NAC1(AT1G56010): global tree
File name: NAC1.mltree0.svg
144 aa residues 81 OTU included.
Results of 0 partial trees included
Assemble Job Id: niX5dN
Tree job id: 0bJkNk
BP values out of 100
Analyzed by t_aoyama@nibb.ac.jp
Date: Wed Jul 05 17:16:15 JST 2006
ABA91268 Oryza sativa (japonica cultivar-group) hypothetical protein LOC_Os11g03310
ABA91280 Oryza sativa (japonica cultivar-group) No apical meristem protein, putative
Q5CD17 Oryza sativa ABA95706 Oryza sativa (japonica cultivar-group) NAC77_ORYSA NAC domain-containing protein 77 (ONAC077) (ONAC300)/hypothetical protein LOC_Os12g03050
BAD91001 Oryza sativa (japonica cultivar-group) ONAC300
NP_912420 Oryza sativa (japonica cultivar-group) AAN64996 Putative NAM (no apical meristem) protein/Putative NAM (no apical meristem) protein
CAA09372 Triticum sp. GRAB2 protein
XP_506578 Oryza sativa (japonica cultivar-group) XP_479577 BAC83810 PREDICTED OSJNBa0060O17.21 gene product/putative GRAB2 protein(Geminivirus Rep A-binding) /putative GRAB2 protein(Geminivirus Rep A-binding)
CAH56054 Zea mays hypothetical protein
AAP86221 Zea mays NAM-related protein 1
NP_914157 Oryza sativa (japonica cultivar-group) BAD68974 OsNAC4-like protein/putative OsNAC2
NP_908359 Oryza sativa (japonica cultivar-group) BAB16335 putative NAM protein/putative NAM protein
AT3G04060 = NP_187056 AAF05864 AAK32791 AAL87404 AAM61417 ANAC046; transcription factor/NAM-like protein (no apical meristem) /AT3g04060/T11I18_17 /AT3g04060/T11I18_17 /NAM-like protein (no apical meristem)
AT5G18270 = NP_974800 ANAC087
AT5G18270 = NP_197328 BAB09485 AAO64133 AAP04055 ANAC087; transcription factor/NAM (no apical meristem)-like protein /putative NAM (no apical meristem) protein /putative NAM (no apical meristem) protein
XP_472753 Oryza sativa (japonica cultivar-group) BAC53811 Oryza sativa CAD40985 Oryza sativa (japonica cultivar-group) OSJNBa0072F16.10/OsNAC2 protein /OSJNBa0072F16.10
BAD29568 Oryza sativa (japonica cultivar-group) putative OsNAC1 protein
BAC53810 Oryza sativa OsNAC1 protein
AT5G39610 = NP_198777 BAB08893 AAL07176 AAM14130 AAM61198 AAO41710 ANAC092; transcription factor/unnamed protein product /putative NAM /CUC2 protein /putative NAM/CUC2 protein /NAM /CUC2-like protein /no apical meristem-like protein
AT3G29035 = NP_189546 AAL32716 AAP42729 ANAC059; transcription factor/Unknown protein /At3g29035
AT5G61430 = NP_200951 BAB08499 AAK96835 AAM10058 AAM61656 ANAC100; transcription factor/NAM (no apical meristem)-like protein /NAM (no apical meristem)-like protein /NAM (no apical meristem)-like protein /NAM, no apical meristem,-like protein
AT5G07680 = NP_568182 NP_850789 BAB11446 AAM65237 AAP21227 ANAC079/ANAC080; transcription factor /ANAC079/ANAC080; transcription factor /NAM (no apical meristem)-like protein /NAM (no apical meristem)-like protein /At5g07680
AAM50521 Petunia x hybrida nam-like protein 18
AAN03466 Glycine max AAX85982 AAY46125 no apical meristem-like protein/NAC5 protein /NAC domain protein NAC5
AT5G53950 = NP_200206 BAA19529 BAB10725 CUC2 (CUP-SHAPED COTYLEDON 2); transcription factor/CUC2 /CUC2
AT1G76420 = NP_177768 AAF16659 AAG51953 AAP82630 CUC3 (CUP SHAPED COTYLEDON3); transcription factor/unknown protein; 31626-33432 /unknown protein; 10137-8331 /cup-shaped cotyledon 3
CAH56059 Zea mays hypothetical protein
XP_483299 Oryza sativa (japonica cultivar-group) BAC57407 putative cup-shaped cotyledon/putative cup-shaped cotyledon
BAD61787 Oryza sativa (japonica cultivar-group) putative NAM
CAH56058 Zea mays hypothetical protein
CAH56057 Zea mays hypothetical protein
AT3G15170 = NP_188135 BAB02571 BAB20598 CUC1 (CUP-SHAPED COTYLEDON1); transcription factor/unnamed protein product /CUC1
CAA63101 Petunia x hybrida CAA63102 NAM/NAM
AT2G24430 = NP_850054 NP_180019 AAD18114 transcription factor/transcription factor /NAM (no apical meristem)-like protein
XP_463226 Oryza sativa (japonica cultivar-group) AAR89042 putative OsNAC1 protein/putative OsNAC1 protein
AT3G18400 = NP_188469 BAB01106 ANAC058; transcription factor/unnamed protein product
Smo_NAC1_7_1 Smo_NAC1_18_1
Smo_NAC1_27_1
Smo_NAC1_15_1 Smo_NAC1_25_1
Ppa_NAC1_15_1
Ppa_NAC1_28_1
Ppa_NAC1_2_1
Smo_NAC1_4_1
AT2G46770 = NP_182200 AAC33506 AAO22745 AAV97804 ANAC043/EMB2301; transcription factor /NAM (no apical meristem)-like protein /putative NAM (no apical meristem) protein /At2g46770
AT3G61910 = NP_191750 CAB71898 ANAC066; transcription factor/NAM-like protein
AT1G71930 = NP_177338 AAG52219 AAM62651 ANAC030; transcription factor/NAM-like protein; 48543-50167 /NAM-like protein
XP_473322 Oryza sativa (japonica cultivar-group) CAE04781 OSJNBb0020O11.1/OSJNBb0020O11.1
XP_467007 Oryza sativa (japonica cultivar-group) BAD25783 putative OsNAC7 protein/putative OsNAC7 protein
AT2G18060 = NP_179397 AAD20120 ANAC037; transcription factor/putative NAM (no apical meristem)-like protein
NP_912453 Oryza sativa (japonica cultivar-group) AAO15294 Hypothetical protein/Hypothetical protein
AT1G54330 = NP_175835 AAD25613 AAQ62866 BAD44041 ANAC020; transcription factor/Unknown protein /At1g54330 /hypothetical protein
Smo_NAC1_14_1 Smo_NAC1_34_1
Ppa_NAC1_17_1
Ppa_NAC1_35_1
Smo_NAC1_29_1 Smo_NAC1_32_1
Smo_NAC1_12_1
Smo_NAC1_20_1
XP_475238 Oryza sativa (japonica cultivar-group) AAT44250 putative no apical meristem (NAM) protein/putative no apical meristem (NAM) protein
CAA09371 Triticum sp. GRAB1 protein
NP_911548 Oryza sativa (japonica cultivar-group) Q7EZT1 Oryza sativa BAA89797 BAC10231 Oryza sativa (japonica cultivar-group) BAD31538 OsNAC3 protein/NAC67_ORYSA NAC domain-containing protein 67 (ONAC067) /OsNAC3 protein /OsNAC3 protein /OsNAC3 p
ABB72842 Elaeis guineensis ABB72844 NAC protein 1/NAC protein 1 splice variant 2
XP_470088 Oryza sativa (japonica cultivar-group) AAR89838 AAT02360 ABD52007 putative NAC-domain protein/putative NAC-domain protein /NAC transcription factor /stress-induced transcription factor NAC1
NP_912551 Oryza sativa (japonica cultivar-group) AAN62790 Putative NAM (no apical meristem) protein/Putative NAM (no apical meristem) protein
XP_468336 Oryza sativa (japonica cultivar-group) XP_507036 BAD21589 BAD22026 putative NAC domain protein NAC1/PREDICTED OJ1116_E04.11 gene product /putative NAC domain protein NAC1 /putative NAC domain protein NAC1
XP_473174 Oryza sativa (japonica cultivar-group) CAE02350 CAD41119 OSJNBb0070J16.15/OSJNBb0072M01.11 /OSJNBb0070J16.15
NP_921992 Oryza sativa (japonica cultivar-group) AAK13151 AAP54279 putative NAM (no apical meristem) protein/putative NAM (no apical meristem) protein /NAM protein, putative
AAM65014 Arabidopsis thaliana NAM/CUC2-like protein
AT4G28530 = NP_567811 AAO00822 AAP42754 ANAC074; transcription factor/NAM /CUC2 -like protein /At4g28530
ABA99367 Oryza sativa (japonica cultivar-group) salicylic acid-induced protein 19
CAB51838 Oryza sativa (indica cultivar-group) I1332.9
XP_473911 Oryza sativa (japonica cultivar-group) CAD41743 OSJNBa0058K23.9/OSJNBa0058K23.9
XP_480565 Oryza sativa (japonica cultivar-group) BAD03222 BAD03589 putative NAC domain protein NAC1/putative NAC domain protein NAC1 /putative NAC domain protein NAC1
XP_464228 Oryza sativa (japonica cultivar-group) BAD26221 BAD25552 putative NAC domain protein NAC1/putative NAC domain protein NAC1 /putative NAC domain protein NAC1
BAD45909 Oryza sativa (japonica cultivar-group) putative NAC domain protein NAC1
AAK84883 Phaseolus vulgaris NAC domain protein NAC1
AAQ75123 Capsicum annuum salicylic acid-induced protein 19
BAB02506 Arabidopsis thaliana NAM (no apical meristem) protein-like
AT1G56010 = NP_175997 Q84TE6 AAF21437 AAF79328 AAM63206 NAC1; transcription factor/NAC22_ARATH NAC domain-containing protein 21/22 (ANAC021) (ANAC022)/NAC1 /F14J16.32 /NAC1
Ppa_NAC1_8_1
Ppa_NAC1_29_1
Ppa_NAC1_13_1
Ppa_NAC1_12_1
10.000