

COMT(At1g21100): partial tree
File name: COMT.pt3.svg
107 aa residues 139 OTU included.
Results of Partial tree including ABA96096.1| O-methyltransferase, putative [Oryza sativa (japonica cultivar-group)].
Assemble Job Id: eRqsET selectNalign job id: 0V6KQY(200)
Tree job id: kw9lZt
BP values out of 100

AT1G21100 = NP_173534 AAF80651 AAK06867 AAL09769 AAN28913 O-methyltransferase/Contains similarity to O-Methyltransferase 1 from Arabidopsis thaliana gb|U70424. It is a member of O-methyltransferase family. ESTs gb|AI994826, gb|N65066 and gb|N38589
AT1G21110 = NP_173535 AAF80650 BAC43382 AAO63966 O-methyltransferase/Contains similarity to O-Methyltransferase 1 from Arabidopsis thaliana gb|U70424. It is a member of O-methyltransferase family. ESTs gb|AI993288 and gb|Z18076 come from this gene
AT1G21120 = NP_173536 AAF80649 AAK06866 BAD94958 O-methyltransferase/Contains similarity to O-Methyltransferase 1 from Arabidopsis thaliana gb|U70424. It is a member of O-methyltransferase family. ESTs gb|AI993288 and gb|Z18076 come from this gene
AT1G21130 = NP_173537 NP_849693 AAF80648 O-methyltransferase/O-methyltransferase /Contains similarity to caffeic acid 3-O-Methyltransferase from Saccharum officinarum gb|AJ231133. It is a member of O-methyltransferase family. ESTs gb|AI994592 and AAM67269 Arabidopsis thaliana O-methyltransferase, putative

