun Edun sb10661178 | OG0 | _[•Horcal_Hcv1.av93.c9.g857.i1 | OG0 | •Pbac_sb2653115 | OG0 | ¹⁰⁰ Pbac sb2640023 | OG0 | •Coeast Cast sb791656 | OG0 | •Lampan_Lpan_9892_01 | OG0 | Bolinf_Binf_sb1261172 | OG0 | Bolinf_Binf_sb1261170 | OG0 | 66 Bolinf Binf sb1261171 | OG0 | Mertsp Mesp sb578298 | OG0 | 981 Mertsp Mesp sb578299 | OG0 | Beraby Baby sb341870 | OG0 | Beraby_Baby_sb341872 | OG0 | ⁷⁵¹-Beraby_Baby_sb341871 | OG0 | Drygla Dgla sb298537 | OG0 | ¹⁰⁰ Drygla_Dgla_sb298538 | OG0 | •Mlei_ML0240.g3.i1 | OG0 | Bolinf_Binf_sb1234941 | OG0 | ¹⁰⁰ Bolinf_Binf_sb1234942 | OG0 | Popea EP00118 Oscarella pearsei P009084 | OG0 | ¹⁰⁰ Ocar_g10081 | OG0 | Plajan_Pjan_scaffold21082_01 | OG0 | ¹⁰⁰ Plajan_Pjan_scaffold21083_01 | OG0 | •Clacor_Ccor_scaffold14871_01 | OG0 | Lcom | lcpid18641 | OG0 | Gracom_Gcom_scaffold86_01 | OG0 | -Scil_scict020596.1 | OG0 | 100 Leuniv_Lniv_C809698_01 | OG0 | Aque Aqu2.1.37449 001 | OG0 | ¹⁰⁰ Aque Aqu2.1.15698_001 | OG0 | Stycar maker-SC scaffold45955-augustus-gene-0.3-mRNA-1 | OG0 | Stycar_maker-SC_scaffold45268-snap-gene-0.3-mRNA-1 | OG0 | •Tetwil g3922.t1 | OG0 | •Plespi_Pspi_scaffold15424_01 | OG0 | •Emue Em0001g1705a.t1 | OG0 | Spolac_c103497_g1_i3_m.34548c103497-g1 | OG0 | Spoiac_c103497_g1_i5_m.34556c103497-g1 | OG0 | Spolac_c103497_g1_i8_m.34558c103497-g1 | OG0 | Spolac_c103497_g1_i1_m.34545c103497-g1 | OG0 | Spolac_c103497_g1_i4_m.34552c103497-g1 | OG0 | spolac_c103497_g1_i3_m.34549c103497-g1 | OG0 | ¹ Spolac_c103497_g1_i5_m.34554c103497-g1 | OG0 | ·Cscu_XM_023363401.1 | OG0 | -Hsap ENST00000232888 | OG0 | •Dpul mRNA26164 | OG0 | -Tcas TC014174 001 | OG0 | •Bralan_BLAG05000846_1 | OG0 | Spur XM 011678350.2 | OG0 | Skow XM 006823432.1 | OG0 | Cgig XM 011443738.3 | OG0 | •Acagra_model.g7673.t1 | OG0 | •Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA122475 | OG0 | Exapal XM 028661222.1 | OG0 | Nvec v1g246398 | OG0 | •Amil Amillepora07002-RA | OG0 | Gfas_gfas1.m1.17866.m1 | OG0 | •Gasp_gasp1.m3.30212.m1 | OG0 | •Fspp_ffun1.m4.15606.m1 | OG0 | -Spis_XP_022801314_1 | OG0 | ¹⁰⁰¹-Pocdam XM 027182107.1 | OG0 | •Xesp_014622-T1 | OG0 | •Dgig_XM_028546106.1 | OG0 | •Morvir_scaffold28.g25.t1 | OG0 | •Sanmal Sma 010838-T1|OG0| •Aaur_scaffold800.g6.t2 | OG0 | •Nemnom_g15818.t1 | OG0 | •Rhoesc_Res_000443-T1 | OG0 | Chem TCONS 00054806 | OG0 | •Hvul_g15990_1 | OG0 | TrH2 TrispH2 008675-RA | OG0 | ¹⁰⁰ Tadh_TriadT26083 | OG0 | _"Hhon_g09825.t1 | OG0 | ¹⁰⁰ HoiH04 mRNA12859 | OG0 | •HoiH06 mRNA5717 | OG0 | ¹⁰⁰¹•HoiH23_PIH23_007848-RA | OG0 |