TCP3(At1g53230): global tree
File name:TCP3.mltree0.svg
72 aa residues 60 OTU included.
Results of 0 partial trees included
Assemble Job Id: c5lmH4
Tree job id: N8WhbV
BP values out of 100
Analyzed by t_aoyama@nibb.ac.jp
Date: Wed May 03 00:06:42 JST 2006

CAJ44128 *Misopates orontium* cycloidea protein
AAF07225 *Antirrhinum graniticum* AAP46512 *Antirrhinum ovatum* AAP46513 *Antirrhinum subcordatum* cyc1A protein/CYCLOIDEA-I
AAP46508 *Antirrhinum cornutum* AAP46509 *Antirrhinum coulterianum* AAP46510 *Antirrhinum multiflorum* AAP46511 *Antirrhinum*
AAP46501 *Antirrhinum multiflorum* CYCLOIDEA-like protein
AAP03348 *Chaenorhinum villosum* CYCLOIDEA-like protein
AAP84102 *Chaenorhinum origanifolium* CYCLOIDEA-like 1
AAP03349 *Nuttallanthus canadensis* CYCLOIDEA-like protein
AAD45359 *Linaria vulgaris* AAP03350 cycloidea-like protein/CYCLOIDEA-like protein
AAP03342 *Kickxia spuria* CYCLOIDEA-like protein
AAP03344 *Maurandya antirrhiniflora* CYCLOIDEA-like protein
AAP03345 *Asarina procumbens* CYCLOIDEA-like protein
AAP03343 *Lophospermum* sp. *91/97* CYCLOIDEA-like protein
AAP03338 *Chaenorhinum villosum* DICHOTOMA-like protein
AAP03340 *Lophospermum* sp. *91/97* DICHOTOMA-like protein
AAP03339 *Asarina procumbens* DICHOTOMA-like protein
AAP46528 *Antirrhinum coulterianum* DICHOTOMA-like protein
AAP46534 *Mohavea confertiflora* DICHOTOMA-like protein
AAP46526 *Mohavea confertiflora* AAP46529 *Antirrhinum multiflorum* AAP46531 *Antirrhinum ovatum* AAP46532 *Antirrhinum subco*
AAP46527 *Antirrhinum cornutum* DICHOTOMA-like protein
AAF07265 *Antirrhinum majus subsp. cirrhigerum* AAF07266 AAF07267 *Digitalis purpurea* AAF07268 *Antirrhinum graniticum* AAF0
AAP46530 *Antirrhinum nuttallianum* AAP84105 DICHOTOMA-like protein/DICHOTOMA-like
AAP46524 *Antirrhinum virga* DICHOTOMA-like protein
AAP46517 *Antirrhinum cornutum* DICHOTOMA-like protein
AAP46523 *Antirrhinum subcordatum* DICHOTOMA-like protein
AAP46525 *Antirrhinum vexillocalyculatum* DICHOTOMA-like protein
AAP84104 *Misopates orontium* DICHOTOMA-like
AAP03341 *Misopates orontium* DICHOTOMA-like protein
AAP46522 *Antirrhinum ovatum* DICHOTOMA-like protein
AAF07263 *Misopates orontium* AAF07264 AAP46518 *Antirrhinum coulterianum* AAP46519 *Antirrhinum leptaleum* AAP46520 *Antirrh*
AAP03346 *Cymbalaria muralis* CYCLOIDEA-like protein
AAP03337 *Linaria vulgaris* DICHOTOMA-like protein
AAG45501 *Populus trichocarpa* teosinte-branched-like protein
AAK60248 *Zea mays* AAL17096 AAL17099 AAL17100 AAL17131 AAL17138 AAS21486*Zea mays subsp. parviglumis* AAS21501 teosinte br
AAK37493 *Arundinella hirta* teosinte branched1 protein
AT2G31070 = NP_565712 AAC63845 AAL24427 AAN65081 transcription factor/expressed protein /Unknown protein /Unknown protein
NP_916166 *Oryza sativa* (japonica cultivar-group) BAB92954 P0503C12.19/transcription factor PCF7
AAO43102 *Antirrhinum majus* mutant cincinnata
AAG43412 *Lycopersicon esculentum* cycloidea
NP_913318 *Oryza sativa* (japonica cultivar-group) BAB92952 BAD73155 putative TCP3/transcription factor PCF5 /transcription
CAD19990 *Lupinus albus* TCP1 protein
AT3G15030 = NP_850589 NP_188121 BAA97066 AAM19816 AAN31111 TCP4; transcription factor/TCP4; transcription factor /unnamed
AT1G53230 = NP_564624 AAC24010 AAF69550 AAM20529 AAN15585 TCP3; transcription factor/TCP3 /F12M16.13 /putative flower dev
NP_916237 *Oryza sativa* (japonica cultivar-group) BAB92377 P0512C01.32/putative transcription factor
XP_475704 *Oryza sativa* (japonica cultivar-group) AAV59297 putative TCP transcription factor/putative TCP transcription fac
AT5G08070 = NP_196424 CAB93708 transcription factor/putative helix-loop-helix DNA binding protein
AT5G60970 = NP_200905 BAB10646 transcription factor/unnamed protein product
AT3G21510 = AT5G39340 = AT3G29350 = AT3G02150 = NP_850501 NP_566164 AAF02122 AAF14823 BAA88760 AAM62743 AAN71993 AAP68278
BAC01119 *Oryza sativa* (japonica cultivar-group) transcription factor PCF8
XP_476577 *Oryza sativa* (japonica cultivar-group) BAC83480 transcription factor PCF8-like/transcription factor PCF8-like
ABA99859 *Oryza sativa* (japonica cultivar-group) transposon protein, putative, unclassified
CAD91129 *Solanum tuberosum* putative transcription factor
AT1G30210 = NP_564351 NP_849730 AAG50562 AAK59513 AAL15303 AAM44946 transcription factor/transcription factor /hypothetic
AT4G18390 = NP_567553 CAA16719 AAC26786 CAB78841 AAK56259 AAK56280 AAM26708 transcription factor/teosinte branched1 - lik
XP_476664 *Oryza sativa* (japonica cultivar-group) BAC79520 BAD31918 putative transcription factor PCF6/putative transcripti
BAB92953 *Oryza sativa* (japonica cultivar-group) transcription factor PCF6
Smo_TCP3_4_1 Smo_TCP3_6_1
Smo_TCP3_1_1 Smo_TCP3_3_1 Smo_TCP31167272820_1
Smo_TCP3_2_1 Smo_TCP3_5_1
Ppa_TCP3_1_1
Ppa_TCP3_2_1

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