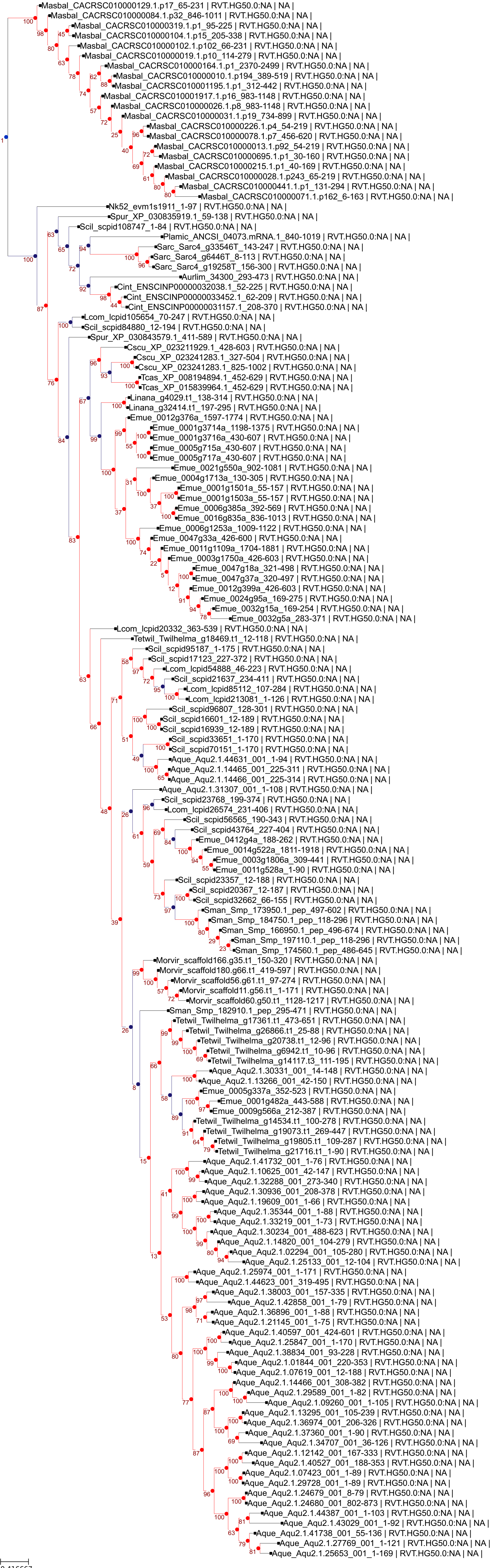
Masbal_CACRSC010000129.1.p17_65-231 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000084.1.p32_846-1011 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000319.1.p1_95-225 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000104.1.p15_205-338 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000102.1.p102_66-231 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000019.1.p10_114-279 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000164.1.p1_2370-2499 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000010.1.p194_389-519 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010001195.1.p1_312-442 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010001917.1.p16_983-1148 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000026.1.p8_983-1148 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000031.1.p19_734-899 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000226.1.p4_54-219 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000078.1.p7_456-620 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000013.1.p92_54-219 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000695.1.p1_30-160 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000215.1.p1_40-169 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000028.1.p243_65-219 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000441.1.p1_131-294 | RVT.HG50.0:NA | NA |
Masbal_CACRSC010000071.1.p162_6-163 | RVT.HG50.0:NA | NA |
Nk52_evm1s1911_1-97 | RVT.HG50.0:NA | NA |
Spur_XP_030835919.1_59-138 | RVT.HG50.0:NA | NA |
Scil_scpid108747_1-84 | RVT.HG50.0:NA | NA |
Plamic_ANCSI_04073.mRNA.1_840-1019 | RVT.HG50.0:NA | NA |
Sarc_Sarc4_g33546T_143-247 | RVT.HG50.0:NA | NA |
Sarc_Sarc4_g6446T_8-113 | RVT.HG50.0:NA | NA |
Sarc_Sarc4_g19258T_156-300 | RVT.HG50.0:NA | NA |
Aurlim_34300_293-473 | RVT.HG50.0:NA | NA |
Cint_ENSCINP00000032038.1_52-225 | RVT.HG50.0:NA | NA |
Cint_ENSCINP00000033452.1_62-209 | RVT.HG50.0:NA | NA |
Cint_ENSCINP00000031157.1_208-370 | RVT.HG50.0:NA | NA |
Lcom_lcpid105654_70-247 | RVT.HG50.0:NA | NA |
Scil_scpid84880_12-194 | RVT.HG50.0:NA | NA |
Spur_XP_030843579.1_411-589 | RVT.HG50.0:NA | NA |
Cscu_XP_023211929.1_428-603 | RVT.HG50.0:NA | NA |
Cscu_XP_023241283.1_327-504 | RVT.HG50.0:NA | NA |
Cscu_XP_023241283.1_825-1002 | RVT.HG50.0:NA | NA |
Tcas_XP_008194894.1_452-629 | RVT.HG50.0:NA | NA |
Tcas_XP_015839964.1_452-629 | RVT.HG50.0:NA | NA |
Linana_g4029.t1_138-314 | RVT.HG50.0:NA | NA |
Linana_g32414.t1_197-295 | RVT.HG50.0:NA | NA |
Emue_0012g376a_1597-1774 | RVT.HG50.0:NA | NA |
Emue_0001g3714a_1198-1375 | RVT.HG50.0:NA | NA |
Emue_0001g3716a_430-607 | RVT.HG50.0:NA | NA |
Emue_0005g715a_430-607 | RVT.HG50.0:NA | NA |
Emue_0005g717a_430-607 | RVT.HG50.0:NA | NA |
Emue_0021g550a_902-1081 | RVT.HG50.0:NA | NA |
Emue_0004g1713a_130-305 | RVT.HG50.0:NA | NA |
Emue_0001g1501a_55-157 | RVT.HG50.0:NA | NA |
Emue_0001g1503a_55-157 | RVT.HG50.0:NA | NA |
Emue_0006g385a_392-569 | RVT.HG50.0:NA | NA |
Emue_0016g835a_836-1013 | RVT.HG50.0:NA | NA |
Emue_0006g1253a_1009-1122 | RVT.HG50.0:NA | NA |
Emue_0047g33a_426-600 | RVT.HG50.0:NA | NA |
Emue_0011g1109a_1704-1881 | RVT.HG50.0:NA | NA |
Emue_0003g1750a_426-603 | RVT.HG50.0:NA | NA |
Emue_0047g18a_321-498 | RVT.HG50.0:NA | NA |
Emue_0047g37a_320-497 | RVT.HG50.0:NA | NA |
Emue_0012g399a_426-603 | RVT.HG50.0:NA | NA |
Emue_0024g95a_169-275 | RVT.HG50.0:NA | NA |
Emue_0032g15a_169-254 | RVT.HG50.0:NA | NA |
Emue_0032g5a_283-371 | RVT.HG50.0:NA | NA |
Lcom_lcpid20332_363-539 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g18469.t1_12-118 | RVT.HG50.0:NA | NA |
Scil_scpid95187_1-175 | RVT.HG50.0:NA | NA |
Scil_scpid17123_227-372 | RVT.HG50.0:NA | NA |
Lcom_lcpid54888_46-223 | RVT.HG50.0:NA | NA |
Scil_scpid21637_234-411 | RVT.HG50.0:NA | NA |
Lcom_lcpid85112_107-284 | RVT.HG50.0:NA | NA |
Lcom_lcpid213081_1-126 | RVT.HG50.0:NA | NA |
Scil_scpid96807_128-301 | RVT.HG50.0:NA | NA |
Scil_scpid16601_12-189 | RVT.HG50.0:NA | NA |
Scil_scpid16939_12-189 | RVT.HG50.0:NA | NA |
Scil_scpid33651_1-170 | RVT.HG50.0:NA | NA |
Scil_scpid70151_1-170 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.44631_001_1-94 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.14465_001_225-311 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.14466_001_225-314 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.31307_001_1-108 | RVT.HG50.0:NA | NA |
Scil_scpid23768_199-374 | RVT.HG50.0:NA | NA |
Lcom_lcpid26574_231-406 | RVT.HG50.0:NA | NA |
Scil_scpid56565_190-343 | RVT.HG50.0:NA | NA |
Scil_scpid43764_227-404 | RVT.HG50.0:NA | NA |
Emue_0412g4a_188-262 | RVT.HG50.0:NA | NA |
Emue_0014g522a_1811-1918 | RVT.HG50.0:NA | NA |
Emue_0003g1806a_309-441 | RVT.HG50.0:NA | NA |
Emue_0011g528a_1-90 | RVT.HG50.0:NA | NA |
Scil_scpid23357_12-188 | RVT.HG50.0:NA | NA |
Scil_scpid20367_12-187 | RVT.HG50.0:NA | NA |
Scil_scpid32662_66-155 | RVT.HG50.0:NA | NA |
Sman_Smp_173950.1_pep_497-602 | RVT.HG50.0:NA | NA |
Sman_Smp_184750.1_pep_118-296 | RVT.HG50.0:NA | NA |
Sman_Smp_166950.1_pep_496-674 | RVT.HG50.0:NA | NA |
Sman_Smp_197110.1_pep_118-296 | RVT.HG50.0:NA | NA |
Sman_Smp_174560.1_pep_486-645 | RVT.HG50.0:NA | NA |
Morvir_scaffold166.g35.t1_150-320 | RVT.HG50.0:NA | NA |
Morvir_scaffold180.g66.t1_419-597 | RVT.HG50.0:NA | NA |
Morvir_scaffold56.g61.t1_97-274 | RVT.HG50.0:NA | NA |
Morvir_scaffold11.g56.t1_1-171 | RVT.HG50.0:NA | NA |
Morvir_scaffold60.g50.t1_1128-1217 | RVT.HG50.0:NA | NA |
Sman_Smp_182910.1_pep_295-471 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g17361.t1_473-651 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g26866.t1_25-88 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g20738.t1_12-96 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g6942.t1_10-96 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g14117.t3_111-195 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.30331_001_14-148 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.13266_001_42-150 | RVT.HG50.0:NA | NA |
Emue_0005g337a_352-523 | RVT.HG50.0:NA | NA |
Emue_0001g482a_443-588 | RVT.HG50.0:NA | NA |
Emue_0009g566a_212-387 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g14534.t1_100-278 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g19073.t1_269-447 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g19805.t1_109-287 | RVT.HG50.0:NA | NA |
Tetwil_Twilhelma_g21716.t1_1-90 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.41732_001_1-76 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.10625_001_42-147 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.32288_001_273-340 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.30936_001_208-378 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.19609_001_1-66 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.35344_001_1-88 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.33219_001_1-73 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.30234_001_488-623 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.14820_001_104-279 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.02294_001_105-280 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.25133_001_12-104 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.25974_001_1-171 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.44623_001_319-495 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.38003_001_157-335 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.42858_001_1-79 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.36896_001_1-88 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.21145_001_1-75 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.40597_001_424-601 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.25847_001_1-170 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.38834_001_93-228 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.01844_001_220-353 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.07619_001_12-188 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.14466_001_308-382 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.29589_001_1-82 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.09260_001_1-105 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.13295_001_105-239 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.36974_001_206-326 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.37360_001_1-90 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.34707_001_36-126 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.12142_001_167-333 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.40527_001_188-353 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.07423_001_1-89 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.29728_001_1-89 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.24679_001_8-79 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.24680_001_802-873 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.44387_001_1-103 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.43029_001_1-92 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.41738_001_55-136 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.27769_001_1-121 | RVT.HG50.0:NA | NA |
Aque_Aqu2.1.25653_001_1-169 | RVT.HG50.0:NA | NA |
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