

Intol- <i>Arabidopsis</i>	MAVNFTPSSGGQLTVSASLRKQDLGRSVHLANLEIRLTHTGAGIPEFLNQMFGEEDVSE	1081
Intol- <i>Populus</i>	MSVNFTPSSGGLLTVSASFSDQLGQSVYLVHLELRIRHPGAGIPEALLDQMYGEDTGASV	1080
Intol- <i>Eucalyptus</i>	VSINYTPTGSQVLVIAAHLTKDHLGESVQLAHLELRITYAGGGISEALLNQMFENDGEISE	1080
Intol- <i>Glycine max</i>	ISINFTPNGGQVVVAGSLTKEQLGKSVHLVKLELSITHGGSGVPEVLLNQMFGNNGLESE	1087
Intol-Scots pine	TVIQFTPAEKWVRIKVSPTKRHLGGGVYVIRIEFRITHPGPGLPEKLVQQMFNFDPDMSQ	806
Tol- <i>Picea glauca</i>	-----	
Tol-Norway spruce	NVIQFTPAEKWVRIKVSSTKRHLGGGVYVIRVEFRITHPGPGLPEKLVQQMFNLDPDMSQ	1120

Intol- <i>Arabidopsis</i>	EGLSLMVSRLVKLMNGDVQYLRQAGKSSFIITAELAAANK-----	1122
Intol- <i>Populus</i>	EGISLVISRKLVKLMNGDVRYMREAGKSS-----	1109
Intol- <i>Eucalyptus</i>	EGISLLISRKLVKLMNGDVQYLRQAGKSTFIISVELAVANKSGSSLKGKN	1130
Intol- <i>Glycine max</i>	EGISLLIRAKLLKLMNGDVRYLRQAGKSAFILSAELAAAHNLKA-----	1131
Intol-Scots pine	EGFGLLICRKLVRMLMNGDVQYLRGGGTSSFVIVVELATGQNEEC-----	850
Tol- <i>Picea glauca</i>	-----	
Tol-Norway spruce	EGFGLFICRKLVRMLMNGDVQYLRGGGTSSFIVVVELASGQTDEC-----	1164

Histidine kinase domain
902 - 1,119

Details of the sequences used for alignment

- Intol-*Arabidopsis* ***Arabidopsis thaliana*** - AT1G09570.1
- Intol-*Populus* ***Populus tremula*** - gi|340007753|gb|AEK26583.1| phytochrome A, partial [*Populus tremula*]
- Intol-*Eucalyptus* ***Eucalyptus grandis*** - gi|702263244|ref|XP_010037943.1| PREDICTED: phytochrome A [*Eucalyptus grandis*]
- Intol-*Glycine max* ***Glycine max*** - gi|351722971|ref|NP_001238286.1| phytochrome A [*Glycine max*]
- Intol-Scots pine ***Pinus sylvestris*** - gi|410718340|gb|AFV79546.1| phytochrome N, partial [*Pinus sylvestris*]
- Tol-*Picea glauca* ***Picea glauca*** - Congenei - 167818246 670 bp
- Tol-Norway spruce ***Picea abies*** - Congenei - gi|543906193|gb|AGV77083.1| PHYN [*Picea abies*]

Arabidopsis-Shade Intolerant	Nucleotide	Spruce-Shade Tolerant	Nucleotide
E-Glutamate Acidic, Polar, Hydrophilic	GAA	K-Lysine Basic, Polar, Hydrophilic	AAG
I-Isoleucine Basic, Non-polar, Hydrophobic	ATT	M-Methionine Non-polar, Hydrophobic	ATG
K-Lysine Basic, Polar, Hydrophilic	AAA	E-Glutamic Acid Acidic, Polar, Hydrophilic	GAA
L-Leucine Basic, Non-polar, Hydrophobic	TTA	F-Phenylalanine Non-polar, hydrophobic	TTC
Q-Glutamine Neutral, Polar, Hydrophilic	CAA	L-Leucine Basic, Non-polar, Hydrophobic	CTG
T-Tyrosine Neutral, Polar, Hydrophilic	ACC	A-Alanine Non-polar, Hydrophobic	GCG
G-Glycine Neutral, Non-polar, Hydrophilic	GGT	E-Glutamic Acid Acidic, Polar, Hydrophilic	GAA
A-Alanine Non-polar, Hydrophobic	GCA	S-Serine Neutral, Polar, Hydrophilic	TCT
K-Lysine Basic, Polar, Hydrophilic	AAA	T-Threonine Neutral, Polar, Hydrophilic	ACA
L-Leucine Basic, Non-polar, Hydroph	TTG	V-Valine Non-polar, Hydrophobic	GTC

Shade Tolerant species

Picea abies (Niinemets, 2010)

Picea glauca (Durzan, 2010)

Shade Intolerant species

Pinus sylvestris (Niinemets, 2010)

Populus tremula (Niinemets *et al.*, 1998)

Eucalyptus grandis (Niinemets, 1997)

Glycine max (Sawada *et al.*, 2002)

Arabidopsis thaliana (Mullen *et al.*, 2006)

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- Mullen JL, Weinig C, Hangarter RP.** 2006. Shade avoidance and the regulation of leaf inclination in Arabidopsis. *Plant Cell and Environment* **29**, 1099-1106.
- Niinemets U.** 1997. Role of foliar nitrogen in light harvesting and shade tolerance of four temperate deciduous woody species. *Functional Ecology* **11**, 518-531.
- Niinemets U.** 2010. A review of light interception in plant stands from leaf to canopy in different plant functional types and in species with varying shade tolerance. *Ecological Research* **25**, 693-714.
- Niinemets U, Kull O, Tenhunen JD.** 1998. An analysis of light effects on foliar morphology, physiology, and light interception in temperate deciduous woody species of contrasting shade tolerance. *Tree Physiology* **18**, 681-696.
- Sawada S, Kato T, Sato M, Kasai M.** 2002. Characteristics of gas exchange and morphology of a spring ephemeral, *Erythronium japonicum*, in comparison with a sun plant, *Glycine max*. *Ecological Research* **17**, 97-108.

[illegible]

Tol-SprucePhyB	YNADARLLQVFEQSGESGKSF	YRSIKSTTE--SVPEQQITAYLSRIQ	RGGR	IQPF	QGV	61
Tol-PsitchensisPhyp	YNADARLLQVFEQSGESGKSF	YRSIKSTTE--SVPEQQITAYLSRIQ	RGGR	IQPF	QGV	94
Intol-Pparviflora	YNADARLLEVFEQSGESGKSF	YRSIKSTTE--SVPEQQITAYLSRIQ	RGGR	IQPF	QGV	94
Intol-PineCon	YNSDARLLQVFEQSGESGKSF	YRSIQVHNR--AVPEQQITAYLSRIQ	RGGR	IQPF	QGV	94
Intol-Ptaeda	YNADARLLQVFEQSGESGKSF	YRSIKSTTE--SVPEQQITAYLSRIQ	RGGR	IQPF	QGV	94
Tol-Pglauca	YNADARLLQVFEQSGESGKSF	YRSIKSTTE--SVPEQQITAYLSRIQ	RGGR	IQPF	QGV	94
Intol-PineGenB	-----GESGKSF	YRSIKSTTE--SVPEQQITAYLSRIQ	RGGR	IQPF	QGV	44
Intol-PopulusB1	YTVDAQLHAVFEQSGGSGKSF	YSQSVRTTSQ--SVPEEQITAYLSKI	QRGG	HIQPF	QGM	101
Intol-PopulusB2	YTVDAQLHAVFEQSGGTGKSF	YSKSVRTPNQ--SVPEEQITAYLSKI	QRGG	HIQPF	QGM	99
Intol-Eucalyptus	YTQDARLHAVFEQSGESGRSF	YSQSVRTTTQ--SVPEQQITAYLSKI	QRGG	HIQPF	QGM	92
Intol-Glycine-max	YTEDARLHAVFEQSGESGRSF	YSESIRIASE--SVPEQQITAYLVKI	QRGG	FIQPF	GSM	102
Intol-Arabidopsis	YTVDARLHAVFEQSGESGKSF	YSQSLKTTTYGSSVPEQQITAYLSRI	QGGY	IQPF	QGM	120
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Tol-SprucePhyB	LAVEETTFRIIAYSENAREMLDLPQSVSPSEMQPQLEVLITIGTDVRTLFTAASAHSLEKA	121
Tol- <i>PsitchensisPhyp</i>	LAVEETTFRIIAYSENAREMLDLPQSVSPSEMQPQLEVLITIGTDVRTLFTAASAHSLEKA	154
Intol-Pparviflora	LAVEETTFRIIAYSENAREMLDLPQSVSPSEMSQLLEVLTIGTDVRTLFTAASSSHSLVKA	154
Intol-PineCon	LAVEETTFRIIAYSENE - EMLDLGAQSVSPSEMKPQQDVLITIGTDVRTLFTAASAHSLEKA	153
Intol-Ptaeda	LAVEETTFRIIAYSENAREMLDLPQSVSPSEMPKQQDVLITIGTDVRTLFTAASAHSLEKA	154
Tol-Pglauca	LAVEETTFRIIAYSENAREMLDLPQSVSPSEMQPQLEVLITIGTDVRTLFTAASAHSLEKA	154
Intol-PineGenB	LAVEETTFRIIAYSENAREMLDLPQSVSPSEMKPQQDVLITIGTDVRTLFTAASAHSLEKA	104
Intol-PopulusB1	IADDEGSFRVIAYSENAREMLGFTTPQSVPSLD - - KQEILSDGTDVRTLFRPSSSAMLEKA	159
Intol-PopulusB2	IADDEQSFRGIAYNENAKNMRLTTPQSVPSLE - - KQEILFVGGDVRILFRPSSAVLLEKT	157
Intol-Eucalyptus	IADDESTFRILAYSENARDMLGLTTPQSVPSLE - - KPEVLGIGTDVRTLFTPTSSGALLEKA	150
Intol-Glycine-max	IADDEPSFRILGYSDNARDMLGITTPQSVPSLDKDNDAAFALGTDRALFTHSSALLLEKA	162
Intol-Arabidopsis	IADDESFRIIGYSENAREMLGITTPQSVPTLE - - KPEILAMGTDVRSFLTSSSILLERA	178
	* * * * *	

Tol-SprucePhyB	AVAQ EISLMNP IWH C KNSR KP FYAI V HRIDVGMVID FEPL KTGD AFVSAAGAVQSQKLA	181
Tol-PsitchensisPhyp	AVAQ EISLMNP IWH C KNSR KP FYAI V HRIDVGMVID FEPL KTGD AFVSAAGAVQSQ---	211
Intol-Pparviflora	AVTQ EISLMNP IWH C KNSR KP FYAI V HRIDVGMVID LEPL RTGD AFMSAAGAVQSQ---	211
Intol-PineCon	AVAQ EISLMNP IWH C KNSR KP FYAI V HRIDVGMVID LEPL RTGD AFMSAAGAVQSQ---	210
Intol-Ptaeda	AVAQ EISLMNP IWH C KNSR KP FYAI V HRIDVGMVID LEPL RTGD AFMSAAGAVQSQ---	211
Tol-Pglauca	AVAQ EISLMNP IWH C KNSR KP FYAI V HRIDVGMVID FEPL KTGD AFVSAAGAVQSQ---	211
Intol-PineGenB	AVAQ EISLMNP IWH C KNSR KP FYAI V HRIDVGMVID LEPL RTGD AFMSAAGAVQSQ---	161
Intol-PopulusB1	FGAREITLLNP IWH I HSKNSG KP FYAIL HRIDVGIVID LEPARTEDPALSIAGAVQSQ---	216
Intol-PopulusB2	FGAREITLLNP IWH I HSKNSG KP FYAIL HRIDVGIVID LEPARTEDPALSIAGAVQSQ---	214
Intol-Eucalyptus	FGAREIMLLNPLWI HSKNSG KAFYAIL HRIDVGIVID LEPT RTEDPALSIAGAVQSQ---	207
Intol-Glycine-max	FSAREISLMNP IWH I HSRTSG KP FYGIL HRIDVGIVID LEPARTEDPALSIAGAVQSQ---	219
Intol-Arabidopsis	FVAREITLLNPVWI HSKNTG KP FYAIL HRIDVGIVID LEPARTEDPALSIAGAVQSQ---	235

PAS fold-2
103-219

Tol-SprucePhyB	VRAISRLQSLPCGDIELLCDTVVENVREL	TGYDRVMVYKFHSAAGAVQSQKLAVRAISRL	241
Tol-PsitchensisPhyp	-----	-----KLAVRAISRL	221
Intol-Pparviflora	-----	-----KLAVRAISRL	221
Intol-PineCon	-----	-----KLAVRAISRL	220
Intol-Ptaeda	-----	-----KLAVRAISRL	221
Tol-Pglauca	-----	-----KLAVRAISRL	221
Intol-PineGenB	-----	-----KLAVRAISRL	171
Intol-PopulusB1	-----	-----KLAVRSISQL	226
Intol-PopulusB2	-----	-----KLAVRAISQL	224
Intol-Eucalyptus	-----	-----KLAVRAISHL	217
Intol-Glycine-max	-----	-----KLAVRAISQL	229
Intol-Arabidopsis	-----	-----KLAVRAISQL	245
		*****:***:	

Tol-SprucePhyB	QSLPCGDIELLCDTVVENVREL	TGYDRVMVYKFHEDEHGEVVAEIR	RRSDLEPYLGLHYPA	301
Tol-PsitchensisPhyp	QSLPCGDIELLCDTVVENVREL	TGYDRVMVYKFHEDEHGEVVAEIR	RRSDLEPYLGLHYPA	281
Intol-Pparviflora	QSLPCGDIIGLLCDTVVENVREL	TGYDRVMVYKFHEDEHGEVVSEIR	RRSDLEPYLGLHYPA	281
Intol-PineCon	QSLPCGDIIGLLCDTVVENVREL	TGYDRVMVYKFHEDEHGEVVSEIR	RRSDLEPYLGLHYPA	280
Intol-Ptaeda	QSLPCGDIIGLLCDTVVENVREL	TGYDRVMVYKFHEDEHGEVVSEIR	RRSDLEPYLGLHYPA	281
Tol-Pglauca	QSLPCGDIIGLLCDTVVENVREL	TGYDRVMVYKFHEDEHGEVVSEIR	RRSDLEPYLGLHYPA	281
Intol-PineGenB	QSLPCGDIIGLLCDTVVENVREL	TGYDRVMVYKFHEDEHGEVVSEIR	RRSDLEPYLGLHYPA	281
Intol-PopulusB1	QSLPGGDIKLLCDTVVESVREL	TGYDRVMVYKFHEDEHGEVVAEN	KRADLEPYLGLHYPS	286
Intol-PopulusB2	QSLPGGDIKLLCDTVVESVREL	TGYDRVMVYKFHEDEHGEVVAEN	KRDLEPYLGLHYPS	284
Intol-Eucalyptus	QSLPGGDIKLLCDTVVESVREL	TGYDRVMVYKFHEDEHGEVVSEK	RADLEPYLGLHYPA	277
Intol-Glycine-max	QSLPGGDIKLLCDTVVESVREL	TGYDRVMVYKFHEDEHGEVVSEK	RPDLEPYLGLHYPA	289
Intol-Arabidopsis	QALPGGDIKLLCDTVVESVREL	TGYDRVMVYKFHEDEHGEVVAESK	RDDLEPYLGLHYPA	305
	. **:	*****:***:	*.* *****:*****:	

Tol-SprucePhyB	TDIPQASRFLFMQNRVRMICDCMATPVQVIO	SEELMQPLCLVGSTLRAPHGCHAQYMANM	361
Tol-PsitchensisPhyp	TDIPQASRFLFMQNRVRMICDCMATPVQVIO	SEELMQPLCLVGSTLRAPHGCHAQYMANM	341
Intol-Pparviflora	TDIPQASRFLFMQNRVRMICDCMATPVQVIO	SEELMQPLCLVGSTLRAPHGCHAQYMANM	341
Intol-PineCon	TDIPQASRFLFMQNRVRMICDCMATPVQVIO	SEELMQPLCLVGSTPSAPHGCHAQYMANM	340
Intol-Ptaeda	TDIPQASRFLFMQNRVRMICDCMATPVQVIO	SEELMQPLCLVGSTLRAPHGCHAQYMANM	341
Tol-Pglauca	TDIPQASRFLFMQNRVRMICDCMATPVQVIO	SEELMQPLCLVGSTLRAPHGCHAQYMANM	341
Intol-PineGenB	TDIPQASRFLFMQNRVRMICDCMATPVQVIO	SEELMQPLCLVGSTLRAPHGCHAQYMANM	291
Intol-PopulusB1	TDIPQASRFLFKQNRVRMIVDCHATPVRVIO	DEALMQPLCLVGSTLRAPHGCHAQYMANM	346
Intol-PopulusB2	TDIPQASRFLFKQNRVRMIVDCHAIPVSVIO	DEGLMQPLCLVGSTLRAPHGCHAQYMANM	344
Intol-Eucalyptus	TDIPQASRFLFKQNRVRMIADCHAMPVHVVO	DESLMQPLCLVGSTLRAPHGCHAQYMANM	337
Intol-Glycine-max	TDIPQASRFLFKQNRVRMIVDCHASAVRVVO	DEALVQPLCLVGSTLRAPHGCHAQYMANM	349
Intol-Arabidopsis	TDIPQASRFLFKQNRVRMIVDCNATPVLVVO	DDRLTQSMCLVGSTLRAPHGCHSQYMANM	365
	*****	*****	*****:*****

Tol-SprucePhyB	GSIASLVMIAVIINGNDDEGA--GSGRNSMKLWGLVVCHHTSPRAVPFPL	-----	408
Tol-PsitchensisPhyp	GSIASLVMIAVIINGNDDEGA--GSGRNSMKLWGLVVCHHTSPRAVPFPLRYACEFL	MQAF	399
Intol-Pparviflora	GSIASLVMIAVIINGNDDEGG--GSGRNSMKLWGLVVCHHTSPRAVPFPLRYACEFL	MQAF	399
Intol-PineCon	GSIRSLMAVIINGNDDEGG--GSGRNSMKLWGLVVCHHTSPRAVPFPLRYACEFL	MQAL	398
Intol-Ptaeda	GSIASLVMIAVIINGNDDEEG--GSGRNSMKLWGLVVCHHTSPRAVPFPLRYACEFL	MQAL	399
Tol-Pglauca	GSIASLVMIAVIINGNDDEGA--GSGRNSMKLWGLVVCHHTSPRAVPFPLRYACEFL	MQAF	399
Intol-PineGenB	GSIASLVMIAVIINGNDDEGG--GSGRNSMKLWGLVVCHHTSPRAVPFPLRYACEFL	MQAL	349
Intol-PopulusB1	GSIASMAMIAVIINGNEEEA---IGGRNSTRLWGLVVCHHTSARCIPFPLRYACEFL	MQAF	403
Intol-PopulusB2	GSIASLAMIAVIINGNDEEA---IGGRNSMRLWGLVVCHHTSARCIPFPLRYACEFL	MQAF	401
Intol-Eucalyptus	GSIASLAMIAVIINGNDEEA---VGGRNSMRLWGLVVCHHTSARCIPFPLRYACEFL	MQAF	394
Intol-Glycine-max	GSIASLVMIAVIINGNDEEG--VGGRSSMRLWGLVVCHHTSARCIPFPLRYACEFL	MQAF	406
Intol-Arabidopsis	GSIASLAMIAVIINGNEDDGSNVASGRSSMRLWGLVVCHHTSSRCIPFPLRYACEFL	MQAF	425
	*** *:	*****	*****:*****

GAF domain
252-443

Tol-SprucePhyB	VVLEFRITHPGPG-----LPAELVQDLFDRSRWTTQEGVG-LSMCR-KLLK	1104
Tol-PsitchensisPhyp	VVLEFRITHPGPG-----LPAELVQDLFDRSRWTTQEGVG-LSMCR-KLLK	1097
Intol-Pparviflora	VVHLEFRITHPGPG-----LPAELVQDLFDRSRWATQEGVG-LSMCR-KLLK	1097
Intol-PineCon	VVHLEFRITHPGLG-----LPAELVQDLFDRSQWATQEGVG-LSMCR-KLLK	1096
Intol-Ptaeda	-----	
Tol-Pglauca	-----	
Intol-PineGenB	-----	
Intol-PopulusB1	LVHMEFXXXXXXXXX-----XLPELVQDMFHSSRWVTQEGLG-LSMCR-KILK	1103
Intol-PopulusB2	LVHTEFRYLRLVQLTRMLGLQHPLRSFLNCKLTLVMLHREGMVPKIQFQGLSVCQGRLIW	1112
Intol-Eucalyptus	LVHNEFRIVCPGEG-----LPPELVQDMFHSSRWMTTEGLG-LSMCR-KILR	1092
Intol-Glycine-max	LLHAEFRMVCPGEG-----LPPELIQDMFNNSRWGTQEGLG-LSMSR-KILK	1104
Intol-Arabidopsis	AIRTEFRMACPGEG-----LPPELVQDMFHSSRWTSPEGLG-LSVCR-KILK	1126

Tol-SprucePhyB	LMNGDVQYIRESGKCYFLVNVELPLAQREDASSMK-----	1139
Tol-PsitchensisPhyp	LMNGDVQYVRESGKCYFLVNVELPLAQREDASSMK-----	1132
Intol-Pparviflora	LMNGDVQYIRESGKCYFLVNVEFPFAQREDAASSMK-----	1132
Intol-PineCon	LMNGDVRYIRESGICYFLVNVEFPMAQREDAASIK-----	1131
Intol-Ptaeda	-----	
Tol-Pglauca	-----	
Intol-PineGenB	-----	
Intol-PopulusB1	LMNGEVQYIRESERCYFLVILEVPM-----	1128
Intol-PopulusB2	LSFPENTTVALYPLTIWFLLLHRHQSRISITH-----	1144
Intol-Eucalyptus	LMSGEVQYIRESERCYFIITLELPMPQRSKDID-----	1125
Intol-Glycine-max	LMNGEVQYIREAERCYFYVLELPLVTRRSSKKC-----	1137
Intol-Arabidopsis	LMNGEVQYIRESERSYFLIILELPVPRKRPLSTASGSGDMMLMMPY	1172

Details of the sequences used for alignment

- Tol-SprucePhyB *Picea abies* - Congenei - MA_10435530g0010, 1139 residues PhyB
- Tol-PsitchensisPhyp *Picea sitchensis* - Congenei - PUT-183a-*Picea sitchensis*-8221 PlantGDB-assembled Unique Transcript-fragment derived from *Picea sitchensis* mRNAs Apr_26_2011 (based on GenBank release 183).
- Intol-Pparviflora *Pinus parviflora* - gi|635149027|gb|AHZ63958.1| phytochrome [*Pinus parviflora*]
- Intol-PineCon *Pinus sylvestris* - Congenei - PUT-187a-*Pinus sylvestris*-67937 PlantGDB-assembled Unique Transcript-fragment derived from *Pinus sylvestris* mRNAs Feb_1_2012 (based on GenBank release 187)
- Intol-Ptaeda *Pinus taeda* - Congenei - PITA_000008046[961 residues]
- Tol-Pglauca *Picea glauca* - Congenei - Sequence maker-Seq167953850-snap-gene-0.12 class=Sequence position=167953850:12666..15909 (+ strand) *P. glauca* seq from congenei 1 ORF:1..2241 Frame +1 sequence got by joining 2 orfs
- Intol-PineGenB *Pinus sylvestris* - gi|158322002|gb|ABW33480.1| phytochrome P, partial [*Pinus sylvestris*]
- Intol-PopulusB1 *Populus tremula* - gi|340007755|gb|AEK26584.1| phytochrome B1, partial [*Populus tremula*]
- Intol-PopulusB2 *Populus tremula* - gi|78643950|emb|CAJ21338.1| phytochrome B2 [*Populus tremula*]
- Intol-Eucalyptus *Eucalyptus grandis* - XP_010065155.1 PREDICTED: phytochrome B [*Eucalyptus grandis*]
- Intol-Glycine-max *Glycine max* - NP_001240097.1 phytochrome B [*Glycine max*]
- Intol-Arabidopsis *Arabidopsis thaliana* - AT2G18790.1

Arabidopsis-Shade Intolerant	Nucleotide	Spruce-Shade Tolerant	Nucleotide
L-Leucine Non polar, Hydrophobic	TTA	F-Phenylalanine Non polar, Hydrophobic	TTT
R-Arginine Basic, Polar, Hydrophilic	AGA	K-Lysine Basic, Polar, Hydrophilic	AAG
L-Leucine Non polar, Hydrophobic	TTG	M-Methionine Non polar, Hydrophobic	ATG
Q-Glutamine Neutral, Polar, Hydrophilic	CAA	H-Histidine Basic, Polar, Hydrophilic	CAT
Q-Glutamine Neutral, Polar, Hydrophilic	CAG	R-Arginine Basic, Polar, Hydrophilic	CGG
G-Glycine Non polar, Hydrophilic	GGA	A-Alanine Non polar, Hydrophobic	GCT

Shade Tolerant species

Picea abies (Niinemets, 2010)

Picea sitchensis (Kennedy *et al.*, 2007)

Picea glauca (Durzan, 2010)

Shade Intolerant species

Pinus parviflora (Watano *et al.*, 2004)

Pinus sylvestris (Niinemets, 2010)

Pinus taeda (Ewers *et al.*, 2001)

Populus tremula (Niinemets *et al.*, 1998)

Eucalyptus grandis (Niinemets, 1997)

Glycine max (Sawada *et al.*, 2002)

Arabidopsis thaliana (Mullen *et al.*, 2006)

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- Ewers BE, Oren R, Phillips N, Stromgren M, Linder S.** 2001. Mean canopy stomatal conductance responses to water and nutrient availabilities in *Picea abies* and *Pinus taeda*. *Tree Physiology* **21**, 841-850.
- Filiault DL, Wessinger CA, Dinneny JR, Lutes J, Borevitz JO, Weigel D, Chory J, Maloof JN.** 2008. Amino acid polymorphisms in *Arabidopsis* phytochrome B cause differential responses to light. *Proceedings of the National Academy of Sciences of the United States of America* **105**, 3157-3162.
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- Niinemets U.** 2010. A review of light interception in plant stands from leaf to canopy in different plant functional types and in species with varying shade tolerance. *Ecological Research* **25**, 693-714.
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