AT1G68920 = NP_849863 NP_177058 AAG51583 AAM91106 AAN72270 transcription factor/transcription factor /putative DNA-binding protein /At1g68920/T6L1_10 /At1g68920/T6L1_10

AT1G10120 = NP_172483 AAT06474 BAD93978 DNA binding/transcription factor /At1g10120 /bHLH transcription factor like protein

XP_470616 Oryza sativa (japonica cultivar-group) AAO00689 Hypothetical protein/Hypothetical protein

Ppa_ALC_21_1

XP_466283 Oryza sativa (japonica cultivar-group) BAD15821 putative bHLH transcription factor PTF1/putative bHLH transcription factor PTF1

AAO73566 Oryza sativa BAD35275 Oryza sativa (japonica cultivar-group) BAD35276 bHLH transcription factor PTF1/bHLH transcription factor PTF1 /putative bHLH transcription factor PTF1

AT4G30980 = NP_194827 CAA18195 CAB79816 AAM10956 DNA binding/transcription factor /putative protein /putative protein /putative bHLH transcription factor

BAD72512 Oryza sativa (japonica cultivar-group) bHLH transcription factor PTF1-like

AT5G58010 = NP_200609 BAA97525 DNA binding/transcription factor /unnamed protein product

Smo_ALC_4_1

AT4G02590 = NP_567245 NP_001031577 AAC78259 CAB80752 AAK32915 AAM10948 AAM65759 AAN28890 DNA binding/transcription factor /DNA binding /hypothetical protein /hypothetical protein /AT4g02590/T10P11_13 /putative bHLH transcription factor /putative lip

AT1G03040 = NP_563672 AAD25805 AAK96661 AAM47952 AAL55714 DNA binding/transcription factor /Contains PF|00010 helix-loop-helix DNA-binding domain. ESTs gb|T45640 and gb|T22783 come from this gene. /Unknown protein /unknown protein /putative transcr

ABB16991 Solanum tuberosum unknown

NP_909951 Oryza sativa (japonica cultivar-group) AAP44685 unknown protein/unknown protein

AAM10955 Arabidopsis thaliana putative bHLH transcription factor

AT2G24260 = NP_180003 AAD03387 BAD44133 BAD44153 DNA binding/transcription factor /unknown protein /putative bHLH transcription factor (bHLH066) /putative bHLH transcription factor (bHLH066)

XP_468258 Oryza sativa (japonica cultivar-group) BAD19075 BAD19276 BAD29274 basic helix-loop-helix (bHLH) -like/basic helix-loop-helix (bHLH) -like /basic helix-loop-helix (bHLH) -like /bHLH transcription factor PTF1-like protein

Smo_ALC_11_1

Smo_ALC_40_1

Ppa_ALC_35_1 Ppa_ALC_30_1

AAF24948 Arabidopsis thaliana T22C5.19

Ppa_ALC_3_1

Ppa_ALC_22_1

Ppa_ALC_31_1

Smo_ALC_15_1

Smo_ALC_10_1

Smo_ALC_1_2

AT5G41315 = NP_680372 BAB08503 AAL36964 GL3 (GLABRA 3); transcription factor/bHLH transcription factor-like protein /bHLH-transcription factor

AT1G63650 = NP_176552 NP_974080 D96661 Q9CAD0 AAG52418 AAL55709 EGL3 (ENHANCER OF GLABRA3); DNA binding/transcription factor /EGL3 (ENHANCER OF GLABRA3); DNA binding /D96661 probable transcription factor F24D7.16 /EGL1_ARATH Transcription factor EGL

AAB72192 Arabidopsis thaliana bHLH protein

ABA99246 Oryza sativa (japonica cultivar-group) Helix-loop-helix DNA-binding domain, putative

AT1G02340 = NP_563650 Q9FE22 AAG40617 AAG45733 AAK15282 HFR1 (LONG HYPOCOTYL IN FAR-RED); DNA binding/transcription factor /HFR1_ARATH Long hypocotyl in far-red 1 (bHLH-like protein HFR1) (Reduced phytochrome signaling) (Basic helix-loop-helix FBI1

AT5G61270 = NP_200935 NP_001032117 BAB08482 AAP86213 AAS79546 CAG25857 BAD95106 DNA binding/transcription factor /DNA binding /transcription factor /unnamed protein product /putative bHLH transcription factor /At5g61270 /hypothetical protein /bHLH t

AT4G00050 = NP_191916 AAM10933 BAC41905 DNA binding/transcription factor /putative bHLH transcription factor /putative bHLH transcription factor bHLH016

AT4G28790 = NP_974634 unknown protein

AT4G28790 = NP_194608 CAA22973 CAB81467 transcription factor/putative protein /putative protein

CAE12173 Arabidopsis thaliana putative bHLH127 transcription factor

AT2G20180 = NP_849996 NP_179608 AAD24380 AAN78308 BAC56979 PIL5 (PHYTOCHROME INTERACTING FACTOR 3-LIKE 5); transcription factor/PIL5 (PHYTOCHROME INTERACTING FACTOR 3-LIKE 5); transcription factor /unknown protein /putative bHLH transcription factor

AT3G59060 = NP_191465 NP_001030889 NP_001030890 NP_851021 CAB86934 AAM10954 BAC56978 PIL6 (PHYTOCHROME-INTERACTING FACTOR 5); DNA binding/transcription factor /PIL6 (PHYTOCHROME-INTERACTING FACTOR 5); DNA binding /transcription factor /PIL6 (PHYTOCH

AT2G43010 = NP_565991 NP_850381 Q8W2F3 H84860 AAL55716 CAD29449 PIF4 (PHYTOCHROME INTERACTING FACTOR 4); DNA binding/transcription factor /PIF4 (PHYTOCHROME INTERACTING FACTOR 4); DNA binding /transcription factor /PIF4_ARATH Phytochrome-interacting

CAD32238 Oryza sativa BP-5 protein

NP_911152 Oryza sativa (japonica cultivar-group) BAC19953 BAC21408 transcription factor BHLH9-like protein/transcription factor BHLH9-like protein /transcription factor BHLH9-like protein

ABA99362 Oryza sativa (japonica cultivar-group) ABA99363 ABA99364 BP-5 protein/BP-5 protein /BP-5 protein

XP_473908 Oryza sativa (japonica cultivar-group) CAE02150 OSJNBa0058K23.6/OSJNBa0058K23.6

ABD33186 Medicago truncatula Basic helix-loop-helix dimerisation region bHLH

AAC99771 Arabidopsis thaliana phytochrome-associated protein 3

AT1G09530 = NP_849626 NP_172424 O80536 AAC33213 AAC95156 AAL55715 BAC41930 PIF3 (PHYTOCHROME INTERACTING FACTOR 3); DNA binding/transcription factor/transcription regulator /PIF3 (PHYTOCHROME INTERACTING FACTOR 3); DNA binding /transcription factor/

AAT77368 Oryza sativa (japonica cultivar-group) AAU10768 hypothetical protein/hypothetical protein

AT3G62090 = NP_191768 BAC10690 PIL2 (PHYTOCHROME INTERACTING FACTOR 3-LIKE 2); transcription factor/PIF3 like basic Helix Loop Helix protein 2

AT2G46970 = NP_182220 BAC10689 AAO37214 AAX55166 PIL1 (PHYTOCHROME INTERACTING FACTOR 3-LIKE 1); transcription factor/PIF3 like basic Helix Loop Helix protein /hypothetical protein /hypothetical protein At2g46970

NP_912624 Oryza sativa (japonica cultivar-group) BAD81566 P0498A12.24/putative BP-5 protein

Smo_ALC_54_1

Smo_ALC_2_1

Ppa_ALC_23_1

Ppa_ALC_18_1

Smo_ALC_25_1

Smo_ALC_45_1

Smo_ALC_14_1 Smo_ALC_35_1

AAV33474 Fragaria x ananassa basic helix-loop-helix protein

AAK98706 Oryza sativa (japonica cultivar-group) Putative SPATULA

BAD67851 Oryza sativa (japonica cultivar-group) basic helix-loop-helix protein SPATULA-like

AT4G36930 = NP_568010 Q9FUA4 CAB16798 CAB80359 AAG33640 SPT (SPATULA); DNA binding/transcription factor /SPT_ARATH Protein SPATULA/putative protein /putative protein /SPATULA

BAD43719 Arabidopsis thaliana putative bHLH transcription factor (bHLH073/ALCATRAZ)

AT5G67110 = NP_201512 BAB10945 AAM61195 ALC (ALCATRAZ); DNA binding/transcription factor /unnamed protein product /unknown

Ppa_ALC_5_1

Ppa_ALC_19_1

10.000