

ANT (AT4G37750): global tree
File name:ANT.njtree0.svg
122 aa residues 80 OTU included.
Results of 19 partial trees included.
Assemble Job Id: MZu87A selectNalign jobid: CD3gUz
Tree job id: YKALcX
BP values out of 100
Analyzed by nonodera@nibb.ac.jp
Date: Thu Aug 31 17:59:22 JST 2006

Short fragmental sequences
Smo_At4g37750_6_1(Pt11)
Smo_At4g37750_4_1(Pt14)
Smo_At4g37750_42_1(Pt14)
AAG52255(Pt14)



BAD33427 *Oryza sativa* (japonica cultivar-group) WRINKLED1-like protein(Pt10)
(Nipponbare)chromosome9

Smo_At4g37750_2_1(Pt1)

Smo_At4g37750_73_1(Pt18) **Smo_At4g37750_63_1(Pt15)**

Smo_At4g37750_3_1 **Smo_At4g37750_1_1(Pt2)**

(cultivar=?)chromosome8

AAW78374 *Oryza sativa* (japonica cultivar-group) **AAW95492** ABA92780 transcription factor AP2D17/AP2/EREBP transcription factor BABY BOOM1 /ovule development aintegumenta-like protein BNM3, putative(Pt3) (cultivar=?)chromosome11
Nipponbare chromosome11

AAW78368 *Oryza sativa* (japonica cultivar-group) transcription factor AP2D8 (Dongjin)chromosome2

AT2G41710 = NP_565957 NP_850355 AAC02777 AAM65663 BAC43380 DNA binding/transcription factor /DNA binding /transcription factor /putative AP2 domain transcription factor /putative AP2 domain transcription factor /putative AP2 domain transcription fac (pt19) Possibly 1 domain possessing

Smo_At4g37750_31_1 **Smo_At4g37750_5_1** 98% similar to Smo_At4g37750_31_1(High quality mismatch=7)
(Nipponbare)chromosome6

AAW78373 (Pt6) (cultivar=?)chromosome1

10382_Cre 5992_Cre(Pt13) Possibly 1 domain possessing