AT3G15170 = NP_188135 BAB02571 BAB20598 CUC1 (CUP-SHAPED COTYLEDON1); transcription factor/unnamed protein product /CUC1
CAA63101 *Petunia x hybrida* CAA63102 NAM/NAM
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
XP_483299 *Oryza sativa* (japonica cultivar-group) BAC57407 putative cup-shaped cotyledon/putative cup-shaped cotyledon
CAH56059 *Zea mays* hypothetical protein
CAH56057 *Zea mays* hypothetical protein
BAD61787 *Oryza sativa* (japonica cultivar-group) putative NAM
AAP86221 *Zea mays* NAM-related protein 1
CAH56054 *Zea mays* hypothetical protein
NP_914157 *Oryza sativa* (japonica cultivar-group) BAD68974 OsNAC4-like protein/putative OsNAC2   Os01g0393100
NP_908359 *Oryza sativa* (japonica cultivar-group) BAB16335 putative NAM protein/putative NAM protein   Os01g0104500
ABA91268 *Oryza sativa* (japonica cultivar-group) hypothetical protein LOC_Os11g03310
ABA91280 *Oryza sativa* (japonica cultivar-group) No apical meristem protein, putative   LOC_Os11g03370
BAD91001 *Oryza sativa* (japonica cultivar-group) ONAC300   Os12g0123800
Q5CD17 *Oryza sativa* ABA95706 *Oryza sativa* (japonica cultivar-group) NAC77_ORYSA NAC domain-containing protein 77 (ONAC077) (ONAC300)/hypothetical protein LOC_Os12g03050
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)
AAM50520 *Petunia x hybrida* nam-like protein 17
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_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
AT5G18270 = NP_974800 ANAC087
BAC53810 *Oryza sativa* BAD29568 *Oryza sativa* (japonica cultivar-group) OsNAC1 protein/putative OsNAC1 protein
XP_472753 *Oryza sativa* (japonica cultivar-group) BAC53811 *Oryza sativa* CAD40985 *Oryza sativa* (japonica cultivar-group) OSJNBa0072F16.10/OsNAC2 protein /OSJNBa0072F16.10
AAM50521 *Petunia x hybrida* nam-like protein 18
AT3G29035 = NP_189546 AAL32716 AAP42729 ANAC059; transcription factor/Unknown protein /At3g29035
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
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
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
AAN03466 *Glycine max* AAX85982 AAY46125 no apical meristem-like protein/NAC5 protein /NAC domain protein NAC5
AT3G18400 = NP_188469 BAB01106 ANAC058; transcription factor/unnamed protein product
XP_463226 *Oryza sativa* (japonica cultivar-group) AAR89042 putative OsNAC1 protein/putative OsNAC1 protein
AT2G24430 = NP_850054 NP_180019 AAD18114 transcription factor/transcription factor /NAM (no apical meristem)-like protein

Smo_CUC1_35_1
Smo_CUC1_16_1
Smo_CUC1_26_1
Ppa_CUC1_13_1   Ppa_CUC1_1_1
Ppa_CUC1_12_1
Ppa_CUC1_39_1
Ppa_CUC1_3_1
Ppa_CUC1_10_1
Ppa_CUC1_19_1
Ppa_CUC1_9_1
Ppa_CUC1_2_1
Smo_CUC1_4_1
Smo_CUC1_14_1

Contig1 is a short clone of Contig13 with the vector sequence

ABA99367 *Oryza sativa* (japonica cultivar-group) salicylic acid-induced protein 19
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
AAQ75123 *Capsicum annuum* salicylic acid-induced protein 19
AT1G56010 = NP_175997 Q84TE6 AAF21437 AAF79328 AAM63206 NAC1; transcription factor/NAC22_ARATH NAC domain-containing protein 21/22 (ANAC021) (ANAC022)/NAC1 /F14J16.32 /NAC1
BAB02506 *Arabidopsis thaliana* NAM (no apical meristem) protein-like
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
NP_921992 *Oryza sativa* (japonica cultivar-group) AAK13151 AAP54279 putative NAM (no apical meristem) protein/putative NAM (no apical meristem) protein /NAM protein, putative
XP_473174 *Oryza sativa* (japonica cultivar-group) CAE02350 CAD41119 OSJNBb0070J16.15/OSJNBb0072M01.11 /OSJNBb0070J16.15
AT1G69490 = NP_564966 O49255 CAA10955 AAG60108 AAK25911 AAG43692 AAM63330 NAP (NAC-LIKE, ACTIVATED BY AP3/PI); transcription factor /NAC29_ARATH NAC domain-containing protein 29 (ANAC029) (NAC2) (NAC-LIKE, ACTIVATED BY AP3/PI protein) (NAP)/unnamed
AAU90314 *Solanum demissum* putative NAC domain protein NAC2
AAX85978 *Glycine max* AAY46121 NAC1 protein/NAC domain protein NAC1
AAK84884 *Phaseolus vulgaris* NAC domain protein NAC2
AAF05865 *Arabidopsis thaliana* NAM-like protein (no apical meristem)
Smo_CUC1_24_1
AT4G27410 = NP_567773 AAL00817 AAL16305 AAM14367 AAM65308 AAN60296 RD26; transcription factor/unknown protein /AT4g27410/F27G19_10 /unknown protein /unknown /unknown
AAX85980 *Glycine max* AAY46123 NAC3 protein/NAC domain protein NAC3
Smo_CUC1_10_1   Smo_CUC1_1_1
Ppa_CUC1_35_1
Ppa_CUC1_14_1
Ppa_CUC1_11_1
Ppa_CUC1_28_1
AT5G46590 = NP_199471 BAA97530 AAV85660 AAW70401 ANAC096; transcription factor/NAM-like /At5g46590 /At5g46590
AT1G54330 = NP_175835 AAD25613 AAQ62866 BAD44041 ANAC020; transcription factor/Unknown protein /At1g54330 /hypothetical protein
AT1G33280 = NP_174598 AAG51291 ANAC015; transcription factor/unknown protein
AT4G10350 = NP_192773 CAB39788 CAB78158 AAM91615 ANAC070; transcription factor/NAM/NAP like protein /NAM/NAP like protein /putative NAM/NAP
Smo_CUC1_2_1 Smo_CUC1_28_1
Smo_CUC1_8_1
Smo_CUC1_5_1 Smo_CUC1_12_1
AAM60909 *Arabidopsis thaliana* NAM-like protein
AAQ06260 *Sorghum bicolor* putative NAM (no apical meristem) protein
XP_493710 *Oryza sativa* (japonica cultivar-group) BAA84803 BAB19365 AAO33144 putative OsNAC7 protein/putative OsNAC7 protein /putative OsNAC7 protein /putative NAM (no apical meristem) protein

0.100