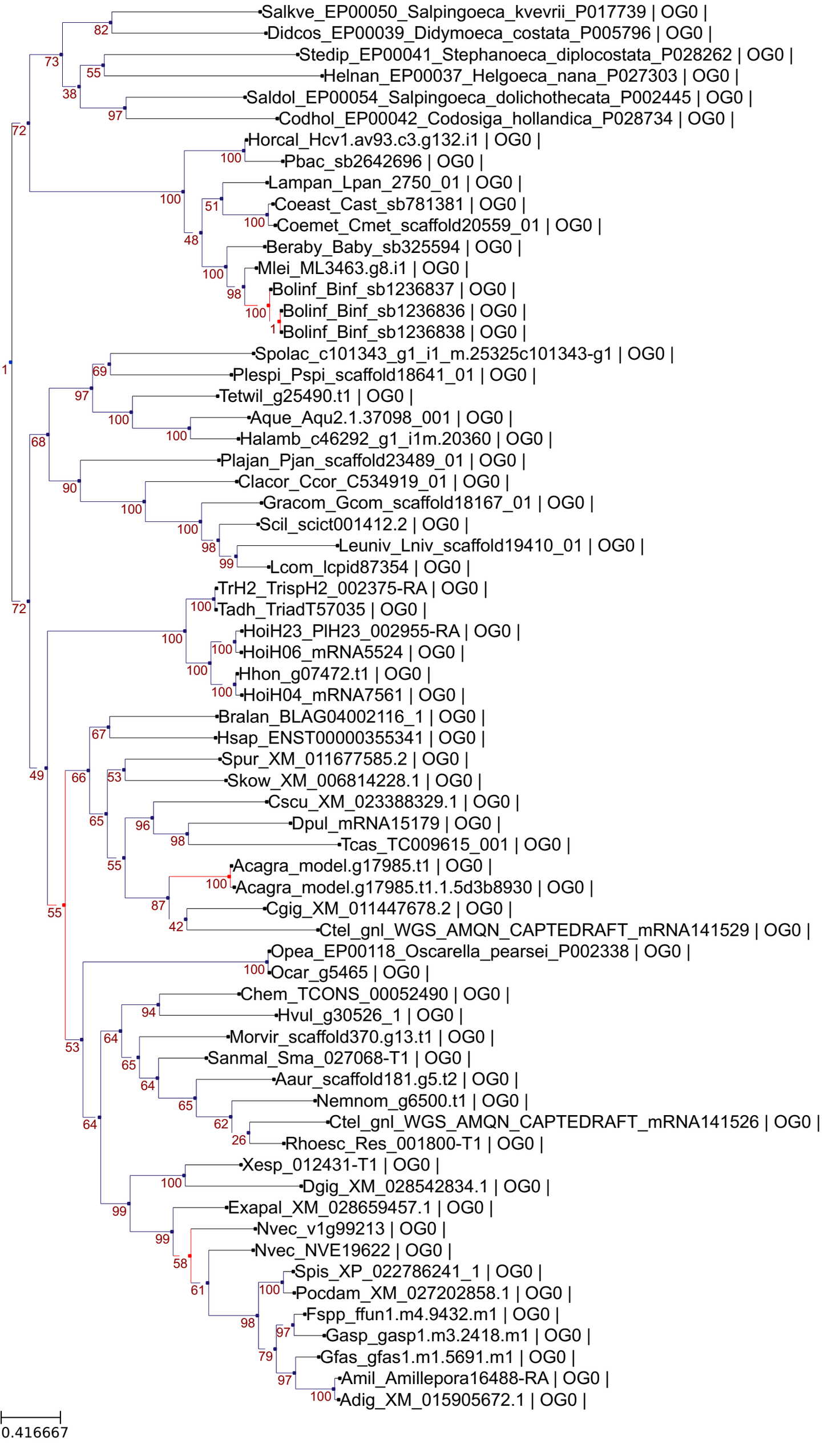
Salkve_EP00050_Salpingoeca_kvevrii_P017739 | OG0 |
Didcos_EP00039_Didymoeca_costata_P005796 | OG0 |
Stedip_EP00041_Stephanoeca_diplocostata_P028262 | OG0 |
Helnan_EP00037_Helgoeca_nana_P027303 | OG0 |
Saldol_EP00054_Salpingoeca_dolichothecata_P002445 | OG0 |
Codhol_EP00042_Codosiga_hollandica_P028734 | OG0 |
Horcal_Hcv1.av93.c3.g132.i1 | OG0 |
Pbac_sb2642696 | OG0 |
Lampan_Lpan_2750_01 | OG0 |
Coeast_Cast_sb781381 | OG0 |
Coemet_Cmet_scaffold20559_01 | OG0 |
Beraby_Baby_sb325594 | OG0 |
Mlei_ML3463.g8.i1 | OG0 |
Bolinf_Binf_sb1236837 | OG0 |
Bolinf_Binf_sb1236836 | OG0 |
Bolinf_Binf_sb1236838 | OG0 |
Spolac_c101343_g1_i1_m.25325c101343-g1 | OG0 |
Plespi_Pspi_scaffold18641_01 | OG0 |
Tetwil_g25490.t1 | OG0 |
Aque_Aqu2.1.37098_001 | OG0 |
Halamb_c46292_g1_i1m.20360 | OG0 |
Plajan_Pjan_scaffold23489_01 | OG0 |
Clacor_Ccor_C534919_01 | OG0 |
Gracom_Gcom_scaffold18167_01 | OG0 |
Scil_scict001412.2 | OG0 |
Leuniv_Lniv_scaffold19410_01 | OG0 |
Lcom_lcpid87354 | OG0 |
TrH2_TrispH2_002375-RA | OG0 |
Tadh_TriadT57035 | OG0 |
HoiH23_PlH23_002955-RA | OG0 |
HoiH06_mRNA5524 | OG0 |
Hhon_g07472.t1 | OG0 |
HoiH04_mRNA7561 | OG0 |
Bralan_BLAG04002116_1 | OG0 |
Hsap_ENST00000355341 | OG0 |
Spur_XM_011677585.2 | OG0 |
Skow_XM_006814228.1 | OG0 |
Cscu_XM_023388329.1 | OG0 |
Dpul_mRNA15179 | OG0 |
Tcas_TC009615_001 | OG0 |
Acagra_model.g17985.t1 | OG0 |
Acagra_model.g17985.t1.1.5d3b8930 | OG0 |
Cgig_XM_011447678.2 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA141529 | OG0 |
Opea_EP00118_Oscarella_pearsei_P002338 | OG0 |
Ocar_g5465 | OG0 |
Chem_TCONS_00052490 | OG0 |
Hvul_g30526_1 | OG0 |
Morvir_scaffold370.g13.t1 | OG0 |
Sanmal_Sma_027068-T1 | OG0 |
Aaur_scaffold181.g5.t2 | OG0 |
Nemnom_g6500.t1 | OG0 |
Ctel_gnl_WGS_AMQN_CAPTEDRAFT_mRNA141526 | OG0 |
Rhoesc_Res_001800-T1 | OG0 |
Xesp_012431-T1 | OG0 |
Dgig_XM_028542834.1 | OG0 |
Exapal_XM_028659457.1 | OG0 |
Nvec_v1g99213 | OG0 |
Nvec_NVE19622 | OG0 |
Spis_XP_022786241_1 | OG0 |
Pocdam_XM_027202858.1 | OG0 |
Fspp_ffun1.m4.9432.m1 | OG0 |
Gasp_gasp1.m3.2418.m1 | OG0 |
Gfas_gfas1.m1.5691.m1 | OG0 |
Amil_Amillepora16488-RA | OG0 |
Adig_XM_015905672.1 | OG0 |
0.416667