SUR(JX74031000): global tree
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Number of sequences:1 (1 selected)
Assembly job id: 4Xu4C2_oeecfhaJgn Job id: g2X3ZN
Tree job id: aZkeF
BP values cut at: 100
Analyzed by: j.aoyama@nibb.ac.jp
Date: Tue Jul 25 01:07:45 JST 2006

AAF05621 Papaver somniferum (S)-N-methylcoclaurine 3'-hydroxylase
AAF14032 Papaver somniferum (S)-N-methylcoclaurine 3'-hydroxylase
AAF14033 Papaver somniferum (S)-N-methylcoclaurine 3'-hydroxylase
BAC40982 Arabidopsis thaliana putative cytochrome P450
BAC53991 Petunia x hybrida cytochrome P450
BAC53992 Petunia x hybrida cytochrome P450
P37132 Solanum melongena CAA50648 C19A2, SOLME Cytochrome P450 76A2 (CYPLXXVIA2) (P-450E03)/P450 hydroxylase
P37131 Solanum melongena CAA50649 C19A1, SOLME Cytochrome P450 76A1 (CYPLXXVIA1) (P-450E02)/cytochrome P450
NP_505008 Oryza sativa (japonica cultivar-group) AAK74370 AAN18475 putative cytochrome P450/Putative cytochrome P450/ Putative cytochrome P450
BAE71262 Triticum aestivum putative cytochrome P450
CAB54741 Cicer arietinum cytochrome P450 monooxygenase
BAE72857 Verbena x hybrida cytochrome P450
AB89365 Vitis vinifera putative flowering-related P-450 enzyme
AAQ56025 Paulownia sativa cytochrome P450
CAD37616 Hordeum lechleri Bx9 cytochrome P-450 monooxygenase
CAA71054 C79B1 Helianthus tuberosus CAA71054 C79B1 ; HELTU Cytochrome P450 76B1 (7-ethoxycoumarin O-deethylase) (ECOD) (Phenylurea dealkylase)/7-ethoxycoumarin O-deethylase
BAC53993 Petunia x hybrida cytochrome P450
CAC93687 Catharanthus roseus geraniol 10-hydroxylase
AT3G61040 = NP_191083 CAB40140 CYP79C7; hence bindingfrom ion binding monooxygenase/oxygen binding lysotochrome P450 monooxygenase-like protein
AT5G05690 = NP_196262 CAB40139 AAC06139 CYP79C6; hence bindingfrom ion binding monooxygenase/oxygen binding C79C2, ARATH Cytochrome P450 76C2(putative cytochrome P450 lysotative cytochrome P450
AAP87087 Arabidopsis thaliana Putative cytochrome P450
AT5G05700 = NP_196263 CAB40138 AAC06138 CYP79C5; hence bindingfrom ion binding monooxygenase/oxygen binding Putative cytochrome P450
AT5G05690 = NP_196262 CAB40139 AAC06139 CYP79C6; hence bindingfrom ion binding monooxygenase/oxygen binding C79C2, ARATH Cytochrome P450 76C2(putative cytochrome P450 lysotative cytochrome P450
AT5G42580 = NP_196981 O64637 AAC08152 AAL32079 Putative cytochrome P450/Putative cytochrome P450/ Putative cytochrome P450
AT5G42590 = NP_196982 O64638 AAC08153 AAL32080 AAL85445 AAN79565 CYP79C1; hence bindingfrom ion binding monooxygenase/oxygen binding C79C1, ARATH Cytochrome P450 79C1(T52158 cytochrome P450 monooxygenase (Arg/Gln55665F1TK2.9 (Arg/Gln5560
NP_501091 Oryza sativa (japonica cultivar-group) putative flavonoid 3',5'-hydroxylase
AAN77484 Sorghum bicolor putative cytochrome P450
AAL27058 Arabidopsis thaliana putative cytochrome P450
AT4G22710 = NP_194002 NP_567665 CAA16554 CAB10556 CAB70326 BAB42538 BAD43108 CYP75B4(2); hence bindingfrom ion binding monooxygenase/oxygen binding CYP75B4(1), hence binding from ion binding monooxygenase/oxygen binding lysotochrome P450
AT5G42400 = NP_196775 BAB10113 AAL14103 AAL91602 CYP75A(1); hence bindingfrom ion binding monooxygenase/oxygen binding Flavonoid 3', 5'-hydroxylase-like; putative cytochrome P450 ,ST/Rg(Gln/Arg)4H2DKTC23, 8 (Arg/Gln4H2DKTC23)
AT5G13210 = NP_196968 CAB40976 CAB70776 CYP75A(4); hence bindingfrom ion binding monooxygenase/oxygen binding flavonoid 3', 5'-hydroxylase-like protein flavonoid 3', 5'-hydroxylase-like protein (Arg/Gln12310
AT5G13220 = NP_196969 CAB40975 CAB70775 AAD55534 CYP75A(5); hence bindingfrom ion binding monooxygenase/oxygen binding hydroxylase P450 homolog (Arg/Gln12320
AT5G13230 = NP_196967 CAB40977 CAB70773 BAC43375 AAO63874 CYP75A(6); hence bindingfrom ion binding monooxygenase/oxygen binding flavonoid 3', 5'-hydroxylase-like protein flavonoid 3', 5'-hydroxylase-like protein flavonoid 3', 5'-hydroxylase
AAZ79541 Phalaenopsis hybrid cultivar flavonoid 3',5'-hydroxylase
O04773 Campanula medium BAA03542 C75A5_CAMME Flavonoid 3',5'-hydroxylase (F3'5'H) (Cytochrome P450 75A5)/flavonoid 3',5'-hydroxylase
AAX11796 Delphinium grandiflorum flavonoid 3',5'-hydroxylase
BAE72857 Verbena x hybrida flavonoid 3',5'-hydroxylase
AF349974 Verbena x hybrida AAO43493 BAE72857 CYP75A(3); hence bindingfrom ion binding
P52418 Petunia x hybrida BAA03542 C75A1_PETHY Flavonoid 3',5'-hydroxylase (F3'5'H) (Cytochrome P450 75A1) (CYPLXXVIA1) (P450E01)/flavonoid 3',5'-hydroxylase
CAD14004 Petunia x hybrida P52418 CAA68230 C75A1_PETHY Flavonoid 3',5'-hydroxylase (F3'5'H) (Cytochrome P450 75A1) (CYPLXXVIA1) (P450E01)/flavonoid
AAO49000 Lycianthes rantonnei flavonoid 3',5'-hydroxylase
AAV85472 Solanum tuberosum flavonoid 3',5'-hydroxylase
AAV85470 Solanum tuberosum AAV85471 flavonoid 3',5'-hydroxylase/flavonoid 3',5'-hydroxylase
P52419 Solanum melongena BAA03543 C75A2_SOLME Flavonoid 3',5'-hydroxylase (F3'5'H) (Cytochrome P450 75A2) (CYPLXXVIA2) (P-450E01)/flavonoid hydroxylase (P450)
AC36640 Vitis vinifera flavonoid 3',5'-hydroxylase
CAB42777 Vitis vinifera ABC70066 flavonoid 3',5'-hydroxylase/flavonoid 3',5'-hydroxylase
CAD17607 Vitis vinifera ABC36641 flavonoid 3',5'-hydroxylase/flavonoid 3',5'-hydroxylase
AAP31058 Gossypium hirsutum flavonoid 3,5'-hydroxylase
AB04503 Camellia sinensis flavonoid 3',5'-hydroxylase
BAB78255 Catharanthus roseus flavonoid 3',5'-hydroxylase
AB81561 Vinca major Flavonoid 3',5'-hydroxylase
AB81562 Vinca major Flavonoid 3',5'-hydroxylase
BAB78255 Gerbera hybrida flavonoid 3',5'-hydroxylase
AAM51564 Glycine max flavonoid 3',5'-hydroxylase
O04790 Eustoma grandiflorum BAA03539 BAD04480 C75A7_EUSGR Flavonoid 3',5'-hydroxylase (F3'5'H) (Cytochrome P450 75A7)/flavonoid 3',5'-hydroxylase/ flavonoid 3',5'-hydroxylase
O99416 Eustoma grandiflorum AAB71562 C75A3_EUSGR Flavonoid 3',5'-hydroxylase (F3'5'H) (Cytochrome P450 75A3)/flavonoid 3',5'-hydroxylase
AAB71561 Eustoma grandiflorum AAB71563 C75A4_EUSGR Flavonoid 3',5'-hydroxylase (F3'5'H) (Cytochrome P450 75A4)/flavonoid 3',5'-hydroxylase
AAD56671 Gentiana sedifolia flavonoid 3',5'-hydroxylase
NP_505099 Oryza sativa (japonica cultivar-group) AAM74364 putative cytochrome 450/Putative cytochrome 450
NP_505027 Oryza sativa (japonica cultivar-group) AAM55948 AAN05137 AAP52014 putative flavonoid 3',5'-hydroxylase/Putative flavonoid 3' hydroxylase /Cytochrome P450
AAY71494 Sorghum bicolor AAY74195 flavonoid 3,5'-hydroxylase/flavonoid 3'-hydroxylase
AAG45029 Catharanthus chinensis flavonoid 3,5'-hydroxylase
ABB43051 Osteospermum hybrid cultivar flavonoid 3'5'-hydroxylase
ABB43052 Osteospermum hybrid cultivar flavonoid 3'5'-hydroxylase
ABB43050 Pericallis cruenta flavonoid 3'5'-hydroxylase
BAE71321 Trifolium pratense putative flavonoid 3,5'-hydroxylase
BAB82501 Glycine max AAD67966 AAD67935 flavonoid 3'5'-hydroxylase
AT5G07990 = NP_196416 Q9SD85 CAB82611 AAF60189 AAF72053 AAG10745 AAG10746 TT7 (TRANSPARENT TESTA 7); flavonoid 3'-monooxygenase/oxygen binding /F3'H, ARATH Flavonoid 3'-monooxygenase (Flavonoid 3'-dydroxylase) (AP3'H) (Cytochrome P450 75B1) (TRANSPA
AAD46851 Matthiola incana flavonoid 3'-hydroxylase
AAD46315 Petunia x hybrida flavonoid 3'-hydroxylase
AB05264 Gentiana sedifolia flavonoid 3'-hydroxylase
AB06383 Anthirrhinum majus flavonoid 3'-hydroxylase
BAE72874 Verbena x hybrida flavonoid 3'-hydroxylase
CAD42278 Vitis vinifera flavonoid 3'-hydroxylase
BAE47005 Vitis vinifera BAE47006 flavonoid 3'-hydroxylase
AB81560 Vinca major Flavonoid 3'-hydroxylase
AB89869 Osteospermum hybrid cultivar flavonoid 3'-hydroxylase
AAG45026 Catharanthus chinensis putative flavonoid 3'-hydroxylase
AB04502 Gerbera hybrida flavonoid 3'-hydroxylase
BAD05189 Ipomoea tricolor BAD05192 flavonoid 3'-hydroxylase/flavonoid 3'-hydroxylase
AB05187 Ipomoea sp BAD05190 flavonoid 3'-hydroxylase/flavonoid 3'-hydroxylase
AB05229 Ipomoea purpurea BAD05188 BAD05191 flavonoid 3'-hydroxylase/flavonoid 3'-hydroxylase
AB04503 Gerbera hybrida flavonoid 3'-hydroxylase
AB86839 Torenia hybrida BAD91806 Gentiana triflora flavonoid 3'-hydroxylase/ Cytochrome P450 75B2/flavonoid 3'-hydroxylase
AB86839 Torenia hybrida BAD91808 Gentiana triflora flavonoid 3'-hydroxylase/ flavonoid 3'-hydroxylase
AAD38930 Glycine max cytochrome P450 monooxygenase CYP92D1
ABC95985 Medicago truncatula cytochrome P450 monooxygenase CYP93A
AB05263 Petunia x hybrida flavonoid 3',5'-hydroxylase cytochrome P450
NP_015946 Orza sativa (japonica cultivar-group) CAE03126 CAE70984 B1160F02.15;OSJNBa0006L06.2 B1160F02.15
BAD05561 Oryza sativa (japonica cultivar-group) putative cytochrome P450
BAA36302 Lotus japonicus cytochrome P450
CAB50768 Cicer arietinum cytochrome P450
AAQ10282 Pisum sativum isoflavone synthase
P93149 Glycyrrhiza echinata BAA20423 CS5B1_GLYEC Cytochrome P450 93B1 (Licodione synthase) (I2S) flavanone 2-hydroxylase (Flavone synthase I) (CYP GE-5)/cytochrome P450
AAD33849 Glycine max cytochrome P450 monooxygenase
AAD35049 Gerbera hybrida cultivar flavone synthase II
AAF64115 Callistephus chinensis flavone synthase II
AAD61908 Gerbera hybrida cultivar flavone synthase II
AAA64072 Torenia hybrida cytochrome P450
AB04504 Gerbera hybrida cultivar flavone synthase II
BAA64071 Antirrhinum majus var. cytochrome P450
AT5G08090 = NP_196387 BAB11147 CYP93D1; hence bindingfrom ion binding monooxygenase/oxygen binding lysotochrome P450
O81973 Glycine max CAA71318 CAB43_SOYBN Cytochrome P450 93A3 (P450 CP3)/cytochrome P450
Q42799 Glycine max BAA13076 CS3A2_SOYBN Cytochrome P450 93A2/cytochrome P-450 (CYP93A2)
Q42798 Glycine max 2058818 BAA13075 CS3A1_SOYBN Cytochrome P450 93A1/P450 CD3(93A1)/cytochrome P-450 (CYP93A1)
AT1G05020 = NP_175485 AAB97871 AAG51197 BAC43325 AAD63960 CYP705A27; hence bindingfrom ion binding monooxygenase/oxygen binding Putative cytochrome P450 /lysotochrome P450, putative cytochrome cytochrome P450 putative cytochrome p450
AT5G52400 = NP_568158 BAB11146 CYP705A2; hence bindingfrom ion binding monooxygenase/oxygen binding lysotochrome-like protein
AT5G42550 = NP_197176 AAB86849 CYP705A1; hence bindingfrom ion binding monooxygenase/oxygen binding lysotochrome P450
Pya_SURZ_60_1
Pya_SURZ_2_1
Pya_SURZ_69_1
BAA52594 Petunia x hybrida cytochrome P450
Smo_SURZ_177_1
Smo_SURZ_120_1
AAN31543 Pinus taeda cytochrome P450 CYPC
BAD15331 Panax ginseng cytochrome P450
AAL96194 Pyrus communis cytochrome P450
AAD13591 Oryza sativa (japonica cultivar-group) Hypothetical protein
ATGG0990 = NP_188311 CYP98A3; hence bindingfrom ion binding monooxygenase/oxygen binding
AAR06877 Brassicanapus paprifera AAR06918 ferulate-5-hydroxylase/ferulate-5-hydroxylase
AAS06825 Centaurium erythrae coniferylaldehyd 5-hydroxylase
AAD14962 Brassica napus cytochrome p450-dependent monooxygenase
AAD14960 Brassica napus cytochrome p450-dependent monooxygenase
AT4G36220 = NP_195345 Q9D60 AAC40389 CAA19128 AAB11500 CAB86263 CAC20022 CAC20033 CAC20034 CAC20035 CAC20036 CAC20037 CAC20038 CAC20039 CAC20040 CAC20041 CAC20042 CAC20043 CAC20044 CAC20045 CAC20046 CAC20047 CAC20048 CAC20049 CAC20050 CAC20051 CAC20052 CAC20053 CAC20054 CAC20055 FAH1 (FERULATE-5-HYDROXYLASE 1); ferulate 5-hydroxylase/C9
AAD05225 Populus tremuloides AAD05226 CAC20059 CAC20060 CAC20061 ferulate-5-hydroxylase/ferulate-5-hydroxylase/ ferulate-5-hydroxylase/ferulate-5-hydroxylase/ferulate-5-hydroxylase
AAD48912 Liquidambar styraciflua aldehyde 5-hydroxylase
AAD37410 Lycopersicon esculentum x Lycopersicon pennavianum ferulate-5-hydroxylase
AAT36511 Camptotheca acuminata ferulate 5-hydroxylase
ABC86896 Sorghum bicolor cytochrome P450 monooxygenase CYP84A
AB05261 Medicago sativa ferulate 5-hydroxylase
AB80216 Medicago sativa ferulate 5-hydroxylase
AB05262 Medicago sativa ferulate 5-hydroxylase
AB04375 Oryza sativa (japonica cultivar-group) Cytochrome P450
AB40304 Triticum aestivum BAD80370 cytochrome P450/P450
AB80619 Triticum aestivum P450
AB80620 Triticum aestivum P450
AB40305 Hordeum lechleri Bx6 P-450
P93103 Zea mays T03050 CAA72207 CAA57424 AAL96768 AAL86769 C71C3_MAIZE Cytochrome P450 71C3 (Benzoxazinoless 5)/T03050 (cytochrome P450 - maize (fragment)/cytochrome p450 /cytochrome P450 monooxygenase CYP71C3v2 /cytochrome P450 monox
T03059 Zea mays CAA57421 T03059 cytochrome P450 - maize/cytochrome P450
Q43350 Zea mays CAA57422 C71C1_MAIZE Cytochrome P450 71C1 (Benzoxazinoless 4)/cytochrome P450
Q43349 Zea mays CAA57423 CAA72208 C71C2_MAIZE Cytochrome P450 71C2 (Benzoxazinoless 3)/cytochrome P450 /cytochrome p450
AB40343 Hordeum lechleri Bx3-like protein
AB40344 Hordeum lechleri P450-like protein
AB80618 Triticum aestivum P450
AB80683 Triticum aestivum BAD83306 cytochrome P450/P450
AB40306 Hordeum lechleri BAD83307 putative cytochrome P450 71C4/cytochrome cytochrome P450 71C4 /cytochrome cytochrome P450 71C4
AAK96807 Lolium rigidum putative cytochrome P450
AB80684 Triticum aestivum BAD83308 putative cytochrome P450/ putative cytochrome P450
AB80685 Triticum aestivum BAD83309 putative cytochrome P450/cytochrome P450
XP_479692 Orza sativa (japonica cultivar-group) BAD83310 putative cytochrome P450 71C1/cytochrome cytochrome P450 71C1 /cytochrome cytochrome P450 71C1
XP_479696 Orza sativa (japonica cultivar-group) BAD83311 putative cytochrome P450 71C2/cytochrome cytochrome P450 71C2 /cytochrome cytochrome P450 71C2
XP_479698 Orza sativa (japonica cultivar-group) BAD83312 putative cytochrome P450 71C3/cytochrome cytochrome P450 71C3 /cytochrome cytochrome P450 71C3
CAD3257 Zea mays CAA57425 CAA72196 C71C4_MAIZE Cytochrome P450 71C4 (Benzoxazinoless 2)/cytochrome P450 /cytochrome p450
BAB87820 Triticum aestivum AAS49242 Hordeum lechleri P450/Bx2-like protein
BAB87821 Triticum aestivum P450
BAB87822 Triticum aestivum P450
BAD16879 Muscari armeniacum BAD16880 cytochrome P450
BAD33773 Oryza sativa (japonica cultivar-group) putative cytochrome P450
BAD33774 Oryza sativa (japonica cultivar-group) putative cytochrome P450
AAU63111 Oryza sativa (japonica cultivar-group) putative cytochrome P450
XP_513462 Orza sativa (japonica cultivar-group) BAB87819 putative cytochrome P-450(LXXXIA1) (cyp71A1) family/putative Cytochrome P450 71A1
XP_513461 Orza sativa (japonica cultivar-group) BAB87818 putative cytochrome P-450(LXXXIA1) (cyp71A1) family/putative cytochrome P450 71A1
XP_513460 Orza sativa (japonica cultivar-group) BAB87817 putative cytochrome P-450(LXXXIA1) (cyp71A1) family/putative cytochrome P450 71A1
XP_513459 Orza sativa (japonica cultivar-group) BAB87816 putative cytochrome P-450(LXXXIA1) (cyp71A1) family/putative Cytochrome P450 71A1
AT4G39260 = NP_195671 hence bindingfrom ion binding monooxygenase/oxygen binding
O65458 Arabidopsis thaliana CYA98, ARATH Cytochrome P450 71A27
AT4G39310 = NP_195677 CAB78700 CAB40766 CAB73373 CYP71A25; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A1/lysotochrome p450-like protein lysotochrome p450-like protein lysotochrome p450 pro
AT4G31310 = NP_195067 Q9FIH2 CAB40766 CAB73373 CYP71A20; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A20/lysotochrome p450 like protein /cytochrome P450 71A20
AT4G26280 = NP_194973 Q9FIH0 BAD43010 BAD43011 CYP71A18; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A18/cytochrome P450 /cytochrome P450 71A18
P08647 Arabidopsis thaliana CYA98, ARATH Cytochrome P450 71A28
AT4G31500 = NP_195069 CAB40767 CAB73374 CYP71A21; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A1/E5E249 hypothetical protein /fibrosg similarity to g2380052 T11J7.14 putative cytochrome P450 from
AT5G07130 = NP_196377 CAB40768 CAB73375 CYP71A26; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A26/cytochrome P450
AT4G37370 = NP_195917 P05183 CAA30040 AAA16000 AAB40748 AAB11600 AAB40800 CYP71A12; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A12/putative cytochrome P450 (Arg/Gln3705/T11J7.14 (Arg/Gln3705/T11J7.14 (Arg/Gln3705/T11J7.14
AT4G37380 = NP_195918 P05184 CAA30041 AAB40749 AAB11601 AAB40801 CYP71A13; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A13/cytochrome P450
AT5G24960 = NP_197878 P05840 CYP71A14; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A14
AT3G48300 = NP_189150 CAB71187 CAB71188 CYP71A30; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A30/cytochrome p450 like protein
AT3G48280 = NP_189157 Q9STK0 CAB71170 CYP71A29; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A29/Cytochrome P450-like protein
AT4G37410 = NP_195921 P05185 CAA30042 AAB40750 AAB11602 AAB40802 CYP71A15; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A15/cytochrome P450 /cytochrome P450 71A15
AT4G37420 = NP_195922 CAB40769 CAB73376 CYP71A16; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A16/cytochrome P450 /cytochrome P450 71A16
AT4G37430 = NP_195923 CAB40770 CAB73377 CYP71A17; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A17/cytochrome P450 /cytochrome P450 71A17
AT4G37440 = NP_195924 CAB40771 CAB73378 CYP71A31; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A31/cytochrome P450 /cytochrome P450 71A31
AT4G37450 = NP_195925 CAB40772 CAB73379 CYP71A32; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A32/cytochrome P450 /cytochrome P450 71A32
AT4G37460 = NP_195926 CAB40773 CAB73380 CYP71A33; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A33/cytochrome P450 /cytochrome P450 71A33
AT5G04660 = NP_196579 CAB40774 CAB73381 CYP71A34; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A34/cytochrome P450 /cytochrome P450 71A34
AT5G04670 = NP_196580 CAB40775 CAB73382 CYP71A35; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A35/cytochrome P450 /cytochrome P450 71A35
AT5G42930 = NP_197001 CAB40776 CAB73383 CYP71A36; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A36/cytochrome P450 /cytochrome P450 71A36
AT5G42940 = NP_197002 CAB40777 CAB73384 CYP71A37; hence bindingfrom ion binding monooxygenase/oxygen binding CYA98, ARATH Cytochrome P450 71A37/cytochrome P450 /cytochrome P450 71A37
AT5G42950 = NP_197003