

NP_849693.1| O-methyltransferase [Arabidopsis thaliana] (identical to At1g21130)
NP_176502.0| O-methyltransferase/S-adenosylmethionine-dependent methyltransferase [Arabidopsis thaliana] (identical to At1g63140)
NP_198533.1| O-methyltransferase [Arabidopsis thaliana] (identical to At5g37170)
NP_176478.1| O-methyltransferase/S-adenosylmethionine-dependent methyltransferase [Arabidopsis thaliana] (identical to At1g62900)
NP_195241.1| O-methyltransferase/S-adenosylmethionine-dependent methyltransferase [Arabidopsis thaliana] (identical to At4g35150)
BAB11374.1| caffeic acid O-methyltransferase-like protein [Arabidopsis thaliana] (identical to At5g37170)
CA178904.1| putative ortho methyl transferase [Nicotiana tabacum]
AAA87043.1| O-methyltransferase [Hordeum vulgare]
AAQ07451.1| caffeic acid O-methyltransferase [Triticum aestivum]
AAD50439.1| caffeic acid O-methyltransferase [Eucalyptus globulus]
AAD50440.1| caffeic acid O-methyltransferase [Eucalyptus globulus]
CAE51883.1| putative caffeate O-methyltransferase [Lolium multiflorum]
Results of 3 partial trees included.
Assemble Job Id: eRqsET, SelectNalign Job Id: oV6KQY (200)
BP values out of 100
Analyzed by tomoakin@kenroku.kanazawa-u.ac.jp Date: Tue Apr 24 15:09:39 JST 2007

COMT (At1g21100): global tree
File name: COMT.njtree2.svg
248 as residues 147 OTU included.
Tree job id: joQUJ
Assemble Job Id: eRqsET, SelectNalign Job Id: oV6KQY (200)
BP values out of 100

BAD14923 *Oryza sativa* (pt1) partial cds from /cultivar="Fujiminori" originally by M. Kubo

gi|21617938|gb|AAM66988.1| putative catechol O-methyltransferase [Arabidopsis thaliana]
gi|13399464|pdb|1FPQ|A Chain A, Crystal Structure Analysis Of Selenomethionine Substituted Chalcone O-Methyltransferase were removed from this tree.