

YP735A1.(At5g38450): global tree
File name: CYP735A1.njtree0.svg
269 aa residues 111 OTU included.
Results of 2 partial trees included.
Assemble Job Id: 7KRKNU, SelectN Assemble JobId: Bln2IL(140)
Tree job id: vUvHKv
BP values out of 100
Analyzed by ku-bo@nibb.ac.jp
Date: Thu Jan 18 16:05:53 JST 2007

NP_173149.1; CYP72C1; heme binding / iron ion binding / monooxygenase/oxygen binding [Arabidopsis thaliana] (identical o AT1G17060)
AAA31430.1; cytochrome P450 3c protein
CAB77247.2; putative cytochrome P-450 [Persea americana]
AAL66766.1; cytochrome P450 monooxygenase CYP72A5 [Zea mays]
AAF64303.1; putative cytochrome P450 [Lycopersicon esculentum]
BAB86912.1; putative cytochrome P450 [Solanum tuberosum]
AAB05376.3; putative cytochrome P-450 [Nicotiana glauca]
NP_001015810.1; MGC97602 protein [Xenopus tropicalis]
XP_541984.2; PREDICTED: similar to cytochrome P450, family 2, subfamily E, polypeptide 2 homolog [Canis familiaris]
BAD02914.1; Cytochrome P450 [Xenopus laevis]
AAL42573.1; cytochrome P450 [Agrobacterium tumefaciens str. C58]
O70537;CP3AV MESAU Cytochrome P450 3A31 (CYP11A31) (P450 SH3A-1)
NP_796281.1; cytochrome P450, family 2, subfamily E, polypeptide 2 homolog [Mus musculus]

were removed from this tree.

