Aque_Aqu2.1.36374_001_201-342 | RVT.HG58.0:NA | NA |
Aque_Aqu2.1.07710_001_754-911 | RVT.HG58.0:NA | NA |
Perkma_EER12830_95-255 | RVT.HG58.0:NA | NA |
Perkma_EEQ97278_123-265 | RVT.HG58.0:NA | NA |
Emue_0008g164a_860-1010 | RVT.HG58.0:NA | NA |
Emue_0017g723a_458-608 | RVT.HG58.0:NA | NA |
Emue_0007g727a_798-948 | RVT.HG58.0:NA | NA |
Auraur_scaffold2413.g4.t1_144-294 | RVT.HG58.0:NA | NA |
Auraur_scaffold1909.g1.t1_517-666 | RVT.HG58.0:NA | NA |
Auraur_scaffold243.g8.t1_924-1070 | RVT.HG58.0:NA | NA |
Morvir_scaffold318.g23.t1_375-520 | RVT.HG58.0:NA | NA |
Morvir_scaffold958.g2.t1_26-175 | RVT.HG58.0:NA | NA |
Morvir_scaffold157.g44.t1_766-911 | RVT.HG58.0:NA | NA |
Morvir_scaffold2.g98.t1_593-738 | RVT.HG58.0:NA | NA |
Morvir_scaffold45.g85.t1_593-738 | RVT.HG58.0:NA | NA |
Morvir_scaffold5.g146.t1_105-246 | RVT.HG58.0:NA | NA |
Morvir_scaffold58.g30.t1_841-985 | RVT.HG58.0:NA | NA |
Morvir_scaffold71.g2.t1_520-662 | RVT.HG58.0:NA | NA |
Auraur_scaffold334.g15.t1_185-360 | RVT.HG58.2:NA | NA |
Clyhem_TCONS_00060908-protein_99-243 | RVT.HG58.1:NA | NA |
Auraur_scaffold3396.g2.t1_342-484 | RVT.HG58.1:NA | NA |
Adig_XP_015776304.1_734-889 | RVT.HG58.1:NA | NA |
Auraur_scaffold1534.g3.t1_487-635 | RVT.HG58.1:NA | NA |
Auraur_scaffold273.g2.t1_121-264 | RVT.HG58.1:NA | NA |
Auraur_scaffold6677.g1.t1_110-240 | RVT.HG58.1:NA | NA |
Auraur_scaffold1062.g8.t1_523-669 | RVT.HG58.1:NA | NA |
Morvir_scaffold14.g11.t1_803-951 | RVT.HG58.1:NA | NA |
Morvir_scaffold322.g3.t1_38-184 | RVT.HG58.1:NA | NA |
Morvir_scaffold1.g62.t1_110-258 | RVT.HG58.1:NA | NA |
Morvir_scaffold272.g4.t1_1188-1336 | RVT.HG58.1:NA | NA |
Morvir_scaffold89.g13.t1_1342-1490 | RVT.HG58.1:NA | NA |
Morvir_scaffold6.g51.t1_882-1030 | RVT.HG58.1:NA | NA |
Morvir_scaffold14.g42.t1_1173-1321 | RVT.HG58.1:NA | NA |
Morvir_scaffold89.g13.t3_1058-1206 | RVT.HG58.1:NA | NA |
Scil_scpid107402_5-163 | RVT.HG58.1:NA | NA |
Scil_scpid11757_698-839 | RVT.HG58.1:NA | NA |
Scil_scpid20451_647-788 | RVT.HG58.1:NA | NA |
Scil_scpid41734_697-843 | RVT.HG58.1:NA | NA |
Scil_scpid41376_56-199 | RVT.HG58.1:NA | NA |
Scil_scpid6188_700-842 | RVT.HG58.1:NA | NA |
Lcom_lcpid36985_68-203 | RVT.HG58.1:NA | NA |
Lcom_lcpid105263_262-404 | RVT.HG58.1:NA | NA |
Scil_scpid17799_675-843 | RVT.HG58.1:NA | NA |
Lcom_lcpid28914_375-514 | RVT.HG58.1:NA | NA |
Scil_scpid84063_350-485 | RVT.HG58.1:NA | NA |
Morvir_scaffold198.g3.t1_779-935 | RVT.HG58.1:NA | NA |
Linana_g741.t1_381-510 | RVT.HG58.1:NA | NA |
Notgen_g16792.t1_109-296 | RVT.HG58.1:NA | NA |
Auraur_scaffold1832.g3.t1_94-231 | RVT.HG58.1:NA | NA |
Auraur_scaffold763.g7.t1_196-337 | RVT.HG58.1:NA | NA |
Scil_scpid7544_961-1102 | RVT.HG58.1:NA | NA |
Scil_scpid28305_738-883 | RVT.HG58.1:NA | NA |
Scil_scpid3998_1255-1402 | RVT.HG58.1:NA | NA |
Notgen_g25529.t1_6-147 | RVT.HG58.1:NA | NA |
Notgen_g38770.t1_84-232 | RVT.HG58.1:NA | NA |
Notgen_g6701.t1_94-232 | RVT.HG58.1:NA | NA |
Notgen_g12700.t1_942-1080 | RVT.HG58.1:NA | NA |
Scil_scpid14339_919-1056 | RVT.HG58.1:NA | NA |
Scil_scpid15785_778-914 | RVT.HG58.1:NA | NA |
Scil_scpid49170_137-281 | RVT.HG58.1:NA | NA |
Lcom_lcpid77365_108-253 | RVT.HG58.1:NA | NA |
Lcom_lcpid79344_108-253 | RVT.HG58.1:NA | NA |
Scil_scpid16991_261-407 | RVT.HG58.1:NA | NA |
Morvir_scaffold192.g5.t1_777-916 | RVT.HG58.1:NA | NA |
Nvec_NVE5622_764-893 | RVT.HG58.1:NA | NA |
Adig_XP_015767845.1_45-184 | RVT.HG58.1:NA | NA |
Adig_XP_015773044.1_89-234 | RVT.HG58.1:NA | NA |
Adig_XP_015757953.1_848-987 | RVT.HG58.1:NA | NA |
Adig_XP_015747436.1_230-372 | RVT.HG58.1:NA | NA |
Exapal_XP_020899647.1_523-660 | RVT.HG58.1:NA | NA |
Auraur_scaffold1077.g2.t1_824-961 | RVT.HG58.1:NA | NA |
Notgen_g27431.t1_146-294 | RVT.HG58.1:NA | NA |
Phoaus_g19691.t1_815-951 | RVT.HG58.1:NA | NA |
Phoaus_g3407.t1_281-418 | RVT.HG58.1:NA | NA |
Morvir_scaffold174.g40.t1_868-1040 | RVT.HG58.1:NA | NA |
Morvir_scaffold77.g77.t1_868-1040 | RVT.HG58.1:NA | NA |
Lcom_lcpid93504_184-320 | RVT.HG58.1:NA | NA |
Lcom_lcpid7428_1152-1288 | RVT.HG58.1:NA | NA |
Scil_scpid5460_1163-1300 | RVT.HG58.1:NA | NA |
Adig_XP_015760854.1_350-497 | RVT.HG58.1:NA | NA |
Exapal_XP_020898652.1_838-1003 | RVT.HG58.1:NA | NA |
Phoaus_g11884.t1_247-424 | RVT.HG58.1:NA | NA |
Notgen_g6408.t1_459-605 | RVT.HG58.1:NA | NA |
Skow_XP_006816890.1_309-457 | RVT.HG58.1:NA | NA |
Skow_XP_006824596.1_56-190 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g30295.t1_74-221 | RVT.HG58.1:NA | NA |
Skow_XP_002734902.1_820-967 | RVT.HG58.1:NA | NA |
Skow_XP_006816675.1_2-141 | RVT.HG58.1:NA | NA |
Skow_XP_006813576.1_73-199 | RVT.HG58.1:NA | NA |
Skow_XP_006814829.1_836-978 | RVT.HG58.1:NA | NA |
Skow_XP_006812926.1_425-573 | RVT.HG58.1:NA | NA |
Skow_XP_006821304.1_364-514 | RVT.HG58.1:NA | NA |
Skow_XP_006817016.1_265-405 | RVT.HG58.1:NA | NA |
Skow_XP_006820435.1_336-512 | RVT.HG58.1:NA | NA |
Skow_XP_006822449.1_84-212 | RVT.HG58.1:NA | NA |
Skow_XP_006818999.1_4-152 | RVT.HG58.1:NA | NA |
Skow_XP_006815892.1_74-252 | RVT.HG58.1:NA | NA |
Skow_XP_006815261.1_28-221 | RVT.HG58.1:NA | NA |
Notgen_g21304.t1_105-251 | RVT.HG58.1:NA | NA |
Skow_XP_006812892.1_758-910 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g9844.t1_238-385 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g10359.t1_606-752 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g7560.t1_239-385 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g23881.t1_558-705 | RVT.HG58.1:NA | NA |
Notgen_g5642.t1_460-607 | RVT.HG58.1:NA | NA |
Skow_XP_006816667.1_214-392 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g996.t1_429-566 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g8202.t1_11-149 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g13747.t1_298-444 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g10612.t1_73-205 | RVT.HG58.1:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g17406.t1_57-192 | RVT.HG58.1:NA | NA |
Notgen_g37030.t1_564-736 | RVT.HG58.1:NA | NA |
Notgen_g10564.t1_772-917 | RVT.HG58.1:NA | NA |
Skow_XP_006815871.1_56-191 | RVT.HG58.1:NA | NA |
Skow_XP_002732171.1_742-905 | RVT.HG58.1:NA | NA |
Skow_XP_006813289.1_1-156 | RVT.HG58.1:NA | NA |
0.416667