Retfil_ETO07158_85-252 | RVT.HG48.0:NA | NA |
Exapal_XP_020900039.1_22-152 | RVT.HG48.0:NA | NA |
Lcom_lcpid98052_184-380 | RVT.HG48.0:NA | NA |
Lcom_lcpid57306_339-546 | RVT.HG48.0:NA | NA |
Spis_XP_022786007_1_19-181 | RVT.HG48.0:NA | NA |
Emue_1045g2a_549-762 | RVT.HG48.0:NA | NA |
Emue_1045g3a_549-762 | RVT.HG48.0:NA | NA |
Emue_0009g26a_645-792 | RVT.HG48.0:NA | NA |
Emue_0034g30a_193-340 | RVT.HG48.0:NA | NA |
Aque_Aqu2.1.14984_001_200-367 | RVT.HG48.0:NA | NA |
Aque_Aqu2.1.26661_001_26-139 | RVT.HG48.0:NA | NA |
Aque_Aqu2.1.05977_001_1-89 | RVT.HG48.0:NA | NA |
Tetwil_Twilhelma_g14963.t1_20-170 | RVT.HG48.0:NA | NA |
Auraur_scaffold1983.g3.t1_156-364 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00043472-protein_321-491 | RVT.HG48.0:NA | NA |
Auraur_scaffold1390.g4.t1_16-176 | RVT.HG48.0:NA | NA |
Lpol_XP_013782192.1_121-279 | RVT.HG48.0:NA | NA |
Auraur_scaffold858.g4.t1_267-461 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00041331-protein_102-256 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00038487-protein_19-132 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00037187-protein_107-330 | RVT.HG48.0:NA | NA |
Auraur_scaffold64.g23.t1_95-317 | RVT.HG48.0:NA | NA |
Auraur_scaffold1844.g5.t1_22-234 | RVT.HG48.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g16971.t1_20-229 | RVT.HG48.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g18119.t1_146-303 | RVT.HG48.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g17599.t1_494-645 | RVT.HG48.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g33576.t1_362-527 | RVT.HG48.0:NA | NA |
Ocar_g7298.t1_669-890 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00026815-protein_313-469 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00039426-protein_241-443 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00041609-protein_202-406 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00050836-protein_137-323 | RVT.HG48.0:NA | NA |
Tetwil_Twilhelma_g19174.t1_1-196 | RVT.HG48.0:NA | NA |
Tetwil_Twilhelma_g232.t1_113-322 | RVT.HG48.0:NA | NA |
Emue_0803g1a_246-454 | RVT.HG48.0:NA | NA |
Emue_0034g30a_1407-1615 | RVT.HG48.0:NA | NA |
Emue_0016g6a_471-675 | RVT.HG48.0:NA | NA |
Emue_0002g1714a_364-570 | RVT.HG48.0:NA | NA |
Emue_0010g351a_365-571 | RVT.HG48.0:NA | NA |
Emue_0024g71a_296-441 | RVT.HG48.0:NA | NA |
Emue_0024g5a_202-348 | RVT.HG48.0:NA | NA |
Emue_0209g6a_202-348 | RVT.HG48.0:NA | NA |
Cscu_XP_023235814.1_328-520 | RVT.HG48.0:NA | NA |
Cscu_XP_023230812.1_154-354 | RVT.HG48.0:NA | NA |
Cscu_XP_023240079.1_207-310 | RVT.HG48.0:NA | NA |
Tcas_XP_008201306.2_389-569 | RVT.HG48.0:NA | NA |
Cscu_XP_023243917.1_203-341 | RVT.HG48.0:NA | NA |
Cscu_XP_023212940.1_1-128 | RVT.HG48.0:NA | NA |
Cscu_XP_023210518.1_1-115 | RVT.HG48.0:NA | NA |
Cscu_XP_023230160.1_1-117 | RVT.HG48.0:NA | NA |
Cscu_XP_023241007.1_1-122 | RVT.HG48.0:NA | NA |
Cscu_XP_023213830.1_1-108 | RVT.HG48.0:NA | NA |
Cscu_XP_023231613.1_725-936 | RVT.HG48.0:NA | NA |
Cscu_XP_023212783.1_281-444 | RVT.HG48.0:NA | NA |
Cscu_XP_023244782.1_1-142 | RVT.HG48.0:NA | NA |
Cscu_XP_023210871.1_90-293 | RVT.HG48.0:NA | NA |
Cscu_XP_023225628.1_1-147 | RVT.HG48.0:NA | NA |
Cscu_XP_023223261.1_1-131 | RVT.HG48.0:NA | NA |
Cscu_XP_023238093.1_159-358 | RVT.HG48.0:NA | NA |
Cscu_XP_023211578.1_8-153 | RVT.HG48.0:NA | NA |
Cscu_XP_023209413.1_144-323 | RVT.HG48.0:NA | NA |
Cscu_XP_023239508.1_1-129 | RVT.HG48.0:NA | NA |
Cscu_XP_023225421.1_1-126 | RVT.HG48.0:NA | NA |
Cscu_XP_023236949.1_1-202 | RVT.HG48.0:NA | NA |
Cscu_XP_023223814.1_20-165 | RVT.HG48.0:NA | NA |
Cscu_XP_023209493.1_1-203 | RVT.HG48.0:NA | NA |
Cscu_XP_023213085.1_206-386 | RVT.HG48.0:NA | NA |
Cscu_XP_023219622.1_316-485 | RVT.HG48.0:NA | NA |
Cscu_XP_023235630.1_153-259 | RVT.HG48.0:NA | NA |
Cscu_XP_023235632.1_113-224 | RVT.HG48.0:NA | NA |
Cscu_XP_023238336.1_114-224 | RVT.HG48.0:NA | NA |
Cscu_XP_023214216.1_110-233 | RVT.HG48.0:NA | NA |
Cscu_XP_023222921.1_43-165 | RVT.HG48.0:NA | NA |
Cscu_XP_023238597.1_350-509 | RVT.HG48.0:NA | NA |
Cscu_XP_023213254.1_267-387 | RVT.HG48.0:NA | NA |
Cscu_XP_023238597.1_565-689 | RVT.HG48.0:NA | NA |
Cscu_XP_023213574.1_205-352 | RVT.HG48.0:NA | NA |
Cscu_XP_023232189.1_7-121 | RVT.HG48.0:NA | NA |
Cscu_XP_023213573.1_201-378 | RVT.HG48.0:NA | NA |
Cscu_XP_023213577.1_201-347 | RVT.HG48.0:NA | NA |
Cscu_XP_023212126.1_1-117 | RVT.HG48.0:NA | NA |
Cscu_XP_023212631.1_16-184 | RVT.HG48.0:NA | NA |
Cscu_XP_023213575.1_166-336 | RVT.HG48.0:NA | NA |
Cscu_XP_023222203.1_122-247 | RVT.HG48.0:NA | NA |
Cscu_XP_023212221.1_270-425 | RVT.HG48.0:NA | NA |
Cscu_XP_023212227.1_193-348 | RVT.HG48.0:NA | NA |
Cscu_XP_023222202.1_130-285 | RVT.HG48.0:NA | NA |
Cscu_XP_023212222.1_242-397 | RVT.HG48.0:NA | NA |
Cscu_XP_023222200.1_130-285 | RVT.HG48.0:NA | NA |
Cscu_XP_023222198.1_130-285 | RVT.HG48.0:NA | NA |
Cscu_XP_023212226.1_221-376 | RVT.HG48.0:NA | NA |
Cscu_XP_023222201.1_102-257 | RVT.HG48.0:NA | NA |
Cscu_XP_023212223.1_270-425 | RVT.HG48.0:NA | NA |
Cscu_XP_023212219.1_270-425 | RVT.HG48.0:NA | NA |
Cscu_XP_023236747.1_51-206 | RVT.HG48.0:NA | NA |
Exapal_XP_020914493.1_73-215 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00043692-protein_266-421 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00059660-protein_72-236 | RVT.HG48.0:NA | NA |
Morvir_scaffold506.g11.t1_252-447 | RVT.HG48.0:NA | NA |
Morvir_scaffold428.g7.t1_191-386 | RVT.HG48.0:NA | NA |
Morvir_scaffold65.g53.t1_327-522 | RVT.HG48.0:NA | NA |
Morvir_scaffold122.g22.t1_391-525 | RVT.HG48.0:NA | NA |
Morvir_scaffold573.g1.t1_92-278 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00063239-protein_19-192 | RVT.HG48.0:NA | NA |
Mlei_ML0513_g18_t1_846-1048 | RVT.HG48.0:NA | NA |
Mlei_ML0622_g3_t1_73-197 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00034510-protein_40-142 | RVT.HG48.0:NA | NA |
Adig_XP_015764052.1_4-187 | RVT.HG48.0:NA | NA |
Adig_XP_015763782.1_4-191 | RVT.HG48.0:NA | NA |
Clyhem_TCONS_00027061-protein_1-196 | RVT.HG48.0:NA | NA |
Exapal_XP_020895056.1_4-198 | RVT.HG48.0:NA | NA |
Exapal_XP_020906008.1_75-269 | RVT.HG48.0:NA | NA |
Scil_scpid46514_346-548 | RVT.HG48.0:NA | NA |
Spis_Spis24110_1_1-147 | RVT.HG48.0:NA | NA |
Adig_XP_015770368.1_32-192 | RVT.HG48.0:NA | NA |
Exapal_XP_020911960.1_18-114 | RVT.HG48.0:NA | NA |
Spis_XP_022802357_1_74-280 | RVT.HG48.0:NA | NA |
Spis_Spis12664_1_59-265 | RVT.HG48.0:NA | NA |
Spis_XP_022805640_1_74-280 | RVT.HG48.0:NA | NA |
Adig_XP_015777507.1_11-171 | RVT.HG48.0:NA | NA |
Adig_XP_015776012.1_142-320 | RVT.HG48.0:NA | NA |
Spis_XP_022805774_1_1-173 | RVT.HG48.0:NA | NA |
Spis_Spis6644_1_232-341 | RVT.HG48.0:NA | NA |
Spis_Spis21821_1_467-654 | RVT.HG48.0:NA | NA |
Spis_Spis11611_1_98-286 | RVT.HG48.0:NA | NA |
Spis_Spis12704_1_254-442 | RVT.HG48.0:NA | NA |
Apla_XP_022111224.1_123-332 | RVT.HG48.0:NA | NA |
Apla_XP_022104795.1_1-208 | RVT.HG48.0:NA | NA |
Morvir_scaffold341.g11.t1_628-840 | RVT.HG48.0:NA | NA |
Morvir_scaffold453.g10.t1_328-540 | RVT.HG48.0:NA | NA |
Morvir_scaffold210.g14.t1_328-540 | RVT.HG48.0:NA | NA |
Morvir_scaffold512.g2.t1_328-540 | RVT.HG48.0:NA | NA |
Bralan_BL02516_cuf0_32-136 | RVT.HG48.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g22530.t1_84-286 | RVT.HG48.0:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g21778.t1_67-177 | RVT.HG48.0:NA | NA |
Bralan_BL26352_evm0_119-216 | RVT.HG48.0:NA | NA |
Bralan_BL70677_cuf0_67-178 | RVT.HG48.0:NA | NA |
Spis_XP_022796700_1_42-241 | RVT.HG48.0:NA | NA |
Spis_XP_022783320_1_1-158 | RVT.HG48.0:NA | NA |
Spis_Spis20803_1_276-471 | RVT.HG48.0:NA | NA |
Spis_Spis24737_1_82-277 | RVT.HG48.0:NA | NA |
Spis_Spis20695_1_24-187 | RVT.HG48.0:NA | NA |
Spis_Spis8415_1_24-187 | RVT.HG48.0:NA | NA |
Bralan_BL25520_evm0_156-375 | RVT.HG48.0:NA | NA |
Bralan_BL31775_evm0_22-204 | RVT.HG48.0:NA | NA |
Apla_XP_022104796.1_259-364 | RVT.HG48.0:NA | NA |
Apla_XP_022096535.1_31-131 | RVT.HG48.0:NA | NA |
Apla_XP_022103511.1_20-139 | RVT.HG48.0:NA | NA |
Apla_XP_022090187.1_20-140 | RVT.HG48.0:NA | NA |
Apla_XP_022110786.1_1-102 | RVT.HG48.0:NA | NA |
Apla_XP_022087613.1_89-209 | RVT.HG48.0:NA | NA |
Apla_XP_022105762.1_113-195 | RVT.HG48.0:NA | NA |
Spur_XP_003724886.1_1-139 | RVT.HG48.0:NA | NA |
Spur_XP_030834186.1_1-139 | RVT.HG48.0:NA | NA |
Spur_XP_030842334.1_342-532 | RVT.HG48.0:NA | NA |
Spur_XP_030842331.1_332-532 | RVT.HG48.0:NA | NA |
Spur_XP_030842332.1_82-282 | RVT.HG48.0:NA | NA |
Spur_XP_003730115.2_332-532 | RVT.HG48.0:NA | NA |
Spur_XP_030849739.1_71-272 | RVT.HG48.0:NA | NA |
Spur_XP_030838373.1_9-193 | RVT.HG48.0:NA | NA |
Spur_XP_030855968.1_335-531 | RVT.HG48.0:NA | NA |
Spur_XP_030855974.1_388-584 | RVT.HG48.0:NA | NA |
Spur_XP_030838458.1_222-423 | RVT.HG48.0:NA | NA |
Spur_XP_011666638.2_338-529 | RVT.HG48.0:NA | NA |
Spur_XP_030841919.1_318-509 | RVT.HG48.0:NA | NA |
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