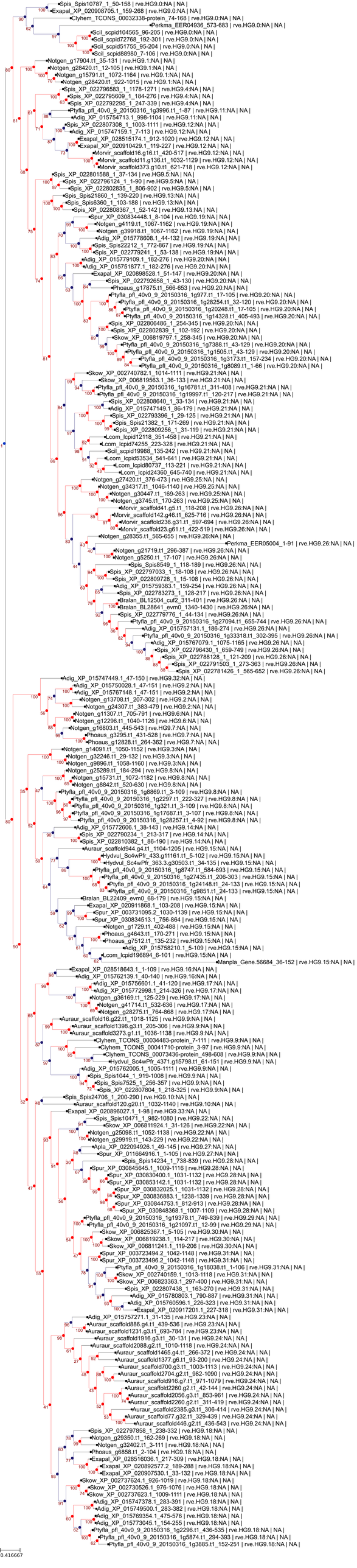
Spis_Spis10787_1_50-158 | rve.HG9.0:NA | NA |
Exapal_XP_020908705.1_159-268 | rve.HG9.0:NA | NA |
Clyhem_TCONS_00032338-protein_74-168 | rve.HG9.0:NA | NA |
Perkma_EER04936_573-683 | rve.HG9.0:NA | NA |
Scil_scpid104565_96-205 | rve.HG9.0:NA | NA |
Scil_scpid72768_192-301 | rve.HG9.0:NA | NA |
Scil_scpid51755_95-204 | rve.HG9.0:NA | NA |
Scil_scpid88980_7-106 | rve.HG9.0:NA | NA |
Notgen_g17904.t1_35-131 | rve.HG9.1:NA | NA |
Notgen_g28420.t1_12-105 | rve.HG9.1:NA | NA |
Notgen_g15791.t1_1072-1164 | rve.HG9.1:NA | NA |
Notgen_g28420.t1_922-1015 | rve.HG9.1:NA | NA |
Spis_XP_022796583_1_1178-1271 | rve.HG9.4:NA | NA |
Spis_XP_022795609_1_184-276 | rve.HG9.4:NA | NA |
Spis_XP_022792295_1_247-339 | rve.HG9.4:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g3996.t1_1-87 | rve.HG9.11:NA | NA |
Adig_XP_015754713.1_998-1104 | rve.HG9.11:NA | NA |
Spis_XP_022807308_1_1003-1111 | rve.HG9.12:NA | NA |
Adig_XP_015747159.1_7-113 | rve.HG9.12:NA | NA |
Exapal_XP_028515174.1_912-1020 | rve.HG9.12:NA | NA |
Exapal_XP_020910429.1_119-227 | rve.HG9.12:NA | NA |
Morvir_scaffold16.g16.t1_420-517 | rve.HG9.12:NA | NA |
Morvir_scaffold11.g136.t1_1032-1129 | rve.HG9.12:NA | NA |
Morvir_scaffold373.g10.t1_621-718 | rve.HG9.12:NA | NA |
Spis_XP_022801588_1_37-134 | rve.HG9.5:NA | NA |
Spis_XP_022796124_1_1-90 | rve.HG9.5:NA | NA |
Spis_XP_022802835_1_806-902 | rve.HG9.5:NA | NA |
Spis_Spis21860_1_139-220 | rve.HG9.13:NA | NA |
Spis_Spis6360_1_103-188 | rve.HG9.13:NA | NA |
Spis_XP_022808367_1_52-142 | rve.HG9.13:NA | NA |
Spur_XP_030834448.1_8-104 | rve.HG9.19:NA | NA |
Notgen_g4119.t1_1067-1162 | rve.HG9.19:NA | NA |
Notgen_g39918.t1_1067-1162 | rve.HG9.19:NA | NA |
Adig_XP_015778608.1_44-132 | rve.HG9.19:NA | NA |
Spis_Spis22212_1_772-867 | rve.HG9.19:NA | NA |
Spis_XP_022779241_1_53-138 | rve.HG9.19:NA | NA |
Adig_XP_015779109.1_182-276 | rve.HG9.20:NA | NA |
Adig_XP_015751877.1_182-276 | rve.HG9.20:NA | NA |
Exapal_XP_020898528.1_51-147 | rve.HG9.20:NA | NA |
Spis_XP_022792658_1_43-130 | rve.HG9.20:NA | NA |
Phoaus_g17875.t1_566-653 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g977.t1_17-105 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g28254.t1_32-120 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g20248.t1_17-105 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g14328.t1_405-493 | rve.HG9.20:NA | NA |
Spis_XP_022806486_1_254-345 | rve.HG9.20:NA | NA |
Spis_XP_022802839_1_102-192 | rve.HG9.20:NA | NA |
Skow_XP_006819797.1_258-345 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g7388.t1_43-129 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g1505.t1_43-129 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g3173.t1_157-234 | rve.HG9.20:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g8089.t1_1-66 | rve.HG9.20:NA | NA |
Skow_XP_002740782.1_1014-1111 | rve.HG9.21:NA | NA |
Skow_XP_006819563.1_36-133 | rve.HG9.21:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g16781.t1_311-408 | rve.HG9.21:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g19997.t1_120-217 | rve.HG9.21:NA | NA |
Spis_XP_022808640_1_33-134 | rve.HG9.21:NA | NA |
Adig_XP_015747149.1_86-179 | rve.HG9.21:NA | NA |
Spis_XP_022793396_1_29-125 | rve.HG9.21:NA | NA |
Spis_Spis21382_1_171-269 | rve.HG9.21:NA | NA |
Spis_XP_022809256_1_31-119 | rve.HG9.21:NA | NA |
Lcom_lcpid12118_351-458 | rve.HG9.21:NA | NA |
Lcom_lcpid74255_223-328 | rve.HG9.21:NA | NA |
Scil_scpid19988_135-242 | rve.HG9.21:NA | NA |
Lcom_lcpid53534_541-641 | rve.HG9.21:NA | NA |
Lcom_lcpid80737_113-221 | rve.HG9.21:NA | NA |
Lcom_lcpid24360_645-740 | rve.HG9.21:NA | NA |
Notgen_g27420.t1_376-473 | rve.HG9.25:NA | NA |
Notgen_g34317.t1_1046-1140 | rve.HG9.25:NA | NA |
Notgen_g30447.t1_169-263 | rve.HG9.25:NA | NA |
Notgen_g3745.t1_170-263 | rve.HG9.25:NA | NA |
Morvir_scaffold41.g5.t1_118-208 | rve.HG9.26:NA | NA |
Morvir_scaffold142.g46.t1_625-716 | rve.HG9.26:NA | NA |
Morvir_scaffold236.g31.t1_597-694 | rve.HG9.26:NA | NA |
Morvir_scaffold223.g61.t1_422-519 | rve.HG9.26:NA | NA |
Notgen_g28355.t1_565-655 | rve.HG9.26:NA | NA |
Perkma_EER05004_1-91 | rve.HG9.26:NA | NA |
Notgen_g21719.t1_296-387 | rve.HG9.26:NA | NA |
Notgen_g5250.t1_17-107 | rve.HG9.26:NA | NA |
Spis_Spis8549_1_118-189 | rve.HG9.26:NA | NA |
Spis_XP_022797033_1_18-108 | rve.HG9.26:NA | NA |
Spis_XP_022809728_1_15-108 | rve.HG9.26:NA | NA |
Adig_XP_015759383.1_159-254 | rve.HG9.26:NA | NA |
Spis_XP_022783273_1_128-217 | rve.HG9.26:NA | NA |
Bralan_BL12504_cuf2_311-401 | rve.HG9.26:NA | NA |
Bralan_BL28641_evm0_1340-1430 | rve.HG9.26:NA | NA |
Spis_XP_022779776_1_44-134 | rve.HG9.26:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g2709.t1_655-744 | rve.HG9.26:NA | NA |
Adig_XP_015757131.1_186-274 | rve.HG9.26:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g33318.t1_302-395 | rve.HG9.26:NA | NA |
Adig_XP_015767079.1_1075-1165 | rve.HG9.26:NA | NA |
Spis_XP_022796430_1_659-749 | rve.HG9.26:NA | NA |
Spis_XP_022788128_1_121-209 | rve.HG9.26:NA | NA |
Spis_XP_022791503_1_273-363 | rve.HG9.26:NA | NA |
Spis_XP_022781426_1_565-652 | rve.HG9.26:NA | NA |
Adig_XP_015747449.1_47-150 | rve.HG9.32:NA | NA |
Adig_XP_015750028.1_47-151 | rve.HG9.2:NA | NA |
Adig_XP_015767148.1_47-151 | rve.HG9.2:NA | NA |
Notgen_g13708.t1_207-302 | rve.HG9.2:NA | NA |
Notgen_g24307.t1_383-479 | rve.HG9.2:NA | NA |
Notgen_g11307.t1_705-791 | rve.HG9.6:NA | NA |
Notgen_g12296.t1_1040-1126 | rve.HG9.6:NA | NA |
Notgen_g16803.t1_445-543 | rve.HG9.7:NA | NA |
Phoaus_g3295.t1_431-528 | rve.HG9.7:NA | NA |
Phoaus_g12828.t1_264-362 | rve.HG9.7:NA | NA |
Notgen_g14091.t1_1050-1152 | rve.HG9.3:NA | NA |
Notgen_g32246.t1_29-132 | rve.HG9.3:NA | NA |
Notgen_g9896.t1_1058-1160 | rve.HG9.3:NA | NA |
Notgen_g25289.t1_184-294 | rve.HG9.8:NA | NA |
Notgen_g15731.t1_1072-1182 | rve.HG9.8:NA | NA |
Notgen_g8842.t1_520-630 | rve.HG9.8:NA | NA |
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Ptyfla_pfl_40v0_9_20150316_1g2297.t1_222-327 | rve.HG9.8:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g321.t1_3-109 | rve.HG9.8:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g17687.t1_3-107 | rve.HG9.8:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g28257.t1_4-92 | rve.HG9.8:NA | NA |
Adig_XP_015772606.1_38-143 | rve.HG9.14:NA | NA |
Spis_XP_022790234_1_213-317 | rve.HG9.14:NA | NA |
Spis_XP_022810382_1_86-190 | rve.HG9.14:NA | NA |
Aurour_scaffold944.g4.t1_1104-1205 | rve.HG9.15:NA | NA |
Hydvul_Sc4wPfr_433.g11161.t1_5-102 | rve.HG9.15:NA | NA |
Hydvul_Sc4wPfr_363.3.g30503.t1_34-135 | rve.HG9.15:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g8747.t1_584-693 | rve.HG9.15:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g27435.t1_206-303 | rve.HG9.15:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g24148.t1_24-133 | rve.HG9.15:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g9851.t1_24-133 | rve.HG9.15:NA | NA |
Bralan_BL22409_evm0_68-179 | rve.HG9.15:NA | NA |
Exapal_XP_020911868.1_103-208 | rve.HG9.15:NA | NA |
Spur_XP_003731095.2_1030-1139 | rve.HG9.15:NA | NA |
Spur_XP_030834513.1_756-864 | rve.HG9.15:NA | NA |
Notgen_g1729.t1_402-488 | rve.HG9.15:NA | NA |
Phoaus_g4643.t1_170-271 | rve.HG9.15:NA | NA |
Phoaus_g7512.t1_135-232 | rve.HG9.15:NA | NA |
Adig_XP_015758210.1_5-109 | rve.HG9.15:NA | NA |
Lcom_lcpid196894_6-101 | rve.HG9.15:NA | NA |
Manpla_Gene.56684_36-152 | rve.HG9.15:NA | NA |
Exapal_XP_028518643.1_1-109 | rve.HG9.16:NA | NA |
Adig_XP_015762139.1_40-140 | rve.HG9.16:NA | NA |
Adig_XP_015756601.1_41-120 | rve.HG9.17:NA | NA |
Adig_XP_015772998.1_214-326 | rve.HG9.17:NA | NA |
Notgen_g36169.t1_125-229 | rve.HG9.17:NA | NA |
Notgen_g41714.t1_532-636 | rve.HG9.17:NA | NA |
Notgen_g28275.t1_764-868 | rve.HG9.17:NA | NA |
Aurour_scaffold16.g22.t1_1018-1125 | rve.HG9.9:NA | NA |
Aurour_scaffold1398.g3.t1_205-306 | rve.HG9.9:NA | NA |
Aurour_scaffold3273.g1.t1_1036-1138 | rve.HG9.9:NA | NA |
Clyhem_TCONS_00034483-protein_7-111 | rve.HG9.9:NA | NA |
Clyhem_TCONS_00041710-protein_3-97 | rve.HG9.9:NA | NA |
Clyhem_TCONS_00073436-protein_498-608 | rve.HG9.9:NA | NA |
Hydvul_Sc4wPfr_4371.g15798.t1_61-151 | rve.HG9.9:NA | NA |
Adig_XP_015762005.1_1005-1111 | rve.HG9.9:NA | NA |
Spis_Spis1044_1_919-1008 | rve.HG9.9:NA | NA |
Spis_Spis7525_1_256-357 | rve.HG9.9:NA | NA |
Spis_XP_022807804_1_218-325 | rve.HG9.9:NA | NA |
Spis_Spis24706_1_200-290 | rve.HG9.10:NA | NA |
Aurour_scaffold120.g20.t1_1032-1140 | rve.HG9.10:NA | NA |
Exapal_XP_020896027.1_1-98 | rve.HG9.33:NA | NA |
Spis_Spis10471_1_982-1080 | rve.HG9.22:NA | NA |
Skow_XP_006811924.1_31-126 | rve.HG9.22:NA | NA |
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Notgen_g29919.t1_143-229 | rve.HG9.22:NA | NA |
Apla_XP_022094926.1_49-145 | rve.HG9.27:NA | NA |
Spur_XP_011664916.1_1-105 | rve.HG9.27:NA | NA |
Spis_Spis14234_1_738-839 | rve.HG9.28:NA | NA |
Spur_XP_030845645.1_1009-1116 | rve.HG9.28:NA | NA |
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Spur_XP_030853142.1_1031-1132 | rve.HG9.28:NA | NA |
Spur_XP_030832025.1_1031-1132 | rve.HG9.28:NA | NA |
Spur_XP_030836883.1_1238-1339 | rve.HG9.28:NA | NA |
Spur_XP_030844753.1_812-913 | rve.HG9.28:NA | NA |
Spur_XP_030848368.1_1007-1109 | rve.HG9.28:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g19378.t1_749-839 | rve.HG9.29:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g21097.t1_12-99 | rve.HG9.29:NA | NA |
Skow_XP_006825367.1_5-105 | rve.HG9.30:NA | NA |
Skow_XP_006819238.1_114-217 | rve.HG9.30:NA | NA |
Skow_XP_006811241.1_119-206 | rve.HG9.30:NA | NA |
Spur_XP_003723494.2_1042-1148 | rve.HG9.31:NA | NA |
Spur_XP_003723496.2_1042-1148 | rve.HG9.31:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g18038.t1_1-106 | rve.HG9.31:NA | NA |
Skow_XP_002740159.1_1013-1118 | rve.HG9.31:NA | NA |
Skow_XP_006823363.1_297-400 | rve.HG9.31:NA | NA |
Spis_XP_022807438_1_163-270 | rve.HG9.31:NA | NA |
Adig_XP_015780803.1_790-887 | rve.HG9.31:NA | NA |
Adig_XP_015760596.1_226-323 | rve.HG9.31:NA | NA |
Exapal_XP_020917201.1_227-318 | rve.HG9.31:NA | NA |
Adig_XP_015757271.1_31-135 | rve.HG9.23:NA | NA |
Aurour_scaffold886.g4.t1_439-536 | rve.HG9.23:NA | NA |
Aurour_scaffold1231.g3.t1_693-784 | rve.HG9.23:NA | NA |
Aurour_scaffold1916.g3.t1_30-131 | rve.HG9.24:NA | NA |
Aurour_scaffold2088.g2.t1_1010-1118 | rve.HG9.24:NA | NA |
Aurour_scaffold1465.g4.t1_266-372 | rve.HG9.24:NA | NA |
Aurour_scaffold1377.g6.t1_93-200 | rve.HG9.24:NA | NA |
Aurour_scaffold700.g3.t1_1003-1113 | rve.HG9.24:NA | NA |
Aurour_scaffold2704.g2.t1_982-1090 | rve.HG9.24:NA | NA |
Aurour_scaffold916.g7.t1_971-1079 | rve.HG9.24:NA | NA |
Aurour_scaffold2260.g2.t1_42-144 | rve.HG9.24:NA | NA |
Aurour_scaffold2056.g3.t1_853-961 | rve.HG9.24:NA | NA |
Aurour_scaffold2260.g2.t1_311-419 | rve.HG9.24:NA | NA |
Aurour_scaffold2385.g3.t1_306-414 | rve.HG9.24:NA | NA |
Aurour_scaffold77.g32.t1_329-439 | rve.HG9.24:NA | NA |
Aurour_scaffold446.g2.t1_436-543 | rve.HG9.24:NA | NA |
Spis_XP_022797858_1_238-332 | rve.HG9.18:NA | NA |
Notgen_g29350.t1_162-269 | rve.HG9.18:NA | NA |
Notgen_g32402.t1_3-111 | rve.HG9.18:NA | NA |
Phoaus_g6858.t1_2-104 | rve.HG9.18:NA | NA |
Exapal_XP_028516036.1_217-309 | rve.HG9.18:NA | NA |
Exapal_XP_020892577.2_189-288 | rve.HG9.18:NA | NA |
Exapal_XP_020907530.1_33-132 | rve.HG9.18:NA | NA |
Skow_XP_002737624.1_926-1019 | rve.HG9.18:NA | NA |
Skow_XP_002730526.1_976-1076 | rve.HG9.18:NA | NA |
Skow_XP_002737623.1_1009-1111 | rve.HG9.18:NA | NA |
Adig_XP_015747378.1_283-391 | rve.HG9.18:NA | NA |
Adig_XP_015749500.1_283-382 | rve.HG9.18:NA | NA |
Adig_XP_015769354.1_475-576 | rve.HG9.18:NA | NA |
Adig_XP_015773045.1_154-255 | rve.HG9.18:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g2296.t1_436-535 | rve.HG9.18:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g5874.t1_294-393 | rve.HG9.18:NA | NA |
Ptyfla_pfl_40v0_9_20150316_1g3885.t1_152-251 | rve.HG9.18:NA | NA |
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