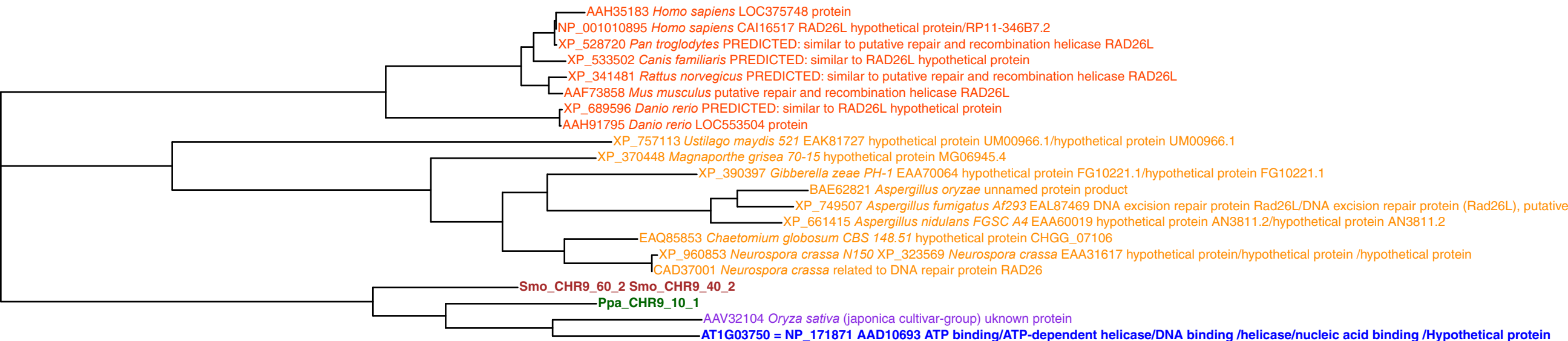CHR9(At1g03750): global tree
File name: CHR9.mltree2.svg
478 aa residues 21 OTU included.
Assemble Job Id: FJI4Wt
Tree job id: Niexdo
BP values out of 100
Analyzed by yoguri@nibb.ac.jp
Date: Fri Aug 25 10:45:08 JST 2006

AAH35183 *Homo sapiens* LOC375748 protein
NP_001010895 *Homo sapiens* CAI16517 RAD26L hypothetical protein/RP11-346B7.2
XP_528720 *Pan troglodytes* PREDICTED: similar to putative repair and recombination helicase RAD26L
XP_533502 *Canis familiaris* PREDICTED: similar to RAD26L hypothetical protein
XP_341481 *Rattus norvegicus* PREDICTED: similar to putative repair and recombination helicase RAD26L
AAF73858 *Mus musculus* putative repair and recombination helicase RAD26L
XP_689596 *Danio rerio* PREDICTED: similar to RAD26L hypothetical protein
AAH91795 *Danio rerio* LOC553504 protein
XP_757113 *Ustilago maydis 521* EAK81727 hypothetical protein UM00966.1/hypothetical protein UM00966.1
XP_370448 *Magnaporthe grisea 70-15* hypothetical protein MG06945.4
XP_390397 *Gibberella zeae PH-1* EAA70064 hypothetical protein FG10221.1/hypothetical protein FG10221.1
BAE62821 *Aspergillus oryzae* unnamed protein product
XP_749507 *Aspergillus fumigatus Af293* EAL87469 DNA excision repair protein Rad26L/DNA excision repair protein (Rad26L), putative
XP_661415 *Aspergillus nidulans FGSC A4* EAA60019 hypothetical protein AN3811.2/hypothetical protein AN3811.2
EAQ85853 *Chaetomium globosum CBS 148.51* hypothetical protein CHGG_07106
XP_960853 *Neurospora crassa N150* XP_323569 *Neurospora crassa* EAA31617 hypothetical protein/hypothetical protein /hypothetical protein
CAD37001 *Neurospora crassa* related to DNA repair protein RAD26
Smo_CHR9_60_2 Smo_CHR9_40_2
Ppa_CHR9_10_1
AAV32104 *Oryza sativa* (japonica cultivar-group) uknown protein
AT1G03750 = NP_171871 AAD10693 ATP binding/ATP-dependent helicase/DNA binding /helicase/nucleic acid binding /Hypothetical protein

10.000