

# Final AFB phylogeny analysis

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2022-09-20

## Load libraries

```
#BiocManager::install("Rcpi")  
library(devtools)
```

```
## Loading required package: usethis
```

```
library(phangorn)
```

```
## Loading required package: ape
```

```
library(ggtree)
```

```
## Registered S3 methods overwritten by 'treeio':
```

```
##   method      from  
##   MRCA.phylo   tidytree  
##   MRCA.treedata tidytree  
##   Nnode.treedata tidytree  
##   Ntip.treedata tidytree  
##   ancestor.phylo tidytree  
##   ancestor.treedata tidytree  
##   child.phylo   tidytree  
##   child.treedata tidytree  
##   full_join.phylo tidytree  
##   full_join.treedata tidytree  
##   groupClade.phylo tidytree  
##   groupClade.treedata tidytree  
##   groupOTU.phylo tidytree  
##   groupOTU.treedata tidytree  
##   inner_join.phylo tidytree  
##   inner_join.treedata tidytree  
##   is.rooted.treedata tidytree  
##   nodeid.phylo   tidytree  
##   nodeid.treedata tidytree  
##   nodelab.phylo  tidytree  
##   nodelab.treedata tidytree  
##   offspring.phylo tidytree  
##   offspring.treedata tidytree  
##   parent.phylo   tidytree  
##   parent.treedata tidytree  
##   root.treedata  tidytree  
##   rootnode.phylo tidytree  
##   sibling.phylo   tidytree
```

```

## ggtree v3.8.2 For help: https://yulab-smu.top/treedata-book/
##
## If you use the ggtree package suite in published research, please cite
## the appropriate paper(s):
##
## Guangchuang Yu, David Smith, Huachen Zhu, Yi Guan, Tommy Tsan-Yuk Lam.
## ggtree: an R package for visualization and annotation of phylogenetic
## trees with their covariates and other associated data. Methods in
## Ecology and Evolution. 2017, 8(1):28-36. doi:10.1111/2041-210X.12628
##
## Shuangbin Xu, Lin Li, Xiao Luo, Meijun Chen, Wenli Tang, Li Zhan, Zehan
## Dai, Tommy T. Lam, Yi Guan, Guangchuang Yu. Ggtree: A serialized data
## object for visualization of a phylogenetic tree and annotation data.
## iMeta 2022, 1(4):e56. doi:10.1002/imt2.56
##
## Guangchuang Yu. Using ggtree to visualize data on tree-like structures.
## Current Protocols in Bioinformatics. 2020, 69:e96. doi:10.1002/cpbi.96
##
## Attaching package: 'ggtree'

## The following object is masked from 'package:ape':
##
##     rotate

library(treeio)

## treeio v1.24.3 For help: https://yulab-smu.top/treedata-book/
##
## If you use the ggtree package suite in published research, please cite
## the appropriate paper(s):
##
## LG Wang, TTY Lam, S Xu, Z Dai, L Zhou, T Feng, P Guo, CW Dunn, BR
## Jones, T Bradley, H Zhu, Y Guan, Y Jiang, G Yu. treeio: an R package
## for phylogenetic tree input and output with richly annotated and
## associated data. Molecular Biology and Evolution. 2020, 37(2):599-603.
## doi: 10.1093/molbev/msz240
##
## Guangchuang Yu. Data Integration, Manipulation and Visualization of
## Phylogenetic Trees (1st edition). Chapman and Hall/CRC. 2022,
## doi:10.1201/9781003279242
##
## Guangchuang Yu. Using ggtree to visualize data on tree-like structures.
## Current Protocols in Bioinformatics. 2020, 69:e96. doi:10.1002/cpbi.96
##
## Attaching package: 'treeio'

## The following object is masked from 'package:ape':
##
##     drop.tip

library(ggplot2)
library(tidyverse)

## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr      1.1.3      v readr      2.1.4

```

```

## v forcats 1.0.0      v stringr 1.5.0
## v lubridate 1.9.2    v tibble 3.2.1
## v purrr 1.0.2       v tidyr 1.3.0

## -- Conflicts ----- tidyverse_conflicts() --
## x tidyr::expand() masks ggtree::expand()
## x dplyr::filter() masks stats::filter()
## x dplyr::lag() masks stats::lag()
## x dplyr::where() masks ape::where()
## i Use the conflicted package (<http://conflicted.r-lib.org/>) to force all conflicts to become errors

library(BSgenome)

## Loading required package: BiocGenerics
##
## Attaching package: 'BiocGenerics'
##
## The following objects are masked from 'package:lubridate':
##
##   intersect, setdiff, union
##
## The following objects are masked from 'package:dplyr':
##
##   combine, intersect, setdiff, union
##
## The following objects are masked from 'package:stats':
##
##   IQR, mad, sd, var, xtabs
##
## The following objects are masked from 'package:base':
##
##   anyDuplicated, aperm, append, as.data.frame, basename, cbind,
##   colnames, dirname, do.call, duplicated, eval, evalq, Filter, Find,
##   get, grep, grepl, intersect, is.unsorted, lapply, Map, mapply,
##   match, mget, order, paste, pmax, pmax.int, pmin, pmin.int,
##   Position, rank, rbind, Reduce, rownames, sapply, setdiff, sort,
##   table, tapply, union, unique, unsplit, which.max, which.min
##
## Loading required package: S4Vectors
## Loading required package: stats4
##
## Attaching package: 'S4Vectors'
##
## The following objects are masked from 'package:lubridate':
##
##   second, second<-
##
## The following objects are masked from 'package:dplyr':
##
##   first, rename
##
## The following object is masked from 'package:tidyr':
##
##   expand
##

```

```

## The following object is masked from 'package:ggtree':
##
##     expand
##
## The following object is masked from 'package:utils':
##
##     findMatches
##
## The following objects are masked from 'package:base':
##
##     expand.grid, I, unname
##
## Loading required package: IRanges
##
## Attaching package: 'IRanges'
##
## The following object is masked from 'package:lubridate':
##
##     %within%
##
## The following objects are masked from 'package:dplyr':
##
##     collapse, desc, slice
##
## The following object is masked from 'package:purrr':
##
##     reduce
##
## The following object is masked from 'package:ggtree':
##
##     collapse
##
## Loading required package: GenomeInfoDb
## Loading required package: GenomicRanges
## Loading required package: Biostrings
## Loading required package: XVector
##
## Attaching package: 'XVector'
##
## The following object is masked from 'package:purrr':
##
##     compact
##
## Attaching package: 'Biostrings'
##
## The following object is masked from 'package:treeio':
##
##     mask
##
## The following object is masked from 'package:ape':
##
##     complement
##

```

```
## The following object is masked from 'package:base':
##
##      strsplit
##
## Loading required package: rtracklayer
library(Biostrings)
library(XVector)
library(rtracklayer)
library(GenomicFeatures)

## Loading required package: AnnotationDbi
## Loading required package: Biobase
## Welcome to Bioconductor
##
##      Vignettes contain introductory material; view with
##      'browseVignettes()'. To cite Bioconductor, see
##      'citation("Biobase")', and for packages 'citation("pkgname)".
##
##
## Attaching package: 'AnnotationDbi'
##
## The following object is masked from 'package:dplyr':
##
##      select
library(biomaRt)
library(ape)
library(DECIPHER)

## Loading required package: RSQLite
## Loading required package: parallel
library(Rcpi)

##
```

From Previous analysis

```
setwd("~/GitHub/SoyArcs_manuscript/final_trees/")

GmAFB <- Biostrings::readAAStringSet(filepath = "AFB_input/GmAFB.fasta")
unique(GmAFB)

## AAStringSet object of length 19:
##      width seq                      names
## [1] 573 MMNYFPDEVIEHIFDYVVS HDR...MYLYRTLAKRRKDAPEYVWTL* Glyma.16G050500|G...
## [2] 574 MNCFPDEVIGHIFGCVTSQRDRN...MYLYRTLAKRRKDAPELVWTL* Glyma.16G146400|G...
## [3] 584 MNQIDDGRTLSPDPQVLE NVLE...YMYRSLDRPRDDAPKVV TILC* Glyma.04G093500|G...
## [4] 574 MNFFPDEVIEHIFDYVTSQRDRN...TYLYRTL VGRRKDAPEHVWTL* Glyma.02G065300|G...
## [5] 631 MRDKSESH PSTNSEDDHRSSPLL...YMYRSLDGPRDDAPRFV TILQ* Glyma.02G211800|G...
## ... ..
## [15] 588 MKQIDDGDDEQRSLSP LPEQVLE...YMYRSLDGPRDDAPKVV TILC* Glyma.06G095400|G...
## [16] 587 MQKMA YTFSPPEEVLEHVFSFIW...PRLDMPGFVW TMEDDSSLRL* Glyma.03G209400|G...
## [17] 573 MMNYFPDEVIEHIFDYVVS HDR...MYLYRTLAKRRKDAPEYVWTL* Glyma.19G100200|G...
## [18] 588 MQRMA YTFSPPEEVLEHVFSFIW...PRLDMPGFVW TMEDDSSLRL* Glyma.19G206800|G...
## [19] 591 MTEERNVRKTRVVDVVLDCVIPY...GQRSDFPD TVVPLDTATCVD T* Glyma.18G030200|G...
```

```
names(GmAFB)
```

```
## [1] "Glyma.16G050500|Glyma.16G050500.1" "Glyma.16G146400|Glyma.16G146400.1"
## [3] "Glyma.04G093500|Glyma.04G093500.1" "Glyma.02G065300|Glyma.02G065300.1"
## [5] "Glyma.02G211800|Glyma.02G211800.1" "Glyma.02G254300|Glyma.02G254300.1"
## [7] "Glyma.02G152800|Glyma.02G152800.2" "Glyma.02G152800|Glyma.02G152800.3"
## [9] "Glyma.10G021500|Glyma.10G021500.2" "Glyma.08G059500|Glyma.08G059500.6"
## [11] "Glyma.07G189800|Glyma.07G189800.5" "Glyma.11G227300|Glyma.11G227300.1"
## [13] "Glyma.14G179500|Glyma.14G179500.1" "Glyma.14G062100|Glyma.14G062100.1"
## [15] "Glyma.06G095400|Glyma.06G095400.1" "Glyma.03G209400|Glyma.03G209400.1"
## [17] "Glyma.19G100200|Glyma.19G100200.1" "Glyma.19G206800|Glyma.19G206800.1"
## [19] "Glyma.18G030200|Glyma.18G030200.1"
```

```
phylotools::rm.sequence.fasta(infile = "AFB_input/GmAFB.fasta", to.rm = "Glyma.02G152800|Glyma.02G152800.2|Glyma.02G152800.3")
```

```
## AFB_input/GmAFB_removed.fasta has been saved to /Users/deisianyneres/GitHub/SoyArcs_manuscript/final
```

```
GmAFB <- Biostrings::readAAStringSet(filepath = "AFB_input/GmAFB_removed.fasta")
unique(GmAFB)
```

```
## AAStringSet object of length 18:
```

##		width	seq	names
##	[1]	573	MMNYFPDEVIEHIFDYVVS HDR...MYLYRTLAKRKDAPEYVWTL*	Glyma.16G050500 G...
##	[2]	574	MNCFPDEVIGHIFGCVTSQRDRN...MYLYRTLAKRRKDAPELVWTL*	Glyma.16G146400 G...
##	[3]	584	MNQIDDGRTLSPDPQVLE NVLE...YMYRSLDRPRDDAPKVV TILC*	Glyma.04G093500 G...
##	[4]	574	MNFFPDEVIEHIFDYVTSQRDRN...TYLYRTLVGRRKDAPEHVWTL*	Glyma.02G065300 G...
##	[5]	631	MRDKSESH PSTNSEDDHRSSPLL...YMYRSLDGPRDDAPRFV TILQ*	Glyma.02G211800 G...
##	...	...	...	...
##	[14]	588	MKQIDDGDDEQRSLSPLEQVLE...YMYRSLDGPRDDAPKVV TILC*	Glyma.06G095400 G...
##	[15]	587	MQKMAYTFSFPEEVLEHVFSFIW...PRLDMPGFVWTMEDDSSLRLE*	Glyma.03G209400 G...
##	[16]	573	MMNYFPDEVIEHIFDYVVS HDR...MYLYRTLAKRKDAPEYVWTL*	Glyma.19G100200 G...
##	[17]	588	MQR MAYTFSFPEEVLEHVFSFIW...PRLDMPGFVWTMEDDSSLRLE*	Glyma.19G206800 G...
##	[18]	591	MTEERNVRKTRVVDVLD CVIPY...GQRSDFPDTVVPLDTATC VDT*	Glyma.18G030200 G...

```
head(GmAFB, n = 10)
```

```
## AAStringSet object of length 10:
```

##		width	seq	names
##	[1]	573	MMNYFPDEVIEHIFDYVVS HDR...MYLYRTLAKRKDAPEYVWTL*	Glyma.16G050500 G...
##	[2]	574	MNCFPDEVIGHIFGCVTSQRDRN...MYLYRTLAKRRKDAPELVWTL*	Glyma.16G146400 G...
##	[3]	584	MNQIDDGRTLSPDPQVLE NVLE...YMYRSLDRPRDDAPKVV TILC*	Glyma.04G093500 G...
##	[4]	574	MNFFPDEVIEHIFDYVTSQRDRN...TYLYRTLVGRRKDAPEHVWTL*	Glyma.02G065300 G...
##	[5]	631	MRDKSESH PSTNSEDDHRSSPLL...YMYRSLDGPRDDAPRFV TILQ*	Glyma.02G211800 G...
##	[6]	585	MTARRLSDVVLDCVMPYI HDSKD...GPR TDFPDTVIPLDPGT YVDT*	Glyma.02G254300 G...
##	[7]	586	MRPRVAYSFPEEVLEHVFSFIEC...PRLDMPGYVWRMQD DSALRIS*	Glyma.02G152800 G...
##	[8]	586	MRPRVNYSFPEEVLEHVFSFIEC...PRLDMPGYVWRMQD DSALRIS*	Glyma.10G021500 G...
##	[9]	580	MMECR RKKENQNP NSTFPDEVLE...VYVYRSVAGPRRDAPPFV LTL*	Glyma.08G059500 G...
##	[10]	579	MECR RKKENQKS NSTFPDEVLER...VYVYRSVAGPRRDAPPFV LTL*	Glyma.07G189800 G...

```
tail(GmAFB, n=10)
```

```
## AAStringSet object of length 10:
```

##		width	seq	names
##	[1]	580	MMECR RKKENQNP NSTFPDEVLE...VYVYRSVAGPRRDAPPFV LTL*	Glyma.08G059500 G...
##	[2]	579	MECR RKKENQKS NSTFPDEVLER...VYVYRSVAGPRRDAPPFV LTL*	Glyma.07G189800 G...
##	[3]	591	MTEDR NVKTRVVDLVLDCVIPY...GQRSDFPDTVVPLDTATC VDT*	Glyma.11G227300 G...
##	[4]	641	MREKENHPSTNSEDDHRSS LLDL...YMYRSLDGPRDDAPRFV TILQ*	Glyma.14G179500 G...

```
## [5] 592 MEDRDAKRMATRLSDVVLDVCMP...GPRTDFPDTVIPLDPATYVDT* Glyma.14G062100|G...
## [6] 588 MKQIDGDEQRSLSPLPEQVLE...YMYRSLDGPRDDAPKVVITLC* Glyma.06G095400|G...
## [7] 587 MQKMAYTFSFPEEVLEHVFSFIW...PRLDMPGFVWTMEDDSSLRLE* Glyma.03G209400|G...
## [8] 573 MMNYFPDEVIEHIFDYVVS HDR...MYLYRTLAKRKDAPEYVWTL* Glyma.19G100200|G...
## [9] 588 MQRMAYTFSFPEEVLEHVFSFIW...PRLDMPGFVWTMEDDSSLRLE* Glyma.19G206800|G...
## [10] 591 MTEERNVRKTRVVDVLDVIPY...GQRSDFPDTVVPLDTATCVD* Glyma.18G030200|G...
```

```
# make alignment
```

```
(AFB_aligned <- DECIPHER::AlignSeqs(GmAFB))
```

```
## Determining distance matrix based on shared 5-mers:
```

```
## =====
##
```

```
## Time difference of 0.02 secs
```

```
##
```

```
## Clustering into groups by similarity:
```

```
## =====
##
```

```
## Time difference of 0.01 secs
```

```
##
```

```
## Aligning Sequences:
```

```
## =====
##
```

```
## Time difference of 0.27 secs
```

```
##
```

```
## Iteration 1 of 2:
```

```
##
```

```
## Determining distance matrix based on alignment:
```

```
## =====
##
```

```
## Time difference of 0 secs
```

```
##
```

```
## Reclustering into groups by similarity:
```

```
## =====
##
```

```
## Time difference of 0.01 secs
```

```
##
```

```
## Realigning Sequences:
```

```
## =====
##
```

```
## Time difference of 0.01 secs
```

```
##
```

```
## Alignment converged - skipping remaining iteration.
```

```
## AASTringSet object of length 18:
```

```
##      width seq                                     names
## [1] 670 -----...KRKDAPEYVWTL-----* Glyma.16G050500|G...
## [2] 670 -----...RRKDAPELVWTL-----* Glyma.16G146400|G...
## [3] 670 -----...PRDDAPKVVITL-----C* Glyma.04G093500|G...
## [4] 670 -----...RRKDAPEHVWTL-----* Glyma.02G065300|G...
## [5] 670 ---MRDKSESHPSTNSEDDHRSS...PRDDAPRFVITL-----Q* Glyma.02G211800|G...
## ...    ...
## [14] 670 -----...PRDDAPKVVITL-----C* Glyma.06G095400|G...
## [15] 670 -----...PRLDMPGFVWTMEDDSSLRLE* Glyma.03G209400|G...
## [16] 670 -----...KRKDAPEYVWTL-----* Glyma.19G100200|G...
```

```
## [17] 670 -----...PRLDMPGFVWTMEDDSSLRL* Glyma.19G206800|G...
## [18] 670 -----...QRSDFPDTPVPLDTATCVD-T* Glyma.18G030200|G...
```

```
# Fix names, because characters such as | are not accepted in MrBayes
names(AFB_aligned) <- stringr::str_remove(names(AFB_aligned), ".*\\|") # remove everything before .*\\|

# write.csv(names(AFB_aligned), "AFB_IDs_all.csv")

# CUT low alignment regions by masking alignment
(AFB_masked <- DECIPHER::MaskAlignment(AFB_aligned,
  type = "sequences",
  windowSize = 6, # same as aliscore default value.
  threshold = 1,
  maxFractionGaps = 0.2,
  includeTerminalGaps = FALSE,
  correction = FALSE,
  showPlot = FALSE))
```

```
## AAMultipleAlignment with 18 rows and 670 columns
```

```
##      aln                                     names
## [1] ----#####...LAGKRKDAPEYVWTL##### Glyma.16G050500.1
## [2] ----#####...LAGRRKDAPELVWTL##### Glyma.16G146400.1
## [3] ----#####...LDRPRDDAPKVVTIL##### Glyma.04G093500.1
## [4] ----#####...LVGRRKDAPEHVWTL##### Glyma.02G065300.1
## [5] ---MR#####...LDGPRDDAPRFVTIL##### Glyma.02G211800.1
## [6] ----#####...LAGPRTDFPDTVIPL##### Glyma.02G254300.1
## [7] ----#####...VSGPRLDMPGYVWRM##### Glyma.02G152800.2
## [8] ----#####...VSGPRLDMPGYVWRM##### Glyma.10G021500.2
## [9] MMECR#####...VAGPRRDAPPFVLT##### Glyma.08G059500.6
## [10] -MECR#####...VAGPRRDAPPFVLT##### Glyma.07G189800.5
## [11] ----#####...LAGQRSDFPDTPVPL##### Glyma.11G227300.1
## [12] ---MR#####...LDGPRDDAPRFVTIL##### Glyma.14G179500.1
## [13] ----#####...LAGPRTDFPDTVIPL##### Glyma.14G062100.1
## [14] ----#####...LDGPRDDAPKVVTIL##### Glyma.06G095400.1
## [15] ----#####...VAGPRLDMPGFVWTM##### Glyma.03G209400.1
## [16] ----#####...LAGKRKDAPEYVWTL##### Glyma.19G100200.1
## [17] ----#####...IAGPRLDMPGFVWTM##### Glyma.19G206800.1
## [18] ----#####...LAGQRSDFPDTPVPL##### Glyma.18G030200.1
```

```
# make it a AAStringSet
AAstr_AFB <- as(AFB_masked, "AAStringSet")
BrowseSeqs(AAstr_AFB) # A few sequences seems really small, however I believe they will probably be dropped

writeXStringSet(AAstr_AFB, file="AFB_input//masked_AFB.fasta" )

#Create a matrix so we can create the nexus file next
(matrix_AFB <- as.matrix(AAstr_AFB, use.names=TRUE))
```

```
##      [,1] [,2] [,3] [,4] [,5] [,6] [,7] [,8] [,9] [,10] [,11]
## Glyma.16G050500.1 "-" "-" "-" "-" "-" "-" "-" "-" "-" "-"
## Glyma.16G146400.1 "-" "-" "-" "-" "-" "-" "-" "-" "-" "-"
## Glyma.04G093500.1 "-" "-" "-" "-" "-" "-" "-" "R" "T" "L" "S"
## Glyma.02G065300.1 "-" "-" "-" "-" "-" "-" "-" "-" "-" "-"
## Glyma.02G211800.1 "-" "-" "-" "M" "R" "P" "S" "E" "P" "Q" "A"
## Glyma.02G254300.1 "-" "-" "-" "-" "-" "-" "-" "-" "-" "-"
## Glyma.02G152800.2 "-" "-" "-" "-" "-" "-" "-" "-" "M" "R" "P"
```



```

## Glyma.10G021500.2 "-" "-" "-" "-" "-" "-" "-" "-" "M" "R" "P"
## Glyma.08G059500.6 "M" "M" "E" "C" "R" "R" "K" "K" "E" "N" "Q"
## Glyma.07G189800.5 "-" "M" "E" "C" "R" "R" "K" "K" "E" "N" "Q"
## Glyma.11G227300.1 "-" "-" "-" "-" "-" "M" "T" "E" "D" "R" "N"
## Glyma.14G179500.1 "-" "-" "-" "M" "R" "T" "S" "E" "P" "Q" "A"
## Glyma.14G062100.1 "-" "-" "-" "-" "-" "E" "D" "R" "D" "A" "K"
## Glyma.06G095400.1 "-" "-" "-" "-" "-" "E" "Q" "R" "S" "L" "S"
## Glyma.03G209400.1 "-" "-" "-" "-" "-" "-" "-" "M" "Q" "K" "M"
## Glyma.19G100200.1 "-" "-" "-" "-" "-" "-" "-" "-" "-" "-" "-"
## Glyma.19G206800.1 "-" "-" "-" "-" "-" "-" "-" "M" "Q" "R" "M"
## Glyma.18G030200.1 "-" "-" "-" "-" "-" "M" "T" "E" "E" "R" "N"
##
## Glyma.16G050500.1 [,12] [,13] [,14] [,15] [,16] [,17] [,18] [,19] [,20] [,21]
## Glyma.16G146400.1 "-" "M" "M" "N" "Y" "F" "P" "D" "E" "V"
## Glyma.16G146400.1 "-" "-" "M" "N" "C" "F" "P" "D" "E" "V"
## Glyma.04G093500.1 "P" "L" "P" "D" "Q" "V" "L" "E" "N" "V"
## Glyma.02G065300.1 "-" "-" "M" "N" "F" "F" "P" "D" "E" "V"
## Glyma.02G211800.1 "P" "F" "P" "D" "Q" "V" "L" "E" "N" "V"
## Glyma.02G254300.1 "M" "T" "A" "R" "R" "L" "S" "D" "V" "V"
## Glyma.02G152800.2 "R" "V" "A" "Y" "S" "F" "P" "E" "E" "V"
## Glyma.10G021500.2 "R" "V" "N" "Y" "S" "F" "P" "E" "E" "V"
## Glyma.08G059500.6 "N" "P" "N" "S" "T" "F" "P" "D" "E" "V"
## Glyma.07G189800.5 "K" "S" "N" "S" "T" "F" "P" "D" "E" "V"
## Glyma.11G227300.1 "V" "R" "K" "T" "R" "V" "V" "D" "L" "V"
## Glyma.14G179500.1 "P" "F" "P" "D" "Q" "V" "L" "E" "N" "V"
## Glyma.14G062100.1 "R" "M" "A" "T" "R" "L" "S" "D" "V" "V"
## Glyma.06G095400.1 "P" "L" "P" "E" "Q" "V" "L" "E" "N" "V"
## Glyma.03G209400.1 "A" "Y" "T" "F" "S" "F" "P" "E" "E" "V"
## Glyma.19G100200.1 "-" "M" "M" "N" "Y" "F" "P" "D" "E" "V"
## Glyma.19G206800.1 "A" "Y" "T" "F" "S" "F" "P" "E" "E" "V"
## Glyma.18G030200.1 "V" "R" "K" "T" "R" "V" "V" "D" "V" "V"
##
## Glyma.16G050500.1 [,22] [,23] [,24] [,25] [,26] [,27] [,28] [,29] [,30] [,31]
## Glyma.16G146400.1 "I" "E" "H" "I" "F" "D" "Y" "V" "V" "S"
## Glyma.16G146400.1 "I" "G" "H" "I" "F" "G" "C" "V" "T" "S"
## Glyma.04G093500.1 "L" "E" "S" "V" "L" "H" "F" "L" "T" "S"
## Glyma.02G065300.1 "I" "E" "H" "I" "F" "D" "Y" "V" "T" "S"
## Glyma.02G211800.1 "L" "E" "N" "V" "L" "H" "F" "L" "P" "S"
## Glyma.02G254300.1 "L" "D" "C" "V" "M" "P" "Y" "I" "H" "D"
## Glyma.02G152800.2 "L" "E" "H" "V" "F" "S" "F" "I" "E" "C"
## Glyma.10G021500.2 "L" "E" "H" "V" "F" "S" "F" "I" "E" "C"
## Glyma.08G059500.6 "L" "E" "R" "I" "S" "G" "M" "L" "K" "S"
## Glyma.07G189800.5 "L" "E" "R" "I" "L" "G" "M" "L" "K" "S"
## Glyma.11G227300.1 "L" "D" "C" "V" "I" "P" "Y" "I" "D" "D"
## Glyma.14G179500.1 "L" "E" "N" "V" "L" "H" "F" "L" "S" "S"
## Glyma.14G062100.1 "L" "D" "C" "V" "M" "P" "Y" "I" "H" "D"
## Glyma.06G095400.1 "L" "E" "S" "V" "L" "H" "F" "L" "T" "S"
## Glyma.03G209400.1 "L" "E" "H" "V" "F" "S" "F" "I" "W" "S"
## Glyma.19G100200.1 "I" "E" "H" "I" "F" "D" "Y" "V" "V" "S"
## Glyma.19G206800.1 "L" "E" "H" "V" "F" "S" "F" "I" "W" "N"
## Glyma.18G030200.1 "L" "D" "C" "V" "I" "P" "Y" "I" "D" "D"
##
## Glyma.16G050500.1 [,32] [,33] [,34] [,35] [,36] [,37] [,38] [,39] [,40] [,41]
## Glyma.16G146400.1 "H" "S" "D" "R" "N" "A" "L" "S" "L" "V"
## Glyma.16G146400.1 "Q" "R" "D" "R" "N" "A" "V" "S" "L" "V"
## Glyma.04G093500.1 "R" "R" "D" "R" "N" "A" "A" "S" "L" "V"
## Glyma.02G065300.1 "Q" "R" "D" "R" "N" "D" "L" "S" "L" "V"

```

## Glyma.02G211800.1	"R"	"R"	"D"	"R"	"N"	"A"	"A"	"S"	"L"	"V"
## Glyma.02G254300.1	"S"	"K"	"D"	"R"	"D"	"A"	"V"	"S"	"Q"	"V"
## Glyma.02G152800.2	"D"	"K"	"D"	"R"	"G"	"S"	"I"	"S"	"L"	"V"
## Glyma.10G021500.2	"D"	"K"	"D"	"R"	"G"	"S"	"I"	"S"	"L"	"V"
## Glyma.08G059500.6	"R"	"K"	"D"	"K"	"S"	"T"	"V"	"S"	"L"	"V"
## Glyma.07G189800.5	"R"	"K"	"D"	"K"	"S"	"T"	"V"	"S"	"L"	"V"
## Glyma.11G227300.1	"P"	"K"	"D"	"R"	"D"	"A"	"V"	"S"	"Q"	"V"
## Glyma.14G179500.1	"R"	"R"	"D"	"R"	"N"	"A"	"A"	"S"	"L"	"V"
## Glyma.14G062100.1	"S"	"K"	"D"	"R"	"D"	"A"	"V"	"S"	"Q"	"V"
## Glyma.06G095400.1	"R"	"R"	"D"	"R"	"N"	"A"	"A"	"S"	"L"	"V"
## Glyma.03G209400.1	"E"	"R"	"D"	"R"	"N"	"A"	"I"	"S"	"L"	"V"
## Glyma.19G100200.1	"H"	"S"	"D"	"R"	"N"	"A"	"L"	"S"	"L"	"V"
## Glyma.19G206800.1	"E"	"R"	"D"	"R"	"N"	"A"	"I"	"S"	"L"	"V"
## Glyma.18G030200.1	"P"	"K"	"D"	"R"	"D"	"A"	"V"	"S"	"Q"	"V"
##	[,42]	[,43]	[,44]	[,45]	[,46]	[,47]	[,48]	[,49]	[,50]	[,51]
## Glyma.16G050500.1	"C"	"K"	"S"	"W"	"Y"	"R"	"I"	"E"	"R"	"C"
## Glyma.16G146400.1	"C"	"K"	"N"	"W"	"H"	"R"	"L"	"E"	"R"	"C"
## Glyma.04G093500.1	"C"	"K"	"S"	"W"	"Y"	"R"	"A"	"E"	"A"	"L"
## Glyma.02G065300.1	"C"	"K"	"N"	"W"	"H"	"R"	"L"	"E"	"R"	"C"
## Glyma.02G211800.1	"C"	"R"	"S"	"W"	"Y"	"R"	"A"	"E"	"A"	"L"
## Glyma.02G254300.1	"C"	"R"	"R"	"W"	"Y"	"E"	"L"	"D"	"S"	"L"
## Glyma.02G152800.2	"C"	"K"	"S"	"W"	"Y"	"E"	"I"	"E"	"R"	"W"
## Glyma.10G021500.2	"C"	"K"	"S"	"W"	"Y"	"E"	"I"	"E"	"R"	"W"
## Glyma.08G059500.6	"C"	"K"	"E"	"W"	"Y"	"N"	"A"	"E"	"R"	"W"
## Glyma.07G189800.5	"C"	"K"	"E"	"W"	"F"	"N"	"A"	"E"	"R"	"W"
## Glyma.11G227300.1	"C"	"R"	"R"	"W"	"Y"	"E"	"L"	"D"	"S"	"L"
## Glyma.14G179500.1	"C"	"R"	"S"	"W"	"Y"	"R"	"A"	"E"	"A"	"L"
## Glyma.14G062100.1	"C"	"R"	"R"	"L"	"Y"	"E"	"L"	"D"	"S"	"L"
## Glyma.06G095400.1	"C"	"K"	"S"	"W"	"Y"	"R"	"A"	"E"	"A"	"L"
## Glyma.03G209400.1	"C"	"K"	"S"	"W"	"Y"	"E"	"I"	"E"	"R"	"W"
## Glyma.19G100200.1	"C"	"K"	"S"	"W"	"Y"	"R"	"I"	"E"	"R"	"C"
## Glyma.19G206800.1	"C"	"K"	"S"	"W"	"Y"	"E"	"I"	"E"	"R"	"W"
## Glyma.18G030200.1	"C"	"R"	"R"	"W"	"Y"	"E"	"L"	"D"	"S"	"L"
##	[,52]	[,53]	[,54]	[,55]	[,56]	[,57]	[,58]	[,59]	[,60]	[,61]
## Glyma.16G050500.1	"T"	"R"	"Q"	"R"	"V"	"F"	"I"	"G"	"N"	"C"
## Glyma.16G146400.1	"C"	"R"	"K"	"S"	"L"	"F"	"I"	"G"	"N"	"C"
## Glyma.04G093500.1	"T"	"R"	"T"	"E"	"L"	"F"	"I"	"G"	"N"	"C"
## Glyma.02G065300.1	"S"	"R"	"K"	"S"	"L"	"F"	"I"	"G"	"N"	"C"
## Glyma.02G211800.1	"T"	"R"	"S"	"E"	"L"	"F"	"I"	"G"	"N"	"C"
## Glyma.02G254300.1	"T"	"R"	"K"	"H"	"V"	"T"	"I"	"A"	"L"	"C"
## Glyma.02G152800.2	"C"	"R"	"R"	"R"	"V"	"F"	"V"	"G"	"N"	"C"
## Glyma.10G021500.2	"C"	"R"	"R"	"R"	"V"	"F"	"V"	"G"	"N"	"C"
## Glyma.08G059500.6	"S"	"R"	"R"	"S"	"V"	"F"	"I"	"G"	"N"	"C"
## Glyma.07G189800.5	"S"	"R"	"R"	"S"	"V"	"F"	"I"	"G"	"N"	"C"
## Glyma.11G227300.1	"T"	"R"	"K"	"H"	"V"	"T"	"I"	"A"	"L"	"C"
## Glyma.14G179500.1	"T"	"R"	"S"	"E"	"L"	"F"	"I"	"G"	"N"	"C"
## Glyma.14G062100.1	"T"	"R"	"K"	"H"	"V"	"T"	"I"	"A"	"L"	"C"
## Glyma.06G095400.1	"T"	"R"	"P"	"D"	"L"	"F"	"I"	"G"	"N"	"C"
## Glyma.03G209400.1	"C"	"R"	"R"	"K"	"V"	"F"	"V"	"G"	"N"	"C"
## Glyma.19G100200.1	"T"	"R"	"Q"	"R"	"V"	"F"	"I"	"G"	"N"	"C"
## Glyma.19G206800.1	"C"	"R"	"R"	"K"	"V"	"F"	"V"	"G"	"N"	"C"
## Glyma.18G030200.1	"T"	"R"	"K"	"H"	"V"	"T"	"I"	"A"	"L"	"C"
##	[,62]	[,63]	[,64]	[,65]	[,66]	[,67]	[,68]	[,69]	[,70]	[,71]
## Glyma.16G050500.1	"Y"	"S"	"I"	"T"	"P"	"E"	"R"	"L"	"I"	"Q"

## Glyma.16G146400.1	"Y"	"T"	"I"	"S"	"P"	"E"	"R"	"V"	"I"	"E"
## Glyma.04G093500.1	"Y"	"A"	"V"	"S"	"P"	"R"	"R"	"A"	"T"	"G"
## Glyma.02G065300.1	"Y"	"S"	"I"	"S"	"P"	"E"	"R"	"V"	"I"	"E"
## Glyma.02G211800.1	"Y"	"A"	"L"	"S"	"P"	"T"	"R"	"A"	"T"	"A"
## Glyma.02G254300.1	"Y"	"T"	"T"	"T"	"P"	"D"	"R"	"L"	"R"	"R"
## Glyma.02G152800.2	"Y"	"A"	"V"	"S"	"P"	"A"	"T"	"V"	"V"	"N"
## Glyma.10G021500.2	"Y"	"A"	"V"	"S"	"P"	"A"	"T"	"V"	"V"	"N"
## Glyma.08G059500.6	"Y"	"S"	"V"	"S"	"P"	"E"	"I"	"L"	"T"	"R"
## Glyma.07G189800.5	"Y"	"S"	"V"	"S"	"P"	"E"	"I"	"L"	"T"	"R"
## Glyma.11G227300.1	"Y"	"T"	"T"	"T"	"P"	"A"	"R"	"L"	"R"	"R"
## Glyma.14G179500.1	"Y"	"A"	"L"	"S"	"P"	"T"	"R"	"A"	"T"	"A"
## Glyma.14G062100.1	"Y"	"T"	"T"	"T"	"P"	"D"	"R"	"L"	"R"	"R"
## Glyma.06G095400.1	"Y"	"A"	"V"	"S"	"P"	"R"	"R"	"A"	"T"	"A"
## Glyma.03G209400.1	"Y"	"A"	"V"	"S"	"P"	"L"	"M"	"V"	"I"	"K"
## Glyma.19G100200.1	"Y"	"S"	"I"	"T"	"P"	"E"	"R"	"L"	"I"	"Q"
## Glyma.19G206800.1	"Y"	"A"	"V"	"S"	"P"	"L"	"M"	"V"	"V"	"K"
## Glyma.18G030200.1	"Y"	"T"	"T"	"T"	"P"	"A"	"R"	"L"	"R"	"R"
##	[,72]	[,73]	[,74]	[,75]	[,76]	[,77]	[,78]	[,79]	[,80]	[,81]
## Glyma.16G050500.1	"R"	"F"	"P"	"G"	"L"	"K"	"S"	"L"	"T"	"L"
## Glyma.16G146400.1	"R"	"F"	"P"	"E"	"L"	"R"	"S"	"L"	"T"	"L"
## Glyma.04G093500.1	"R"	"F"	"P"	"R"	"V"	"R"	"S"	"V"	"T"	"I"
## Glyma.02G065300.1	"R"	"F"	"P"	"E"	"L"	"K"	"S"	"L"	"T"	"L"
## Glyma.02G211800.1	"R"	"F"	"T"	"R"	"V"	"M"	"S"	"V"	"T"	"V"
## Glyma.02G254300.1	"R"	"F"	"P"	"H"	"L"	"E"	"S"	"L"	"K"	"L"
## Glyma.02G152800.2	"R"	"F"	"P"	"K"	"V"	"R"	"S"	"I"	"A"	"I"
## Glyma.10G021500.2	"R"	"F"	"P"	"K"	"V"	"R"	"S"	"I"	"A"	"I"
## Glyma.08G059500.6	"R"	"F"	"P"	"N"	"I"	"R"	"S"	"V"	"T"	"L"
## Glyma.07G189800.5	"R"	"F"	"P"	"N"	"I"	"R"	"S"	"V"	"T"	"L"
## Glyma.11G227300.1	"R"	"F"	"P"	"H"	"L"	"E"	"S"	"L"	"K"	"L"
## Glyma.14G179500.1	"R"	"F"	"T"	"R"	"A"	"R"	"S"	"V"	"T"	"V"
## Glyma.14G062100.1	"R"	"F"	"P"	"H"	"L"	"E"	"S"	"L"	"N"	"L"
## Glyma.06G095400.1	"R"	"F"	"P"	"R"	"V"	"R"	"S"	"L"	"T"	"I"
## Glyma.03G209400.1	"R"	"F"	"P"	"E"	"L"	"R"	"S"	"I"	"A"	"L"
## Glyma.19G100200.1	"R"	"F"	"P"	"G"	"L"	"K"	"S"	"L"	"T"	"L"
## Glyma.19G206800.1	"R"	"F"	"P"	"E"	"V"	"R"	"S"	"I"	"A"	"L"
## Glyma.18G030200.1	"R"	"F"	"P"	"H"	"L"	"E"	"S"	"L"	"K"	"L"
##	[,82]	[,83]	[,84]	[,85]	[,86]	[,87]	[,88]	[,89]	[,90]	[,91]
## Glyma.16G050500.1	"K"	"G"	"K"	"P"	"H"	"F"	"A"	"D"	"F"	"S"
## Glyma.16G146400.1	"K"	"G"	"K"	"P"	"H"	"F"	"P"	"Y"	"F"	"S"
## Glyma.04G093500.1	"K"	"G"	"K"	"P"	"R"	"F"	"A"	"D"	"F"	"D"
## Glyma.02G065300.1	"K"	"G"	"K"	"P"	"H"	"F"	"A"	"D"	"F"	"N"
## Glyma.02G211800.1	"K"	"G"	"K"	"P"	"R"	"F"	"A"	"D"	"F"	"D"
## Glyma.02G254300.1	"K"	"G"	"K"	"P"	"R"	"A"	"A"	"M"	"F"	"N"
## Glyma.02G152800.2	"K"	"G"	"K"	"P"	"H"	"F"	"A"	"D"	"F"	"N"
## Glyma.10G021500.2	"K"	"G"	"K"	"P"	"H"	"F"	"A"	"D"	"F"	"N"
## Glyma.08G059500.6	"K"	"G"	"K"	"P"	"R"	"F"	"S"	"D"	"F"	"N"
## Glyma.07G189800.5	"K"	"G"	"K"	"P"	"R"	"F"	"S"	"D"	"F"	"N"
## Glyma.11G227300.1	"K"	"G"	"K"	"P"	"R"	"A"	"A"	"M"	"F"	"N"
## Glyma.14G179500.1	"K"	"G"	"K"	"P"	"R"	"F"	"A"	"D"	"F"	"D"
## Glyma.14G062100.1	"K"	"G"	"K"	"P"	"R"	"A"	"A"	"M"	"F"	"N"
## Glyma.06G095400.1	"K"	"G"	"K"	"P"	"R"	"F"	"A"	"D"	"F"	"D"
## Glyma.03G209400.1	"K"	"G"	"K"	"P"	"H"	"F"	"A"	"D"	"F"	"N"
## Glyma.19G100200.1	"K"	"G"	"K"	"P"	"H"	"F"	"A"	"D"	"F"	"S"
## Glyma.19G206800.1	"K"	"G"	"K"	"P"	"H"	"F"	"A"	"D"	"F"	"N"

## Glyma.18G030200.1	"K"	"G"	"K"	"P"	"R"	"A"	"A"	"M"	"F"	"N"
##	[,92]	[,93]	[,94]	[,95]	[,96]	[,97]	[,98]	[,99]	[,100]	[,101]
## Glyma.16G050500.1	"L"	"V"	"P"	"Y"	"D"	"W"	"G"	"G"	"F"	"V"
## Glyma.16G146400.1	"L"	"V"	"P"	"S"	"G"	"W"	"G"	"G"	"F"	"V"
## Glyma.04G093500.1	"L"	"M"	"P"	"L"	"N"	"W"	"G"	"A"	"H"	"F"
## Glyma.02G065300.1	"L"	"V"	"P"	"H"	"G"	"W"	"G"	"G"	"F"	"V"
## Glyma.02G211800.1	"L"	"M"	"P"	"P"	"D"	"W"	"G"	"A"	"H"	"F"
## Glyma.02G254300.1	"L"	"I"	"P"	"E"	"D"	"W"	"G"	"G"	"F"	"V"
## Glyma.02G152800.2	"L"	"V"	"P"	"E"	"G"	"W"	"G"	"A"	"Y"	"V"
## Glyma.10G021500.2	"L"	"V"	"P"	"E"	"G"	"W"	"G"	"A"	"Y"	"V"
## Glyma.08G059500.6	"L"	"V"	"P"	"A"	"N"	"W"	"G"	"A"	"D"	"I"
## Glyma.07G189800.5	"L"	"V"	"P"	"A"	"N"	"W"	"G"	"A"	"D"	"I"
## Glyma.11G227300.1	"L"	"I"	"P"	"E"	"D"	"W"	"G"	"G"	"H"	"V"
## Glyma.14G179500.1	"L"	"M"	"P"	"A"	"D"	"W"	"G"	"A"	"H"	"F"
## Glyma.14G062100.1	"L"	"I"	"P"	"E"	"D"	"W"	"G"	"G"	"F"	"V"
## Glyma.06G095400.1	"L"	"M"	"P"	"L"	"N"	"W"	"G"	"A"	"H"	"F"
## Glyma.03G209400.1	"L"	"V"	"P"	"E"	"G"	"W"	"G"	"G"	"Y"	"V"
## Glyma.19G100200.1	"L"	"V"	"P"	"Y"	"D"	"W"	"G"	"G"	"F"	"V"
## Glyma.19G206800.1	"L"	"V"	"P"	"D"	"G"	"W"	"G"	"G"	"Y"	"V"
## Glyma.18G030200.1	"L"	"I"	"P"	"E"	"D"	"W"	"G"	"G"	"H"	"V"
##	[,102]	[,103]	[,104]	[,105]	[,106]	[,107]	[,108]	[,109]		
## Glyma.16G050500.1	"H"	"P"	"W"	"I"	"E"	"A"	"L"	"A"		
## Glyma.16G146400.1	"A"	"P"	"W"	"I"	"E"	"A"	"L"	"A"		
## Glyma.04G093500.1	"T"	"P"	"W"	"V"	"I"	"A"	"L"	"S"		
## Glyma.02G065300.1	"S"	"P"	"W"	"I"	"E"	"A"	"M"	"V"		
## Glyma.02G211800.1	"G"	"P"	"W"	"A"	"S"	"A"	"L"	"A"		
## Glyma.02G254300.1	"T"	"P"	"W"	"V"	"R"	"V"	"I"	"S"		
## Glyma.02G152800.2	"G"	"P"	"W"	"I"	"K"	"A"	"M"	"A"		
## Glyma.10G021500.2	"G"	"P"	"W"	"I"	"K"	"A"	"M"	"A"		
## Glyma.08G059500.6	"H"	"S"	"W"	"L"	"V"	"V"	"F"	"A"		
## Glyma.07G189800.5	"H"	"S"	"W"	"L"	"V"	"V"	"F"	"A"		
## Glyma.11G227300.1	"T"	"P"	"W"	"V"	"K"	"E"	"I"	"S"		
## Glyma.14G179500.1	"S"	"P"	"W"	"A"	"S"	"A"	"L"	"S"		
## Glyma.14G062100.1	"T"	"P"	"W"	"V"	"R"	"E"	"I"	"S"		
## Glyma.06G095400.1	"T"	"P"	"W"	"A"	"I"	"A"	"L"	"S"		
## Glyma.03G209400.1	"C"	"P"	"W"	"I"	"A"	"A"	"M"	"A"		
## Glyma.19G100200.1	"H"	"P"	"W"	"V"	"E"	"A"	"L"	"A"		
## Glyma.19G206800.1	"C"	"P"	"W"	"I"	"A"	"A"	"M"	"A"		
## Glyma.18G030200.1	"T"	"P"	"W"	"V"	"K"	"E"	"I"	"S"		
##	[,110]	[,111]	[,112]	[,113]	[,114]	[,115]	[,116]	[,117]		
## Glyma.16G050500.1	"K"	"N"	"K"	"V"	"G"	"L"	"E"	"E"		
## Glyma.16G146400.1	"R"	"S"	"R"	"V"	"D"	"L"	"E"	"E"		
## Glyma.04G093500.1	"Q"	"S"	"Y"	"H"	"S"	"L"	"N"	"K"		
## Glyma.02G065300.1	"K"	"S"	"R"	"V"	"D"	"L"	"E"	"E"		
## Glyma.02G211800.1	"Q"	"A"	"Y"	"P"	"W"	"L"	"E"	"K"		
## Glyma.02G254300.1	"Q"	"Y"	"F"	"D"	"C"	"L"	"K"	"S"		
## Glyma.02G152800.2	"A"	"A"	"Y"	"P"	"W"	"L"	"Q"	"E"		
## Glyma.10G021500.2	"A"	"A"	"Y"	"P"	"W"	"L"	"Q"	"E"		
## Glyma.08G059500.6	"G"	"K"	"Y"	"P"	"W"	"L"	"E"	"E"		
## Glyma.07G189800.5	"E"	"K"	"Y"	"P"	"W"	"L"	"E"	"E"		
## Glyma.11G227300.1	"Q"	"Y"	"F"	"D"	"C"	"L"	"K"	"S"		
## Glyma.14G179500.1	"Q"	"A"	"Y"	"P"	"W"	"L"	"E"	"K"		
## Glyma.14G062100.1	"Q"	"Y"	"F"	"D"	"C"	"L"	"K"	"S"		
## Glyma.06G095400.1	"Q"	"S"	"Y"	"P"	"S"	"L"	"N"	"K"		

## Glyma.03G209400.1	"R"	"A"	"F"	"P"	"C"	"L"	"E"	"E"
## Glyma.19G100200.1	"K"	"S"	"R"	"V"	"G"	"L"	"E"	"E"
## Glyma.19G206800.1	"R"	"A"	"F"	"P"	"C"	"L"	"E"	"E"
## Glyma.18G030200.1	"Q"	"Y"	"F"	"D"	"C"	"L"	"K"	"S"
##	[,118]	[,119]	[,120]	[,121]	[,122]	[,123]	[,124]	[,125]
## Glyma.16G050500.1	"L"	"R"	"L"	"K"	"R"	"M"	"V"	"V"
## Glyma.16G146400.1	"L"	"R"	"L"	"K"	"R"	"M"	"V"	"V"
## Glyma.04G093500.1	"L"	"H"	"L"	"K"	"R"	"M"	"S"	"L"
## Glyma.02G065300.1	"L"	"R"	"L"	"K"	"R"	"M"	"V"	"V"
## Glyma.02G211800.1	"L"	"H"	"L"	"K"	"R"	"M"	"L"	"V"
## Glyma.02G254300.1	"L"	"H"	"F"	"R"	"R"	"M"	"I"	"V"
## Glyma.02G152800.2	"I"	"R"	"L"	"K"	"R"	"M"	"V"	"I"
## Glyma.10G021500.2	"I"	"R"	"L"	"K"	"R"	"M"	"V"	"I"
## Glyma.08G059500.6	"L"	"R"	"L"	"K"	"R"	"M"	"T"	"V"
## Glyma.07G189800.5	"L"	"R"	"L"	"K"	"R"	"M"	"T"	"V"
## Glyma.11G227300.1	"L"	"H"	"F"	"R"	"R"	"M"	"I"	"V"
## Glyma.14G179500.1	"L"	"H"	"L"	"K"	"R"	"M"	"L"	"L"
## Glyma.14G062100.1	"L"	"H"	"F"	"R"	"R"	"M"	"I"	"V"
## Glyma.06G095400.1	"L"	"H"	"L"	"K"	"R"	"M"	"S"	"L"
## Glyma.03G209400.1	"I"	"R"	"L"	"K"	"R"	"M"	"V"	"I"
## Glyma.19G100200.1	"L"	"R"	"L"	"K"	"R"	"M"	"V"	"V"
## Glyma.19G206800.1	"I"	"R"	"L"	"K"	"R"	"M"	"V"	"I"
## Glyma.18G030200.1	"L"	"H"	"F"	"R"	"R"	"M"	"I"	"V"
##	[,126]	[,127]	[,128]	[,129]	[,130]	[,131]	[,132]	[,133]
## Glyma.16G050500.1	"S"	"D"	"E"	"S"	"L"	"E"	"L"	"L"
## Glyma.16G146400.1	"S"	"D"	"E"	"S"	"L"	"E"	"L"	"L"
## Glyma.04G093500.1	"T"	"D"	"H"	"D"	"L"	"T"	"L"	"L"
## Glyma.02G065300.1	"T"	"D"	"K"	"S"	"L"	"E"	"L"	"L"
## Glyma.02G211800.1	"T"	"D"	"A"	"D"	"L"	"A"	"L"	"I"
## Glyma.02G254300.1	"R"	"D"	"S"	"D"	"L"	"Q"	"V"	"L"
## Glyma.02G152800.2	"A"	"D"	"E"	"C"	"L"	"E"	"L"	"I"
## Glyma.10G021500.2	"S"	"D"	"E"	"C"	"L"	"E"	"L"	"I"
## Glyma.08G059500.6	"T"	"D"	"E"	"S"	"L"	"E"	"F"	"L"
## Glyma.07G189800.5	"T"	"D"	"E"	"S"	"L"	"E"	"F"	"L"
## Glyma.11G227300.1	"K"	"D"	"S"	"D"	"L"	"R"	"N"	"L"
## Glyma.14G179500.1	"T"	"D"	"A"	"D"	"L"	"A"	"L"	"I"
## Glyma.14G062100.1	"R"	"D"	"S"	"D"	"L"	"Q"	"V"	"L"
## Glyma.06G095400.1	"T"	"D"	"H"	"D"	"L"	"I"	"L"	"L"
## Glyma.03G209400.1	"T"	"D"	"E"	"S"	"L"	"E"	"L"	"I"
## Glyma.19G100200.1	"S"	"D"	"E"	"S"	"L"	"E"	"L"	"L"
## Glyma.19G206800.1	"T"	"D"	"E"	"S"	"L"	"E"	"L"	"I"
## Glyma.18G030200.1	"K"	"D"	"S"	"D"	"L"	"Q"	"N"	"L"
##	[,134]	[,135]	[,136]	[,137]	[,138]	[,139]	[,140]	[,141]
## Glyma.16G050500.1	"S"	"R"	"S"	"F"	"T"	"H"	"F"	"K"
## Glyma.16G146400.1	"S"	"R"	"S"	"F"	"V"	"N"	"F"	"K"
## Glyma.04G093500.1	"S"	"H"	"S"	"L"	"P"	"S"	"F"	"Q"
## Glyma.02G065300.1	"S"	"R"	"S"	"F"	"M"	"N"	"F"	"K"
## Glyma.02G211800.1	"A"	"D"	"S"	"F"	"A"	"G"	"F"	"R"
## Glyma.02G254300.1	"A"	"R"	"S"	"R"	"G"	"H"	"L"	"Q"
## Glyma.02G152800.2	"A"	"K"	"S"	"F"	"K"	"N"	"F"	"Q"
## Glyma.10G021500.2	"A"	"K"	"S"	"F"	"K"	"N"	"F"	"Q"
## Glyma.08G059500.6	"A"	"L"	"Q"	"F"	"P"	"N"	"F"	"K"
## Glyma.07G189800.5	"A"	"L"	"K"	"F"	"P"	"N"	"F"	"K"
## Glyma.11G227300.1	"A"	"R"	"D"	"R"	"G"	"H"	"L"	"H"

## Glyma.14G179500.1	"A"	"D"	"S"	"F"	"A"	"A"	"F"	"R"
## Glyma.14G062100.1	"A"	"R"	"S"	"R"	"G"	"H"	"L"	"Q"
## Glyma.06G095400.1	"S"	"H"	"S"	"F"	"P"	"S"	"F"	"Q"
## Glyma.03G209400.1	"A"	"K"	"S"	"F"	"K"	"N"	"F"	"K"
## Glyma.19G100200.1	"S"	"R"	"S"	"F"	"T"	"H"	"F"	"K"
## Glyma.19G206800.1	"A"	"K"	"S"	"F"	"K"	"N"	"F"	"K"
## Glyma.18G030200.1	"A"	"R"	"D"	"R"	"G"	"H"	"L"	"H"
##	[,142]	[,143]	[,144]	[,145]	[,146]	[,147]	[,148]	[,149]
## Glyma.16G050500.1	"S"	"L"	"V"	"L"	"V"	"S"	"C"	"E"
## Glyma.16G146400.1	"S"	"L"	"V"	"L"	"V"	"R"	"C"	"E"
## Glyma.04G093500.1	"D"	"L"	"L"	"L"	"T"	"C"	"C"	"E"
## Glyma.02G065300.1	"S"	"L"	"V"	"L"	"V"	"S"	"C"	"E"
## Glyma.02G211800.1	"E"	"L"	"V"	"L"	"V"	"C"	"C"	"E"
## Glyma.02G254300.1	"A"	"L"	"R"	"L"	"E"	"N"	"C"	"S"
## Glyma.02G152800.2	"V"	"L"	"V"	"L"	"T"	"S"	"C"	"E"
## Glyma.10G021500.2	"V"	"L"	"V"	"L"	"T"	"S"	"C"	"E"
## Glyma.08G059500.6	"A"	"L"	"S"	"L"	"L"	"S"	"C"	"D"
## Glyma.07G189800.5	"A"	"L"	"S"	"L"	"L"	"S"	"C"	"D"
## Glyma.11G227300.1	"S"	"L"	"K"	"L"	"D"	"K"	"C"	"S"
## Glyma.14G179500.1	"E"	"L"	"V"	"L"	"V"	"C"	"C"	"E"
## Glyma.14G062100.1	"A"	"L"	"K"	"L"	"D"	"K"	"C"	"S"
## Glyma.06G095400.1	"D"	"L"	"V"	"L"	"T"	"C"	"C"	"E"
## Glyma.03G209400.1	"V"	"L"	"V"	"L"	"T"	"S"	"C"	"E"
## Glyma.19G100200.1	"S"	"L"	"V"	"L"	"V"	"S"	"C"	"E"
## Glyma.19G206800.1	"V"	"L"	"V"	"L"	"T"	"S"	"C"	"E"
## Glyma.18G030200.1	"A"	"L"	"K"	"L"	"D"	"K"	"C"	"S"
##	[,150]	[,151]	[,152]	[,153]	[,154]	[,155]	[,156]	[,157]
## Glyma.16G050500.1	"G"	"F"	"S"	"T"	"D"	"G"	"L"	"A"
## Glyma.16G146400.1	"G"	"F"	"T"	"T"	"E"	"G"	"L"	"A"
## Glyma.04G093500.1	"G"	"F"	"G"	"T"	"T"	"A"	"L"	"A"
## Glyma.02G065300.1	"G"	"F"	"T"	"T"	"D"	"G"	"L"	"A"
## Glyma.02G211800.1	"G"	"F"	"G"	"T"	"P"	"G"	"L"	"A"
## Glyma.02G254300.1	"G"	"F"	"S"	"T"	"D"	"G"	"L"	"Y"
## Glyma.02G152800.2	"G"	"F"	"T"	"T"	"D"	"G"	"L"	"A"
## Glyma.10G021500.2	"G"	"F"	"T"	"T"	"D"	"G"	"L"	"A"
## Glyma.08G059500.6	"G"	"F"	"S"	"T"	"D"	"G"	"L"	"A"
## Glyma.07G189800.5	"G"	"F"	"S"	"T"	"D"	"G"	"L"	"A"
## Glyma.11G227300.1	"G"	"F"	"T"	"T"	"D"	"G"	"L"	"F"
## Glyma.14G179500.1	"G"	"F"	"G"	"T"	"P"	"G"	"L"	"A"
## Glyma.14G062100.1	"G"	"F"	"S"	"T"	"D"	"G"	"L"	"Y"
## Glyma.06G095400.1	"G"	"F"	"G"	"T"	"T"	"G"	"L"	"A"
## Glyma.03G209400.1	"G"	"F"	"T"	"A"	"D"	"G"	"L"	"T"
## Glyma.19G100200.1	"G"	"F"	"S"	"T"	"D"	"G"	"L"	"A"
## Glyma.19G206800.1	"G"	"F"	"T"	"T"	"D"	"G"	"L"	"A"
## Glyma.18G030200.1	"G"	"F"	"T"	"T"	"D"	"G"	"L"	"F"
##	[,158]	[,159]	[,160]	[,161]	[,162]	[,163]	[,164]	[,165]
## Glyma.16G050500.1	"A"	"L"	"A"	"A"	"N"	"C"	"R"	"F"
## Glyma.16G146400.1	"A"	"I"	"A"	"A"	"N"	"C"	"R"	"F"
## Glyma.04G093500.1	"A"	"L"	"A"	"S"	"N"	"C"	"R"	"L"
## Glyma.02G065300.1	"A"	"I"	"A"	"A"	"N"	"C"	"R"	"F"
## Glyma.02G211800.1	"A"	"V"	"A"	"S"	"K"	"C"	"R"	"L"
## Glyma.02G254300.1	"Y"	"I"	"G"	"R"	"Y"	"C"	"R"	"N"
## Glyma.02G152800.2	"A"	"I"	"A"	"A"	"N"	"C"	"R"	"N"
## Glyma.10G021500.2	"A"	"I"	"A"	"A"	"N"	"C"	"R"	"N"

## Glyma.08G059500.6	"S"	"I"	"A"	"T"	"N"	"C"	"K"	"N"
## Glyma.07G189800.5	"S"	"I"	"A"	"T"	"N"	"C"	"K"	"N"
## Glyma.11G227300.1	"H"	"I"	"G"	"R"	"F"	"C"	"K"	"S"
## Glyma.14G179500.1	"V"	"V"	"V"	"S"	"K"	"C"	"R"	"L"
## Glyma.14G062100.1	"Y"	"I"	"G"	"R"	"Y"	"C"	"R"	"N"
## Glyma.06G095400.1	"A"	"L"	"T"	"S"	"N"	"C"	"R"	"L"
## Glyma.03G209400.1	"A"	"I"	"A"	"S"	"N"	"C"	"R"	"N"
## Glyma.19G100200.1	"A"	"I"	"A"	"A"	"N"	"C"	"R"	"F"
## Glyma.19G206800.1	"A"	"I"	"A"	"A"	"N"	"C"	"R"	"N"
## Glyma.18G030200.1	"H"	"I"	"G"	"R"	"F"	"C"	"K"	"S"
##	[,166]	[,167]	[,168]	[,169]	[,170]	[,171]	[,172]	[,173]
## Glyma.16G050500.1	"L"	"R"	"E"	"L"	"D"	"L"	"Q"	"E"
## Glyma.16G146400.1	"L"	"K"	"D"	"L"	"D"	"L"	"H"	"E"
## Glyma.04G093500.1	"L"	"R"	"V"	"L"	"E"	"L"	"V"	"E"
## Glyma.02G065300.1	"L"	"K"	"E"	"L"	"D"	"L"	"Q"	"E"
## Glyma.02G211800.1	"L"	"R"	"V"	"L"	"E"	"L"	"V"	"E"
## Glyma.02G254300.1	"L"	"R"	"V"	"L"	"F"	"L"	"E"	"E"
## Glyma.02G152800.2	"L"	"R"	"E"	"L"	"E"	"L"	"R"	"E"
## Glyma.10G021500.2	"L"	"R"	"E"	"L"	"E"	"L"	"R"	"E"
## Glyma.08G059500.6	"L"	"T"	"E"	"L"	"D"	"I"	"Q"	"E"
## Glyma.07G189800.5	"L"	"T"	"E"	"L"	"D"	"I"	"Q"	"E"
## Glyma.11G227300.1	"L"	"R"	"V"	"L"	"F"	"L"	"E"	"E"
## Glyma.14G179500.1	"L"	"R"	"V"	"L"	"E"	"L"	"V"	"E"
## Glyma.14G062100.1	"L"	"R"	"V"	"L"	"F"	"L"	"E"	"E"
## Glyma.06G095400.1	"L"	"R"	"V"	"L"	"E"	"L"	"V"	"E"
## Glyma.03G209400.1	"L"	"R"	"E"	"L"	"D"	"L"	"Q"	"E"
## Glyma.19G100200.1	"L"	"R"	"E"	"L"	"D"	"L"	"Q"	"E"
## Glyma.19G206800.1	"L"	"R"	"E"	"L"	"D"	"L"	"Q"	"E"
## Glyma.18G030200.1	"L"	"R"	"V"	"L"	"F"	"L"	"E"	"E"
##	[,174]	[,175]	[,176]	[,177]	[,178]	[,179]	[,180]	[,181]
## Glyma.16G050500.1	"N"	"E"	"V"	"E"	"D"	"H"	"K"	"G"
## Glyma.16G146400.1	"N"	"V"	"V"	"T"	"D"	"L"	"K"	"G"
## Glyma.04G093500.1	"C"	"V"	"V"	"G"	"D"	"E"	"E"	"L"
## Glyma.02G065300.1	"N"	"E"	"V"	"D"	"D"	"H"	"R"	"G"
## Glyma.02G211800.1	"S"	"V"	"V"	"D"	"D"	"E"	"E"	"V"
## Glyma.02G254300.1	"S"	"S"	"L"	"V"	"E"	"K"	"D"	"G"
## Glyma.02G152800.2	"S"	"E"	"V"	"D"	"D"	"I"	"C"	"G"
## Glyma.10G021500.2	"S"	"E"	"V"	"D"	"D"	"I"	"C"	"G"
## Glyma.08G059500.6	"N"	"G"	"I"	"E"	"D"	"K"	"S"	"G"
## Glyma.07G189800.5	"N"	"G"	"I"	"E"	"D"	"K"	"S"	"G"
## Glyma.11G227300.1	"S"	"S"	"I"	"V"	"E"	"K"	"D"	"G"
## Glyma.14G179500.1	"S"	"E"	"V"	"D"	"D"	"E"	"E"	"V"
## Glyma.14G062100.1	"S"	"S"	"L"	"V"	"E"	"N"	"D"	"G"
## Glyma.06G095400.1	"C"	"V"	"V"	"G"	"D"	"E"	"E"	"M"
## Glyma.03G209400.1	"S"	"E"	"V"	"E"	"D"	"L"	"S"	"G"
## Glyma.19G100200.1	"N"	"E"	"V"	"E"	"D"	"H"	"K"	"G"
## Glyma.19G206800.1	"S"	"E"	"V"	"E"	"D"	"L"	"S"	"G"
## Glyma.18G030200.1	"S"	"S"	"I"	"L"	"E"	"K"	"D"	"G"
##	[,182]	[,183]	[,184]	[,185]	[,186]	[,187]	[,188]	[,189]
## Glyma.16G050500.1	"Q"	"W"	"L"	"S"	"C"	"F"	"P"	"D"
## Glyma.16G146400.1	"Q"	"W"	"L"	"S"	"C"	"F"	"P"	"D"
## Glyma.04G093500.1	"D"	"W"	"I"	"S"	"C"	"F"	"P"	"E"
## Glyma.02G065300.1	"Q"	"W"	"L"	"S"	"C"	"F"	"P"	"D"
## Glyma.02G211800.1	"D"	"W"	"I"	"S"	"C"	"F"	"P"	"E"

## Glyma.02G254300.1	"D"	"W"	"L"	"H"	"E"	"L"	"A"	"L"
## Glyma.02G152800.2	"H"	"W"	"L"	"S"	"H"	"F"	"P"	"D"
## Glyma.10G021500.2	"H"	"W"	"L"	"S"	"H"	"F"	"P"	"D"
## Glyma.08G059500.6	"N"	"W"	"L"	"G"	"C"	"F"	"P"	"D"
## Glyma.07G189800.5	"N"	"W"	"L"	"S"	"C"	"F"	"P"	"D"
## Glyma.11G227300.1	"E"	"W"	"L"	"H"	"E"	"L"	"A"	"L"
## Glyma.14G179500.1	"D"	"W"	"I"	"S"	"C"	"F"	"P"	"E"
## Glyma.14G062100.1	"D"	"W"	"L"	"H"	"E"	"L"	"A"	"L"
## Glyma.06G095400.1	"D"	"W"	"I"	"S"	"C"	"F"	"P"	"E"
## Glyma.03G209400.1	"H"	"W"	"L"	"S"	"H"	"F"	"P"	"D"
## Glyma.19G100200.1	"Q"	"W"	"L"	"S"	"C"	"F"	"P"	"D"
## Glyma.19G206800.1	"H"	"W"	"L"	"S"	"H"	"F"	"P"	"D"
## Glyma.18G030200.1	"E"	"W"	"L"	"H"	"E"	"L"	"A"	"L"
##	[,190]	[,191]	[,192]	[,193]	[,194]	[,195]	[,196]	[,197]
## Glyma.16G050500.1	"N"	"C"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.16G146400.1	"C"	"C"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.04G093500.1	"I"	"Q"	"T"	"Y"	"L"	"E"	"S"	"L"
## Glyma.02G065300.1	"C"	"C"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.02G211800.1	"T"	"Q"	"T"	"N"	"M"	"E"	"S"	"L"
## Glyma.02G254300.1	"N"	"N"	"T"	"V"	"L"	"E"	"T"	"L"
## Glyma.02G152800.2	"S"	"Y"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.10G021500.2	"S"	"Y"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.08G059500.6	"S"	"F"	"T"	"S"	"L"	"E"	"V"	"L"
## Glyma.07G189800.5	"S"	"F"	"T"	"S"	"L"	"E"	"V"	"L"
## Glyma.11G227300.1	"N"	"N"	"T"	"V"	"L"	"E"	"T"	"L"
## Glyma.14G179500.1	"S"	"Q"	"T"	"N"	"L"	"E"	"S"	"L"
## Glyma.14G062100.1	"N"	"N"	"T"	"V"	"L"	"E"	"T"	"L"
## Glyma.06G095400.1	"S"	"Q"	"T"	"H"	"L"	"E"	"S"	"L"
## Glyma.03G209400.1	"S"	"Y"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.19G100200.1	"N"	"C"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.19G206800.1	"S"	"Y"	"T"	"S"	"L"	"V"	"S"	"L"
## Glyma.18G030200.1	"N"	"N"	"T"	"V"	"L"	"E"	"T"	"L"
##	[,198]	[,199]	[,200]	[,201]	[,202]	[,203]	[,204]	[,205]
## Glyma.16G050500.1	"N"	"F"	"A"	"C"	"L"	"K"	"G"	"E"
## Glyma.16G146400.1	"N"	"F"	"A"	"C"	"L"	"K"	"G"	"Q"
## Glyma.04G093500.1	"V"	"F"	"D"	"C"	"V"	"E"	"C"	"P"
## Glyma.02G065300.1	"N"	"F"	"A"	"C"	"L"	"K"	"G"	"Q"
## Glyma.02G211800.1	"V"	"F"	"D"	"C"	"V"	"E"	"V"	"P"
## Glyma.02G254300.1	"N"	"F"	"Y"	"V"	"T"	"D"	"A"	"N"
## Glyma.02G152800.2	"N"	"I"	"S"	"C"	"L"	"G"	"N"	"E"
## Glyma.10G021500.2	"N"	"I"	"S"	"C"	"L"	"G"	"N"	"E"
## Glyma.08G059500.6	"N"	"F"	"A"	"N"	"L"	"H"	"N"	"D"
## Glyma.07G189800.5	"N"	"F"	"A"	"N"	"L"	"H"	"N"	"D"
## Glyma.11G227300.1	"N"	"F"	"Y"	"L"	"T"	"D"	"A"	"V"
## Glyma.14G179500.1	"V"	"F"	"D"	"C"	"V"	"D"	"V"	"P"
## Glyma.14G062100.1	"N"	"F"	"Y"	"L"	"T"	"D"	"A"	"N"
## Glyma.06G095400.1	"V"	"F"	"D"	"C"	"V"	"E"	"C"	"P"
## Glyma.03G209400.1	"N"	"I"	"S"	"C"	"L"	"N"	"H"	"E"
## Glyma.19G100200.1	"N"	"F"	"A"	"C"	"L"	"K"	"G"	"E"
## Glyma.19G206800.1	"N"	"I"	"S"	"C"	"L"	"N"	"N"	"E"
## Glyma.18G030200.1	"N"	"F"	"Y"	"L"	"T"	"D"	"A"	"V"
##	[,206]	[,207]	[,208]	[,209]	[,210]	[,211]	[,212]	[,213]
## Glyma.16G050500.1	"V"	"S"	"L"	"G"	"A"	"L"	"E"	"R"
## Glyma.16G146400.1	"I"	"N"	"A"	"G"	"D"	"L"	"E"	"R"



## Glyma.04G093500.1	"I"	"N"	"L"	"E"	"A"	"L"	"E"	"R"
## Glyma.02G065300.1	"I"	"N"	"L"	"G"	"A"	"L"	"E"	"R"
## Glyma.02G211800.1	"I"	"N"	"F"	"E"	"A"	"L"	"E"	"G"
## Glyma.02G254300.1	"V"	"R"	"I"	"Q"	"D"	"L"	"E"	"L"
## Glyma.02G152800.2	"V"	"N"	"L"	"S"	"A"	"L"	"E"	"R"
## Glyma.10G021500.2	"V"	"N"	"L"	"S"	"A"	"L"	"E"	"R"
## Glyma.08G059500.6	"V"	"N"	"F"	"D"	"A"	"L"	"E"	"K"
## Glyma.07G189800.5	"V"	"N"	"F"	"D"	"A"	"L"	"E"	"K"
## Glyma.11G227300.1	"V"	"K"	"I"	"Q"	"D"	"L"	"E"	"L"
## Glyma.14G179500.1	"I"	"N"	"F"	"E"	"A"	"L"	"E"	"G"
## Glyma.14G062100.1	"V"	"R"	"I"	"Q"	"D"	"L"	"E"	"L"
## Glyma.06G095400.1	"V"	"N"	"F"	"D"	"A"	"L"	"E"	"R"
## Glyma.03G209400.1	"V"	"S"	"L"	"S"	"A"	"L"	"E"	"R"
## Glyma.19G100200.1	"V"	"S"	"L"	"G"	"A"	"L"	"E"	"R"
## Glyma.19G206800.1	"V"	"S"	"L"	"S"	"A"	"L"	"E"	"R"
## Glyma.18G030200.1	"V"	"K"	"I"	"E"	"D"	"L"	"E"	"L"
##	[,214]	[,215]	[,216]	[,217]	[,218]	[,219]	[,220]	[,221]
## Glyma.16G050500.1	"L"	"V"	"A"	"R"	"S"	"P"	"Y"	"L"
## Glyma.16G146400.1	"L"	"V"	"A"	"R"	"S"	"P"	"N"	"L"
## Glyma.04G093500.1	"L"	"V"	"A"	"R"	"S"	"P"	"S"	"L"
## Glyma.02G065300.1	"L"	"V"	"A"	"R"	"S"	"P"	"N"	"L"
## Glyma.02G211800.1	"L"	"V"	"A"	"R"	"S"	"P"	"R"	"L"
## Glyma.02G254300.1	"I"	"A"	"R"	"N"	"C"	"P"	"N"	"L"
## Glyma.02G152800.2	"L"	"V"	"S"	"R"	"C"	"P"	"N"	"L"
## Glyma.10G021500.2	"L"	"V"	"S"	"R"	"C"	"P"	"N"	"L"
## Glyma.08G059500.6	"L"	"V"	"C"	"R"	"C"	"K"	"S"	"L"
## Glyma.07G189800.5	"L"	"V"	"S"	"R"	"C"	"K"	"S"	"L"
## Glyma.11G227300.1	"L"	"A"	"K"	"N"	"C"	"P"	"N"	"L"
## Glyma.14G179500.1	"L"	"V"	"A"	"R"	"S"	"P"	"R"	"L"
## Glyma.14G062100.1	"I"	"A"	"R"	"N"	"C"	"P"	"N"	"L"
## Glyma.06G095400.1	"L"	"V"	"A"	"R"	"S"	"P"	"L"	"L"
## Glyma.03G209400.1	"L"	"L"	"G"	"R"	"C"	"R"	"N"	"L"
## Glyma.19G100200.1	"F"	"V"	"A"	"R"	"S"	"P"	"N"	"L"
## Glyma.19G206800.1	"L"	"L"	"G"	"R"	"C"	"P"	"N"	"L"
## Glyma.18G030200.1	"L"	"A"	"K"	"N"	"C"	"P"	"N"	"L"
##	[,222]	[,223]	[,224]	[,225]	[,226]	[,227]	[,228]	[,229]
## Glyma.16G050500.1	"K"	"S"	"L"	"K"	"L"	"N"	"R"	"S"
## Glyma.16G146400.1	"K"	"S"	"L"	"R"	"L"	"N"	"H"	"T"
## Glyma.04G093500.1	"R"	"K"	"L"	"R"	"L"	"N"	"R"	"Y"
## Glyma.02G065300.1	"K"	"S"	"L"	"R"	"L"	"N"	"H"	"T"
## Glyma.02G211800.1	"K"	"K"	"L"	"R"	"L"	"N"	"Q"	"F"
## Glyma.02G254300.1	"N"	"S"	"V"	"K"	"I"	"T"	"D"	"C"
## Glyma.02G152800.2	"Q"	"T"	"L"	"R"	"L"	"N"	"R"	"A"
## Glyma.10G021500.2	"Q"	"T"	"L"	"R"	"L"	"N"	"R"	"A"
## Glyma.08G059500.6	"K"	"T"	"L"	"K"	"V"	"N"	"K"	"S"
## Glyma.07G189800.5	"K"	"T"	"L"	"K"	"V"	"N"	"K"	"S"
## Glyma.11G227300.1	"V"	"S"	"V"	"K"	"L"	"T"	"D"	"S"
## Glyma.14G179500.1	"K"	"K"	"L"	"R"	"L"	"N"	"R"	"Y"
## Glyma.14G062100.1	"N"	"S"	"V"	"K"	"I"	"T"	"D"	"C"
## Glyma.06G095400.1	"R"	"K"	"L"	"R"	"L"	"N"	"R"	"Y"
## Glyma.03G209400.1	"R"	"T"	"L"	"R"	"L"	"N"	"R"	"A"
## Glyma.19G100200.1	"K"	"S"	"L"	"K"	"L"	"N"	"R"	"S"
## Glyma.19G206800.1	"R"	"T"	"L"	"R"	"L"	"N"	"R"	"A"
## Glyma.18G030200.1	"V"	"S"	"V"	"K"	"L"	"T"	"D"	"C"

	[,230]	[,231]	[,232]	[,233]	[,234]	[,235]	[,236]	[,237]
## Glyma.16G050500.1	"V"	"P"	"F"	"D"	"L"	"Q"	"R"	"I"
## Glyma.16G146400.1	"V"	"P"	"L"	"S"	"L"	"Q"	"R"	"I"
## Glyma.04G093500.1	"V"	"S"	"M"	"S"	"L"	"H"	"R"	"L"
## Glyma.02G065300.1	"V"	"P"	"L"	"N"	"L"	"Q"	"R"	"I"
## Glyma.02G211800.1	"V"	"S"	"M"	"A"	"L"	"Y"	"R"	"L"
## Glyma.02G254300.1	"E"	"V"	"L"	"D"	"L"	"V"	"N"	"F"
## Glyma.02G152800.2	"V"	"P"	"L"	"D"	"L"	"A"	"T"	"L"
## Glyma.10G021500.2	"V"	"P"	"L"	"D"	"L"	"A"	"N"	"L"
## Glyma.08G059500.6	"V"	"T"	"L"	"E"	"L"	"Q"	"R"	"L"
## Glyma.07G189800.5	"V"	"T"	"L"	"E"	"L"	"Q"	"R"	"L"
## Glyma.11G227300.1	"E"	"I"	"L"	"D"	"L"	"V"	"N"	"F"
## Glyma.14G179500.1	"V"	"S"	"M"	"A"	"L"	"Y"	"R"	"L"
## Glyma.14G062100.1	"E"	"V"	"L"	"D"	"L"	"V"	"N"	"F"
## Glyma.06G095400.1	"V"	"S"	"M"	"S"	"L"	"H"	"R"	"L"
## Glyma.03G209400.1	"V"	"P"	"L"	"D"	"L"	"P"	"N"	"L"
## Glyma.19G100200.1	"V"	"P"	"V"	"D"	"L"	"Q"	"R"	"I"
## Glyma.19G206800.1	"V"	"P"	"L"	"D"	"L"	"P"	"N"	"L"
## Glyma.18G030200.1	"E"	"I"	"L"	"D"	"L"	"V"	"N"	"F"
	[,238]	[,239]	[,240]	[,241]	[,242]	[,243]	[,244]	[,245]
## Glyma.16G050500.1	"M"	"M"	"R"	"A"	"P"	"Q"	"L"	"S"
## Glyma.16G146400.1	"L"	"M"	"Q"	"A"	"P"	"Q"	"L"	"V"
## Glyma.04G093500.1	"M"	"H"	"R"	"A"	"P"	"Q"	"L"	"T"
## Glyma.02G065300.1	"L"	"R"	"R"	"A"	"P"	"Q"	"I"	"V"
## Glyma.02G211800.1	"L"	"L"	"R"	"A"	"P"	"Q"	"L"	"T"
## Glyma.02G254300.1	"F"	"Q"	"Y"	"A"	"S"	"A"	"L"	"E"
## Glyma.02G152800.2	"L"	"R"	"G"	"A"	"P"	"Q"	"L"	"V"
## Glyma.10G021500.2	"L"	"R"	"G"	"A"	"P"	"Q"	"L"	"V"
## Glyma.08G059500.6	"L"	"V"	"H"	"V"	"P"	"Q"	"L"	"G"
## Glyma.07G189800.5	"I"	"V"	"H"	"V"	"P"	"Q"	"L"	"G"
## Glyma.11G227300.1	"F"	"K"	"H"	"A"	"S"	"A"	"L"	"E"
## Glyma.14G179500.1	"L"	"L"	"R"	"A"	"P"	"Q"	"L"	"T"
## Glyma.14G062100.1	"F"	"R"	"Y"	"A"	"S"	"A"	"L"	"E"
## Glyma.06G095400.1	"M"	"H"	"R"	"A"	"P"	"Q"	"L"	"T"
## Glyma.03G209400.1	"L"	"L"	"R"	"C"	"P"	"Q"	"L"	"V"
## Glyma.19G100200.1	"M"	"M"	"R"	"A"	"P"	"Q"	"L"	"S"
## Glyma.19G206800.1	"L"	"L"	"Q"	"C"	"P"	"Q"	"L"	"V"
## Glyma.18G030200.1	"F"	"K"	"H"	"A"	"S"	"A"	"L"	"E"
	[,246]	[,247]	[,248]	[,249]	[,250]	[,251]	[,252]	[,253]
## Glyma.16G050500.1	"D"	"L"	"G"	"I"	"G"	"S"	"F"	"V"
## Glyma.16G146400.1	"D"	"L"	"G"	"I"	"G"	"S"	"F"	"V"
## Glyma.04G093500.1	"H"	"L"	"G"	"T"	"G"	"S"	"F"	"S"
## Glyma.02G065300.1	"D"	"L"	"G"	"I"	"G"	"S"	"F"	"I"
## Glyma.02G211800.1	"H"	"L"	"G"	"T"	"G"	"S"	"F"	"S"
## Glyma.02G254300.1	"E"	"F"	"S"	"G"	"G"	"S"	"Y"	"N"
## Glyma.02G152800.2	"E"	"L"	"G"	"T"	"G"	"A"	"Y"	"T"
## Glyma.10G021500.2	"E"	"L"	"G"	"T"	"G"	"T"	"Y"	"T"
## Glyma.08G059500.6	"E"	"L"	"G"	"T"	"G"	"S"	"F"	"S"
## Glyma.07G189800.5	"E"	"L"	"G"	"T"	"G"	"S"	"F"	"S"
## Glyma.11G227300.1	"E"	"F"	"C"	"G"	"G"	"T"	"Y"	"N"
## Glyma.14G179500.1	"H"	"L"	"G"	"T"	"G"	"S"	"F"	"S"
## Glyma.14G062100.1	"E"	"F"	"C"	"G"	"G"	"S"	"Y"	"N"
## Glyma.06G095400.1	"H"	"L"	"G"	"T"	"G"	"S"	"F"	"S"
## Glyma.03G209400.1	"E"	"L"	"G"	"T"	"G"	"V"	"Y"	"S"

## Glyma.19G100200.1	"D"	"L"	"G"	"I"	"G"	"S"	"L"	"V"
## Glyma.19G206800.1	"E"	"L"	"G"	"T"	"G"	"V"	"Y"	"S"
## Glyma.18G030200.1	"E"	"F"	"C"	"G"	"G"	"T"	"Y"	"N"
##	[,254]	[,255]	[,256]	[,257]	[,258]	[,259]	[,260]	[,261]
## Glyma.16G050500.1	"H"	"D"	"P"	"E"	"S"	"E"	"A"	"Y"
## Glyma.16G146400.1	"F"	"D"	"P"	"R"	"S"	"E"	"V"	"Y"
## Glyma.04G093500.1	"A"	"S"	"E"	"L"	"D"	"_"	"Q"	"E"
## Glyma.02G065300.1	"P"	"D"	"P"	"N"	"S"	"N"	"V"	"F"
## Glyma.02G211800.1	"A"	"T"	"E"	"A"	"G"	"V"	"Q"	"E"
## Glyma.02G254300.1	"E"	"E"	"S"	"E"	"K"	"Y"	"S"	"A"
## Glyma.02G152800.2	"T"	"E"	"M"	"R"	"P"	"E"	"V"	"F"
## Glyma.10G021500.2	"T"	"E"	"M"	"R"	"P"	"E"	"V"	"F"
## Glyma.08G059500.6	"Q"	"E"	"L"	"T"	"A"	"Q"	"Q"	"C"
## Glyma.07G189800.5	"Q"	"E"	"L"	"T"	"S"	"Q"	"Q"	"C"
## Glyma.11G227300.1	"E"	"E"	"P"	"E"	"K"	"Y"	"S"	"A"
## Glyma.14G179500.1	"A"	"T"	"E"	"A"	"G"	"A"	"Q"	"E"
## Glyma.14G062100.1	"E"	"E"	"S"	"E"	"K"	"Y"	"S"	"A"
## Glyma.06G095400.1	"A"	"S"	"E"	"L"	"D"	"_"	"Q"	"E"
## Glyma.03G209400.1	"T"	"E"	"M"	"R"	"P"	"E"	"V"	"F"
## Glyma.19G100200.1	"H"	"D"	"P"	"E"	"S"	"E"	"A"	"Y"
## Glyma.19G206800.1	"T"	"E"	"M"	"R"	"P"	"E"	"V"	"F"
## Glyma.18G030200.1	"E"	"E"	"P"	"E"	"R"	"Y"	"S"	"A"
##	[,262]	[,263]	[,264]	[,265]	[,266]	[,267]	[,268]	[,269]
## Glyma.16G050500.1	"I"	"K"	"L"	"K"	"N"	"T"	"I"	"L"
## Glyma.16G146400.1	"N"	"N"	"M"	"K"	"N"	"A"	"I"	"L"
## Glyma.04G093500.1	"L"	"D"	"F"	"A"	"S"	"A"	"F"	"A"
## Glyma.02G065300.1	"I"	"K"	"L"	"M"	"N"	"T"	"I"	"I"
## Glyma.02G211800.1	"P"	"D"	"Y"	"A"	"A"	"A"	"F"	"E"
## Glyma.02G254300.1	"M"	"S"	"L"	"P"	"A"	"K"	"L"	"S"
## Glyma.02G152800.2	"T"	"N"	"L"	"A"	"E"	"A"	"F"	"S"
## Glyma.10G021500.2	"T"	"N"	"L"	"A"	"E"	"A"	"F"	"S"
## Glyma.08G059500.6	"S"	"D"	"L"	"E"	"S"	"A"	"F"	"K"
## Glyma.07G189800.5	"S"	"D"	"L"	"E"	"S"	"A"	"L"	"K"
## Glyma.11G227300.1	"I"	"S"	"L"	"P"	"A"	"K"	"L"	"C"
## Glyma.14G179500.1	"P"	"D"	"Y"	"A"	"A"	"A"	"F"	"E"
## Glyma.14G062100.1	"I"	"S"	"L"	"P"	"A"	"K"	"L"	"S"
## Glyma.06G095400.1	"L"	"D"	"F"	"A"	"S"	"A"	"F"	"A"
## Glyma.03G209400.1	"S"	"N"	"L"	"E"	"A"	"A"	"F"	"S"
## Glyma.19G100200.1	"I"	"K"	"L"	"K"	"N"	"T"	"I"	"L"
## Glyma.19G206800.1	"S"	"N"	"L"	"E"	"A"	"A"	"F"	"S"
## Glyma.18G030200.1	"I"	"S"	"L"	"P"	"A"	"K"	"L"	"C"
##	[,270]	[,271]	[,272]	[,273]	[,274]	[,275]	[,276]	[,277]
## Glyma.16G050500.1	"K"	"S"	"G"	"F"	"L"	"E"	"V"	"A"
## Glyma.16G146400.1	"K"	"S"	"G"	"F"	"F"	"W"	"V"	"Y"
## Glyma.04G093500.1	"A"	"S"	"G"	"F"	"R"	"E"	"I"	"W"
## Glyma.02G065300.1	"E"	"S"	"G"	"F"	"F"	"E"	"V"	"T"
## Glyma.02G211800.1	"A"	"S"	"G"	"F"	"R"	"E"	"I"	"W"
## Glyma.02G254300.1	"R"	"L"	"G"	"L"	"T"	"Y"	"I"	"T"
## Glyma.02G152800.2	"G"	"S"	"G"	"F"	"W"	"D"	"V"	"L"
## Glyma.10G021500.2	"G"	"S"	"G"	"F"	"W"	"D"	"V"	"L"
## Glyma.08G059500.6	"N"	"S"	"G"	"L"	"W"	"V"	"A"	"T"
## Glyma.07G189800.5	"N"	"S"	"G"	"L"	"W"	"V"	"A"	"T"
## Glyma.11G227300.1	"R"	"L"	"G"	"L"	"T"	"Y"	"I"	"G"
## Glyma.14G179500.1	"A"	"S"	"G"	"F"	"R"	"E"	"I"	"W"

## Glyma.14G062100.1	"R"	"L"	"G"	"L"	"T"	"Y"	"I"	"T"
## Glyma.06G095400.1	"S"	"S"	"G"	"F"	"R"	"E"	"F"	"W"
## Glyma.03G209400.1	"G"	"S"	"G"	"F"	"W"	"D"	"V"	"L"
## Glyma.19G100200.1	"K"	"S"	"G"	"F"	"L"	"E"	"V"	"A"
## Glyma.19G206800.1	"G"	"S"	"G"	"F"	"W"	"D"	"V"	"L"
## Glyma.18G030200.1	"R"	"L"	"G"	"L"	"T"	"Y"	"I"	"G"
##	[,278]	[,279]	[,280]	[,281]	[,282]	[,283]	[,284]	[,285]
## Glyma.16G050500.1	"P"	"H"	"C"	"L"	"A"	"A"	"I"	"Y"
## Glyma.16G146400.1	"P"	"H"	"C"	"L"	"S"	"A"	"L"	"Y"
## Glyma.04G093500.1	"P"	"D"	"Y"	"L"	"P"	"A"	"I"	"Y"
## Glyma.02G065300.1	"P"	"R"	"C"	"L"	"P"	"A"	"I"	"Y"
## Glyma.02G211800.1	"A"	"D"	"Y"	"L"	"P"	"A"	"I"	"Y"
## Glyma.02G254300.1	"K"	"N"	"E"	"M"	"P"	"I"	"V"	"F"
## Glyma.02G152800.2	"P"	"S"	"Y"	"L"	"P"	"A"	"V"	"Y"
## Glyma.10G021500.2	"P"	"S"	"Y"	"L"	"P"	"A"	"V"	"Y"
## Glyma.08G059500.6	"A"	"Q"	"Y"	"L"	"P"	"V"	"L"	"Y"
## Glyma.07G189800.5	"A"	"Q"	"Y"	"L"	"P"	"V"	"L"	"Y"
## Glyma.11G227300.1	"K"	"N"	"E"	"L"	"P"	"I"	"V"	"F"
## Glyma.14G179500.1	"A"	"D"	"Y"	"L"	"P"	"A"	"I"	"Y"
## Glyma.14G062100.1	"K"	"N"	"E"	"M"	"P"	"M"	"V"	"F"
## Glyma.06G095400.1	"A"	"D"	"Y"	"L"	"P"	"A"	"I"	"Y"
## Glyma.03G209400.1	"P"	"S"	"Y"	"L"	"P"	"A"	"V"	"Y"
## Glyma.19G100200.1	"P"	"H"	"C"	"L"	"A"	"A"	"I"	"Y"
## Glyma.19G206800.1	"P"	"S"	"Y"	"L"	"P"	"A"	"V"	"Y"
## Glyma.18G030200.1	"K"	"N"	"E"	"L"	"P"	"I"	"V"	"F"
##	[,286]	[,287]	[,288]	[,289]	[,290]	[,291]	[,292]	[,293]
## Glyma.16G050500.1	"P"	"I"	"C"	"P"	"N"	"L"	"T"	"S"
## Glyma.16G146400.1	"P"	"V"	"C"	"M"	"N"	"L"	"T"	"T"
## Glyma.04G093500.1	"P"	"A"	"C"	"A"	"N"	"L"	"I"	"S"
## Glyma.02G065300.1	"P"	"V"	"C"	"M"	"N"	"L"	"T"	"A"
## Glyma.02G211800.1	"P"	"V"	"C"	"A"	"N"	"L"	"T"	"S"
## Glyma.02G254300.1	"P"	"Y"	"A"	"A"	"L"	"L"	"K"	"K"
## Glyma.02G152800.2	"P"	"I"	"C"	"S"	"N"	"L"	"T"	"S"
## Glyma.10G021500.2	"P"	"I"	"C"	"S"	"N"	"L"	"T"	"S"
## Glyma.08G059500.6	"S"	"A"	"C"	"T"	"N"	"L"	"T"	"F"
## Glyma.07G189800.5	"S"	"A"	"C"	"T"	"N"	"L"	"T"	"F"
## Glyma.11G227300.1	"M"	"F"	"A"	"A"	"V"	"L"	"K"	"K"
## Glyma.14G179500.1	"P"	"V"	"C"	"T"	"N"	"L"	"T"	"S"
## Glyma.14G062100.1	"P"	"Y"	"A"	"A"	"L"	"L"	"K"	"K"
## Glyma.06G095400.1	"P"	"A"	"C"	"A"	"N"	"L"	"I"	"S"
## Glyma.03G209400.1	"P"	"I"	"C"	"S"	"R"	"L"	"T"	"S"
## Glyma.19G100200.1	"P"	"I"	"C"	"P"	"N"	"L"	"T"	"S"
## Glyma.19G206800.1	"P"	"I"	"C"	"S"	"R"	"L"	"T"	"S"
## Glyma.18G030200.1	"M"	"F"	"A"	"A"	"V"	"L"	"K"	"K"
##	[,294]	[,295]	[,296]	[,297]	[,298]	[,299]	[,300]	[,301]
## Glyma.16G050500.1	"L"	"N"	"L"	"S"	"Y"	"A"	"A"	"I"
## Glyma.16G146400.1	"L"	"N"	"L"	"R"	"F"	"A"	"A"	"I"
## Glyma.04G093500.1	"L"	"N"	"F"	"S"	"Y"	"A"	"D"	"I"
## Glyma.02G065300.1	"M"	"N"	"L"	"S"	"Y"	"A"	"A"	"I"
## Glyma.02G211800.1	"L"	"N"	"L"	"S"	"Y"	"A"	"D"	"I"
## Glyma.02G254300.1	"L"	"D"	"L"	"L"	"Y"	"A"	"M"	"L"
## Glyma.02G152800.2	"L"	"N"	"L"	"S"	"Y"	"A"	"T"	"I"
## Glyma.10G021500.2	"L"	"N"	"L"	"S"	"Y"	"A"	"T"	"I"
## Glyma.08G059500.6	"L"	"N"	"F"	"S"	"Y"	"A"	"P"	"L"

## Glyma.07G189800.5	"L"	"N"	"F"	"S"	"Y"	"A"	"P"	"L"
## Glyma.11G227300.1	"L"	"D"	"L"	"L"	"Y"	"A"	"M"	"L"
## Glyma.14G179500.1	"L"	"N"	"L"	"S"	"Y"	"A"	"D"	"V"
## Glyma.14G062100.1	"L"	"D"	"L"	"L"	"Y"	"A"	"M"	"L"
## Glyma.06G095400.1	"L"	"N"	"F"	"S"	"F"	"A"	"D"	"I"
## Glyma.03G209400.1	"L"	"N"	"L"	"S"	"Y"	"A"	"I"	"I"
## Glyma.19G100200.1	"L"	"N"	"L"	"S"	"Y"	"A"	"A"	"I"
## Glyma.19G206800.1	"L"	"N"	"L"	"S"	"Y"	"A"	"I"	"I"
## Glyma.18G030200.1	"L"	"D"	"L"	"L"	"Y"	"A"	"M"	"L"
##	[,302]	[,303]	[,304]	[,305]	[,306]	[,307]	[,308]	[,309]
## Glyma.16G050500.1	"Q"	"G"	"S"	"D"	"L"	"I"	"K"	"L"
## Glyma.16G146400.1	"Q"	"N"	"T"	"E"	"L"	"I"	"K"	"L"
## Glyma.04G093500.1	"S"	"A"	"D"	"Q"	"L"	"I"	"S"	"V"
## Glyma.02G065300.1	"Q"	"S"	"R"	"E"	"L"	"I"	"K"	"L"
## Glyma.02G211800.1	"N"	"T"	"D"	"Q"	"L"	"K"	"S"	"V"
## Glyma.02G254300.1	"D"	"T"	"E"	"D"	"H"	"C"	"T"	"L"
## Glyma.02G152800.2	"Q"	"S"	"P"	"D"	"L"	"I"	"K"	"L"
## Glyma.10G021500.2	"Q"	"S"	"P"	"D"	"L"	"I"	"K"	"L"
## Glyma.08G059500.6	"D"	"S"	"D"	"G"	"L"	"A"	"K"	"L"
## Glyma.07G189800.5	"D"	"S"	"D"	"G"	"L"	"T"	"K"	"L"
## Glyma.11G227300.1	"D"	"T"	"E"	"D"	"H"	"C"	"M"	"L"
## Glyma.14G179500.1	"N"	"T"	"D"	"Q"	"L"	"K"	"S"	"V"
## Glyma.14G062100.1	"D"	"T"	"E"	"D"	"H"	"C"	"T"	"L"
## Glyma.06G095400.1	"S"	"A"	"D"	"Q"	"L"	"K"	"S"	"V"
## Glyma.03G209400.1	"Q"	"S"	"S"	"D"	"L"	"I"	"K"	"L"
## Glyma.19G100200.1	"Q"	"G"	"S"	"A"	"L"	"V"	"K"	"L"
## Glyma.19G206800.1	"Q"	"S"	"S"	"D"	"L"	"I"	"K"	"L"
## Glyma.18G030200.1	"D"	"T"	"E"	"D"	"H"	"C"	"M"	"L"
##	[,310]	[,311]	[,312]	[,313]	[,314]	[,315]	[,316]	[,317]
## Glyma.16G050500.1	"I"	"R"	"H"	"C"	"V"	"K"	"L"	"Q"
## Glyma.16G146400.1	"I"	"C"	"C"	"C"	"G"	"K"	"L"	"Q"
## Glyma.04G093500.1	"I"	"R"	"H"	"C"	"H"	"K"	"L"	"Q"
## Glyma.02G065300.1	"I"	"C"	"R"	"C"	"G"	"K"	"L"	"Q"
## Glyma.02G211800.1	"I"	"C"	"H"	"C"	"H"	"K"	"L"	"Q"
## Glyma.02G254300.1	"I"	"Q"	"R"	"C"	"P"	"N"	"L"	"E"
## Glyma.02G152800.2	"V"	"G"	"Q"	"C"	"E"	"S"	"L"	"Q"
## Glyma.10G021500.2	"V"	"G"	"Q"	"C"	"E"	"S"	"L"	"Q"
## Glyma.08G059500.6	"L"	"V"	"H"	"C"	"P"	"K"	"L"	"Q"
## Glyma.07G189800.5	"L"	"V"	"H"	"C"	"P"	"K"	"L"	"Q"
## Glyma.11G227300.1	"I"	"Q"	"K"	"C"	"P"	"N"	"L"	"E"
## Glyma.14G179500.1	"I"	"R"	"H"	"C"	"H"	"K"	"L"	"Q"
## Glyma.14G062100.1	"I"	"Q"	"R"	"C"	"P"	"N"	"L"	"E"
## Glyma.06G095400.1	"I"	"R"	"H"	"C"	"H"	"K"	"L"	"Q"
## Glyma.03G209400.1	"I"	"S"	"Q"	"C"	"P"	"N"	"L"	"L"
## Glyma.19G100200.1	"I"	"H"	"H"	"C"	"V"	"K"	"L"	"Q"
## Glyma.19G206800.1	"I"	"S"	"Q"	"C"	"P"	"N"	"L"	"L"
## Glyma.18G030200.1	"I"	"Q"	"R"	"C"	"P"	"N"	"L"	"E"
##	[,318]	[,319]	[,320]	[,321]	[,322]	[,323]	[,324]	[,325]
## Glyma.16G050500.1	"R"	"L"	"L"	"I"	"M"	"D"	"C"	"I"
## Glyma.16G146400.1	"R"	"L"	"S"	"I"	"M"	"D"	"C"	"I"
## Glyma.04G093500.1	"T"	"F"	"W"	"V"	"L"	"D"	"T"	"I"
## Glyma.02G065300.1	"R"	"L"	"W"	"I"	"M"	"D"	"C"	"I"
## Glyma.02G211800.1	"I"	"F"	"W"	"V"	"L"	"D"	"S"	"I"
## Glyma.02G254300.1	"V"	"L"	"E"	"S"	"R"	"N"	"V"	"I"

## Glyma.02G152800.2	"R"	"L"	"W"	"V"	"L"	"D"	"Y"	"I"
## Glyma.10G021500.2	"R"	"L"	"W"	"V"	"L"	"D"	"Y"	"I"
## Glyma.08G059500.6	"R"	"I"	"W"	"V"	"V"	"D"	"T"	"V"
## Glyma.07G189800.5	"R"	"L"	"W"	"V"	"V"	"D"	"T"	"V"
## Glyma.11G227300.1	"V"	"L"	"E"	"T"	"R"	"N"	"V"	"I"
## Glyma.14G179500.1	"I"	"F"	"W"	"V"	"L"	"D"	"S"	"I"
## Glyma.14G062100.1	"V"	"L"	"E"	"S"	"R"	"N"	"V"	"I"
## Glyma.06G095400.1	"T"	"F"	"W"	"V"	"L"	"D"	"T"	"I"
## Glyma.03G209400.1	"R"	"L"	"W"	"V"	"L"	"D"	"Y"	"I"
## Glyma.19G100200.1	"R"	"L"	"W"	"I"	"M"	"D"	"C"	"I"
## Glyma.19G206800.1	"R"	"L"	"W"	"V"	"L"	"D"	"Y"	"I"
## Glyma.18G030200.1	"V"	"L"	"E"	"T"	"R"	"N"	"V"	"I"
##	[,326]	[,327]	[,328]	[,329]	[,330]	[,331]	[,332]	[,333]
## Glyma.16G050500.1	"G"	"D"	"K"	"G"	"L"	"D"	"V"	"V"
## Glyma.16G146400.1	"G"	"D"	"N"	"G"	"L"	"G"	"V"	"V"
## Glyma.04G093500.1	"C"	"D"	"E"	"G"	"L"	"Q"	"A"	"V"
## Glyma.02G065300.1	"G"	"D"	"F"	"G"	"L"	"H"	"V"	"V"
## Glyma.02G211800.1	"R"	"D"	"E"	"G"	"L"	"Q"	"A"	"V"
## Glyma.02G254300.1	"G"	"D"	"R"	"G"	"L"	"E"	"V"	"L"
## Glyma.02G152800.2	"E"	"D"	"A"	"G"	"L"	"E"	"V"	"I"
## Glyma.10G021500.2	"E"	"D"	"A"	"G"	"L"	"E"	"V"	"I"
## Glyma.08G059500.6	"E"	"D"	"K"	"G"	"L"	"E"	"A"	"V"
## Glyma.07G189800.5	"E"	"D"	"K"	"G"	"L"	"E"	"A"	"V"
## Glyma.11G227300.1	"G"	"D"	"R"	"G"	"L"	"E"	"V"	"L"
## Glyma.14G179500.1	"R"	"D"	"E"	"G"	"L"	"Q"	"A"	"V"
## Glyma.14G062100.1	"G"	"D"	"R"	"G"	"L"	"E"	"V"	"L"
## Glyma.06G095400.1	"C"	"D"	"E"	"G"	"L"	"Q"	"A"	"V"
## Glyma.03G209400.1	"E"	"D"	"A"	"G"	"L"	"Y"	"A"	"L"
## Glyma.19G100200.1	"G"	"D"	"K"	"G"	"L"	"G"	"V"	"V"
## Glyma.19G206800.1	"E"	"D"	"A"	"G"	"L"	"Y"	"A"	"L"
## Glyma.18G030200.1	"G"	"D"	"R"	"G"	"L"	"E"	"V"	"L"
##	[,334]	[,335]	[,336]	[,337]	[,338]	[,339]	[,340]	[,341]
## Glyma.16G050500.1	"A"	"T"	"S"	"C"	"K"	"D"	"L"	"Q"
## Glyma.16G146400.1	"A"	"A"	"T"	"C"	"K"	"D"	"L"	"Q"
## Glyma.04G093500.1	"A"	"E"	"T"	"C"	"K"	"D"	"L"	"R"
## Glyma.02G065300.1	"A"	"S"	"T"	"C"	"K"	"D"	"L"	"Q"
## Glyma.02G211800.1	"A"	"A"	"T"	"C"	"K"	"D"	"L"	"R"
## Glyma.02G254300.1	"A"	"R"	"C"	"C"	"R"	"R"	"L"	"K"
## Glyma.02G152800.2	"A"	"A"	"S"	"C"	"K"	"D"	"L"	"R"
## Glyma.10G021500.2	"A"	"A"	"S"	"C"	"K"	"D"	"L"	"R"
## Glyma.08G059500.6	"G"	"S"	"H"	"C"	"P"	"L"	"L"	"E"
## Glyma.07G189800.5	"G"	"S"	"H"	"C"	"P"	"L"	"L"	"E"
## Glyma.11G227300.1	"G"	"R"	"C"	"C"	"K"	"R"	"L"	"K"
## Glyma.14G179500.1	"A"	"A"	"T"	"C"	"K"	"D"	"L"	"R"
## Glyma.14G062100.1	"A"	"R"	"C"	"C"	"R"	"R"	"L"	"K"
## Glyma.06G095400.1	"A"	"E"	"T"	"C"	"K"	"D"	"L"	"R"
## Glyma.03G209400.1	"A"	"A"	"S"	"C"	"K"	"D"	"L"	"R"
## Glyma.19G100200.1	"A"	"T"	"T"	"C"	"K"	"D"	"L"	"Q"
## Glyma.19G206800.1	"A"	"A"	"S"	"C"	"K"	"D"	"L"	"R"
## Glyma.18G030200.1	"G"	"R"	"C"	"C"	"K"	"R"	"L"	"K"
##	[,342]	[,343]	[,344]	[,345]	[,346]	[,347]	[,348]	[,349]
## Glyma.16G050500.1	"E"	"L"	"R"	"V"	"F"	"P"	"S"	"V"
## Glyma.16G146400.1	"E"	"L"	"R"	"V"	"F"	"P"	"V"	"V"
## Glyma.04G093500.1	"E"	"L"	"R"	"V"	"F"	"P"	"V"	"N"

## Glyma.02G065300.1	"E"	"L"	"R"	"V"	"F"	"P"	"S"	"V"
## Glyma.02G211800.1	"E"	"L"	"R"	"V"	"F"	"P"	"M"	"D"
## Glyma.02G254300.1	"R"	"I"	"R"	"I"	"E"	"R"	"D"	"D"
## Glyma.02G152800.2	"E"	"L"	"R"	"V"	"F"	"P"	"S"	"D"
## Glyma.10G021500.2	"E"	"L"	"R"	"V"	"F"	"P"	"S"	"D"
## Glyma.08G059500.6	"E"	"L"	"R"	"V"	"F"	"P"	"A"	"D"
## Glyma.07G189800.5	"E"	"L"	"R"	"V"	"F"	"P"	"A"	"D"
## Glyma.11G227300.1	"R"	"L"	"R"	"I"	"E"	"R"	"D"	"D"
## Glyma.14G179500.1	"E"	"L"	"R"	"V"	"F"	"P"	"V"	"D"
## Glyma.14G062100.1	"R"	"L"	"R"	"I"	"E"	"R"	"D"	"D"
## Glyma.06G095400.1	"E"	"L"	"R"	"V"	"F"	"P"	"V"	"N"
## Glyma.03G209400.1	"E"	"L"	"R"	"V"	"F"	"P"	"S"	"E"
## Glyma.19G100200.1	"E"	"L"	"R"	"V"	"F"	"P"	"S"	"V"
## Glyma.19G206800.1	"E"	"L"	"R"	"V"	"F"	"P"	"S"	"D"
## Glyma.18G030200.1	"R"	"L"	"R"	"I"	"E"	"R"	"D"	"D"
##	[,350]	[,351]	[,352]	[,353]	[,354]	[,355]	[,356]	[,357]
## Glyma.16G050500.1	"P"	"G"	"_"	"_"	"N"	"P"	"A"	"A"
## Glyma.16G146400.1	"R"	"G"	"G"	"N"	"G"	"P"	"T"	"R"
## Glyma.04G093500.1	"T"	"R"	"E"	"E"	"I"	"E"	"G"	"P"
## Glyma.02G065300.1	"R"	"G"	"R"	"N"	"D"	"P"	"A"	"G"
## Glyma.02G211800.1	"A"	"R"	"E"	"E"	"I"	"D"	"G"	"P"
## Glyma.02G254300.1	"Q"	"M"	"E"	"D"	"E"	"E"	"G"	"V"
## Glyma.02G152800.2	"P"	"G"	"L"	"E"	"P"	"N"	"V"	"A"
## Glyma.10G021500.2	"P"	"G"	"L"	"E"	"P"	"N"	"V"	"A"
## Glyma.08G059500.6	"P"	"D"	"E"	"G"	"I"	"V"	"H"	"G"
## Glyma.07G189800.5	"P"	"D"	"E"	"G"	"I"	"V"	"H"	"G"
## Glyma.11G227300.1	"Q"	"M"	"E"	"D"	"E"	"E"	"G"	"T"
## Glyma.14G179500.1	"A"	"R"	"E"	"E"	"I"	"D"	"G"	"P"
## Glyma.14G062100.1	"Q"	"M"	"E"	"D"	"E"	"E"	"G"	"V"
## Glyma.06G095400.1	"T"	"R"	"E"	"E"	"I"	"E"	"G"	"P"
## Glyma.03G209400.1	"P"	"G"	"L"	"E"	"P"	"N"	"V"	"S"
## Glyma.19G100200.1	"P"	"G"	"_"	"_"	"D"	"P"	"A"	"A"
## Glyma.19G206800.1	"P"	"G"	"L"	"E"	"P"	"N"	"V"	"A"
## Glyma.18G030200.1	"Q"	"M"	"E"	"D"	"E"	"E"	"G"	"T"
##	[,358]	[,359]	[,360]	[,361]	[,362]	[,363]	[,364]	[,365]
## Glyma.16G050500.1	"V"	"T"	"E"	"K"	"G"	"L"	"V"	"A"
## Glyma.16G146400.1	"V"	"T"	"E"	"K"	"G"	"L"	"V"	"A"
## Glyma.04G093500.1	"V"	"S"	"E"	"V"	"G"	"F"	"E"	"A"
## Glyma.02G065300.1	"V"	"T"	"E"	"K"	"G"	"L"	"V"	"A"
## Glyma.02G211800.1	"V"	"S"	"E"	"V"	"G"	"F"	"E"	"A"
## Glyma.02G254300.1	"V"	"S"	"Q"	"R"	"G"	"L"	"I"	"A"
## Glyma.02G152800.2	"L"	"T"	"E"	"Q"	"G"	"L"	"V"	"S"
## Glyma.10G021500.2	"L"	"T"	"E"	"Q"	"G"	"L"	"V"	"S"
## Glyma.08G059500.6	"V"	"T"	"E"	"S"	"G"	"F"	"I"	"A"
## Glyma.07G189800.5	"V"	"T"	"E"	"S"	"G"	"F"	"I"	"A"
## Glyma.11G227300.1	"V"	"S"	"H"	"R"	"G"	"L"	"I"	"A"
## Glyma.14G179500.1	"V"	"S"	"E"	"V"	"G"	"F"	"E"	"A"
## Glyma.14G062100.1	"V"	"S"	"Q"	"R"	"G"	"L"	"I"	"A"
## Glyma.06G095400.1	"V"	"S"	"E"	"V"	"G"	"F"	"E"	"A"
## Glyma.03G209400.1	"L"	"T"	"E"	"Q"	"G"	"L"	"V"	"S"
## Glyma.19G100200.1	"V"	"T"	"E"	"K"	"G"	"L"	"V"	"A"
## Glyma.19G206800.1	"L"	"T"	"E"	"Q"	"G"	"L"	"V"	"S"
## Glyma.18G030200.1	"V"	"S"	"H"	"R"	"G"	"L"	"I"	"A"
##	[,366]	[,367]	[,368]	[,369]	[,370]	[,371]	[,372]	[,373]

## Glyma.16G050500.1	"I"	"S"	"M"	"G"	"C"	"P"	"K"	"L"
## Glyma.16G146400.1	"I"	"S"	"M"	"G"	"C"	"P"	"E"	"L"
## Glyma.04G093500.1	"I"	"S"	"R"	"G"	"C"	"R"	"K"	"L"
## Glyma.02G065300.1	"I"	"S"	"M"	"G"	"C"	"P"	"K"	"L"
## Glyma.02G211800.1	"I"	"S"	"Q"	"G"	"C"	"R"	"K"	"L"
## Glyma.02G254300.1	"L"	"S"	"H"	"G"	"C"	"P"	"E"	"L"
## Glyma.02G152800.2	"V"	"S"	"E"	"G"	"C"	"T"	"K"	"L"
## Glyma.10G021500.2	"V"	"S"	"E"	"G"	"C"	"T"	"K"	"L"
## Glyma.08G059500.6	"V"	"S"	"Q"	"G"	"C"	"P"	"R"	"L"
## Glyma.07G189800.5	"V"	"S"	"Q"	"G"	"C"	"P"	"R"	"L"
## Glyma.11G227300.1	"L"	"S"	"Q"	"G"	"C"	"S"	"E"	"L"
## Glyma.14G179500.1	"I"	"S"	"Q"	"G"	"C"	"R"	"K"	"L"
## Glyma.14G062100.1	"L"	"S"	"H"	"G"	"C"	"P"	"E"	"L"
## Glyma.06G095400.1	"I"	"S"	"R"	"G"	"C"	"R"	"K"	"L"
## Glyma.03G209400.1	"V"	"S"	"E"	"G"	"C"	"P"	"K"	"L"
## Glyma.19G100200.1	"I"	"S"	"M"	"G"	"C"	"P"	"K"	"L"
## Glyma.19G206800.1	"V"	"S"	"E"	"G"	"C"	"P"	"K"	"L"
## Glyma.18G030200.1	"L"	"S"	"Q"	"G"	"C"	"S"	"E"	"L"
##	[,374]	[,375]	[,376]	[,377]	[,378]	[,379]	[,380]	[,381]
## Glyma.16G050500.1	"H"	"S"	"L"	"L"	"Y"	"F"	"C"	"H"
## Glyma.16G146400.1	"H"	"S"	"L"	"L"	"Y"	"F"	"C"	"Q"
## Glyma.04G093500.1	"Q"	"S"	"I"	"L"	"F"	"F"	"C"	"Q"
## Glyma.02G065300.1	"H"	"S"	"L"	"L"	"Y"	"F"	"C"	"Q"
## Glyma.02G211800.1	"E"	"S"	"I"	"L"	"F"	"L"	"C"	"Q"
## Glyma.02G254300.1	"E"	"Y"	"L"	"A"	"V"	"Y"	"V"	"S"
## Glyma.02G152800.2	"Q"	"S"	"V"	"L"	"Y"	"F"	"C"	"R"
## Glyma.10G021500.2	"Q"	"S"	"V"	"L"	"Y"	"F"	"C"	"R"
## Glyma.08G059500.6	"H"	"Y"	"V"	"L"	"Y"	"F"	"C"	"R"
## Glyma.07G189800.5	"H"	"Y"	"V"	"L"	"Y"	"F"	"C"	"R"
## Glyma.11G227300.1	"E"	"Y"	"M"	"A"	"V"	"Y"	"V"	"S"
## Glyma.14G179500.1	"E"	"S"	"I"	"L"	"F"	"F"	"T"	"Q"
## Glyma.14G062100.1	"E"	"Y"	"L"	"A"	"V"	"Y"	"V"	"S"
## Glyma.06G095400.1	"Q"	"S"	"I"	"L"	"F"	"F"	"C"	"Q"
## Glyma.03G209400.1	"Q"	"S"	"V"	"L"	"Y"	"F"	"C"	"R"
## Glyma.19G100200.1	"H"	"S"	"L"	"L"	"Y"	"F"	"C"	"H"
## Glyma.19G206800.1	"Q"	"S"	"V"	"L"	"Y"	"F"	"C"	"R"
## Glyma.18G030200.1	"E"	"Y"	"M"	"A"	"V"	"Y"	"V"	"S"
##	[,382]	[,383]	[,384]	[,385]	[,386]	[,387]	[,388]	[,389]
## Glyma.16G050500.1	"Q"	"M"	"T"	"N"	"A"	"A"	"L"	"I"
## Glyma.16G146400.1	"Q"	"M"	"T"	"N"	"A"	"A"	"L"	"I"
## Glyma.04G093500.1	"R"	"M"	"T"	"N"	"A"	"A"	"V"	"V"
## Glyma.02G065300.1	"Q"	"M"	"T"	"N"	"A"	"A"	"L"	"I"
## Glyma.02G211800.1	"R"	"M"	"T"	"N"	"A"	"A"	"V"	"V"
## Glyma.02G254300.1	"D"	"I"	"T"	"N"	"A"	"S"	"L"	"E"
## Glyma.02G152800.2	"Q"	"M"	"S"	"N"	"A"	"A"	"L"	"D"
## Glyma.10G021500.2	"Q"	"M"	"T"	"N"	"S"	"A"	"L"	"D"
## Glyma.08G059500.6	"Q"	"M"	"T"	"N"	"A"	"A"	"V"	"A"
## Glyma.07G189800.5	"Q"	"M"	"T"	"N"	"A"	"A"	"V"	"A"
## Glyma.11G227300.1	"D"	"I"	"T"	"N"	"A"	"S"	"L"	"E"
## Glyma.14G179500.1	"R"	"M"	"T"	"N"	"A"	"A"	"V"	"V"
## Glyma.14G062100.1	"D"	"I"	"T"	"N"	"A"	"S"	"L"	"E"
## Glyma.06G095400.1	"R"	"M"	"T"	"N"	"A"	"A"	"V"	"V"
## Glyma.03G209400.1	"Q"	"M"	"S"	"N"	"A"	"A"	"L"	"H"
## Glyma.19G100200.1	"Q"	"M"	"T"	"N"	"A"	"A"	"L"	"I"



## Glyma.19G206800.1	"Q"	"M"	"S"	"N"	"A"	"A"	"L"	"H"
## Glyma.18G030200.1	"D"	"I"	"T"	"N"	"A"	"S"	"L"	"E"
##	[,390]	[,391]	[,392]	[,393]	[,394]	[,395]	[,396]	[,397]
## Glyma.16G050500.1	"T"	"V"	"A"	"K"	"N"	"C"	"P"	"N"
## Glyma.16G146400.1	"T"	"V"	"A"	"K"	"N"	"C"	"P"	"N"
## Glyma.04G093500.1	"A"	"M"	"S"	"N"	"N"	"C"	"P"	"D"
## Glyma.02G065300.1	"T"	"V"	"A"	"K"	"N"	"C"	"P"	"N"
## Glyma.02G211800.1	"A"	"M"	"S"	"K"	"N"	"C"	"P"	"D"
## Glyma.02G254300.1	"H"	"I"	"G"	"T"	"H"	"L"	"K"	"N"
## Glyma.02G152800.2	"T"	"I"	"A"	"R"	"S"	"R"	"P"	"N"
## Glyma.10G021500.2	"T"	"I"	"A"	"R"	"N"	"R"	"P"	"N"
## Glyma.08G059500.6	"T"	"V"	"V"	"Q"	"N"	"C"	"P"	"D"
## Glyma.07G189800.5	"T"	"V"	"V"	"Q"	"N"	"C"	"P"	"D"
## Glyma.11G227300.1	"H"	"I"	"G"	"T"	"H"	"L"	"K"	"N"
## Glyma.14G179500.1	"A"	"M"	"S"	"K"	"N"	"C"	"P"	"D"
## Glyma.14G062100.1	"H"	"I"	"G"	"T"	"H"	"L"	"K"	"N"
## Glyma.06G095400.1	"A"	"M"	"S"	"N"	"N"	"C"	"P"	"D"
## Glyma.03G209400.1	"T"	"I"	"A"	"R"	"N"	"R"	"P"	"N"
## Glyma.19G100200.1	"T"	"V"	"A"	"K"	"N"	"C"	"P"	"N"
## Glyma.19G206800.1	"T"	"I"	"A"	"R"	"N"	"R"	"T"	"N"
## Glyma.18G030200.1	"H"	"I"	"G"	"T"	"H"	"L"	"K"	"N"
##	[,398]	[,399]	[,400]	[,401]	[,402]	[,403]	[,404]	[,405]
## Glyma.16G050500.1	"F"	"I"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.16G146400.1	"F"	"I"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.04G093500.1	"L"	"V"	"V"	"F"	"R"	"L"	"C"	"I"
## Glyma.02G065300.1	"F"	"I"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.02G211800.1	"L"	"V"	"V"	"F"	"R"	"L"	"C"	"I"
## Glyma.02G254300.1	"L"	"C"	"D"	"F"	"R"	"L"	"V"	"L"
## Glyma.02G152800.2	"M"	"T"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.10G021500.2	"M"	"T"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.08G059500.6	"F"	"T"	"H"	"F"	"R"	"L"	"C"	"I"
## Glyma.07G189800.5	"F"	"T"	"H"	"F"	"R"	"L"	"C"	"I"
## Glyma.11G227300.1	"L"	"C"	"D"	"F"	"R"	"L"	"V"	"L"
## Glyma.14G179500.1	"L"	"V"	"V"	"F"	"R"	"L"	"C"	"I"
## Glyma.14G062100.1	"L"	"C"	"D"	"F"	"R"	"L"	"V"	"L"
## Glyma.06G095400.1	"L"	"V"	"V"	"F"	"R"	"L"	"C"	"I"
## Glyma.03G209400.1	"L"	"T"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.19G100200.1	"F"	"I"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.19G206800.1	"L"	"T"	"R"	"F"	"R"	"L"	"C"	"I"
## Glyma.18G030200.1	"L"	"C"	"D"	"F"	"R"	"L"	"V"	"L"
##	[,406]	[,407]	[,408]	[,409]	[,410]	[,411]	[,412]	[,413]
## Glyma.16G050500.1	"L"	"D"	"A"	"T"	"K"	"P"	"D"	"T"
## Glyma.16G146400.1	"L"	"D"	"P"	"T"	"K"	"P"	"D"	"T"
## Glyma.04G093500.1	"I"	"G"	"R"	"Y"	"R"	"P"	"E"	"T"
## Glyma.02G065300.1	"L"	"D"	"P"	"T"	"K"	"P"	"D"	"T"
## Glyma.02G211800.1	"I"	"G"	"R"	"Y"	"R"	"P"	"V"	"T"
## Glyma.02G254300.1	"L"	"D"	"R"	"E"	"E"	"K"	"I"	"T"
## Glyma.02G152800.2	"I"	"E"	"P"	"R"	"A"	"Y"	"L"	"T"
## Glyma.10G021500.2	"I"	"E"	"P"	"Q"	"A"	"H"	"L"	"T"
## Glyma.08G059500.6	"M"	"H"	"P"	"G"	"Q"	"Y"	"L"	"T"
## Glyma.07G189800.5	"M"	"H"	"P"	"G"	"Q"	"Y"	"L"	"T"
## Glyma.11G227300.1	"L"	"D"	"H"	"E"	"E"	"K"	"I"	"T"
## Glyma.14G179500.1	"I"	"G"	"R"	"Y"	"R"	"P"	"V"	"T"
## Glyma.14G062100.1	"L"	"D"	"R"	"E"	"E"	"K"	"I"	"T"

## Glyma.06G095400.1	"I"	"G"	"Q"	"Y"	"R"	"P"	"V"	"T"
## Glyma.03G209400.1	"I"	"E"	"P"	"R"	"T"	"Y"	"L"	"T"
## Glyma.19G100200.1	"L"	"D"	"A"	"T"	"K"	"P"	"D"	"T"
## Glyma.19G206800.1	"I"	"E"	"P"	"R"	"T"	"Y"	"L"	"T"
## Glyma.18G030200.1	"L"	"D"	"H"	"E"	"E"	"K"	"I"	"T"
##	[,414]	[,415]	[,416]	[,417]	[,418]	[,419]	[,420]	[,421]
## Glyma.16G050500.1	"M"	"Q"	"P"	"L"	"D"	"E"	"G"	"F"
## Glyma.16G146400.1	"M"	"Q"	"P"	"L"	"N"	"E"	"G"	"F"
## Glyma.04G093500.1	"L"	"E"	"P"	"M"	"D"	"E"	"G"	"F"
## Glyma.02G065300.1	"V"	"Q"	"P"	"L"	"D"	"E"	"G"	"F"
## Glyma.02G211800.1	"Q"	"E"	"P"	"M"	"D"	"E"	"G"	"F"
## Glyma.02G254300.1	"D"	"L"	"P"	"L"	"D"	"N"	"G"	"V"
## Glyma.02G152800.2	"H"	"Q"	"P"	"L"	"D"	"A"	"G"	"F"
## Glyma.10G021500.2	"H"	"Q"	"P"	"L"	"D"	"A"	"G"	"F"
## Glyma.08G059500.6	"Q"	"E"	"S"	"M"	"D"	"E"	"A"	"F"
## Glyma.07G189800.5	"Q"	"E"	"S"	"M"	"D"	"E"	"A"	"F"
## Glyma.11G227300.1	"D"	"L"	"P"	"L"	"D"	"N"	"G"	"V"
## Glyma.14G179500.1	"E"	"E"	"P"	"M"	"D"	"E"	"G"	"F"
## Glyma.14G062100.1	"D"	"L"	"P"	"L"	"D"	"N"	"G"	"V"
## Glyma.06G095400.1	"L"	"E"	"P"	"M"	"D"	"E"	"G"	"F"
## Glyma.03G209400.1	"L"	"E"	"P"	"L"	"D"	"S"	"G"	"F"
## Glyma.19G100200.1	"M"	"Q"	"P"	"L"	"D"	"E"	"G"	"F"
## Glyma.19G206800.1	"H"	"E"	"P"	"L"	"D"	"S"	"G"	"F"
## Glyma.18G030200.1	"D"	"L"	"P"	"L"	"D"	"N"	"G"	"V"
##	[,422]	[,423]	[,424]	[,425]	[,426]	[,427]	[,428]	[,429]
## Glyma.16G050500.1	"G"	"A"	"I"	"V"	"Q"	"S"	"C"	"R"
## Glyma.16G146400.1	"G"	"A"	"I"	"V"	"Q"	"S"	"C"	"K"
## Glyma.04G093500.1	"G"	"A"	"I"	"V"	"M"	"N"	"C"	"K"
## Glyma.02G065300.1	"G"	"A"	"I"	"V"	"Q"	"S"	"C"	"K"
## Glyma.02G211800.1	"G"	"A"	"I"	"V"	"M"	"N"	"C"	"K"
## Glyma.02G254300.1	"R"	"A"	"L"	"L"	"R"	"G"	"C"	"D"
## Glyma.02G152800.2	"G"	"A"	"I"	"V"	"E"	"H"	"C"	"K"
## Glyma.10G021500.2	"G"	"A"	"I"	"V"	"E"	"H"	"C"	"K"
## Glyma.08G059500.6	"G"	"A"	"V"	"V"	"K"	"T"	"C"	"T"
## Glyma.07G189800.5	"G"	"A"	"V"	"V"	"K"	"T"	"C"	"T"
## Glyma.11G227300.1	"R"	"A"	"L"	"L"	"R"	"G"	"C"	"N"
## Glyma.14G179500.1	"G"	"A"	"I"	"V"	"M"	"N"	"C"	"K"
## Glyma.14G062100.1	"R"	"A"	"L"	"L"	"R"	"G"	"C"	"D"
## Glyma.06G095400.1	"G"	"A"	"I"	"V"	"M"	"N"	"C"	"K"
## Glyma.03G209400.1	"G"	"A"	"I"	"V"	"E"	"Q"	"C"	"K"
## Glyma.19G100200.1	"G"	"A"	"I"	"V"	"Q"	"S"	"C"	"R"
## Glyma.19G206800.1	"G"	"A"	"I"	"V"	"E"	"Q"	"C"	"K"
## Glyma.18G030200.1	"R"	"A"	"L"	"L"	"R"	"G"	"C"	"D"
##	[,430]	[,431]	[,432]	[,433]	[,434]	[,435]	[,436]	[,437]
## Glyma.16G050500.1	"R"	"L"	"R"	"R"	"L"	"S"	"L"	"S"
## Glyma.16G146400.1	"Q"	"L"	"R"	"R"	"L"	"S"	"L"	"S"
## Glyma.04G093500.1	"K"	"L"	"T"	"R"	"L"	"A"	"V"	"S"
## Glyma.02G065300.1	"Q"	"L"	"R"	"R"	"L"	"S"	"L"	"S"
## Glyma.02G211800.1	"K"	"L"	"T"	"R"	"L"	"A"	"V"	"S"
## Glyma.02G254300.1	"K"	"L"	"R"	"R"	"F"	"A"	"L"	"P"
## Glyma.02G152800.2	"D"	"L"	"Q"	"R"	"L"	"S"	"L"	"S"
## Glyma.10G021500.2	"D"	"L"	"Q"	"R"	"L"	"S"	"L"	"S"
## Glyma.08G059500.6	"K"	"L"	"Q"	"R"	"L"	"A"	"V"	"S"
## Glyma.07G189800.5	"K"	"L"	"Q"	"R"	"L"	"A"	"V"	"S"

## Glyma.11G227300.1	"K"	"L"	"R"	"R"	"F"	"A"	"L"	"R"
## Glyma.14G179500.1	"K"	"L"	"T"	"R"	"L"	"A"	"M"	"S"
## Glyma.14G062100.1	"K"	"L"	"R"	"R"	"F"	"A"	"L"	"P"
## Glyma.06G095400.1	"K"	"L"	"T"	"R"	"L"	"A"	"V"	"S"
## Glyma.03G209400.1	"D"	"L"	"Q"	"R"	"L"	"S"	"L"	"S"
## Glyma.19G100200.1	"R"	"L"	"R"	"R"	"L"	"S"	"L"	"S"
## Glyma.19G206800.1	"D"	"L"	"Q"	"R"	"L"	"S"	"L"	"S"
## Glyma.18G030200.1	"K"	"L"	"R"	"R"	"F"	"A"	"L"	"R"
##	[,438]	[,439]	[,440]	[,441]	[,442]	[,443]	[,444]	[,445]
## Glyma.16G050500.1	"G"	"Q"	"L"	"T"	"D"	"Q"	"V"	"F"
## Glyma.16G146400.1	"G"	"Q"	"L"	"T"	"D"	"Q"	"V"	"F"
## Glyma.04G093500.1	"G"	"L"	"L"	"T"	"D"	"R"	"A"	"F"
## Glyma.02G065300.1	"G"	"Q"	"L"	"T"	"D"	"Q"	"V"	"F"
## Glyma.02G211800.1	"G"	"L"	"L"	"T"	"D"	"R"	"A"	"F"
## Glyma.02G254300.1	"G"	"G"	"L"	"T"	"D"	"V"	"G"	"L"
## Glyma.02G152800.2	"G"	"L"	"L"	"T"	"D"	"R"	"V"	"F"
## Glyma.10G021500.2	"G"	"L"	"L"	"T"	"D"	"R"	"V"	"F"
## Glyma.08G059500.6	"G"	"Y"	"L"	"T"	"D"	"L"	"T"	"F"
## Glyma.07G189800.5	"G"	"Y"	"L"	"T"	"D"	"L"	"T"	"F"
## Glyma.11G227300.1	"G"	"G"	"L"	"T"	"D"	"V"	"G"	"L"
## Glyma.14G179500.1	"G"	"L"	"L"	"T"	"D"	"R"	"V"	"F"
## Glyma.14G062100.1	"G"	"G"	"L"	"T"	"D"	"V"	"G"	"L"
## Glyma.06G095400.1	"G"	"L"	"L"	"T"	"D"	"R"	"A"	"F"
## Glyma.03G209400.1	"G"	"L"	"L"	"T"	"D"	"R"	"V"	"F"
## Glyma.19G100200.1	"G"	"K"	"L"	"T"	"D"	"Q"	"V"	"F"
## Glyma.19G206800.1	"G"	"L"	"L"	"T"	"D"	"R"	"V"	"F"
## Glyma.18G030200.1	"G"	"G"	"L"	"T"	"D"	"V"	"G"	"L"
##	[,446]	[,447]	[,448]	[,449]	[,450]	[,451]	[,452]	[,453]
## Glyma.16G050500.1	"L"	"Y"	"I"	"G"	"M"	"Y"	"A"	"E"
## Glyma.16G146400.1	"L"	"Y"	"I"	"G"	"M"	"Y"	"A"	"E"
## Glyma.04G093500.1	"N"	"Y"	"I"	"G"	"T"	"Y"	"G"	"K"
## Glyma.02G065300.1	"L"	"Y"	"I"	"G"	"V"	"Y"	"A"	"E"
## Glyma.02G211800.1	"E"	"Y"	"I"	"G"	"T"	"Y"	"G"	"K"
## Glyma.02G254300.1	"G"	"Y"	"V"	"G"	"Q"	"Y"	"S"	"P"
## Glyma.02G152800.2	"E"	"Y"	"I"	"G"	"T"	"Y"	"G"	"K"
## Glyma.10G021500.2	"E"	"Y"	"I"	"G"	"T"	"Y"	"G"	"K"
## Glyma.08G059500.6	"E"	"Y"	"I"	"G"	"K"	"Y"	"A"	"K"
## Glyma.07G189800.5	"E"	"Y"	"I"	"G"	"K"	"Y"	"A"	"K"
## Glyma.11G227300.1	"G"	"Y"	"I"	"G"	"Q"	"Y"	"S"	"P"
## Glyma.14G179500.1	"E"	"Y"	"I"	"G"	"M"	"Y"	"G"	"K"
## Glyma.14G062100.1	"G"	"Y"	"I"	"G"	"Q"	"Y"	"S"	"P"
## Glyma.06G095400.1	"S"	"Y"	"I"	"G"	"T"	"Y"	"G"	"K"
## Glyma.03G209400.1	"E"	"Y"	"I"	"G"	"T"	"Y"	"A"	"K"
## Glyma.19G100200.1	"L"	"Y"	"I"	"G"	"M"	"Y"	"A"	"E"
## Glyma.19G206800.1	"E"	"Y"	"I"	"G"	"T"	"C"	"G"	"K"
## Glyma.18G030200.1	"G"	"Y"	"I"	"G"	"Q"	"Y"	"S"	"P"
##	[,454]	[,455]	[,456]	[,457]	[,458]	[,459]	[,460]	[,461]
## Glyma.16G050500.1	"K"	"L"	"E"	"M"	"L"	"S"	"I"	"A"
## Glyma.16G146400.1	"Q"	"L"	"E"	"M"	"L"	"S"	"V"	"A"
## Glyma.04G093500.1	"L"	"I"	"R"	"T"	"L"	"S"	"V"	"A"
## Glyma.02G065300.1	"Q"	"L"	"E"	"M"	"L"	"S"	"I"	"A"
## Glyma.02G211800.1	"L"	"V"	"R"	"T"	"L"	"S"	"V"	"A"
## Glyma.02G254300.1	"N"	"V"	"R"	"W"	"M"	"L"	"L"	"G"
## Glyma.02G152800.2	"K"	"L"	"E"	"M"	"L"	"S"	"V"	"A"

## Glyma.10G021500.2	"K"	"L"	"E"	"M"	"L"	"S"	"V"	"A"
## Glyma.08G059500.6	"N"	"L"	"E"	"T"	"L"	"S"	"V"	"A"
## Glyma.07G189800.5	"N"	"L"	"E"	"T"	"L"	"S"	"V"	"A"
## Glyma.11G227300.1	"N"	"V"	"R"	"W"	"M"	"L"	"L"	"G"
## Glyma.14G179500.1	"L"	"V"	"R"	"T"	"L"	"S"	"V"	"A"
## Glyma.14G062100.1	"N"	"V"	"R"	"W"	"M"	"L"	"L"	"G"
## Glyma.06G095400.1	"L"	"I"	"R"	"T"	"L"	"S"	"V"	"A"
## Glyma.03G209400.1	"K"	"L"	"E"	"M"	"L"	"S"	"V"	"A"
## Glyma.19G100200.1	"K"	"L"	"E"	"M"	"L"	"S"	"I"	"A"
## Glyma.19G206800.1	"K"	"L"	"E"	"M"	"L"	"S"	"V"	"A"
## Glyma.18G030200.1	"N"	"V"	"R"	"W"	"M"	"L"	"L"	"G"
##	[,462]	[,463]	[,464]	[,465]	[,466]	[,467]	[,468]	[,469]
## Glyma.16G050500.1	"F"	"A"	"G"	"E"	"S"	"D"	"K"	"G"
## Glyma.16G146400.1	"F"	"A"	"G"	"E"	"S"	"D"	"K"	"A"
## Glyma.04G093500.1	"F"	"A"	"G"	"D"	"I"	"D"	"L"	"G"
## Glyma.02G065300.1	"F"	"A"	"G"	"E"	"S"	"D"	"K"	"A"
## Glyma.02G211800.1	"F"	"A"	"G"	"D"	"I"	"D"	"V"	"G"
## Glyma.02G254300.1	"Y"	"V"	"G"	"E"	"I"	"D"	"A"	"G"
## Glyma.02G152800.2	"F"	"A"	"G"	"D"	"S"	"D"	"L"	"G"
## Glyma.10G021500.2	"F"	"A"	"G"	"D"	"S"	"D"	"L"	"G"
## Glyma.08G059500.6	"F"	"A"	"G"	"S"	"S"	"D"	"W"	"G"
## Glyma.07G189800.5	"F"	"A"	"G"	"S"	"S"	"D"	"W"	"G"
## Glyma.11G227300.1	"Y"	"V"	"G"	"E"	"S"	"D"	"A"	"G"
## Glyma.14G179500.1	"F"	"A"	"G"	"D"	"I"	"D"	"V"	"G"
## Glyma.14G062100.1	"Y"	"V"	"G"	"E"	"I"	"D"	"A"	"G"
## Glyma.06G095400.1	"F"	"A"	"G"	"D"	"I"	"D"	"L"	"G"
## Glyma.03G209400.1	"F"	"A"	"G"	"D"	"S"	"D"	"L"	"G"
## Glyma.19G100200.1	"F"	"A"	"G"	"D"	"G"	"D"	"K"	"G"
## Glyma.19G206800.1	"F"	"A"	"G"	"D"	"S"	"D"	"L"	"G"
## Glyma.18G030200.1	"Y"	"V"	"G"	"E"	"S"	"D"	"A"	"G"
##	[,470]	[,471]	[,472]	[,473]	[,474]	[,475]	[,476]	[,477]
## Glyma.16G050500.1	"M"	"L"	"Y"	"V"	"L"	"N"	"G"	"C"
## Glyma.16G146400.1	"M"	"L"	"Y"	"V"	"L"	"N"	"G"	"C"
## Glyma.04G093500.1	"L"	"Q"	"Y"	"V"	"L"	"E"	"G"	"C"
## Glyma.02G065300.1	"M"	"L"	"Y"	"V"	"L"	"N"	"G"	"C"
## Glyma.02G211800.1	"L"	"K"	"Y"	"V"	"L"	"K"	"G"	"C"
## Glyma.02G254300.1	"L"	"L"	"E"	"F"	"S"	"K"	"G"	"C"
## Glyma.02G152800.2	"L"	"H"	"H"	"V"	"L"	"S"	"G"	"C"
## Glyma.10G021500.2	"L"	"H"	"H"	"V"	"L"	"S"	"G"	"C"
## Glyma.08G059500.6	"M"	"R"	"C"	"V"	"L"	"D"	"G"	"C"
## Glyma.07G189800.5	"M"	"R"	"C"	"V"	"L"	"D"	"G"	"C"
## Glyma.11G227300.1	"L"	"L"	"E"	"F"	"S"	"K"	"G"	"C"
## Glyma.14G179500.1	"L"	"K"	"Y"	"V"	"L"	"E"	"G"	"C"
## Glyma.14G062100.1	"L"	"L"	"E"	"F"	"S"	"K"	"G"	"C"
## Glyma.06G095400.1	"L"	"Q"	"Y"	"V"	"L"	"Q"	"G"	"C"
## Glyma.03G209400.1	"L"	"H"	"H"	"V"	"L"	"S"	"G"	"C"
## Glyma.19G100200.1	"M"	"L"	"Y"	"V"	"L"	"N"	"G"	"C"
## Glyma.19G206800.1	"L"	"H"	"H"	"V"	"L"	"S"	"G"	"C"
## Glyma.18G030200.1	"L"	"L"	"E"	"F"	"A"	"K"	"G"	"C"
##	[,478]	[,479]	[,480]	[,481]	[,482]	[,483]	[,484]	[,485]
## Glyma.16G050500.1	"K"	"K"	"L"	"R"	"K"	"L"	"E"	"I"
## Glyma.16G146400.1	"K"	"K"	"I"	"H"	"K"	"L"	"A"	"I"
## Glyma.04G093500.1	"P"	"N"	"L"	"Q"	"K"	"L"	"E"	"I"
## Glyma.02G065300.1	"K"	"K"	"L"	"R"	"K"	"L"	"E"	"I"

## Glyma.02G211800.1	"P"	"N"	"L"	"Q"	"K"	"L"	"E"	"I"
## Glyma.02G254300.1	"P"	"S"	"L"	"Q"	"K"	"L"	"E"	"M"
## Glyma.02G152800.2	"D"	"N"	"L"	"R"	"K"	"L"	"E"	"I"
## Glyma.10G021500.2	"D"	"N"	"L"	"R"	"K"	"L"	"E"	"I"
## Glyma.08G059500.6	"P"	"K"	"L"	"R"	"K"	"L"	"E"	"V"
## Glyma.07G189800.5	"P"	"K"	"L"	"R"	"K"	"L"	"E"	"V"
## Glyma.11G227300.1	"P"	"S"	"L"	"Q"	"K"	"L"	"E"	"M"
## Glyma.14G179500.1	"P"	"N"	"L"	"Q"	"K"	"L"	"E"	"I"
## Glyma.14G062100.1	"P"	"S"	"L"	"Q"	"K"	"L"	"E"	"M"
## Glyma.06G095400.1	"P"	"N"	"L"	"Q"	"K"	"L"	"E"	"I"
## Glyma.03G209400.1	"D"	"N"	"L"	"R"	"K"	"L"	"E"	"I"
## Glyma.19G100200.1	"K"	"K"	"L"	"R"	"K"	"L"	"E"	"I"
## Glyma.19G206800.1	"D"	"N"	"L"	"R"	"K"	"L"	"E"	"I"
## Glyma.18G030200.1	"P"	"S"	"L"	"Q"	"K"	"L"	"E"	"M"
##	[,486]	[,487]	[,488]	[,489]	[,490]	[,491]	[,492]	[,493]
## Glyma.16G050500.1	"R"	"D"	"C"	"P"	"F"	"G"	"N"	"V"
## Glyma.16G146400.1	"R"	"G"	"S"	"P"	"F"	"G"	"D"	"S"
## Glyma.04G093500.1	"R"	"D"	"S"	"P"	"F"	"G"	"D"	"G"
## Glyma.02G065300.1	"R"	"D"	"S"	"P"	"F"	"G"	"D"	"S"
## Glyma.02G211800.1	"R"	"D"	"S"	"P"	"F"	"G"	"D"	"G"
## Glyma.02G254300.1	"R"	"G"	"C"	"S"	"F"	"S"	"E"	"Y"
## Glyma.02G152800.2	"R"	"D"	"C"	"P"	"F"	"G"	"D"	"K"
## Glyma.10G021500.2	"R"	"D"	"C"	"P"	"F"	"G"	"D"	"K"
## Glyma.08G059500.6	"R"	"D"	"C"	"P"	"F"	"G"	"N"	"G"
## Glyma.07G189800.5	"R"	"D"	"C"	"P"	"F"	"G"	"N"	"G"
## Glyma.11G227300.1	"R"	"G"	"C"	"S"	"F"	"S"	"E"	"R"
## Glyma.14G179500.1	"R"	"D"	"S"	"P"	"F"	"G"	"D"	"G"
## Glyma.14G062100.1	"R"	"G"	"C"	"S"	"F"	"S"	"E"	"Y"
## Glyma.06G095400.1	"R"	"D"	"S"	"P"	"F"	"G"	"D"	"G"
## Glyma.03G209400.1	"R"	"D"	"C"	"P"	"F"	"G"	"D"	"K"
## Glyma.19G100200.1	"R"	"D"	"C"	"P"	"F"	"G"	"D"	"M"
## Glyma.19G206800.1	"R"	"D"	"C"	"P"	"F"	"G"	"D"	"K"
## Glyma.18G030200.1	"R"	"G"	"C"	"L"	"F"	"S"	"E"	"R"
##	[,494]	[,495]	[,496]	[,497]	[,498]	[,499]	[,500]	[,501]
## Glyma.16G050500.1	"A"	"L"	"L"	"T"	"D"	"V"	"G"	"K"
## Glyma.16G146400.1	"A"	"L"	"L"	"M"	"D"	"V"	"G"	"K"
## Glyma.04G093500.1	"A"	"L"	"R"	"S"	"G"	"L"	"H"	"H"
## Glyma.02G065300.1	"A"	"L"	"L"	"M"	"D"	"V"	"G"	"K"
## Glyma.02G211800.1	"A"	"L"	"R"	"S"	"G"	"L"	"H"	"H"
## Glyma.02G254300.1	"A"	"L"	"A"	"I"	"A"	"A"	"T"	"Q"
## Glyma.02G152800.2	"A"	"L"	"L"	"A"	"N"	"A"	"A"	"K"
## Glyma.10G021500.2	"A"	"L"	"L"	"A"	"N"	"A"	"A"	"K"
## Glyma.08G059500.6	"A"	"L"	"L"	"S"	"G"	"L"	"G"	"K"
## Glyma.07G189800.5	"A"	"L"	"L"	"S"	"G"	"L"	"G"	"K"
## Glyma.11G227300.1	"A"	"L"	"A"	"V"	"A"	"A"	"T"	"Q"
## Glyma.14G179500.1	"A"	"L"	"R"	"S"	"G"	"L"	"H"	"H"
## Glyma.14G062100.1	"A"	"L"	"A"	"I"	"A"	"A"	"T"	"Q"
## Glyma.06G095400.1	"A"	"L"	"H"	"S"	"G"	"L"	"H"	"H"
## Glyma.03G209400.1	"A"	"L"	"L"	"A"	"N"	"A"	"E"	"K"
## Glyma.19G100200.1	"A"	"L"	"L"	"T"	"D"	"V"	"G"	"K"
## Glyma.19G206800.1	"A"	"L"	"L"	"A"	"N"	"A"	"E"	"K"
## Glyma.18G030200.1	"A"	"L"	"A"	"V"	"A"	"A"	"T"	"Q"
##	[,502]	[,503]	[,504]	[,505]	[,506]	[,507]	[,508]	[,509]
## Glyma.16G050500.1	"Y"	"E"	"T"	"M"	"R"	"S"	"L"	"W"

## Glyma.16G146400.1	"Y"	"E"	"T"	"M"	"Q"	"F"	"L"	"W"
## Glyma.04G093500.1	"F"	"Y"	"N"	"M"	"R"	"F"	"L"	"W"
## Glyma.02G065300.1	"Y"	"E"	"T"	"M"	"R"	"S"	"L"	"W"
## Glyma.02G211800.1	"Y"	"Y"	"N"	"M"	"R"	"F"	"L"	"W"
## Glyma.02G254300.1	"L"	"N"	"S"	"L"	"R"	"Y"	"L"	"W"
## Glyma.02G152800.2	"L"	"E"	"T"	"M"	"R"	"S"	"L"	"W"
## Glyma.10G021500.2	"L"	"E"	"T"	"M"	"R"	"S"	"L"	"W"
## Glyma.08G059500.6	"Y"	"E"	"S"	"M"	"R"	"S"	"L"	"W"
## Glyma.07G189800.5	"Y"	"E"	"S"	"M"	"R"	"S"	"L"	"W"
## Glyma.11G227300.1	"L"	"T"	"S"	"L"	"R"	"Y"	"L"	"W"
## Glyma.14G179500.1	"Y"	"Y"	"N"	"M"	"R"	"F"	"L"	"W"
## Glyma.14G062100.1	"L"	"N"	"S"	"L"	"R"	"Y"	"L"	"W"
## Glyma.06G095400.1	"F"	"Y"	"N"	"M"	"R"	"F"	"L"	"W"
## Glyma.03G209400.1	"L"	"E"	"T"	"M"	"R"	"S"	"L"	"W"
## Glyma.19G100200.1	"Y"	"E"	"T"	"M"	"R"	"S"	"L"	"W"
## Glyma.19G206800.1	"L"	"E"	"T"	"M"	"R"	"S"	"L"	"W"
## Glyma.18G030200.1	"L"	"T"	"S"	"L"	"R"	"Y"	"L"	"W"
##	[,510]	[,511]	[,512]	[,513]	[,514]	[,515]	[,516]	[,517]
## Glyma.16G050500.1	"M"	"S"	"S"	"C"	"E"	"V"	"T"	"V"
## Glyma.16G146400.1	"M"	"T"	"S"	"C"	"N"	"V"	"T"	"V"
## Glyma.04G093500.1	"M"	"S"	"S"	"C"	"K"	"L"	"T"	"R"
## Glyma.02G065300.1	"M"	"S"	"S"	"C"	"D"	"V"	"T"	"I"
## Glyma.02G211800.1	"M"	"S"	"T"	"C"	"K"	"L"	"T"	"L"
## Glyma.02G254300.1	"V"	"Q"	"G"	"Y"	"S"	"A"	"S"	"A"
## Glyma.02G152800.2	"M"	"S"	"S"	"C"	"L"	"V"	"S"	"Y"
## Glyma.10G021500.2	"M"	"S"	"S"	"C"	"L"	"V"	"S"	"Y"
## Glyma.08G059500.6	"M"	"S"	"D"	"C"	"N"	"L"	"T"	"M"
## Glyma.07G189800.5	"M"	"S"	"D"	"C"	"N"	"L"	"T"	"M"
## Glyma.11G227300.1	"V"	"Q"	"G"	"Y"	"G"	"V"	"S"	"P"
## Glyma.14G179500.1	"M"	"S"	"S"	"C"	"K"	"L"	"T"	"R"
## Glyma.14G062100.1	"V"	"Q"	"G"	"Y"	"G"	"A"	"S"	"P"
## Glyma.06G095400.1	"M"	"S"	"S"	"C"	"K"	"L"	"T"	"R"
## Glyma.03G209400.1	"M"	"S"	"S"	"C"	"S"	"V"	"S"	"Y"
## Glyma.19G100200.1	"M"	"S"	"S"	"C"	"E"	"V"	"T"	"V"
## Glyma.19G206800.1	"M"	"S"	"S"	"C"	"S"	"V"	"S"	"Y"
## Glyma.18G030200.1	"V"	"Q"	"G"	"Y"	"G"	"V"	"S"	"P"
##	[,518]	[,519]	[,520]	[,521]	[,522]	[,523]	[,524]	[,525]
## Glyma.16G050500.1	"G"	"A"	"C"	"K"	"L"	"L"	"A"	"K"
## Glyma.16G146400.1	"G"	"A"	"C"	"K"	"A"	"L"	"A"	"E"
## Glyma.04G093500.1	"Q"	"A"	"C"	"R"	"E"	"V"	"A"	"R"
## Glyma.02G065300.1	"G"	"A"	"C"	"K"	"A"	"L"	"A"	"K"
## Glyma.02G211800.1	"Q"	"A"	"C"	"Q"	"E"	"V"	"A"	"R"
## Glyma.02G254300.1	"S"	"G"	"R"	"D"	"L"	"L"	"A"	"M"
## Glyma.02G152800.2	"G"	"A"	"C"	"K"	"L"	"L"	"G"	"Q"
## Glyma.10G021500.2	"G"	"A"	"C"	"K"	"L"	"L"	"G"	"Q"
## Glyma.08G059500.6	"N"	"G"	"V"	"R"	"L"	"L"	"A"	"K"
## Glyma.07G189800.5	"N"	"G"	"V"	"R"	"L"	"L"	"A"	"Q"
## Glyma.11G227300.1	"S"	"G"	"R"	"D"	"L"	"L"	"A"	"M"
## Glyma.14G179500.1	"Q"	"A"	"C"	"Q"	"E"	"V"	"A"	"R"
## Glyma.14G062100.1	"S"	"G"	"R"	"D"	"L"	"L"	"A"	"M"
## Glyma.06G095400.1	"Q"	"A"	"C"	"Q"	"E"	"V"	"A"	"Q"
## Glyma.03G209400.1	"G"	"A"	"C"	"K"	"L"	"L"	"G"	"Q"
## Glyma.19G100200.1	"G"	"A"	"C"	"K"	"L"	"L"	"A"	"K"
## Glyma.19G206800.1	"G"	"A"	"C"	"K"	"L"	"L"	"G"	"Q"

## Glyma.18G030200.1	"S"	"G"	"R"	"D"	"L"	"L"	"V"	"M"
##	[,526]	[,527]	[,528]	[,529]	[,530]	[,531]	[,532]	[,533]
## Glyma.16G050500.1	"K"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.16G146400.1	"K"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.04G093500.1	"M"	"L"	"P"	"H"	"L"	"V"	"L"	"E"
## Glyma.02G065300.1	"K"	"M"	"P"	"G"	"L"	"N"	"V"	"E"
## Glyma.02G211800.1	"V"	"L"	"P"	"N"	"L"	"V"	"F"	"E"
## Glyma.02G254300.1	"A"	"R"	"P"	"Y"	"W"	"N"	"I"	"E"
## Glyma.02G152800.2	"K"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.10G021500.2	"K"	"L"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.08G059500.6	"E"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.07G189800.5	"E"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.11G227300.1	"A"	"R"	"P"	"F"	"W"	"N"	"I"	"E"
## Glyma.14G179500.1	"A"	"L"	"P"	"N"	"L"	"V"	"L"	"E"
## Glyma.14G062100.1	"A"	"R"	"P"	"Y"	"W"	"N"	"I"	"E"
## Glyma.06G095400.1	"T"	"L"	"P"	"H"	"L"	"V"	"L"	"E"
## Glyma.03G209400.1	"K"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.19G100200.1	"K"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.19G206800.1	"K"	"M"	"P"	"R"	"L"	"N"	"V"	"E"
## Glyma.18G030200.1	"A"	"R"	"P"	"F"	"W"	"N"	"I"	"E"
##	[,534]	[,535]	[,536]	[,537]	[,538]	[,539]	[,540]	[,541]
## Glyma.16G050500.1	"I"	"F"	"N"	"E"	"N"	"E"	"Q"	"E"
## Glyma.16G146400.1	"I"	"F"	"N"	"E"	"N"	"K"	"K"	"V"
## Glyma.04G093500.1	"V"	"I"	"N"	"_"	"_"	"S"	"E"	"E"
## Glyma.02G065300.1	"I"	"F"	"N"	"G"	"N"	"E"	"K"	"V"
## Glyma.02G211800.1	"V"	"I"	"N"	"N"	"N"	"S"	"E"	"E"
## Glyma.02G254300.1	"L"	"I"	"P"	"S"	"R"	"S"	"V"	"V"
## Glyma.02G152800.2	"V"	"I"	"D"	"E"	"R"	"G"	"P"	"P"
## Glyma.10G021500.2	"V"	"I"	"D"	"E"	"R"	"G"	"P"	"P"
## Glyma.08G059500.6	"V"	"I"	"K"	"E"	"E"	"T"	"Y"	"E"
## Glyma.07G189800.5	"V"	"I"	"K"	"E"	"E"	"S"	"Y"	"E"
## Glyma.11G227300.1	"L"	"I"	"P"	"S"	"R"	"K"	"V"	"A"
## Glyma.14G179500.1	"V"	"I"	"N"	"N"	"N"	"N"	"E"	"E"
## Glyma.14G062100.1	"L"	"I"	"P"	"S"	"R"	"R"	"V"	"V"
## Glyma.06G095400.1	"V"	"I"	"N"	"_"	"_"	"S"	"E"	"E"
## Glyma.03G209400.1	"V"	"I"	"D"	"E"	"R"	"G"	"P"	"P"
## Glyma.19G100200.1	"I"	"F"	"N"	"E"	"N"	"E"	"Q"	"E"
## Glyma.19G206800.1	"V"	"I"	"D"	"E"	"R"	"G"	"P"	"P"
## Glyma.18G030200.1	"L"	"I"	"P"	"S"	"R"	"K"	"V"	"A"
##	[,542]	[,543]	[,544]	[,545]	[,546]	[,547]	[,548]	[,549]
## Glyma.16G050500.1	"D"	"C"	"S"	"Q"	"K"	"V"	"E"	"K"
## Glyma.16G146400.1	"D"	"R"	"D"	"Q"	"K"	"V"	"E"	"K"
## Glyma.04G093500.1	"D"	"K"	"A"	"D"	"D"	"I"	"E"	"I"
## Glyma.02G065300.1	"D"	"R"	"D"	"Q"	"K"	"V"	"E"	"K"
## Glyma.02G211800.1	"N"	"A"	"G"	"D"	"E"	"V"	"E"	"T"
## Glyma.02G254300.1	"V"	"S"	"N"	"E"	"H"	"L"	"A"	"H"
## Glyma.02G152800.2	"D"	"S"	"R"	"S"	"P"	"V"	"E"	"K"
## Glyma.10G021500.2	"D"	"S"	"R"	"S"	"P"	"V"	"E"	"K"
## Glyma.08G059500.6	"T"	"_"	"_"	"H"	"Q"	"A"	"K"	"K"
## Glyma.07G189800.5	"T"	"_"	"_"	"H"	"Q"	"A"	"K"	"K"
## Glyma.11G227300.1	"M"	"N"	"T"	"E"	"H"	"P"	"A"	"H"
## Glyma.14G179500.1	"N"	"A"	"G"	"D"	"E"	"V"	"E"	"T"
## Glyma.14G062100.1	"V"	"S"	"N"	"E"	"H"	"P"	"A"	"H"
## Glyma.06G095400.1	"D"	"K"	"A"	"D"	"G"	"I"	"E"	"I"

## Glyma.03G209400.1	"D"	"S"	"R"	"C"	"P"	"V"	"E"	"K"
## Glyma.19G100200.1	"D"	"C"	"S"	"Q"	"K"	"V"	"E"	"K"
## Glyma.19G206800.1	"D"	"S"	"R"	"C"	"P"	"V"	"E"	"K"
## Glyma.18G030200.1	"T"	"N"	"T"	"E"	"H"	"P"	"A"	"H"
##	[,550]	[,551]	[,552]	[,553]	[,554]	[,555]	[,556]	[,557]
## Glyma.16G050500.1	"M"	"Y"	"L"	"Y"	"R"	"T"	"L"	"A"
## Glyma.16G146400.1	"M"	"Y"	"L"	"Y"	"R"	"T"	"L"	"A"
## Glyma.04G093500.1	"L"	"Y"	"M"	"Y"	"R"	"S"	"L"	"D"
## Glyma.02G065300.1	"T"	"Y"	"L"	"Y"	"R"	"T"	"L"	"V"
## Glyma.02G211800.1	"L"	"Y"	"M"	"Y"	"R"	"S"	"L"	"D"
## Glyma.02G254300.1	"I"	"L"	"A"	"Y"	"Y"	"S"	"L"	"A"
## Glyma.02G152800.2	"L"	"Y"	"I"	"Y"	"R"	"T"	"V"	"S"
## Glyma.10G021500.2	"L"	"Y"	"M"	"Y"	"R"	"T"	"V"	"S"
## Glyma.08G059500.6	"V"	"Y"	"V"	"Y"	"R"	"S"	"V"	"A"
## Glyma.07G189800.5	"V"	"Y"	"V"	"Y"	"R"	"S"	"V"	"A"
## Glyma.11G227300.1	"I"	"L"	"A"	"Y"	"Y"	"S"	"L"	"A"
## Glyma.14G179500.1	"L"	"Y"	"M"	"Y"	"R"	"S"	"L"	"D"
## Glyma.14G062100.1	"I"	"L"	"A"	"Y"	"Y"	"S"	"L"	"A"
## Glyma.06G095400.1	"L"	"Y"	"M"	"Y"	"R"	"S"	"L"	"D"
## Glyma.03G209400.1	"L"	"Y"	"I"	"Y"	"R"	"T"	"V"	"A"
## Glyma.19G100200.1	"M"	"Y"	"L"	"Y"	"R"	"T"	"L"	"A"
## Glyma.19G206800.1	"L"	"Y"	"I"	"Y"	"R"	"T"	"I"	"A"
## Glyma.18G030200.1	"I"	"L"	"A"	"Y"	"Y"	"S"	"L"	"A"
##	[,558]	[,559]	[,560]	[,561]	[,562]	[,563]	[,564]	[,565]
## Glyma.16G050500.1	"G"	"K"	"R"	"K"	"D"	"A"	"P"	"E"
## Glyma.16G146400.1	"G"	"R"	"R"	"K"	"D"	"A"	"P"	"E"
## Glyma.04G093500.1	"R"	"P"	"R"	"D"	"D"	"A"	"P"	"K"
## Glyma.02G065300.1	"G"	"R"	"R"	"K"	"D"	"A"	"P"	"E"
## Glyma.02G211800.1	"G"	"P"	"R"	"D"	"D"	"A"	"P"	"R"
## Glyma.02G254300.1	"G"	"P"	"R"	"T"	"D"	"F"	"P"	"D"
## Glyma.02G152800.2	"G"	"P"	"R"	"L"	"D"	"M"	"P"	"G"
## Glyma.10G021500.2	"G"	"P"	"R"	"L"	"D"	"M"	"P"	"G"
## Glyma.08G059500.6	"G"	"P"	"R"	"R"	"D"	"A"	"P"	"P"
## Glyma.07G189800.5	"G"	"P"	"R"	"R"	"D"	"A"	"P"	"P"
## Glyma.11G227300.1	"G"	"Q"	"R"	"S"	"D"	"F"	"P"	"D"
## Glyma.14G179500.1	"G"	"P"	"R"	"D"	"D"	"A"	"P"	"R"
## Glyma.14G062100.1	"G"	"P"	"R"	"T"	"D"	"F"	"P"	"D"
## Glyma.06G095400.1	"G"	"P"	"R"	"D"	"D"	"A"	"P"	"K"
## Glyma.03G209400.1	"G"	"P"	"R"	"L"	"D"	"M"	"P"	"G"
## Glyma.19G100200.1	"G"	"K"	"R"	"K"	"D"	"A"	"P"	"E"
## Glyma.19G206800.1	"G"	"P"	"R"	"L"	"D"	"M"	"P"	"G"
## Glyma.18G030200.1	"G"	"Q"	"R"	"S"	"D"	"F"	"P"	"D"
##	[,566]	[,567]	[,568]	[,569]	[,570]	[,571]		
## Glyma.16G050500.1	"Y"	"V"	"W"	"T"	"L"	"*"		
## Glyma.16G146400.1	"L"	"V"	"W"	"T"	"L"	"*"		
## Glyma.04G093500.1	"V"	"V"	"T"	"I"	"L"	"*"		
## Glyma.02G065300.1	"H"	"V"	"W"	"T"	"L"	"*"		
## Glyma.02G211800.1	"F"	"V"	"T"	"I"	"L"	"*"		
## Glyma.02G254300.1	"T"	"V"	"I"	"P"	"L"	"*"		
## Glyma.02G152800.2	"Y"	"V"	"W"	"R"	"M"	"*"		
## Glyma.10G021500.2	"Y"	"V"	"W"	"R"	"M"	"*"		
## Glyma.08G059500.6	"F"	"V"	"L"	"T"	"L"	"*"		
## Glyma.07G189800.5	"F"	"V"	"L"	"T"	"L"	"*"		
## Glyma.11G227300.1	"T"	"V"	"V"	"P"	"L"	"*"		



```
## Glyma.14G179500.1 "F"      "V"      "T"      "I"      "L"      "*"
## Glyma.14G062100.1 "T"      "V"      "I"      "P"      "L"      "*"
## Glyma.06G095400.1 "V"      "V"      "T"      "I"      "L"      "*"
## Glyma.03G209400.1 "F"      "V"      "W"      "T"      "M"      "*"
## Glyma.19G100200.1 "Y"      "V"      "W"      "T"      "L"      "*"
## Glyma.19G206800.1 "F"      "V"      "W"      "T"      "M"      "*"
## Glyma.18G030200.1 "T"      "V"      "V"      "P"      "L"      "*"
```

```
# Also remove all the * from sequences because MrBayes does not like it
matrix_AFB <- sub("\\*", "-", matrix_AFB)
```

```
DECIPHER::BrowseSeqs(AAStringSet(matrix_AFB))
```

```
# Write nex file for MrBayes analysis
ape::write.nexus.data(matrix_AFB, format = "protein",
                      file = "AFB_input/AFB_final.nex",
                      datablock = TRUE,
                      interleaved = TRUE,
                      charsperline = NULL,
                      gap = NULL, missing = NULL)
```

```
# MBayes steps was the same as specified in the AFB_final.Rmd file. Outgroup was GLYMA_11G227300 (GmC01
# mcmc stopval=0.01 stoprule=yes Mcmcdiag=yes nchains=4 printfreq=1000 samplefreq=100 burnin=200 relb
```

## Build tree with multiple family

I will be adding L. japonicus, G. max, G. soja, A. thaliana, Medicago truncatula

```
Mult_AFB <- Biostrings::readAAStringSet(filepath = "AFB_input/AFB_multi.fasta")
Mult_AFB <- unique(Mult_AFB)
names(Mult_AFB)
```

```
## [1] "Medtr3g070140|Medtr3g070140.1"
## [2] "Medtr3g100860|Medtr3g100860.1"
## [3] "Medtr8g098695|Medtr8g098695.2"
## [4] "Medtr7g083610|Medtr7g083610.1"
## [5] "Medtr5g065490|Medtr5g065490.1"
## [6] "Medtr5g081870|Medtr5g081870.1"
## [7] "Medtr1g088950|Medtr1g088950.1"
## [8] "Medtr4g091290|Medtr4g091290.1"
## [9] "AT2G39940|AT2G39940.1"
## [10] "AT4G03190|AT4G03190.1"
## [11] "AT4G24390|AT4G24390.1"
## [12] "AT1G12820|AT1G12820.1"
## [13] "AT3G26810|AT3G26810.1"
## [14] "AT3G62980|AT3G62980.1"
## [15] "AT5G49980|AT5G49980.1"
## [16] "Glyma.16G050500|Glyma.16G050500.1"
## [17] "Glyma.16G146400|Glyma.16G146400.1"
## [18] "Glyma.04G093500|Glyma.04G093500.1"
## [19] "Glyma.02G065300|Glyma.02G065300.1"
## [20] "Glyma.02G211800|Glyma.02G211800.1"
## [21] "Glyma.02G254300|Glyma.02G254300.1"
## [22] "Glyma.02G152800|Glyma.02G152800.2"
```

```

## [23] "Glyma.02G152800|Glyma.02G152800.3"
## [24] "Glyma.10G021500|Glyma.10G021500.2"
## [25] "Glyma.08G059500|Glyma.08G059500.6"
## [26] "Glyma.07G189800|Glyma.07G189800.5"
## [27] "Glyma.11G227300|Glyma.11G227300.1"
## [28] "Glyma.14G179500|Glyma.14G179500.1"
## [29] "Glyma.14G062100|Glyma.14G062100.1"
## [30] "Glyma.06G095400|Glyma.06G095400.1"
## [31] "Glyma.03G209400|Glyma.03G209400.1"
## [32] "Glyma.19G100200|Glyma.19G100200.1"
## [33] "Glyma.19G206800|Glyma.19G206800.1"
## [34] "Glyma.18G030200|Glyma.18G030200.1"
## [35] "GlysoPI483463.19G168300|GlysoPI483463.19G168300.1"
## [36] "GlysoPI483463.03G169700|GlysoPI483463.03G169700.1"
## [37] "GlysoPI483463.14G141800|GlysoPI483463.14G141800.1"
## [38] "GlysoPI483463.02G212600|GlysoPI483463.02G212600.1"
## [39] "GlysoPI483463.02G175100|GlysoPI483463.02G175100.1"
## [40] "Lj1g0006513|Lj1g0006513.1"
## [41] "Lj1g0027142|Lj1g0027142.1"
## [42] "Lj1g0015670|Lj1g0015670.1"
## [43] "Lj3g0010583|Lj3g0010583.1"
## [44] "Lj5g0004781|Lj5g0004781.1"
## [45] "LjContig00357g0020173|LjContig00357g0020173.1"
## [46] "Lj6g0020395|Lj6g0020395.1"
## [47] "Lj4g0012889|Lj4g0012889.1"
## [48] "LOC_0s04g32460|LOC_0s04g32460.1"
## [49] "LOC_0s01g63420|LOC_0s01g63420.1"
## [50] "LOC_0s01g63420|LOC_0s01g63420.2"
## [51] "LOC_0s01g63420|LOC_0s01g63420.3"
## [52] "LOC_0s03g08850|LOC_0s03g08850.1"
## [53] "LOC_0s03g15880|LOC_0s03g15880.4"
## [54] "LOC_0s02g52230|LOC_0s02g52230.1"
## [55] "LOC_0s05g37690|LOC_0s05g37690.1"
## [56] "LOC_0s05g05800|LOC_0s05g05800.1"
## [57] "LOC_0s11g31620|LOC_0s11g31620.1"

phylotools::rm.sequence.fasta(infile = "AFB_input/AFB_multi.fasta", to.rm =c("Glyma.02G152800|Glyma.02G152800.3",
"Warning in readLines(file): incomplete final line found on
'AFB_input/AFB_multi.fasta'

## AFB_input/AFB_multi_removed.fasta has been saved to /Users/deisianyneres/GitHub/SoyArcs_manuscript/

Mult_AFB2 <- Biostrings::readAAStringSet(filepath = "AFB_input/AFB_multi_removed.fasta")
names(Mult_AFB2)

## [1] "Medtr3g070140|Medtr3g070140.1"
## [2] "Medtr3g100860|Medtr3g100860.1"
## [3] "Medtr8g098695|Medtr8g098695.2"
## [4] "Medtr7g083610|Medtr7g083610.1"
## [5] "Medtr5g065490|Medtr5g065490.1"
## [6] "Medtr5g081870|Medtr5g081870.1"
## [7] "Medtr1g088950|Medtr1g088950.1"
## [8] "Medtr4g091290|Medtr4g091290.1"
## [9] "AT2G39940|AT2G39940.1"
## [10] "AT4G03190|AT4G03190.1"

```

```
## [11] "AT4G24390|AT4G24390.1"
## [12] "AT1G12820|AT1G12820.1"
## [13] "AT3G26810|AT3G26810.1"
## [14] "AT3G62980|AT3G62980.1"
## [15] "AT5G49980|AT5G49980.1"
## [16] "Glyma.16G050500|Glyma.16G050500.1"
## [17] "Glyma.16G146400|Glyma.16G146400.1"
## [18] "Glyma.04G093500|Glyma.04G093500.1"
## [19] "Glyma.02G065300|Glyma.02G065300.1"
## [20] "Glyma.02G211800|Glyma.02G211800.1"
## [21] "Glyma.02G254300|Glyma.02G254300.1"
## [22] "Glyma.02G152800|Glyma.02G152800.2"
## [23] "Glyma.10G021500|Glyma.10G021500.2"
## [24] "Glyma.08G059500|Glyma.08G059500.6"
## [25] "Glyma.07G189800|Glyma.07G189800.5"
## [26] "Glyma.11G227300|Glyma.11G227300.1"
## [27] "Glyma.14G179500|Glyma.14G179500.1"
## [28] "Glyma.14G062100|Glyma.14G062100.1"
## [29] "Glyma.06G095400|Glyma.06G095400.1"
## [30] "Glyma.03G209400|Glyma.03G209400.1"
## [31] "Glyma.19G100200|Glyma.19G100200.1"
## [32] "Glyma.19G206800|Glyma.19G206800.1"
## [33] "Glyma.18G030200|Glyma.18G030200.1"
## [34] "GlysoPI483463.19G168300|GlysoPI483463.19G168300.1"
## [35] "GlysoPI483463.03G169700|GlysoPI483463.03G169700.1"
## [36] "GlysoPI483463.14G141800|GlysoPI483463.14G141800.1"
## [37] "GlysoPI483463.02G212600|GlysoPI483463.02G212600.1"
## [38] "GlysoPI483463.02G175100|GlysoPI483463.02G175100.1"
## [39] "Lj1g0006513|Lj1g0006513.1"
## [40] "Lj1g0027142|Lj1g0027142.1"
## [41] "Lj1g0015670|Lj1g0015670.1"
## [42] "Lj3g0010583|Lj3g0010583.1"
## [43] "Lj5g0004781|Lj5g0004781.1"
## [44] "Lj6g0020395|Lj6g0020395.1"
## [45] "Lj4g0012889|Lj4g0012889.1"
## [46] "LOC_0s04g32460|LOC_0s04g32460.1"
## [47] "LOC_0s04g32460|LOC_0s04g32460.2"
## [48] "LOC_0s01g63420|LOC_0s01g63420.1"
## [49] "LOC_0s03g08850|LOC_0s03g08850.1"
## [50] "LOC_0s03g15880|LOC_0s03g15880.4"
## [51] "LOC_0s03g15880|LOC_0s03g15880.5"
## [52] "LOC_0s03g15880|LOC_0s03g15880.3"
## [53] "LOC_0s03g15880|LOC_0s03g15880.2"
## [54] "LOC_0s03g15880|LOC_0s03g15880.1"
## [55] "LOC_0s02g52230|LOC_0s02g52230.1"
## [56] "LOC_0s05g37690|LOC_0s05g37690.1"
## [57] "LOC_0s05g05800|LOC_0s05g05800.1"
## [58] "LOC_0s11g31620|LOC_0s11g31620.1"
```

```
Mult_AFB2 <- unique(Mult_AFB2)
```

```
# Biostrings::writeXStringSet(Mult_AFB2, "AFB_input/multi_AFB_final.fasta")
```

```
# Load from here fasta for alignment without Rice (removed all rice sequences manually)
```

```

Mult_AFB2 <- Biostrings::readAAStringSet(filepath = "AFB_input/")

## Warning in file(fp): 'raw = FALSE' but 'AFB_input/' is not a regular file
## Warning in file(fp): 'raw = FALSE' but 'AFB_input/' is not a regular file
names(Mult_AFB2)

## character(0)
Mult_AFB2 <- unique(Mult_AFB2)

# make alignment
(Mult_AFB_aligned <- DECIPHER::AlignSeqs(Mult_AFB2))

## Error in DECIPHER::AlignSeqs(Mult_AFB2): At least two sequences are required in myXStringSet.
# Fix names, because characters such as / are not accepted in MrBayes
names(Mult_AFB_aligned) <- stringr::str_remove(names(Mult_AFB_aligned), ".*\\|") # remove everything be

## Error in eval(expr, envir, enclos): object 'Mult_AFB_aligned' not found
# Biostrings::writeXStringSet(Mult_AFB2, "AFB_input/multi_AFB_aligned.fasta")

# write.csv(names(Mult_AFB_aligned), "AFB_IDS_all.csv")

# CUT low alignment regions by masking alignment
(multAFB_masked <- DECIPHER::MaskAlignment(Mult_AFB_aligned,
  type = "sequences",
  windowSize = 6, # same as aliscore default value.
  threshold = 1,
  maxFractionGaps = 0.2,
  includeTerminalGaps = FALSE,
  correction = FALSE,
  showPlot = FALSE))

## Error in eval(expr, envir, enclos): object 'Mult_AFB_aligned' not found
# make it a AAStringSet
AAstr_multAFB <- as(multAFB_masked, "AAStringSet")

## Error in eval(expr, envir, enclos): object 'multAFB_masked' not found
BrowseSeqs(AAstr_multAFB) # A few sequences seems really small, however I believe they will probably be

## Error in eval(expr, envir, enclos): object 'AAstr_multAFB' not found
writeXStringSet(AAstr_multAFB, file="AFB_input/masked_multAFB.fasta" )

## Error in eval(expr, envir, enclos): object 'AAstr_multAFB' not found
#Create a matrix so we can create the nexus file next
(matrix_multAFB <- as.matrix(AAstr_multAFB, use.names=TRUE))

## Error in h(simpleError(msg, call)): error in evaluating the argument 'x' in selecting a method for f
# Also remove all the * from sequences because MrBayes does not like it
matrix_multAFB <- sub("\\*", "-", matrix_multAFB)

## Error in h(simpleError(msg, call)): error in evaluating the argument 'x' in selecting a method for f

```

```
DECIPHER::BrowseSeqs(AAStringSet(matrix_multAFB))
```

```
## Error in h(simpleError(msg, call)): error in evaluating the argument 'x' in selecting a method for function h
```

```
# Write nex file for MrBayes analysis
```

```
ape::write.nexus.data(matrix_multAFB, format = "protein",  
  file = "AFB_input/AFBmult_final.nex",  
  datablock = TRUE,  
  interleaved = TRUE,  
  charsperline = NULL,  
  gap = NULL, missing = NULL)
```

```
## Error in eval(expr, envir, enclos): object 'matrix_multAFB' not found
```

```
# Mbayes analysis were done following steps in AFB_Final.Rmd script
```

```
./mb
```

```
execute AFB_final.nex
```

```
# set as the outgroup
```

```
outgroup AT2G39940.1
```

```
lset nucmodel=protein Rates=invgamma
```

```
# or include Nst=6 ngammacat=4 omegavar=M3
```

```
prset aamodelpr=fixed(jones) statefreqpr=fixed(empirical)
```

```
# initial run with aamodelpr=mixed yielded posterior prob of 1 for jones model
```

```
propset ExtTBR$prob=0
```

```
mcmc stopval=0.01 stoprule=yes Mcmcdiag=yes nchains=6 printfreq=1000 samplefreq=100 burnin=200 relburnin=0.01
```

```
## bash: ./mb: No such file or directory
```

```
## bash: line 2: execute: command not found
```

```
## bash: line 4: outgroup: command not found
```

```
## bash: line 6: lset: command not found
```

```
## bash: -c: line 8: syntax error near unexpected token `('
```

```
## bash: -c: line 8: `prset aamodelpr=fixed(jones) statefreqpr=fixed(empirical) '
```

```
mcmc
```

```
sump
```

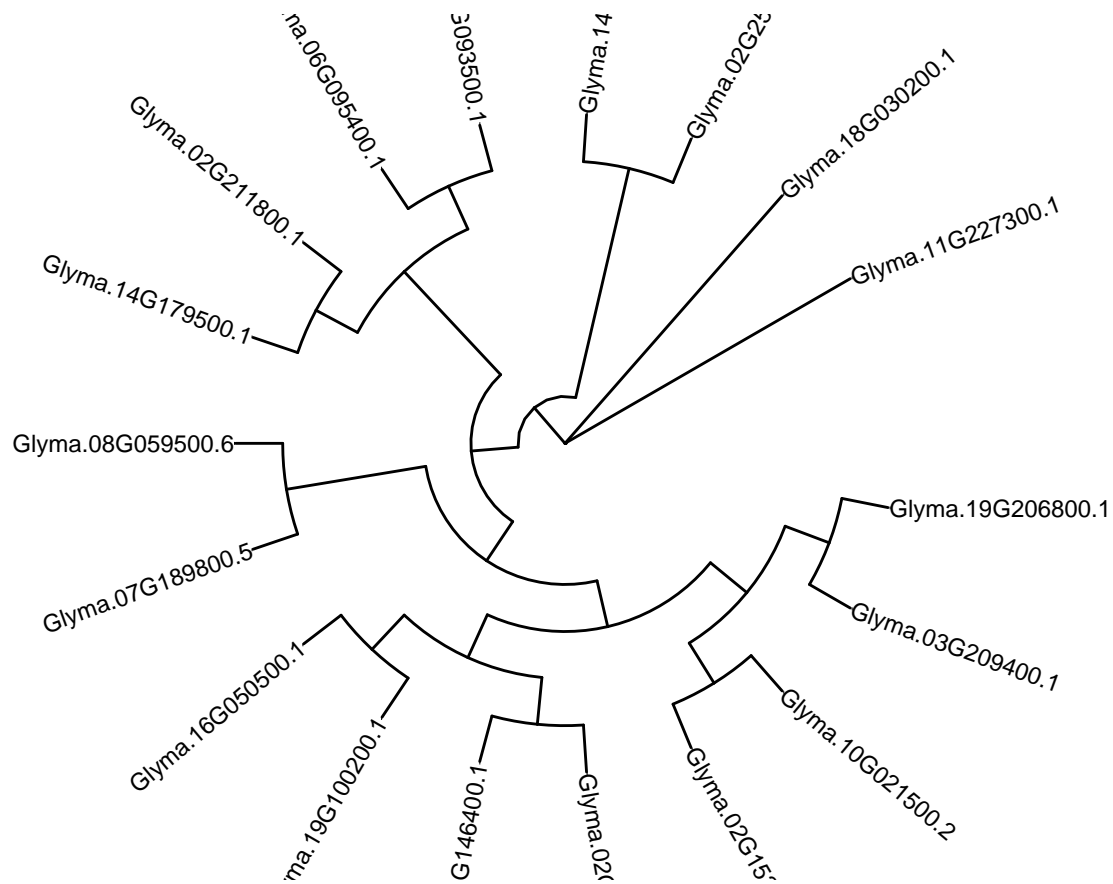
```
sumt
```

```
## Error in running command bash
```

## build G. max tree

```
tree <- treeio::read.mrbayes(file = "AFB_input/mb_results/AFB_final.nex.con.tre")
```

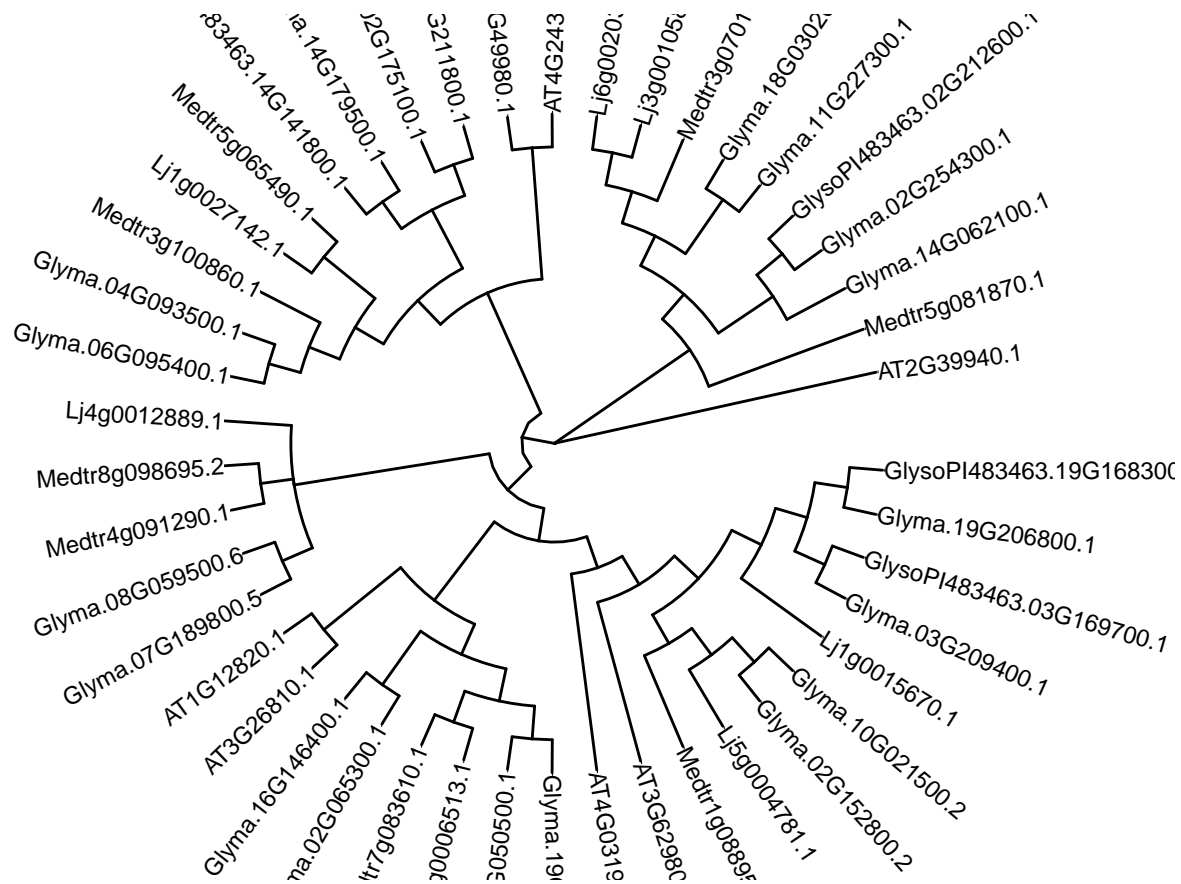
```
ggtree(tree, layout="circular", branch.length = 'none') + geom_tiplab(size=3, align = T)
```



```
# ggsave("AFB_input/figures/g_max_mb_tree.pdf", height = 8, width = 7)
```

## Build G. max tree with multiple species.

```
tree <- treeio::read.mrbayes(file = "AFB_input/mb_results_mult/AFBmult_final.nex.con.tre")
g_tree <- ggtree(tree, layout="circular", branch.length = 'none') + geom_tiplab(size=3, align = T)
g_tree
```



```
ggtree(tree, layout="circular", color = "darkgrey",
        branch.length='prob', ladderize = TRUE) + geom_tiplab(align = F) +
geom_text(aes(label=node), # node numbers
          size=3, hjust=-.3, color="red", fontface=2)
```





```

class_TIR1_AFB1 <- data.frame(get_taxa_name(g_tree, node = 72)) # TIR1_AFB1
class_TIR1_AFB1 <- class_TIR1_AFB1 %>%
  dplyr::rename(tair_locus = `get_taxa_name.g_tree..node...72.`) %>%
  mutate(class = "TIR1/AFB1",
         name = ifelse(grepl("Glyma", .$tair_locus), "GmTIR1/AFB1", ""),
         Clade = "I")

TIR_AFB_class <- rbind(class_TIR1_AFB1, class_AFB2_3, class_AFB4_5, class_AFB6, class_COI1)

# Assign orthology
TIR_AFB_class <- TIR_AFB_class %>% mutate(ensembl_gene_id = str_remove_all(.$tair_locus, "\\.[^.]*$"))

Gm_ortholog <- TIR_AFB_class %>%
  dplyr::filter(str_detect(ensembl_gene_id, "Glyma")) %>%
  # arrange by tree order.
  group_by(name) %>% unique() %>%
  mutate(ortholog_name = dplyr::row_number()) %>%
  mutate(ortholog_name = paste0(name, "_", LETTERS[ortholog_name]))

#write_csv(Gm_ortholog, "AFB_input/AFB_Gm_ortholog.csv")

df4tips <- left_join(TIR_AFB_class, Gm_ortholog)

## Joining with `by = join_by(tair_locus, class, name, Clade, ensembl_gene_id)`
df4tips[c(27:28), 5] <- c("MtAFB6") #Assign AFB6 to IDs
df4tips[29, 5] <- c("LjAFB6")

df4tips2 <- within(df4tips, ortholog_name[is.na(ortholog_name)] <- tair_locus[is.na(ortholog_name)])

```

Now I want to join dataframe with nodes classification and tips and arabidopsis classification according to ViVa tool paper.

```

# change tips according to Arabidopsis classification
names <- as.data.frame(tree@phylo[["tip.label"]]) %>% dplyr::rename(tair_locus = `tree@phylo[["tip.label"]])

IDs <- readxl::read_xls("../Arquive/AFB_files/all_IDs.xls") %>%
  dplyr::rename(tair_locus = gene_IDs)
IDs

```

```

## # A tibble: 47 x 3
##   tair_locus      names      Source
##   <chr>          <chr>      <chr>
## 1 Medtr3g070140.1 Medtr3g070140.1 <NA>
## 2 Medtr3g100860.1 Medtr3g100860.1 <NA>
## 3 Medtr8g098695.2 Medtr8g098695.2/AFB6 https://www.ncbi.nlm.nih.gov/pmc/articl~
## 4 Medtr7g083610.1 Medtr7g083610.1 <NA>
## 5 Medtr5g065490.1 Medtr5g065490.1 <NA>
## 6 Medtr5g081870.1 Medtr5g081870.1 <NA>
## 7 Medtr1g088950.1 Medtr1g088950.1 <NA>
## 8 Medtr4g091290.1 Medtr4g091290.1 <NA>
## 9 AT2G39940.1      AT2G39940/COI1    <NA>
## 10 AT4G03190.1      AT4G03190/AFB1    <NA>
## # i 37 more rows

```

```

all_Tip_IDs <- left_join(names, IDs[,-3]) %>%
  mutate(ensembl_gene_id = str_remove_all(tair_locus, "\\.[^.]*$"))

## Joining with `by = join_by(tair_locus)`
TipLabel <- left_join(all_Tip_IDs, df4tips2)

## Joining with `by = join_by(tair_locus, ensembl_gene_id)`
TipLabel <- TipLabel %>% dplyr::rename(name_ID = names) %>% as.data.frame()

TipLabel2 <- cbind(TipLabel, replicate(1, TipLabel$name_ID)) %>%
  dplyr::rename(Tips_ID = `replicate(1, TipLabel$name_ID)`)

TipLabel2$name_ID <- gsub( ".*/", "At", TipLabel2$name_ID)

TipLabel2 <- within(TipLabel2, class[is.na(class)] <-
  gsub("At", "", name_ID[is.na(class)]))

TipLabel2[c(3, 39),2] <- c("MtAFB6", "LjAFB6") # correct row 3 and 39 back to what they are. they are n

TipLabel2[c(1:2, 4:7, 915, 34:38, 40:45), 7] <- NA #ortholog_name column

# Join ortholog name with name_ID
TipLabel3 <-
  TipLabel2 %>% dplyr::relocate(ortholog_name, .after = name_ID) %>%
  unite("New_label",
    ortholog_name:name_ID,
    sep = "_",
    na.rm = T) %>% .[, -c(5:6)]

# make a clade column for make visualization of tree easier to readers
TipLabel3 <- TipLabel3 %>%
  mutate(
    Clade = ifelse(grepl("COI", .$New_label), "V",
      ifelse(
        grepl("AFB1", .$New_label), "I",
        ifelse(grepl("TIR1", .$New_label), "I",
          ifelse(
            grepl("AFB2", .$New_label), "II",
            ifelse(grepl("AFB3", .$New_label), "II",
              ifelse(
                grepl("AFB4", .$New_label), "IV",
                ifelse(
                  grepl("AFB5", .$New_label),
                    "IV",
                    ifelse(grepl("AFB6", .$New_label), "III",
                      grepl("Medtr4g091290.1", .$New_label), "III", Clade
                    )
                )
              )
            )
          )
        )
      )
    ),
    class = ifelse(

```

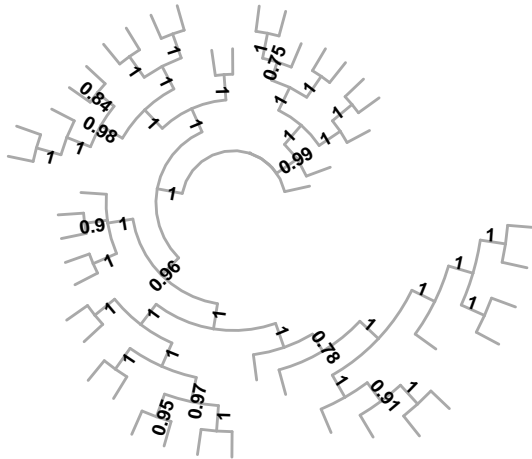


```

    branch.length='prob', ladderize = TRUE) +
  scale_x_continuous(expand = c(0, 1.2)) +
  geom_nodelab(mapping = aes(label=round(
    as.numeric(prob), digits = 2), fontface=2),
    nudge_x = -.9, nudge_y = .2, hjust = -.05, size = 2.5)

```

p

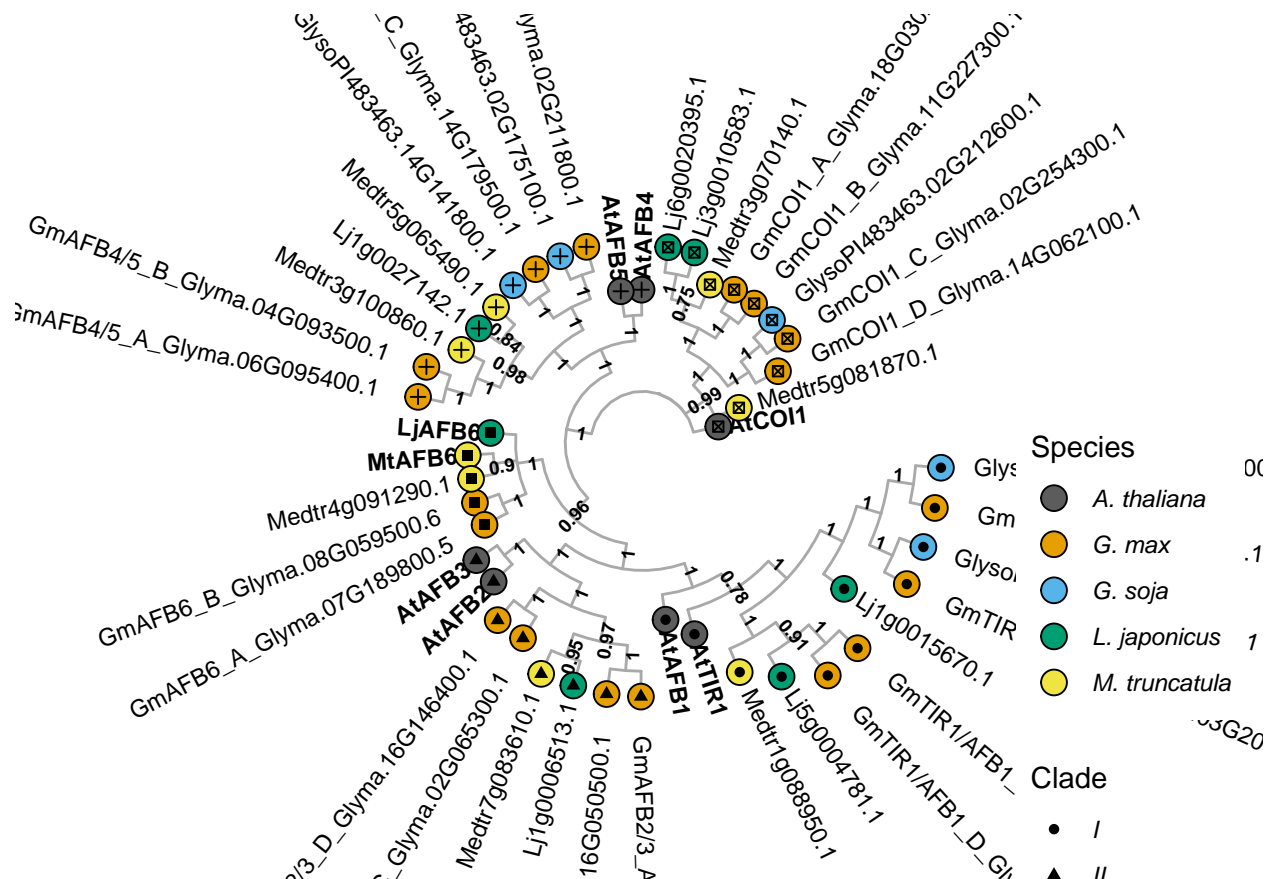


```

(p2 <- p %<+% TipLabel4[1:45,-1] +
  geom_tippoint(aes(fill=factor(Species)),
    size=4.2, shape=21) + geom_tiplab( size=3, hjust=-.15) +
  scale_fill_manual(values =
    c('#000000', '#E69F00', '#56B4E9', '#009E73', '#F0E442')) + #COLORS FROM GGTHERMES
  guides(fill=guide_legend(title="Species"))
)

```





```
ggsave("AFB_input/figures/AFB_FINAL.pdf", height = 11, width = 14, dpi = 1000)
```

```
## Warning: Removed 42 rows containing missing values (`geom_point()`).
```

```
# save in png
p %<+% tipLabel5[1:45,-1] +
  geom_tiplab(aes(subset=tip_color == "T0_not"), size=3, hjust=-.1, fontface=2) + #subset tree in order
  geom_tiplab(aes(subset=tip_color == "T0_color"),
    colour='black', size=4, hjust=-.1) +
  geom_tippoint(aes(fill=factor(Species)),
    size=4.2, shape =21) + #geom_tiplab( size=3, hjust=-.15) +
  scale_fill_manual(values =
    c('grey36', '#E69F00', '#56B4E9', '#009E73', '#F0E442')) + #COLORS FROM GGTHEMES
  guides(fill=guide_legend(title="Species")) +
  theme(legend.position = c(1.10, 0.10)) +
  geom_point(aes(shape = Clade)) +
  theme(legend.text = element_text(face = "italic"))
```

```
## Warning: Removed 42 rows containing missing values (`geom_point()`).
```

