

# R\_markdown\_iLand\_exploratory\_analysis.r

baldo

2023-06-14

```
# R wind plot events random - paper on recovery rate after disturbance (09.11.2022)

# R wind plot event in block - paper on recovery rate after disturbance (before editing)

library(RSQLite)
library(dplyr)

## Warning: package 'dplyr' was built under R version 4.2.3

##
## Attaching package: 'dplyr'

## The following objects are masked from 'package:stats':
##      filter, lag

## The following objects are masked from 'package:base':
##      intersect, setdiff, setequal, union

library(tidyr)
library(ggplot2)
library(gridExtra)

##
## Attaching package: 'gridExtra'

## The following object is masked from 'package:dplyr':
##      combine

library(fields)

## Loading required package: spam

## Spam version 2.9-1 (2022-08-07) is loaded.
## Type 'help( Spam)' or 'demo( spam)' for a short introduction
## and overview of this package.
## Help for individual functions is also obtained by adding the
## suffix '.spam' to the function name, e.g. 'help( chol.spam)'.
```

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##  

## Attaching package: 'spam'

## The following objects are masked from 'package:base':  

##  

##     backsolve, forwardsolve

## Loading required package: viridis

## Warning: package 'viridis' was built under R version 4.2.3

## Loading required package: viridisLite

##  

## Try help(fields) to get started.

# Path to search the data
dataroot <- ("C:/iLand/2023/browsing_revision/db_test/") # Root for the selection of the data

# CREATE NEW EMPTY DATAFRAME # this is needed to create a new database

# removals<-c()

lnd<-c()

aUnit<-c()

bb <-c()

w <- c()

damage.all<-c()

dys <- c()

variables.all <- c()

# FOR CYCLE FOR THE IMPORT AND ANALYSIS OF THE LANDSCAPE VOLUME AND VOLUME HARVESTED

all_v <- list.files(dataroot, ".sqlite") # alternative way to select all the files

for (i in (1:length(all_v))) { # We read in the files in the loop

  # "for" = for argument, "(in"= in, "1"= first element of the for cycle to analysis, ": length(cases))" = number of cases
  # PAY ATTENTION this part HERE is just FOR TESTING

  #i<-1 # to test but remember to don't run also the }

  # ORDINATION OF THE CASE TO IMPORT AS DATABASE see line 45 the loop will be as long as number of cases
}

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# Name of the database
file <- paste0(dataroot, all_v[i])                                # File to read here the case is
# "file"= name of the object, "paste0"+ function to create a NAME for a computer path of selection of
# case<- strsplit(all_v[i], ".s")[[1]][1]                      # Why we used it in the paper analysis? # explanation of
case<-all_v[i]

# Control
print(file)

# connect to the database of clear cut model
sqlite.driver <- dbDriver("SQLite")
db1 <- dbConnect(sqlite.driver, dbname = file) # connect to the file
tables.in.the.file<-dbListTables(db1)          # explore the tables in the file
print(tables.in.the.file)

#-----
# LOAD THE DATABASE

landscape <- dbReadTable(db1, "landscape")
abeUnit <- dbReadTable(db1, "abeUnit")
#abeStandRemoval <- dbReadTable(db1, "abeStandRemoval")
barkbeetle <- dbReadTable(db1, "barkbeetle")
wind <- dbReadTable(db1, "wind")
dynamicstand <- dbReadTable(db1, "dynamicstand")
carbon <- dbReadTable(db1, "carbon")
carbonflow <- dbReadTable(db1, "carbonflow")

dbDisconnect(db1)      # close the file

#-----
# CREATE SHANNON VARIABLE

annual.data<-landscape %>% group_by(year) %>% filter(year>0) %>% summarise(VOL.tot=sum(volume_m3), BA=BA/sum(BA))
annual.spec.data<-landscape %>% group_by(year, species) %>% filter(year>0) %>% summarise(VOL=(volume_m3*prop.BA)/sum(volume_m3*prop.BA))

print(head(annual.data))
print(head(annual.spec.data))

S<-landscape %>% group_by(year) %>% filter(volume_m3>0 & year>0) %>% summarise(n=n()) # number of species per year
print(S$n)

t<-annual.spec.data %>% right_join(annual.data, by="year") %>%
  mutate(prop.VOL=VOL/VOL.tot, prop.BA=BA/BA.tot, prop.count=count/count.tot) %>%
  filter(prop.VOL>0) # here I also filtering them to have only records with m3>0

#https://www.statology.org/shannon-diversity-index/
#https://www.statology.org/shannon-diversity-index-calculator/

```

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# Shannon diversity index (SDI): this is already Shannon.... that extra step by dividing the by S$n is
# so maybe we can make shannon based on BA, VOL and number of trees.... (? can discuss or save all 3)

H.BA<-t %>% group_by(year) %>% summarize(H=-1*sum(prop.BA*log(prop.BA)))
H.VOL<-t %>% group_by(year) %>% summarize(H=-1*sum(prop.VOL*log(prop.VOL)))
H.count<-t %>% group_by(year) %>% summarize(H=-1*sum(prop.count*log(prop.count))) # here I just put
set.panel(3,1)
plot(H.BA, main= "H.BA")
plot(H.VOL, main= "H.vol")
plot(H.count, main="H.count")

#The higher the value of H, the higher the diversity of species in a particular community.
#The lower the value of H, the lower the diversity.
#A value of H = 0 indicates a community that only has one species.

# Shannon Equitability Index (SEI).... maybe we do not need this at all.....
SEI<-H.BA$H/log(S$n)
SEI[which(is.na(SEI)==T)]<-0

#-----
# Pioneer species proportion based on BA
# CSP conifer species proportion

conifers <-c("pini", "pisy", "psme", "lade", "piab", "abal")

conif.spec.prop<-t %>% filter(species %in% conifers) %>% summarize(BA=sum(prop.BA))

csp<-data.frame(year=c(1:max(landscape$year)))
csp<-csp %>% left_join(conif.spec.prop, by="year")
csp$BA[which(is.na(csp$BA)==T)]<-0

# ESP

earlyspegs<-c("lade", "acps", "frex", "cabe", "bepe", "alin", "algl", "acca", "acpl", "soau", "soar", "coav", "alv")
early.spec.prop<-t %>% filter(species %in% earlyspegs) %>% summarize(BA=sum(prop.BA))

esp<-data.frame(year=c(1:max(landscape$year)))
esp<-esp %>% left_join(early.spec.prop, by="year")
esp$BA[which(is.na(esp$BA)==T)]<-0

#-----
# CREATE THE DATA FRAME FOR ADD VARIABLES ABOUT CARBON IN THE FINAL DATA FRAME

ab.tot.c<- data.frame(carbon %>%
                           group_by(year) %>%
                           summarise(tot_carbon=sum(stem_c, branch_c, foliage_c, coarseRoot_c, fineRoot_c)))

#-----
# CREATE NEE VARIABLE

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ab.nee.v<- data.frame(carbonflow %>%
                        group_by(year) %>%
                        summarise(NEE= (-1*(NPP + Rh))))


#-----
# CREATE THE CALCULATION FOR DAMAGES LOOK LINE 370

landscape.area<-landscape$area[1]                                     # CREATE THE VARIABLE F

lnd_volume = landscape %>% group_by(year)  %>%
  summarise(tot.vol = sum(volume_m3),                                         # CREATE THE SUMMARIZAT
            .groups = 'drop')

head(lnd_volume)

# WIND AND BARKBEETLE MERGING

head(barkbeetle)
head(wind)

damage <- data.frame(year=barkbeetle$year,                                # CREATE THE DATA FRAME
                      barkbeetle=barkbeetle$killedVolume)

# ADD WIND IMPACT IN THE DAMAGE DATA FRAME
damage<-left_join(damage,wind[,c(1,8)],by=("year"))
damage<-left_join(damage,lnd_volume,by=("year"))

# LEFT_JOIN IS A FUNCTION
# NB ...wind[,c(1,8)]...
# ADD THE LANDSCAPE VOL

# CREATE VARIABLES AND ADD NEW COLUMNS: IMPACT (BB+WIND), IMPACT % ON TOTAL VOLUME.

# damage$killedVolume[which(is.na(damage$killedVolume)==TRUE)] <- 0          # FOR MAKE THE na = 0
# damage$wind[which(is.na(damage$wind)==TRUE)] <-0                         # FOR MAKE THE na = 0

# TOTAL IMPACT IN M3/ha
impact <- data.frame(year=damage[,1],                                         # "impact <- data.frame
                      impact= (damage[,2] + damage[,4])/landscape.area)
# IT IS THE SAME CODE FOR CREATE THE DATAFRAME IMPACT variable
# Add the variable
damage <- left_join(damage,impact,by=("year"))

# IMPACT RELATIVE (%) AT THE TOTAL VOLUME
rel_imp <- data.frame(year=damage$year,
                       rel_imp=((damage$impact/damage$tot.vol)*100))

damage<-left_join(damage,rel_imp,by=("year"))

# GIVE THE NAME AT EVERY VARIABLE

# colnames(damage)<-c("year", "barkbeetle", "case", "wind", "volume", "impact_m3", "relative_imp")

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# damage <- damage[,-3]

head(damage)

#-----
# CREATE A NEW DATA FRAME TO WORK ON THE SPECIFIC VARIABLES NEEDED

head(landscape)
head(dynamicstand)

# TO UNDERSTAND THE OPERATORS %>% AND %IN% HAVE TO STUDY THEM IN DATACAMP AND IN DPLYR CRAN PACKAGES

ab.lnd.v<- data.frame(landscape %>%
                        group_by(year) %>%
                        filter(year>0) %>%
                        summarise(tot_volume=sum(volume_m3), living_c=sum(total_carbon_kg), count_ha=su

dynamicstand_1 <-dynamicstand %>% filter(year>0) # THIS IS NEEDED FOR MAKE THE na = 0 !

# CREATE THE NEW DATA FRAME FOR VARIABLES

variables <- data.frame(case=case,
                         year=dynamicstand_1$year,
                         h=dynamicstand_1$height_mean,
                         dbh=dynamicstand_1$dbh_mean,
                         age=dynamicstand_1$age_mean,
                         Rh=carbonflow$Rh,
                         NPP=carbonflow$NPP,
                         GPP=carbonflow$GPP,
                         NEP=carbonflow$NEP,
                         H.count=H.count$H,
                         H.VOL=H.VOL$H,
                         H.BA=H.BA$H,
                         SEI=SEI,
                         esp_BA_prop=esp$BA,
                         csp_BA_prop=csp$BA)

# ADD LANDSCAPE VARIABLES AT V DATA FRAME

variables = inner_join(variables, ab.lnd.v, by="year")
variables = inner_join(variables, ab.tot.c, by="year")
variables = inner_join(variables, ab.nee.v, by="year")
variables = inner_join(variables, damage, by="year")

# variables[which(is.na(damage$wind)==TRUE)] <-0 # FOR MAKE THE na = 0 !

head(variables)

```

```

#-----#
# Make the 3 categories of removals:

# activity.names<-unique(abeStandRemoval$activity)
# swcuts<- grep("sw",activity.names)
# activity.names.sw<-activity.names[swcuts]
# activity.names.notsw<-activity.names[!swcuts]

# print(activity.names.sw)
# print(activity.names.notsw)

# Here I filter only the listed activity names and calculate thinning/finalcut values for every year
# (each line is per ha for a stand, so I scale with the area, sum up all the harvest on the landscape

# TO UNDERSTAND THE OPERATORS %>% AND %IN% HAVE TO STUDY THEM IN DATACAMP AND IN DPLER CRAN PACKAGES

# ab.regcuts<- data.frame(abeStandRemoval %>% filter(activity %in% activity.names.sw)      %>%
#                           # group_by(year)    %>%   summarise(volume=sum(volumeThinning*area)/landscape.area, type="regcut", run

# ab.finalcuts<- data.frame(abeStandRemoval %>% filter(activity %in% activity.names.sw)      %>%
#                           # group_by(year)    %>%   summarise(volume=sum(volumeFinal*area)/landscape.area, type="finalcut", run

# ab.thinnig<- data.frame(abeStandRemoval %>% filter(activity %in% activity.names.notsw)      %>%
#                           # group_by(year)    %>%   summarise(volume=sum(volumeThinning*area)/landscape.area, type="thinning",

# ab.salvaged<- data.frame(abeStandRemoval %>% filter(activity %in% activity.names.sw)      %>%
#                           # group_by(year)    %>%   summarise(volume=sum(volumeSalvaged*area)/landscape.area, type="salvager",

# CREATE A FOR CYCLE INTO THE FOR CYCLE

# if (length(activity.names[swcuts])==0) {

#   ab.finalcuts<- data.frame(abeStandRemoval %>% filter(activity %in% activity.names.notsw)      %>%
#                             # group_by(year)    %>%   summarise(volume=sum(volumeFinal*area)/landscap

# CLOSE THE FOR CYCLE

# CREATE THE VARIABLE FOR THE DIFFERENT WOOD REMOVAL ACTIVITY
# removals<-rbind(removals,ab.regcuts,ab.finalcuts,ab.thinnig,ab.salvaged)

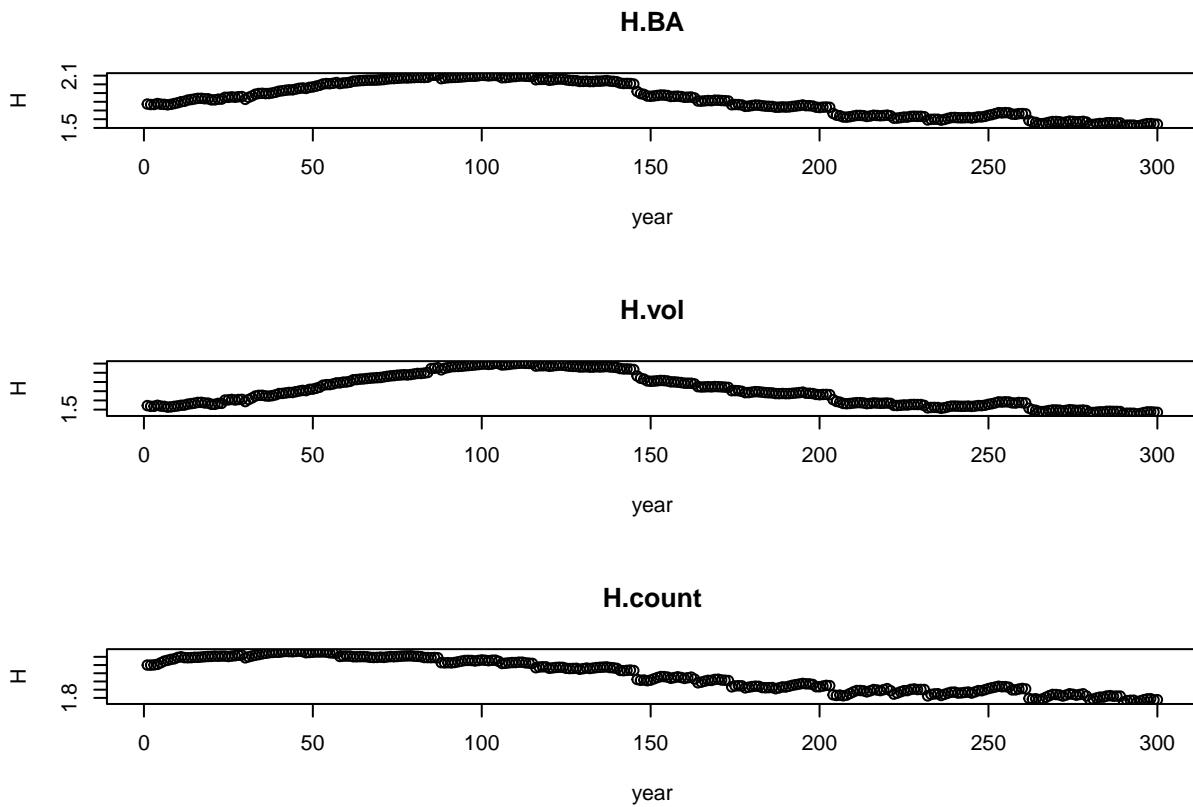
# Collect landscape data FOR CREATE THE VARIABLE LND FOR ALL THE RUNS
landscape<- (landscape %>% mutate(run=case))
lnd<-rbind(lnd, landscape)

# Collect abeUnit data FOR CREATE THE VARIABLE ABEUNIT FOR ALL THE RUNS
abeUnit<-(abeUnit %>% mutate(run=case))
aUnit<-rbind(aUnit, abeUnit)

# Collect barkbeetle data FOR CREATE THE VARIABLE BB FOR ALL THE RUNS
barkbeetle <-(barkbeetle %>% mutate(run=case))
bb <-rbind(bb, barkbeetle)

```





```

## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>    <dbl>    <dbl>     <dbl>
## 1     1     299.    30.6      999.
## 2     2     296.    30.4     1004.
## 3     3     294.    30.3     1010.
## 4     4     284.    29.4      999.
## 5     5     288.    29.9     1024.
## 6     6     297.    30.9     1054.

```

```

## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species    VOL     BA count
##   <int> <chr>     <dbl>   <dbl> <dbl>
## 1     1 abal      6.59  0.553  17.1
## 2     1 acps      7.89  1.05   67.7
## 3     1 algl      0.605 0.0860  3.57
## 4     1 bepe      0.398 0.0940  5.28
## 5     1 cabe      7.92  1.22   65.7
## 6     1 fasy     23.9   2.49   121.

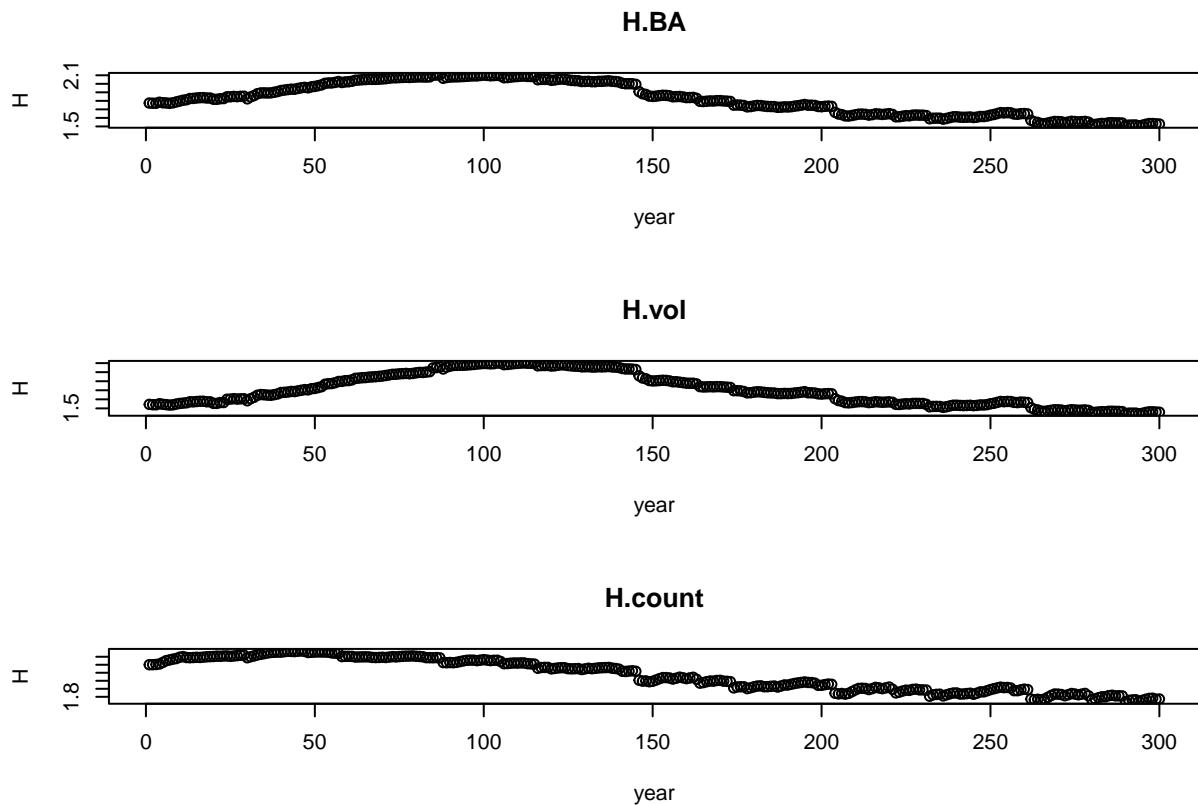
##   [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
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##  [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [101] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [126] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 16 16 16 16 16 16 16 16 16 16 16 16 16
## [151] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16
## [176] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16
## [201] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16
## [226] 16 16 16 16 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15

## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B0._refclim_w3.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"           "soilinput"
## [10] "wind"

## 'summarise()' has grouped output by 'year'. You can override using the
## '.groups' argument.

```



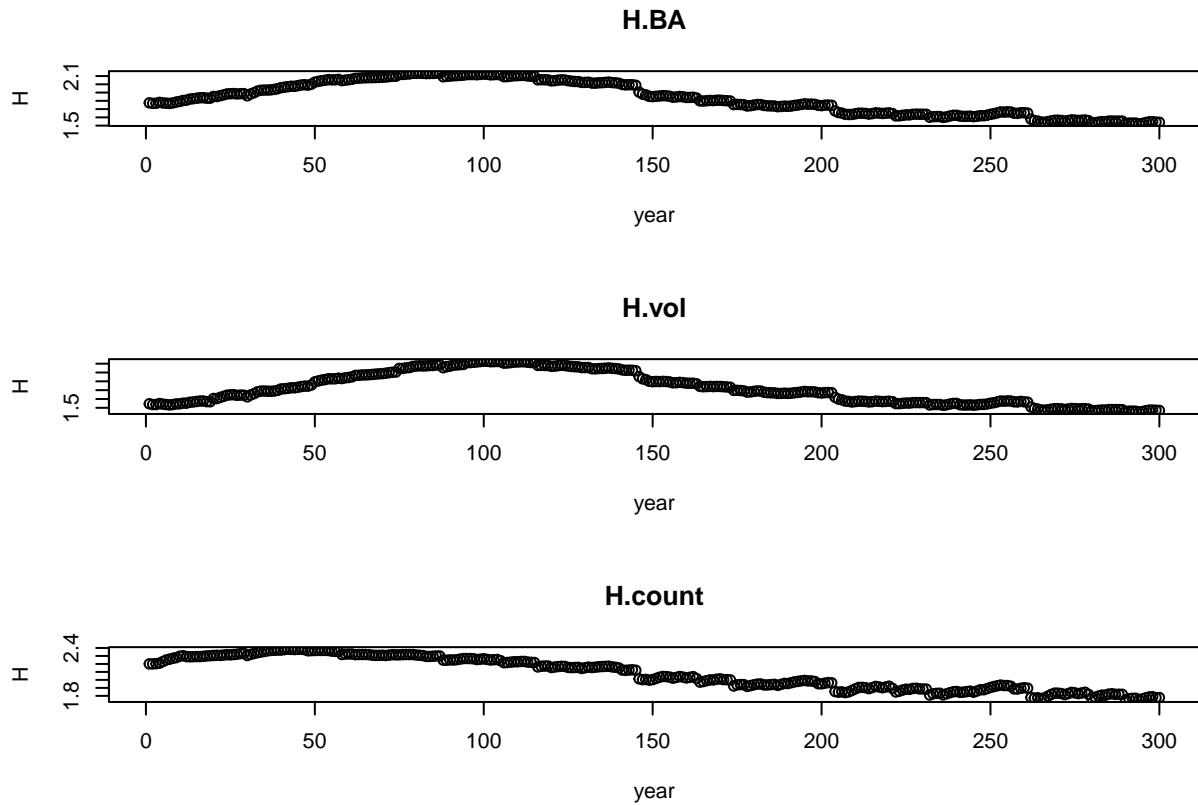
```

## [176] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 15 15 15
## [201] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 15 15 15
## [226] 15 16 16 16 16 16 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B0.5_refclim_w1.sqlite"
## [1] "abeUnit"          "barkbeetle"       "carbon"
## [4] "carbonflow"       "dynamicstand"    "landscape"
## [7] "landscape_removed" "runinfo"          "soilinput"
## [10] "wind"

## `summarise()` has grouped output by 'year'. You can override using the
## `groups` argument.

```



```

## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>   <dbl>   <dbl>     <dbl>
## 1     1    299.    30.6     999.
## 2     2    296.    30.4    1004.
## 3     3    294.    30.3    1007.
## 4     4    284.    29.4     992.
## 5     5    288.    29.8    1007.
## 6     6    297.    30.8    1024.

```

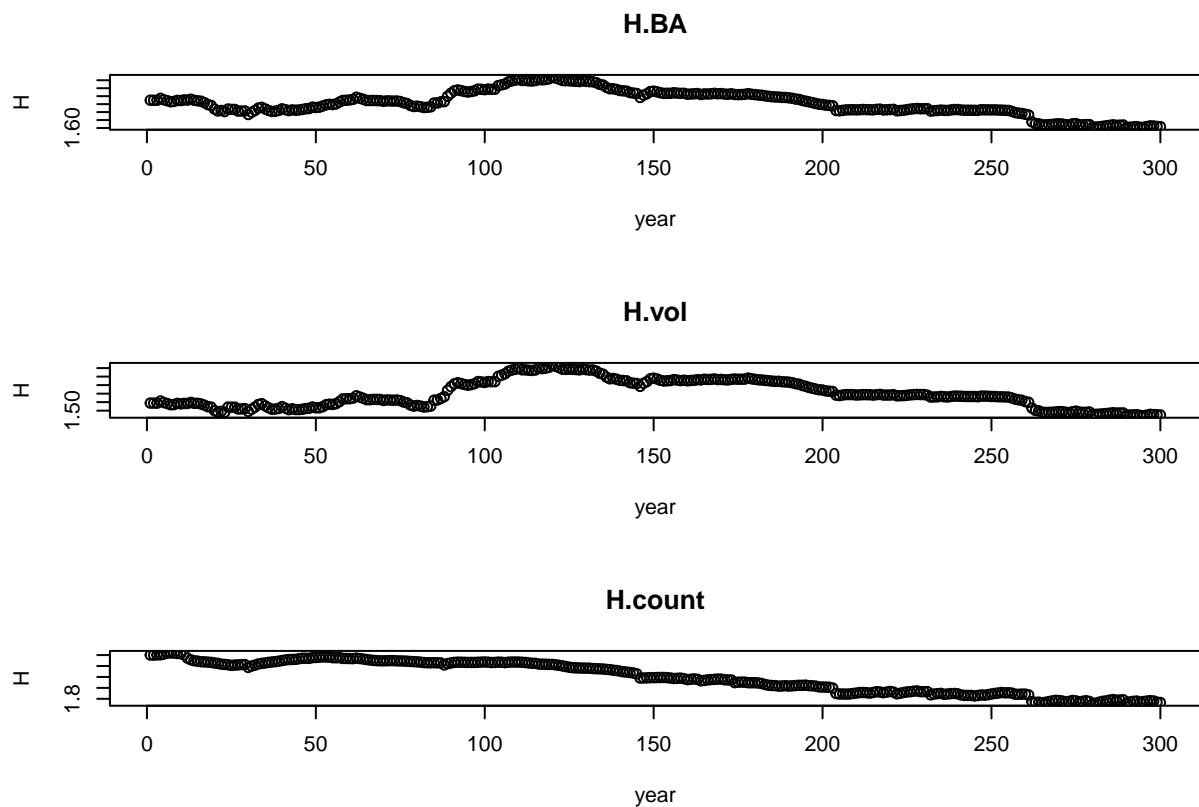
```

## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species    VOL     BA count
##   <int> <chr>     <dbl>   <dbl> <dbl>
## 1     1 abal      6.45  0.546  17.1
## 2     1 acps      7.88  1.05   67.6
## 3     1 alg1      0.603  0.0857  3.57
## 4     1 bepe      0.399  0.0942  5.30
## 5     1 cabe      7.91   1.22   65.7
## 6     1 fasy     24.0    2.50   121.
##   [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [51] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [101] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [126] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 16 16 16 16 16 16 16 16 16 16 16 16 16 16
## [151] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 15 15
## [176] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [201] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [226] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B0.5_refclim_w2.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"           "soilinput"
## [10] "wind"

## 'summarise()' has grouped output by 'year'. You can override using the
## '.groups' argument.

```



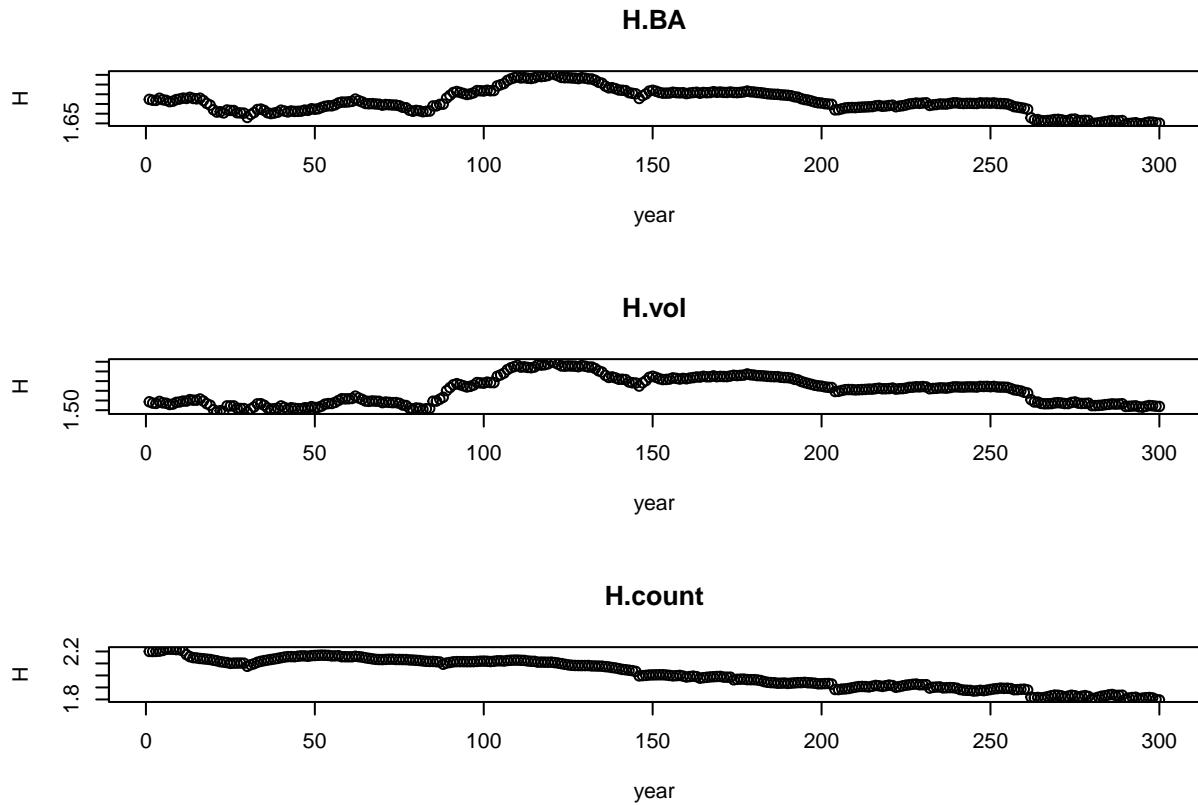
```

## [176] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 16 16 16 15 15 15 15 15 15 15 15
## [201] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [226] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B0.5_refclim_w3.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"          "soilinput"
## [10] "wind"

## `summarise()` has grouped output by 'year'. You can override using the
## `.` argument.

```



```

## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>   <dbl>   <dbl>     <dbl>
## 1     1    299.    30.6     1000.
## 2     2    296.    30.4     1006.
## 3     3    294.    30.3     1011.
## 4     4    284.    29.4      995.
## 5     5    288.    29.9     1009.
## 6     6    297.    30.8     1024.

```

```

## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species     VOL     BA count
##   <int> <chr>    <dbl>  <dbl> <dbl>
## 1     1 abal      6.63  0.556  17.2
## 2     1 acps      7.89  1.05   67.8
## 3     1 algl      0.604 0.0860  3.58
## 4     1 bepe      0.398 0.0940  5.28
## 5     1 cabe      7.92  1.22   65.8
## 6     1 fasy     23.9   2.49   121.

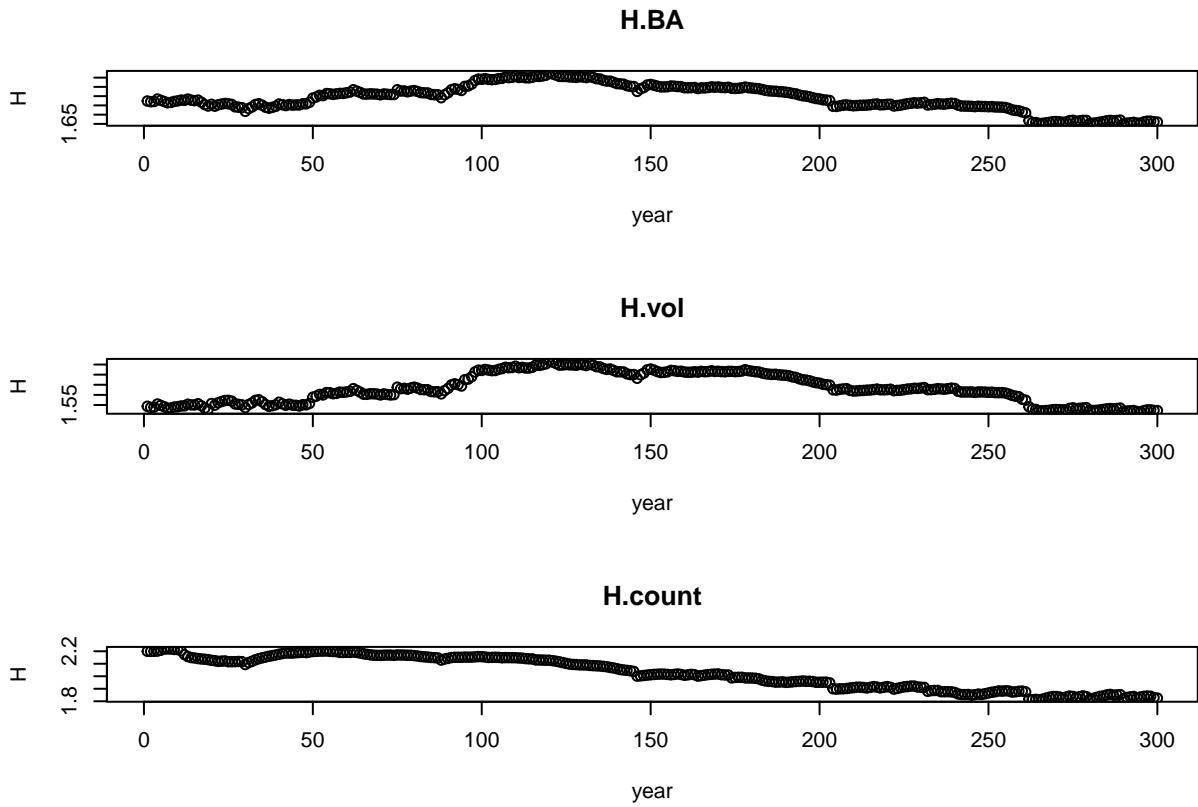
##   [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [51] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
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##  [126] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [151] 17 17 17 17 17 17 17 17 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16
##  [176] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 15 16 16 16 16 16 16 16 16 16 16 16 16 16
##  [201] 16 16 16 16 16 16 16 16 16 16 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
##  [226] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
##  [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
##  [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15

## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B1_refclim_w1.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"           "soilinput"
## [10] "wind"

## 'summarise()' has grouped output by 'year'. You can override using the
## '.groups' argument.

```



```

## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>   <dbl>   <dbl>     <dbl>
## 1     1    299.    30.6     1000.
## 2     2    296.    30.4     1007.
## 3     3    294.    30.3     1010.
## 4     4    284.    29.4      992.
## 5     5    288.    29.9     1003.
## 6     6    297.    30.8     1014.

## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species   VOL     BA   count
##   <int> <chr>    <dbl>   <dbl>   <dbl>
## 1     1  abal     6.63   0.557   17.2
## 2     1  acps     7.88   1.05    67.7
## 3     1  alg1     0.604   0.0859  3.57
## 4     1  bepe     0.400   0.0945  5.30
## 5     1  cabe     7.92   1.22    65.7
## 6     1  fasy    23.6    2.47   121.

## [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [51] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [101] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [126] 17 17 17 16 16 16 16 16 16 16 16 16 16 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [151] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15

```

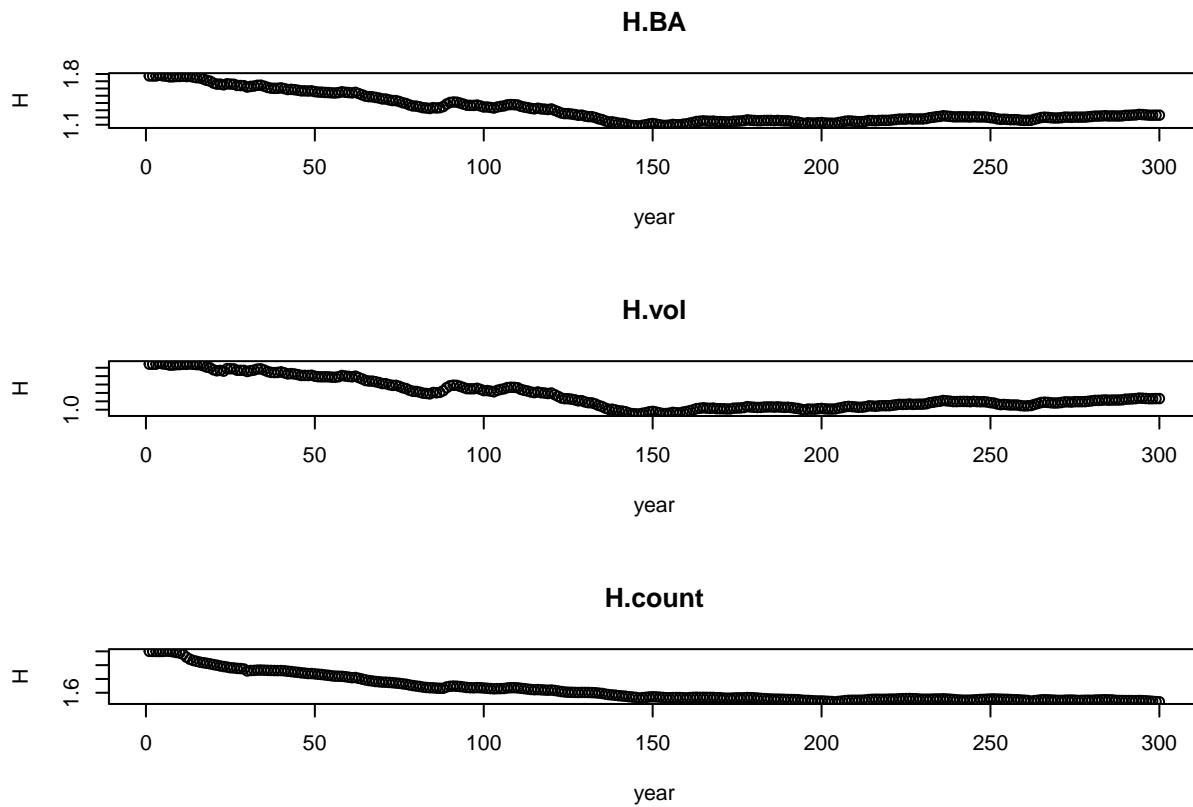
```

## [176] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [201] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [226] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B1_refclim_w2.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"          "soilinput"
## [10] "wind"

## `summarise()` has grouped output by 'year'. You can override using the
## `groups` argument.

```



```

## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>   <dbl>   <dbl>    <dbl>
## 1     1     299.    30.6     1000.
## 2     2     296.    30.4     1005.
## 3     3     294.    30.3     1008.
## 4     4     284.    29.3      990.
## 5     5     288.    29.9     1003.
## 6     6     297.    30.8     1014.

```

```

## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species     VOL     BA count
##   <int> <chr>    <dbl>   <dbl> <dbl>
## 1     1 abal      6.63  0.557  17.2
## 2     1 acps      7.86  1.05   67.7
## 3     1 alg1      0.605  0.0860  3.57
## 4     1 bepe      0.400  0.0945  5.30
## 5     1 cabe      7.91   1.22   65.7
## 6     1 fasy     23.6   2.47   121.

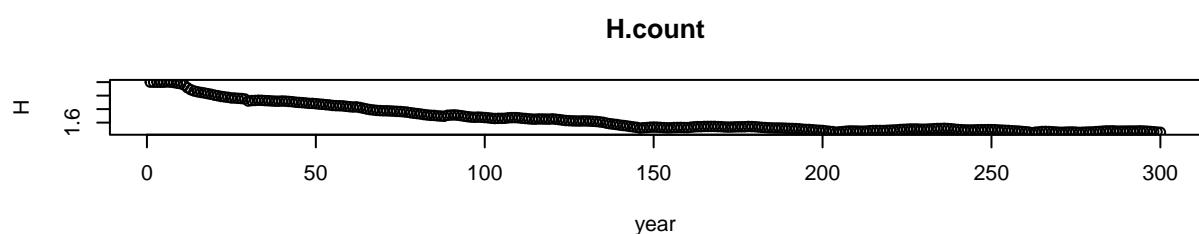
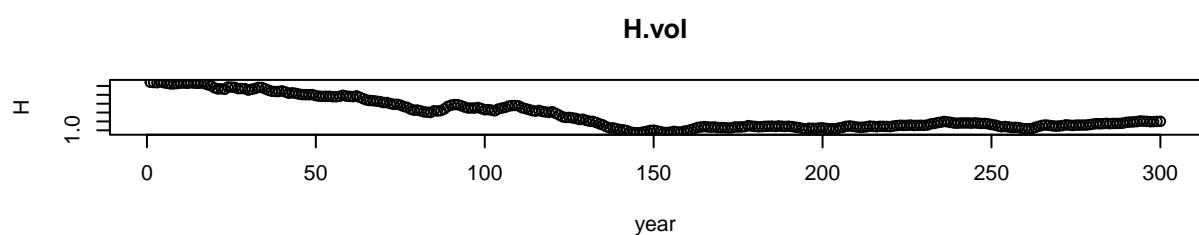
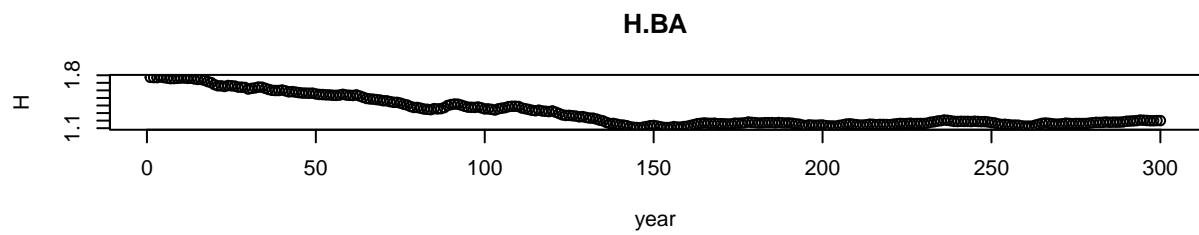
##   [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [51] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [101] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 16 16 16 16 16 16 16 16 16 16 16
## [126] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16
## [151] 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 15 15 15 15 15 15 15 15 15 15 15
## [176] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [201] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [226] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15

## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B1_refclim_w3.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"           "soilinput"
## [10] "wind"

## 'summarise()' has grouped output by 'year'. You can override using the
## '.groups' argument.

```



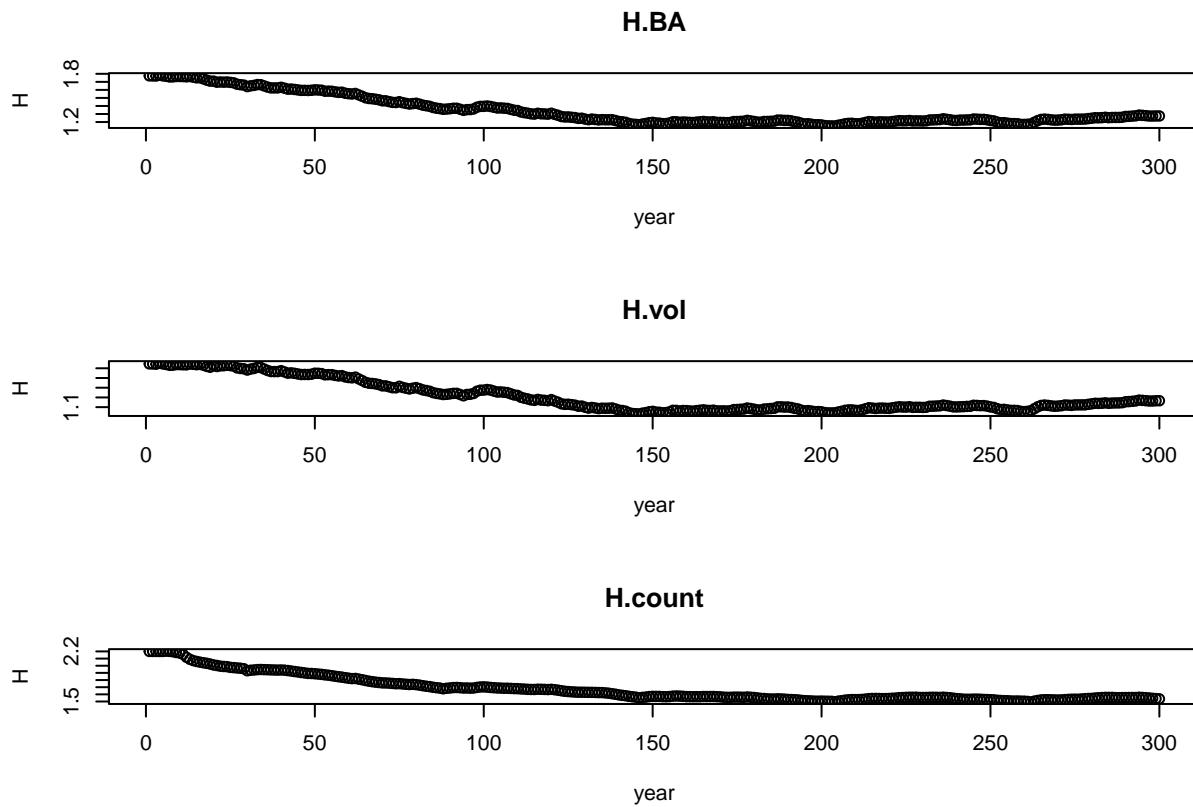
```

## [176] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [201] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [226] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [251] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [276] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B2_refclim_w1.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"          "soilinput"
## [10] "wind"

## `summarise()` has grouped output by 'year'. You can override using the
## `groups` argument.

```



```

## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>   <dbl>   <dbl>     <dbl>
## 1     1    299.    30.6     999.
## 2     2    296.    30.4    1003.
## 3     3    294.    30.3    1007.
## 4     4    284.    29.4     989.
## 5     5    288.    29.9     998.
## 6     6    297.    30.9    1005.

```

```

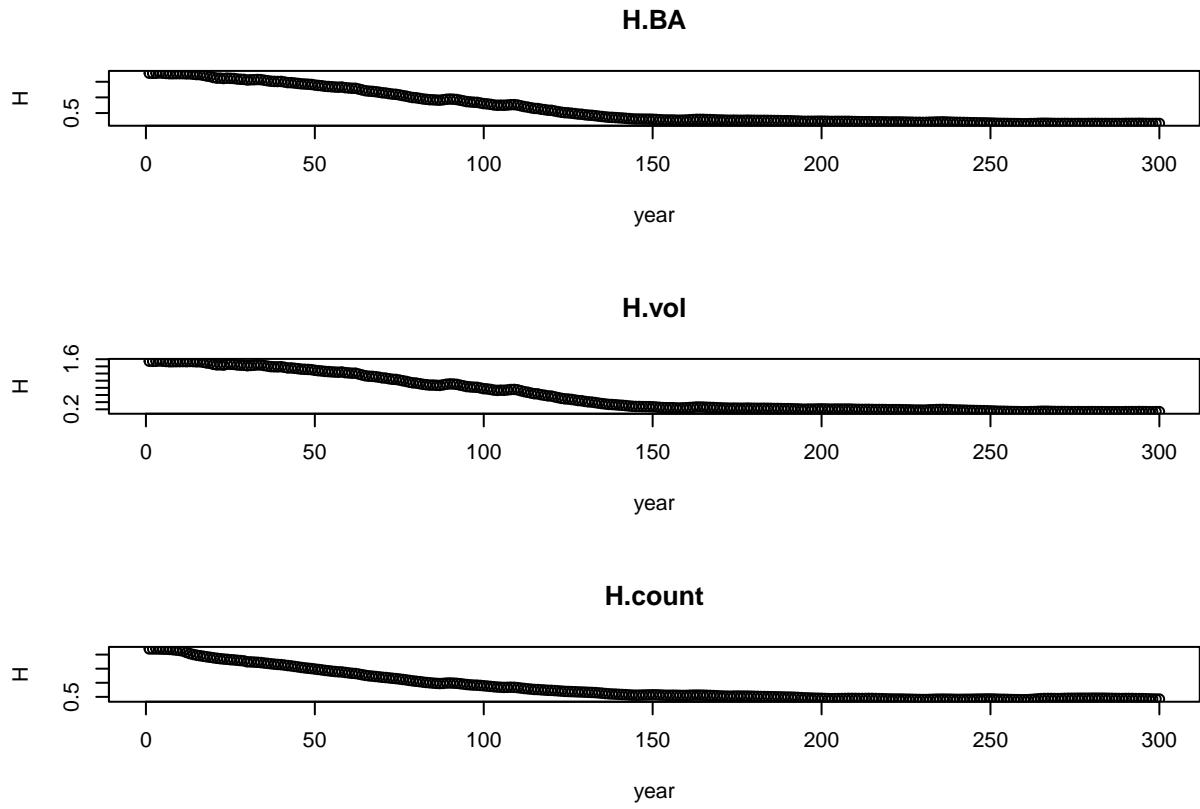
## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species    VOL     BA count
##   <int> <chr>     <dbl>   <dbl> <dbl>
## 1     1 abal      6.54  0.551  17.1
## 2     1 acps      7.91  1.05   67.7
## 3     1 algl      0.606 0.0861  3.58
## 4     1 bepe      0.400 0.0944  5.30
## 5     1 cabe      7.92  1.22   65.7
## 6     1 fasy     23.8   2.49   121.

##   [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [51] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16
## [101] 16 16 16 16 16 16 16 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 14 14 14
## [126] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
## [151] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
## [176] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
## [201] 14 14 14 14 14 14 14 14 14 14 14 14 14 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13
## [226] 13 13 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12
## [251] 11 11 11 11 11 11 11 11 11 11 11 11 11 11 11 11 11 11 10 10 10 10 10 10 10 10 10 10 10 10 10 10 10
## [276] 10 10 10 10 10 10 9 9 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8 8
## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B2_refclim_w2.sqlite"
## [1] "abeUnit"           "barkbeetle"        "carbon"
## [4] "carbonflow"        "dynamicstand"      "landscape"
## [7] "landscape_removed" "runinfo"          "soilinput"
## [10] "wind"

## 'summarise()' has grouped output by 'year'. You can override using the
## '.groups' argument.

```



```
## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>    <dbl>   <dbl>      <dbl>
## 1     1    299.   30.6      999.
## 2     2    296.   30.4     1004.
## 3     3    294.   30.3     1006.
## 4     4    284.   29.4      988.
## 5     5    288.   29.8      995.
## 6     6    297.   30.8     1001.

## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species    VOL     BA   count
##   <int> <chr>    <dbl>   <dbl>   <dbl>
## 1     1  abal     6.63   0.556   17.2
## 2     1  acps     7.90   1.05    67.7
## 3     1  algl     0.605   0.0861   3.58
## 4     1  bepe     0.396   0.0937   5.28
## 5     1  cabe     7.90   1.22    65.7
## 6     1  fasy    24.0    2.50    121.

## [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [51] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
## [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16 16
## [101] 17 17 17 17 17 16 16 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
## [126] 15 15 15 15 15 15 15 15 15 15 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
## [151] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
```

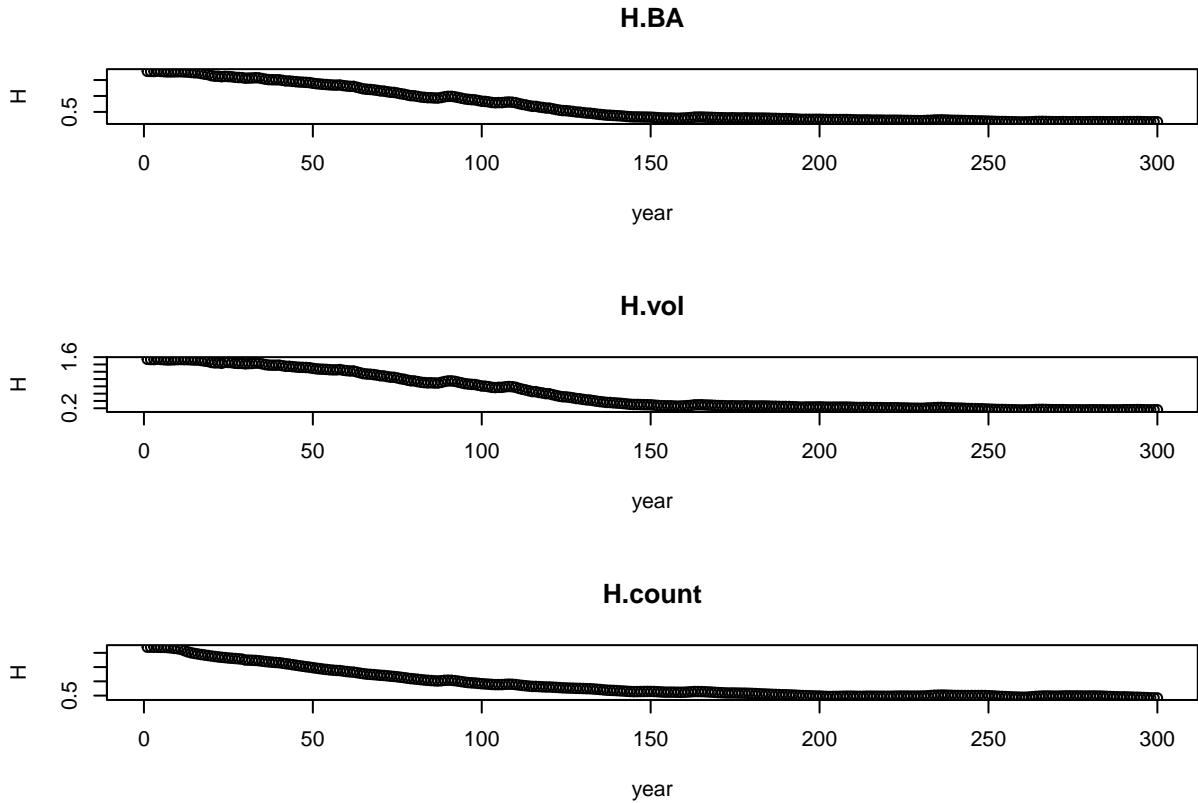
```

## [176] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
## [201] 14 14 14 14 14 14 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13
## [226] 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 13 12 12 12 12
## [251] 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12
## [276] 12 11 11 11 11 11 10 10 10 10 10 10 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9
## plot window will lay out plots in a 3 by 1 matrix

## [1] "C:/iLand/2023/browsing_revision/db_test/B2_refclim_w3.sqlite"
## [1] "abeUnit"          "barkbeetle"       "carbon"
## [4] "carbonflow"       "dynamicstand"    "landscape"
## [7] "landscape_removed" "runinfo"         "soilinput"
## [10] "wind"

## `summarise()` has grouped output by 'year'. You can override using the
## `.` argument.

```



```

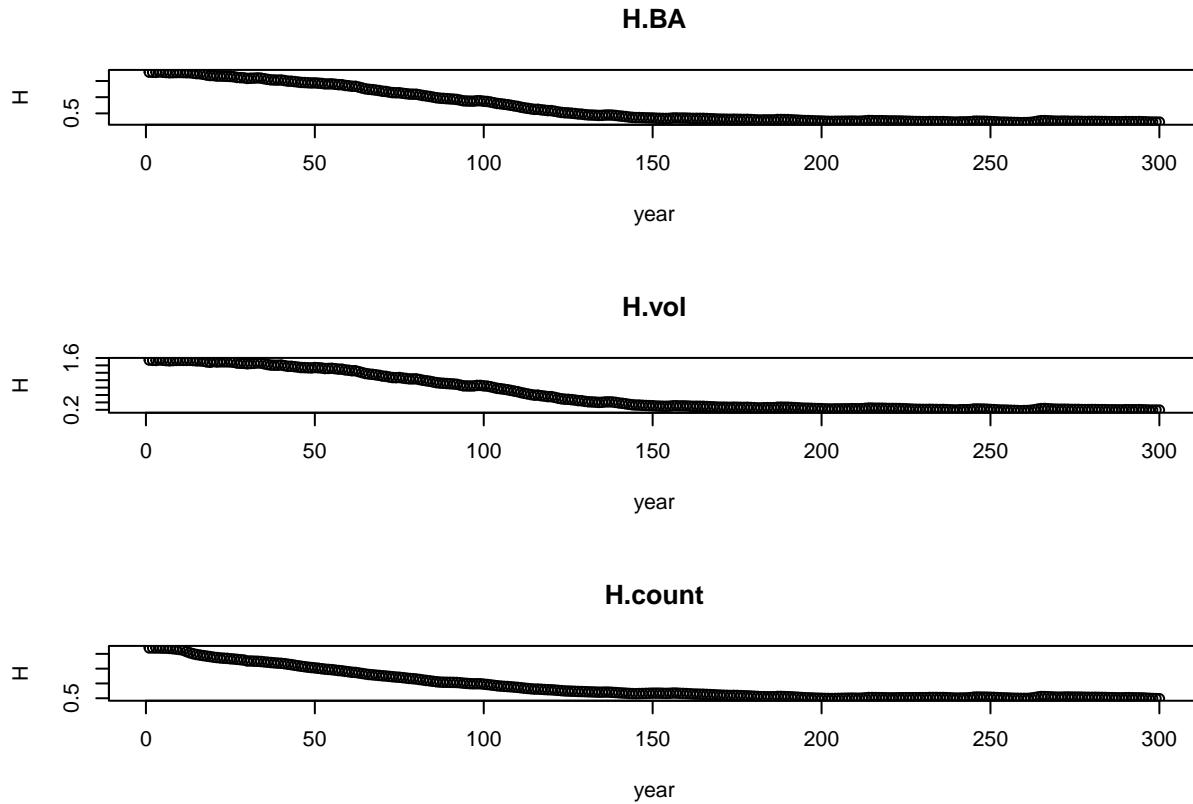
## # A tibble: 6 x 4
##   year VOL.tot BA.tot count.tot
##   <int>   <dbl>   <dbl>     <dbl>
## 1     1     299.    30.6     1000.
## 2     2     296.    30.4     1004.
## 3     3     294.    30.3     1007.
## 4     4     284.    29.3      987.
## 5     5     288.    29.8     996.
## 6     6     297.    30.8     1002.

```

```

## # A tibble: 6 x 5
## # Groups:   year [1]
##   year species    VOL     BA count
##   <int> <chr>     <dbl>   <dbl> <dbl>
## 1     1 abal      6.59  0.555  17.2
## 2     1 acps      7.89  1.05   67.7
## 3     1 alg1      0.604 0.0859  3.57
## 4     1 bepe      0.401 0.0946  5.30
## 5     1 cabe      7.91  1.22   65.7
## 6     1 fasy     23.7   2.47   121.
##   [1] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [26] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [51] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17
##  [76] 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 17 16 16 16 16 16 16 16 16 17 17 17
##  [101] 17 17 17 17 17 17 16 16 16 16 16 16 16 16 16 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15
##  [126] 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 15 14 14 14 14 14 14 14
##  [151] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
##  [176] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14
##  [201] 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 14 13 13 13 13 13 13 13 13
##  [226] 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12
##  [251] 12 12 12 12 12 12 12 12 12 12 12 12 12 12 12 11 11 11 10 10 10 10 10 10 10 10 10 10 10 10
##  [276] 10 10 10 10 10 10 10 10 10 10 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9 9
## plot window will lay out plots in a 3 by 1 matrix

```



```

# This tells the colors:

species.we.have<-unique(lnd$species)                                     # IT IS SAYING WHICH SP

# LIST OF ALL POSSIBLE SPECIES

cols.all=c( "rops"="#e0e0e0", "acpl"="#A9A9A9",     "alin"="#696969", "alvi"="#2e2e2e",
           "bepe"="#fadfad",
           "casa"="#7eeadf", "coav"="#20c6b6",
           "tipl"="#645394", "ulgl"="#311432" ,
           "saca"="#D8BFD8", "soar"="#DDA0DD", "soau"="#BA55D3",
           "pice"="#D27D2D", "pini"="#a81c07",
           "algl"="#2ECBE9", "tico"="#128FC8", "potr"="#00468B", "poni"="#5BAEB7",
           "frex"="#fe9cb5", "cabe"="#fe6181", "acps"="#fe223e",
           "lade"="#FFFE71", "abal"="#FFD800", "pisy"="#A4DE02",
           "fasy"="#76BA1B", "piab"="#006600",
           "quro"="#FF7F00", "qupe"="#FF9900", "qupu"="#CC9900"
)

# COLORATION ORDER FOR ALL THE POSSIBLE SPECIES

new_order_gg.all=c("alvi","alin", "acpl", "rops", "bepe" , "coav", "casa", "ulgl", "tipl", "soau", "soar",
                   "poni", "algl", "tico", "potr", "frex", "cabe", "acps", "lade", "abal", "qupu", "q

# This will show at the end only the species we really have on the landscape.

cols<-cols.all[names(cols.all) %in% species.we.have]
new_order_gg<- new_order_gg.all[new_order_gg.all %in% species.we.have]

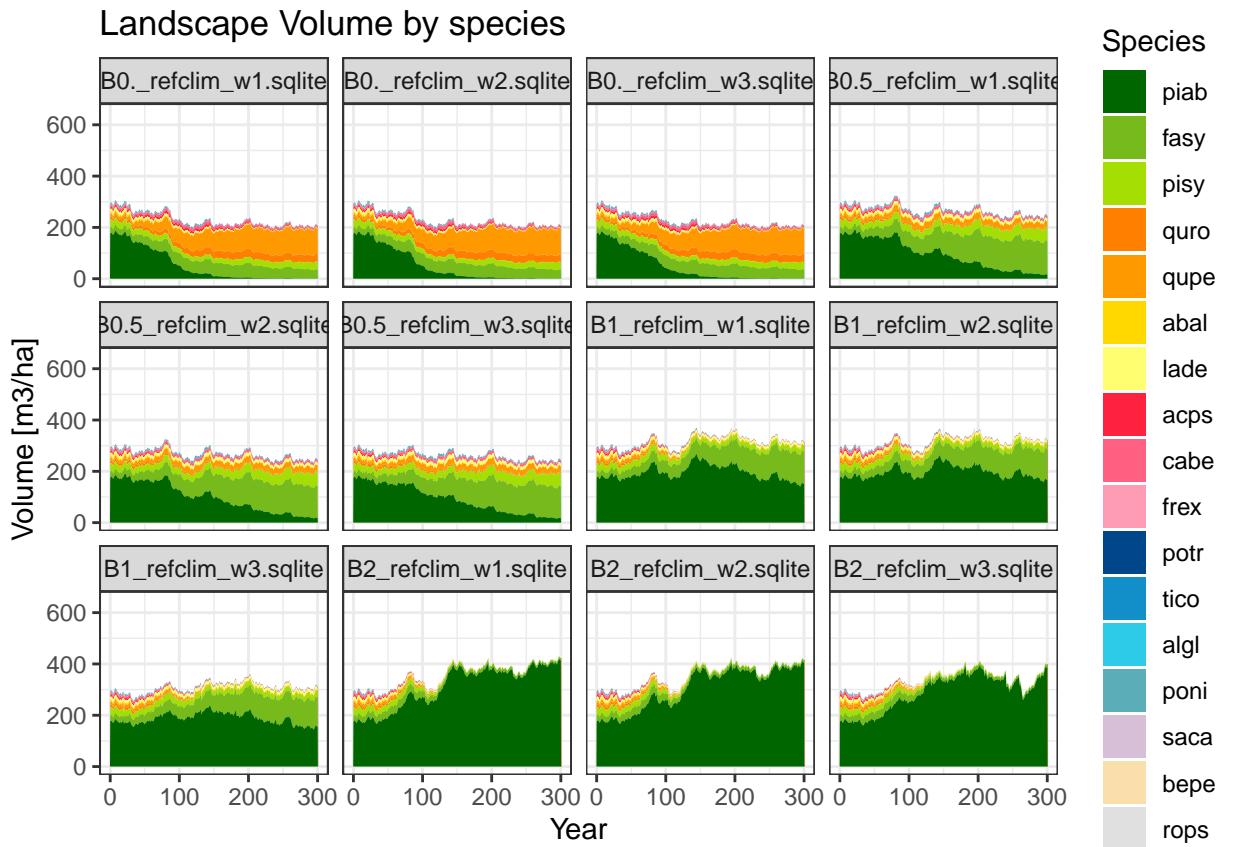
# STARTING PLOTS

# Make a plot with ggplot, volume, colored by species for the transitional period for Clear cut managem

# PLOT LANDSCAPE VOLUME PLOT FOR CASES (GEOM AREA)

ggplot(lnd, aes(year,volume_m3, fill=factor(species, levels=new_order_gg)))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+ 
  ggtitle("Landscape Volume by species")+
  facet_wrap(~run, ncol=4)+ 
  labs(x = "Year",y="Volume [m3/ha]",fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+ 
  ylim(0,650)+ 
  theme_bw()

```



```

# Pie chart year 0

# Filter per the year and create the percentage data frame

B0._refclim_w1 <- lnd %>% filter(run=="B0._refclim_w1.sqlite" & year==300)
B0._refclim_w1_per<-B0._refclim_w1 %>% mutate( sumvol=sum(volume_m3)) %>% mutate(perc.vol=100*volume_m3/sumvol)

B0.5_refclim_w1 <- lnd %>% filter(run=="B0.5_refclim_w1.sqlite" & year==300)
B0.5_refclim_w1_per<-B0.5_refclim_w1 %>% mutate( sumvol=sum(volume_m3)) %>% mutate(perc.vol=100*volume_m3/sumvol)

B1_refclim_w1 <- lnd %>% filter(run=="B1_refclim_w1.sqlite" & year==300)
B1_refclim_w1_per<-B1_refclim_w1 %>% mutate( sumvol=sum(volume_m3)) %>% mutate(perc.vol=100*volume_m3/sumvol)

B2_refclim_w1 <- lnd %>% filter(run=="B2_refclim_w1.sqlite" & year==300)
B2_refclim_w1_per<-B2_refclim_w1 %>% mutate( sumvol=sum(volume_m3)) %>% mutate(perc.vol=100*volume_m3/sumvol)

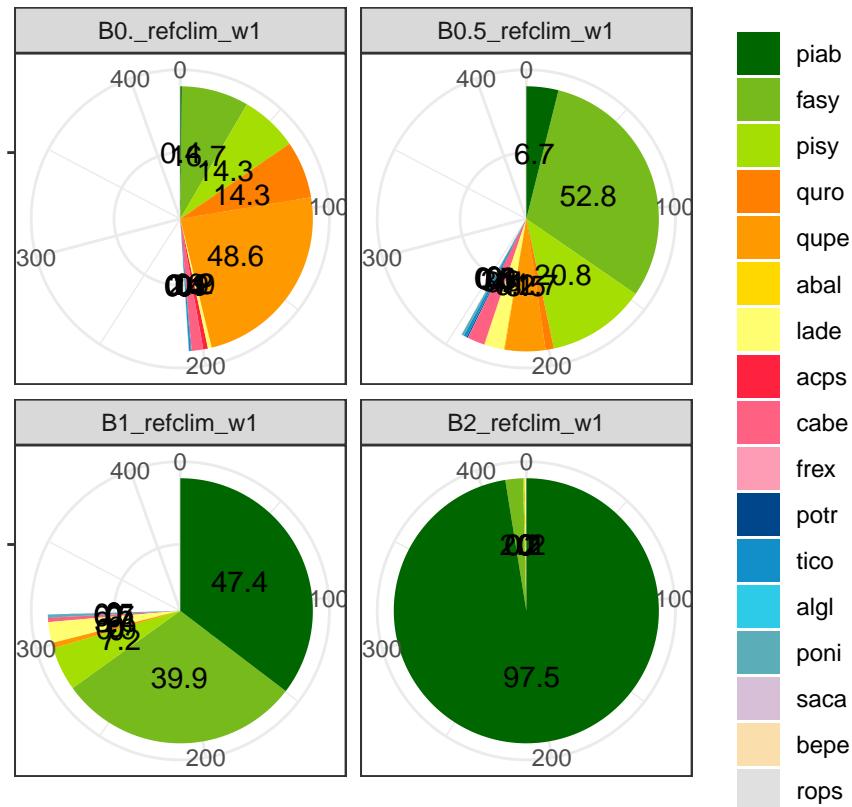
r1wb<-rbind(B0._refclim_w1_per,B0.5_refclim_w1_per, B1_refclim_w1_per, B2_refclim_w1_per)

x7wb <- ggplot(r1wb, aes(x="", y=volume_m3, fill=factor(species,levels=new_order_gg))) +
  geom_bar(stat="identity", width=1, show.legend = T) +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+ 
  facet_wrap(~case, ncol=2)+ 
  coord_polar("y", start=0) + 
  geom_text(aes(label = paste0( round(perc.vol, 1) )), position = position_stack(vjust=0.5)) + 
  labs(x = NULL, y = NULL, fill = NULL)

```

```
ggtitle("Species proportions [%] based on landscape volume [m3/ha] in year 300")+
  theme_bw()
x7wb + theme(plot.title = element_text(hjust = 0.5))
```

Species proportions [%] based on landscape volume [m3/ha] in year 300

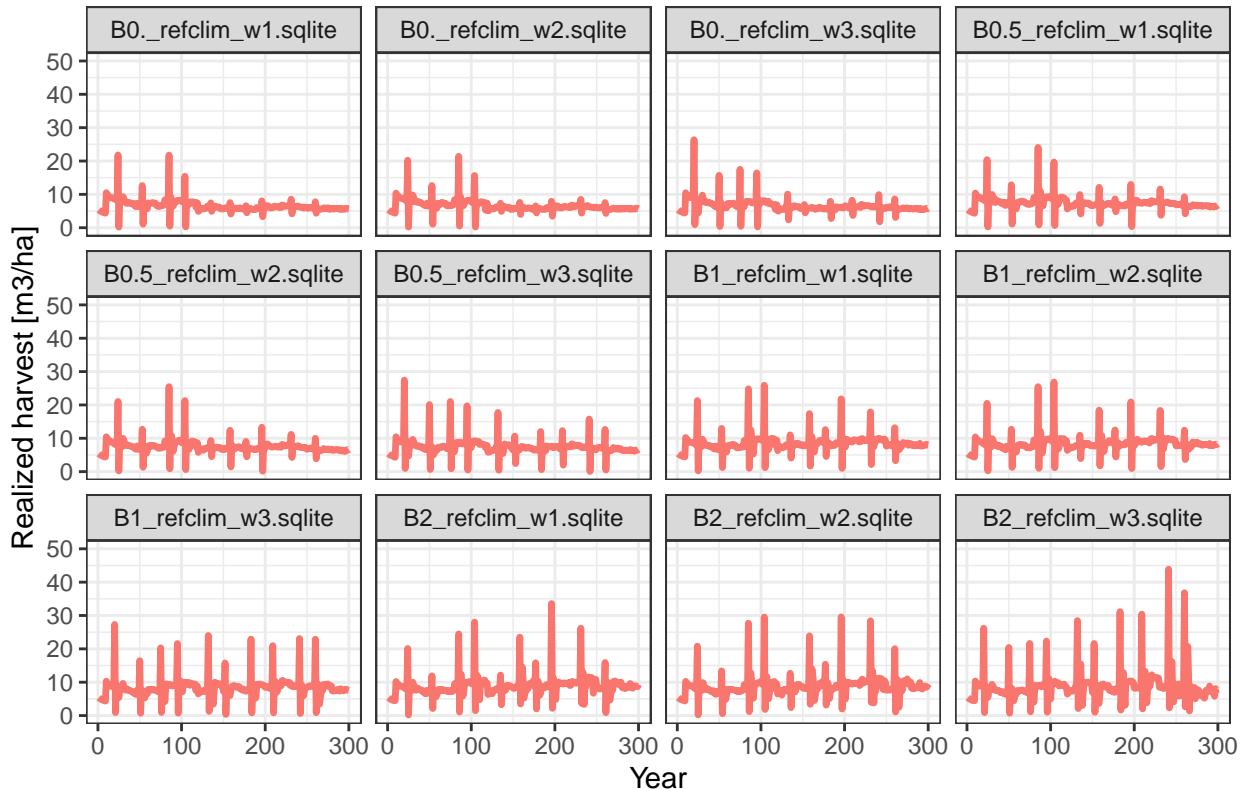


```
# (SHOULD BE REALIZED) PLOT 2 "Y" AXIS WITH relationship between realized harvest and volume increasing
# Total realized harvest at landscape level in average per ha
```

```
ggplot(aUnit, aes(year,realizedHarvest, color=case))+  
  geom_line(size=1.2, show.legend = F)+  
  facet_wrap(~run, ncol=4)+  
  ylim(0,50)+  
  ggtitle("Realized Harvest") +  
  theme(plot.title = element_text(hjust = 0.5))+  
  ylab("Realized harvest [m3/ha]")+  
  xlab("Year") +  
  theme_bw()
```

```
## Warning: Using 'size' aesthetic for lines was deprecated in ggplot2 3.4.0.  
## i Please use 'linewidth' instead.  
## This warning is displayed once every 8 hours.  
## Call 'lifecycle::last_lifecycle_warnings()' to see where this warning was  
## generated.
```

## Realized Harvest



```
# SPECIES specifically BA:

#species.to.keep<-c("piab", "pisy", "fasy", "qupe")

#lnd2 <- lnd %>% filter(species %in% species.to.keep)

#ggplot(data=lnd2, aes(x=year, y=basal_area_m2, colour=species)) +
#  geom_line(size=1.2) +
#  scale_colour_manual(values = c("#76BA1B", "#006600", "#A4DE02", "orange")) +
#  ggtitle("Clearcut management in brow pressure 0") +
#  theme(plot.title = element_text(hjust = 0.5)) +
#  ylab("Basal area [m²/ha]") +
#  theme_bw()
```

```
# PLOT BASAL AREA GEOM_LINE AT LANDSCAPE LEVEL BY SPECIES SELECTED
```

```
# SPECIES specifically BA:

species.to.keep<-c("piab", "fasy", "qupe", "pisy")

lnd2 <- lnd %>% filter(species %in% species.to.keep)

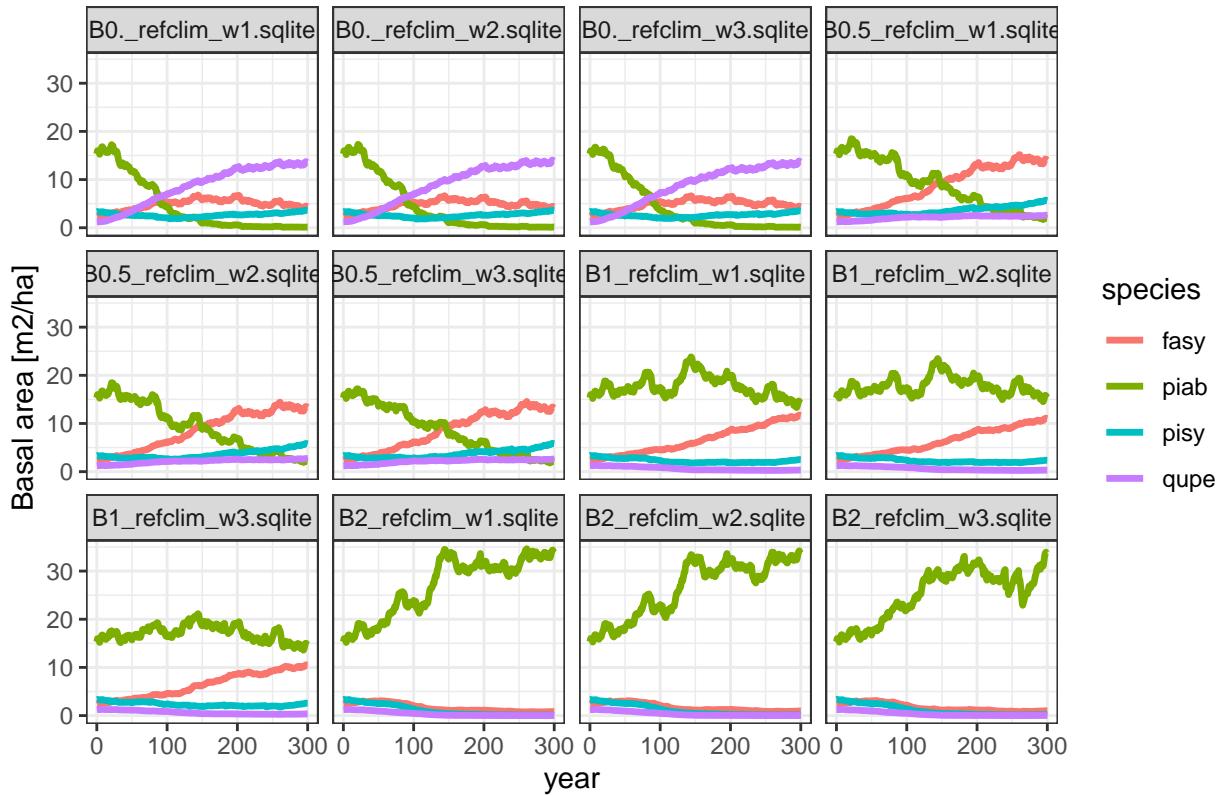
ggplot(data=lnd2, aes(x=year, y=basal_area_m2, color=species)) +
  geom_line(size=1.2) +
  ggtitle("Basal area by species") +
```

```

facet_wrap(~run, ncol=4)+
theme(plot.title = element_text(hjust = 0.5))+
ylab("Basal area [m2/ha]")+
theme_bw()

```

### Basal area by species



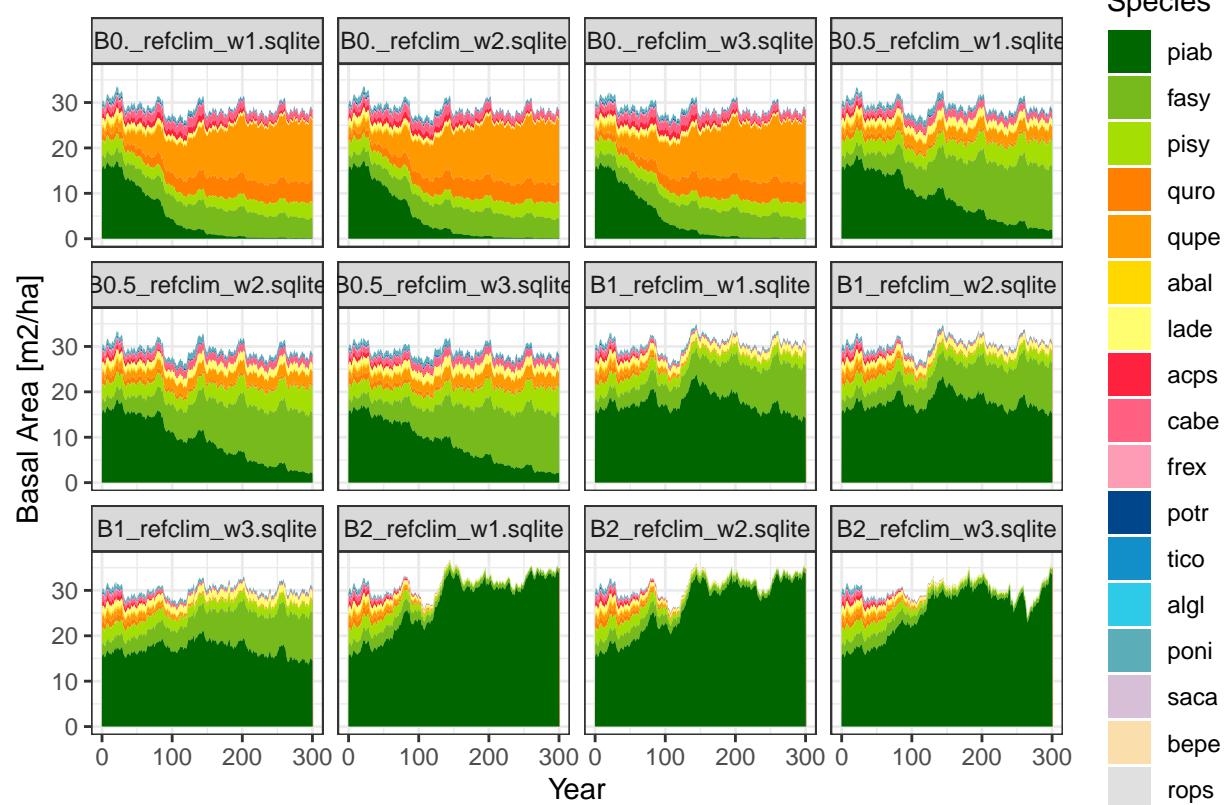
```
# PLOT TOTAL AVG BASAL AREA AT LANDSCAPE LEVEL BY SPECIES
```

```

ggplot(lnd, aes(year, basal_area_m2, fill=factor(species, levels=new_order_gg)))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
  ggtitle("Total Basal Area")+
  facet_wrap(~run, ncol=4)+
  labs(x = "Year", y="Basal Area [m2/ha]", fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+
  theme_bw()

```

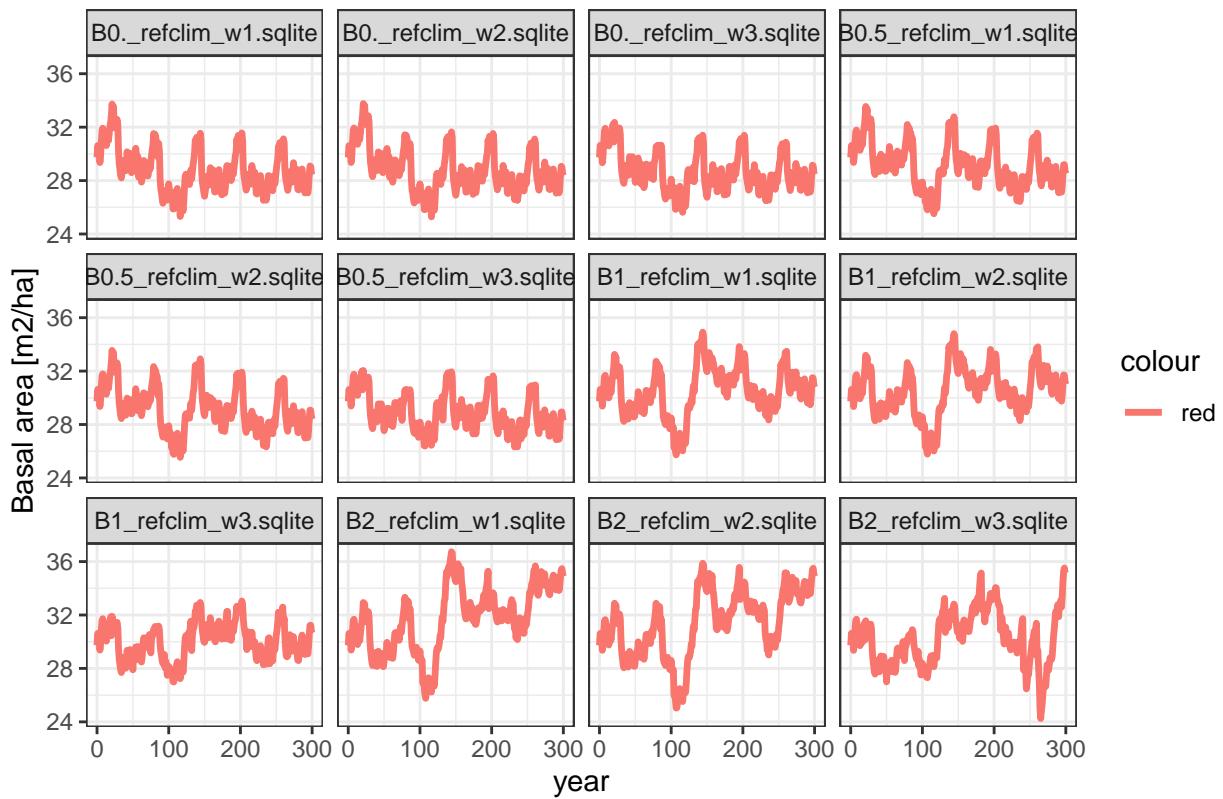
## Total Basal Area



```
# PLOT SUM BASAL AREA AT LANDSCAPE
```

```
ggplot(data=dys, aes(x=year, y=basalarea_sum/landscape.area, color="red")) +
  geom_line(size=1.2) +
  ggtitle("Avarege Basal area") +
  facet_wrap(~run, ncol=4) +
  theme(plot.title = element_text(hjust = 0.5)) +
  ylab("Basal area [m2/ha]") +
  theme_bw()
```

## Average Basal area



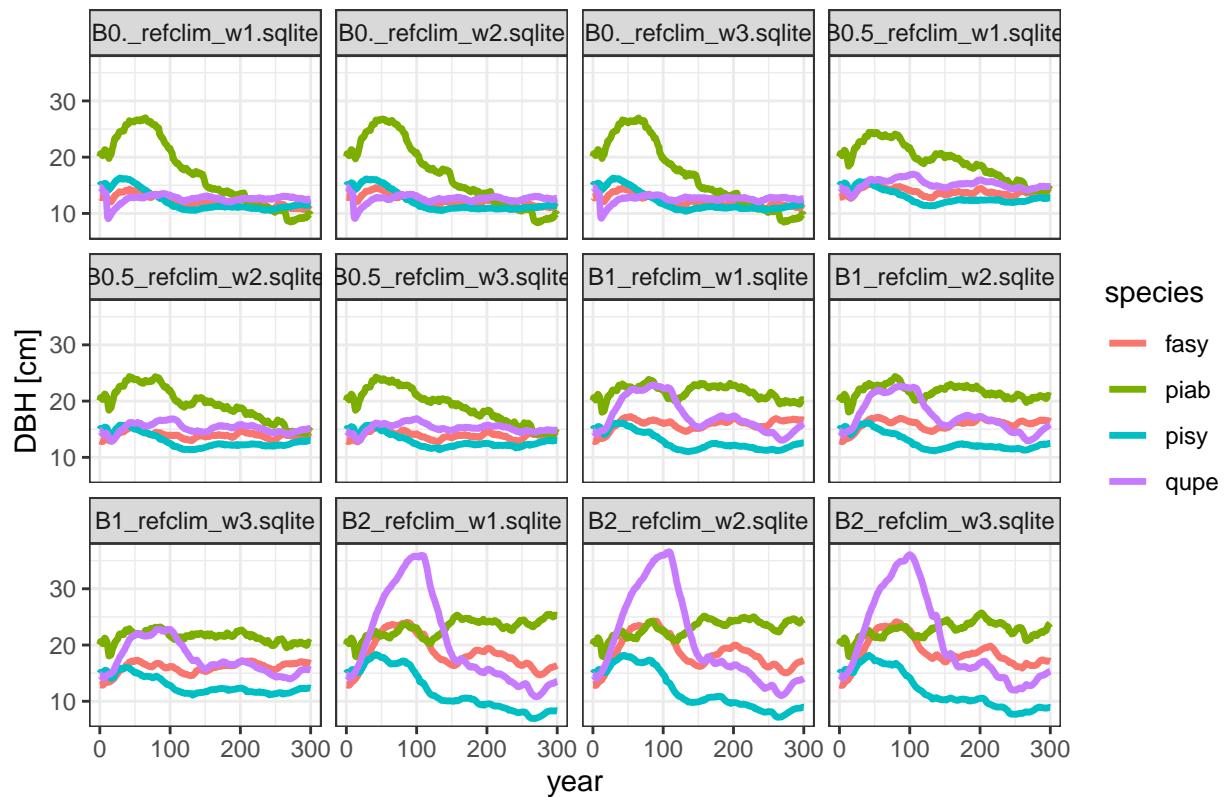
```
# PLOT DBH GEOM_LINE AT LANDSCAPE LEVEL BY SPECIES

species.to.keep<-c("piab", "fasy", "qupe", "pisy")

lnd2 <- lnd %>% filter(species %in% species.to.keep)

ggplot(data=lnd2, aes(x=year, y=dbh_avg_cm, color=species)) +
  geom_line(size=1.2) +
  ggtitle("Average DBH by species") +
  facet_wrap(~run, ncol=4) +
  theme(plot.title = element_text(hjust = 0.5)) +
  ylab("DBH [cm]") +
  theme_bw()
```

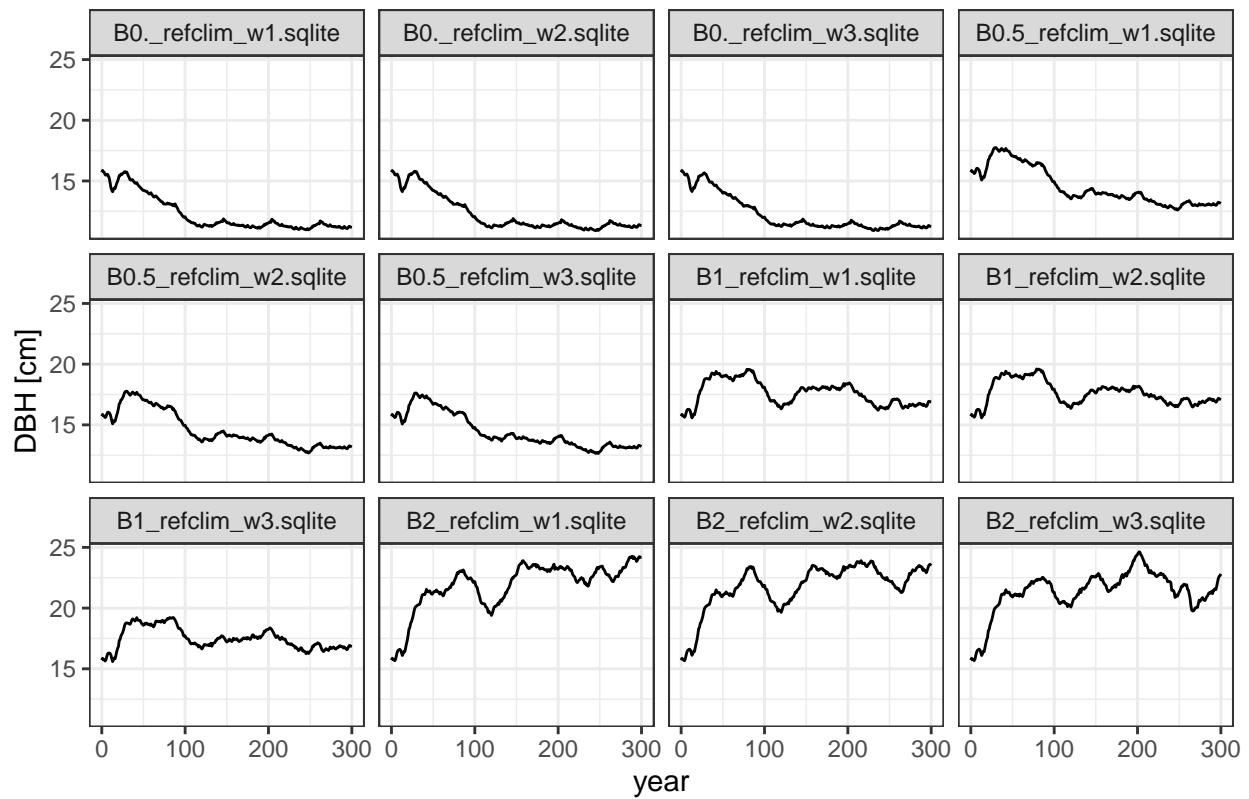
## Average DBH by species



```
# PLOT DBH GEOM_AREA AT LANDSCAPE LEVEL AVERAGE ALL SP TOGETHER
```

```
ggplot(data=dys, aes(x=year, y=dbh_mean)) +
  geom_line()+
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

## Average DBH



```
# SD DBH / try with violin plots

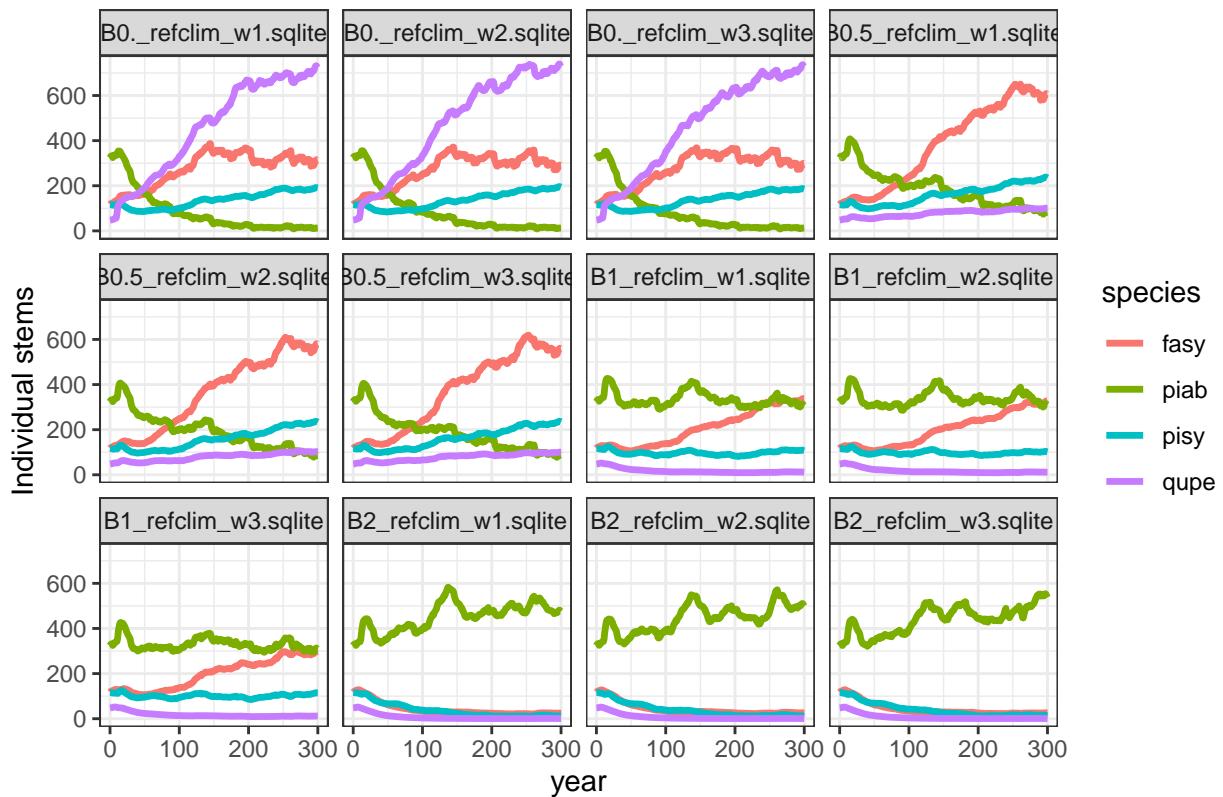
# ggplot(data=dys, aes(x=year, y=dbh_sd)) +
# geom_area(size=1.2) +
# scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE)) +
# ggtitle("Avarage DBH by species") +
# facet_wrap(~run, ncol=1) +
# theme(plot.title = element_text(hjust = 0.5)) +
# ylab("DBH [cm]") +
# theme_bw()

# PLOT NUMBER OF STEMS GEOM_LINE AT LANDSCAPE LEVEL BY SPECIES

# lnd2 <- lnd %>% filter(species %in% species.to.keep)      --- ADD IF YOU WANT SELECT SPECIES TO KEEP

ggplot(data=lnd2, aes(x=year, y=count_ha, color=species)) +
  geom_line(size=1.2) +
  ggtitle("N. individual stems by species") +
  facet_wrap(~run, ncol=4) +
  theme(plot.title = element_text(hjust = 0.5)) +
  ylab("Individual stems") +
  theme_bw()
```

## N. individual stems by species



```
# PLOT NUMBER OF STEMS GEOM_AREA AT LANDSCAPE LEVEL BY SPECIES
```

```
ggplot(lnd, aes(x=year, y=count_ha, fill=factor(species, levels=new_order_gg)))+
  geom_area(size=1.2)+  

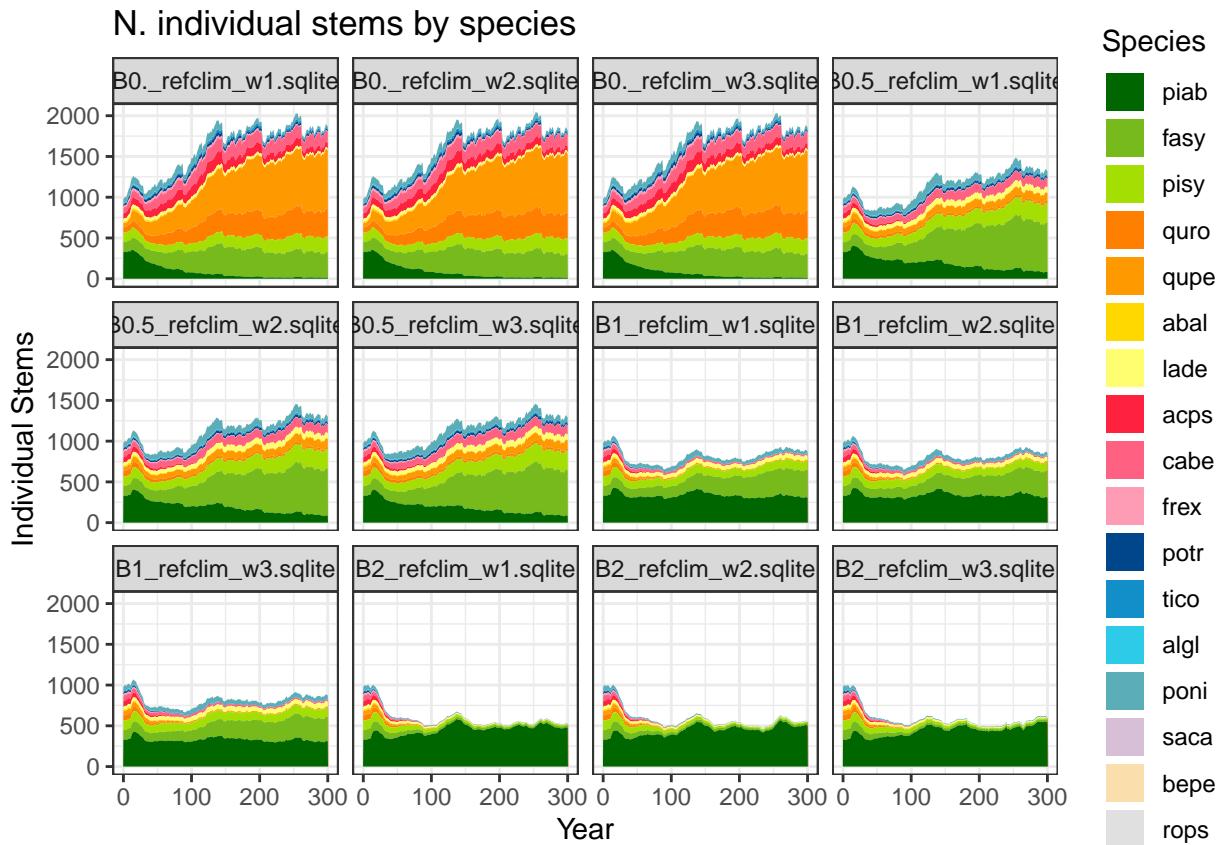
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+  

  ggtitle("N. individual stems by species") +  

  facet_wrap(~run, ncol=4)+  

  labs(x = "Year", y="Individual Stems", fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+  

  theme_bw()
```

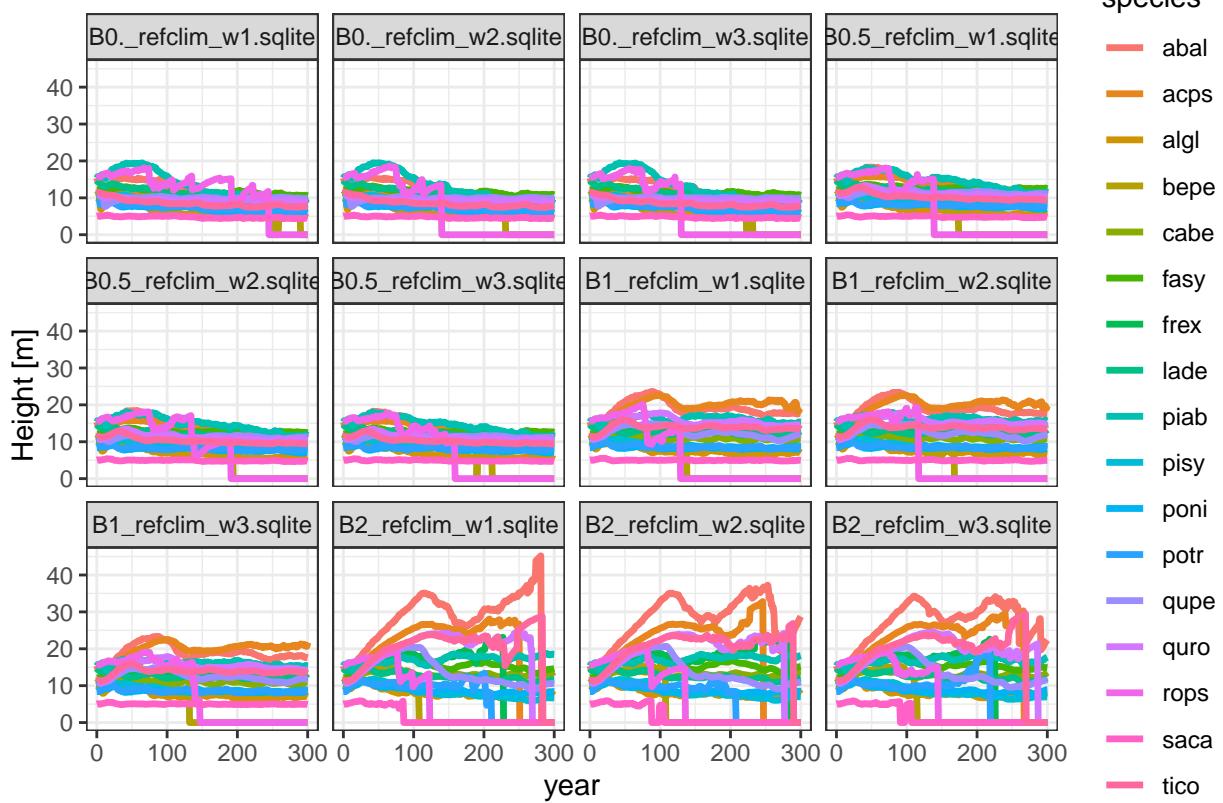


```
# PLOT HEIGHT GEOM_LINE AT LANDSCAPE LEVEL BY SPECIES

# lnd2 <- lnd %>% filter(species %in% species.to.keep)      #--- ADD IF YOU WANT SELECT SPECIES TO KEEP

ggplot(data=lnd, aes(x=year, y=height_avg_m, color=species)) +
  geom_line(size=1.2) +
  ggtitle("Avarage Height by species") +
  facet_wrap(~run, ncol=4) +
  theme(plot.title = element_text(hjust = 0.5)) +
  ylab("Height [m]") +
  theme_bw()
```

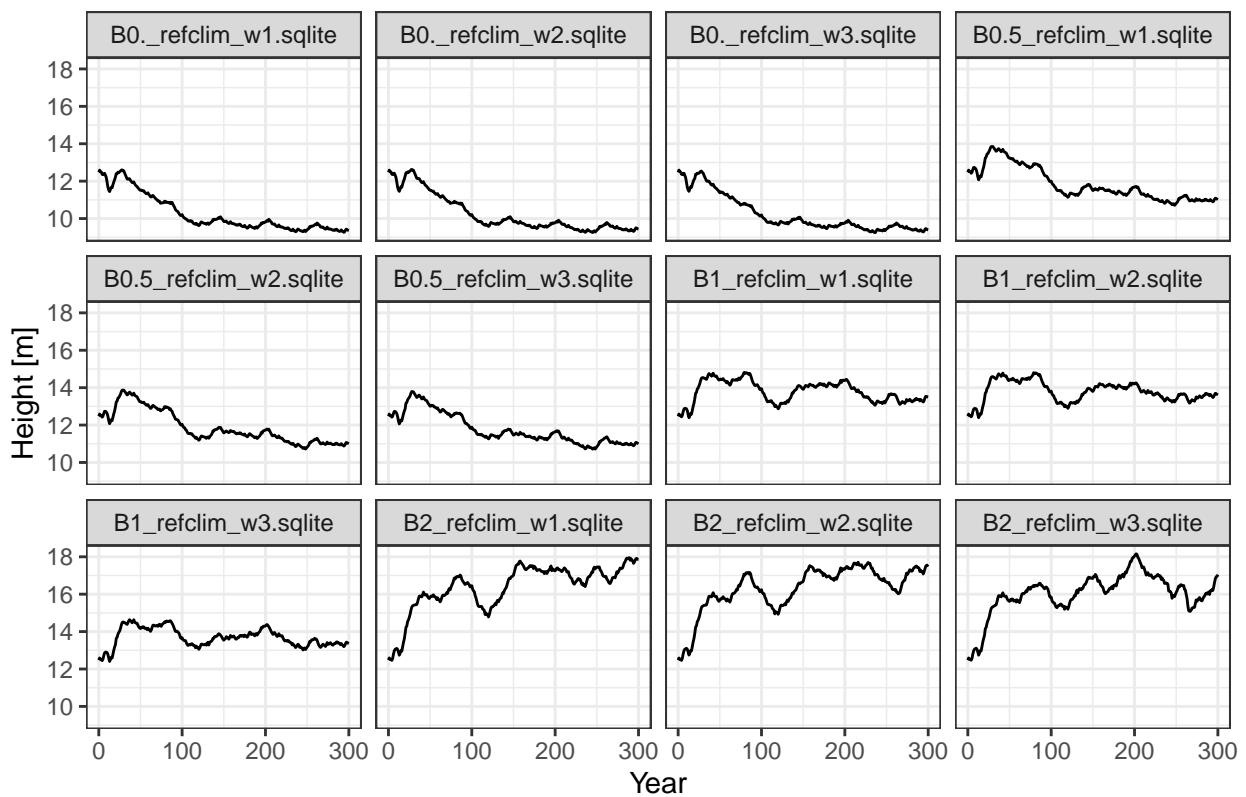
## Average Height by species



```
# PLOT TOTAL HEIGHT GEOM_AREA AT LANDSCAPE LEVEL BY SPECIES
```

```
ggplot(dys, aes(x=year, y=height_mean))+
  geom_line() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

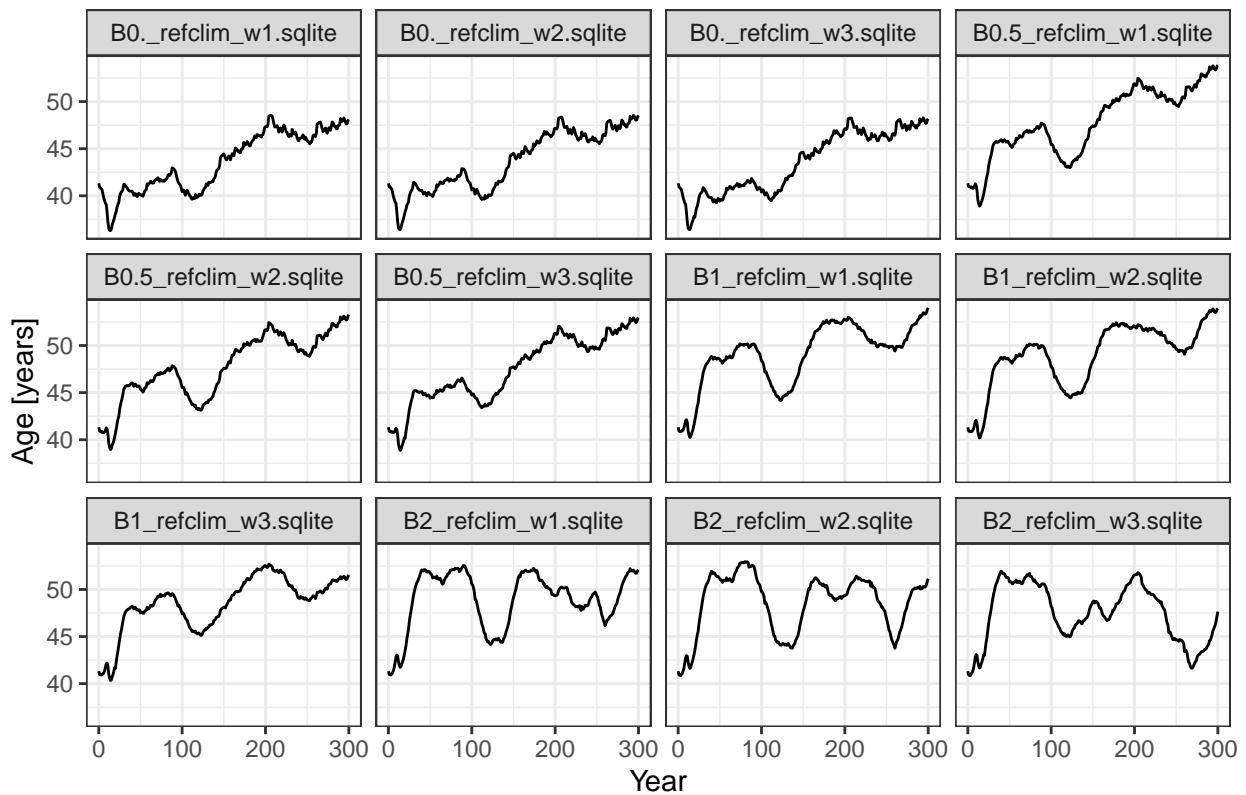
## Total Avarage Height



```
# AGE

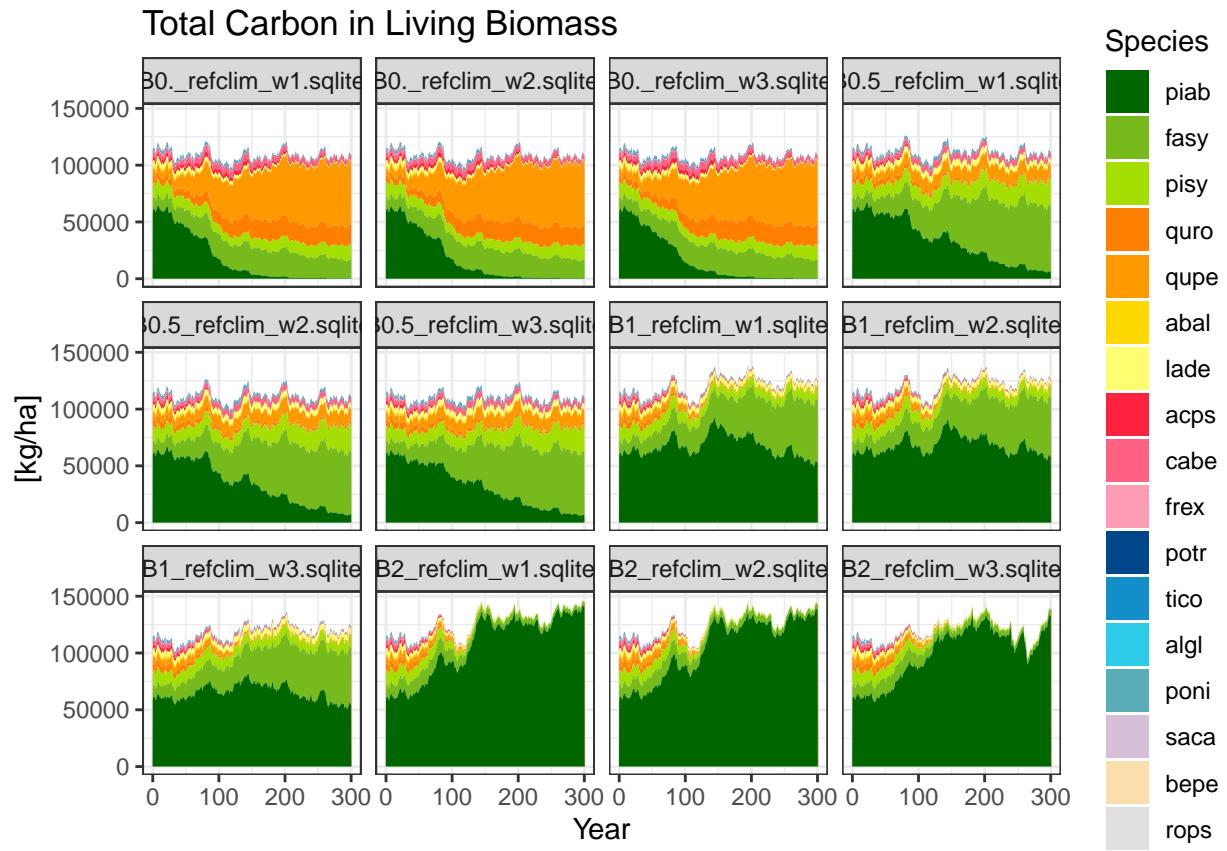
ggplot(dys, aes(x=year, y=age_mean))+
  geom_line() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
  ggtitle("Avarage Tree Age")+
  facet_wrap(~run, ncol=4)+
  labs(x = "Year", y="Age [years]")+
  theme(plot.title = element_text(hjust = 0.5))+
  theme_bw()
```

## Average Tree Age



```
# Total Carbon in Kg (total_carbon_kg)    double  total carbon in living biomass (aboveground compartment)

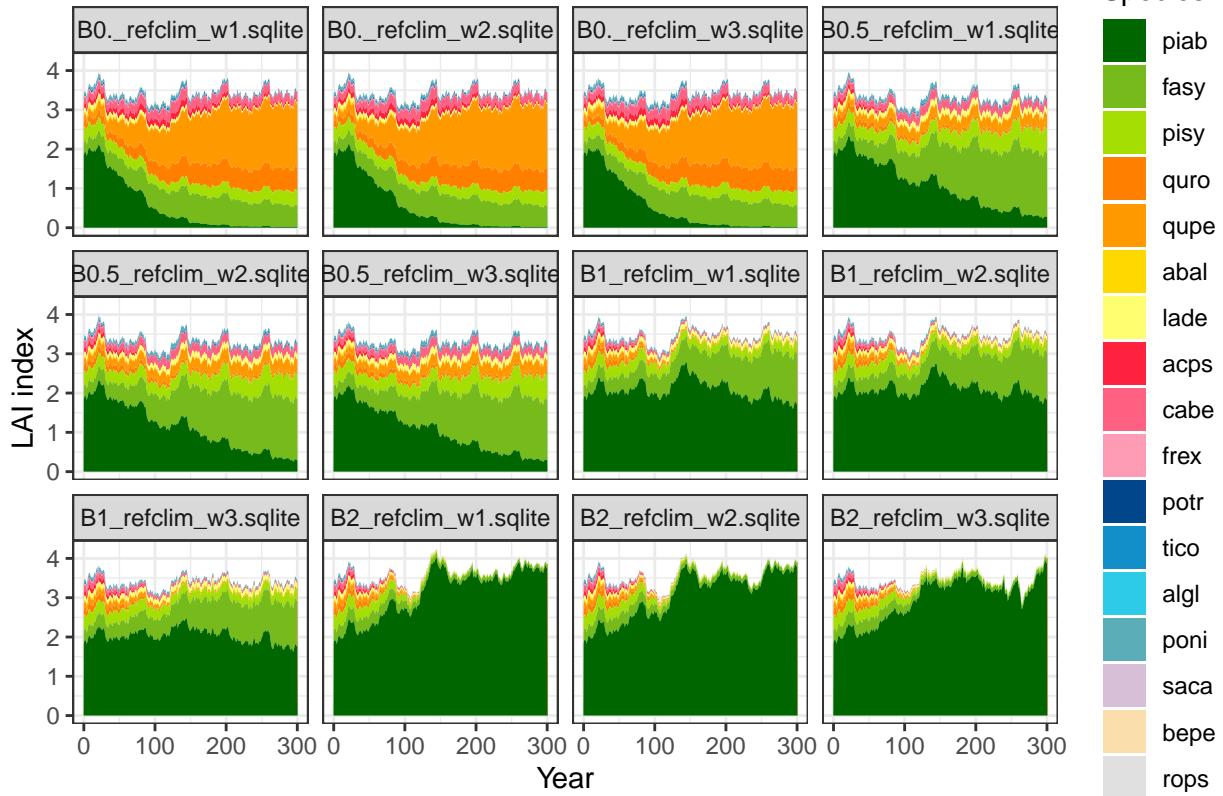
ggplot(lnd, aes(year, total_carbon_kg, fill=factor(species, levels=new_order_gg)))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
  ggtitle("Total Carbon in Living Biomass")+
  facet_wrap(~run, ncol=4)+
  labs(x = "Year", y = "[kg/ha]", fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+
  theme_bw()
```



```
# PLOT LAI AT LANDSCAPE LEVEL BY SPECIES
```

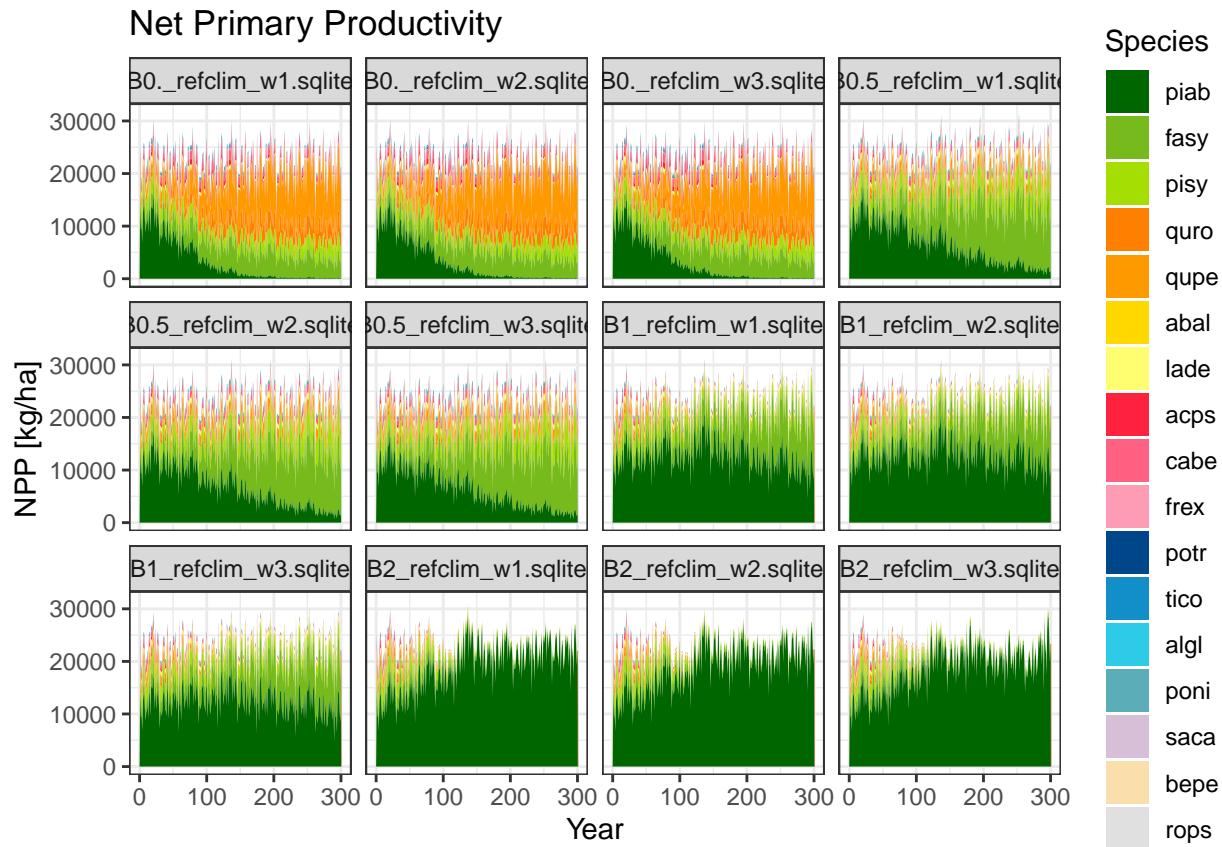
```
ggplot(lnd, aes(year, LAI, fill=factor(species, levels=new_order_gg)))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
  ggtitle("LAI index by species")+
  facet_wrap(~run, ncol=4)+
  labs(x = "Year", y="LAI index", fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+
  theme_bw()
```

## LAI index by species



```
# PLOT NPP AT LANDSCAPE LEVEL BY SPECIES
```

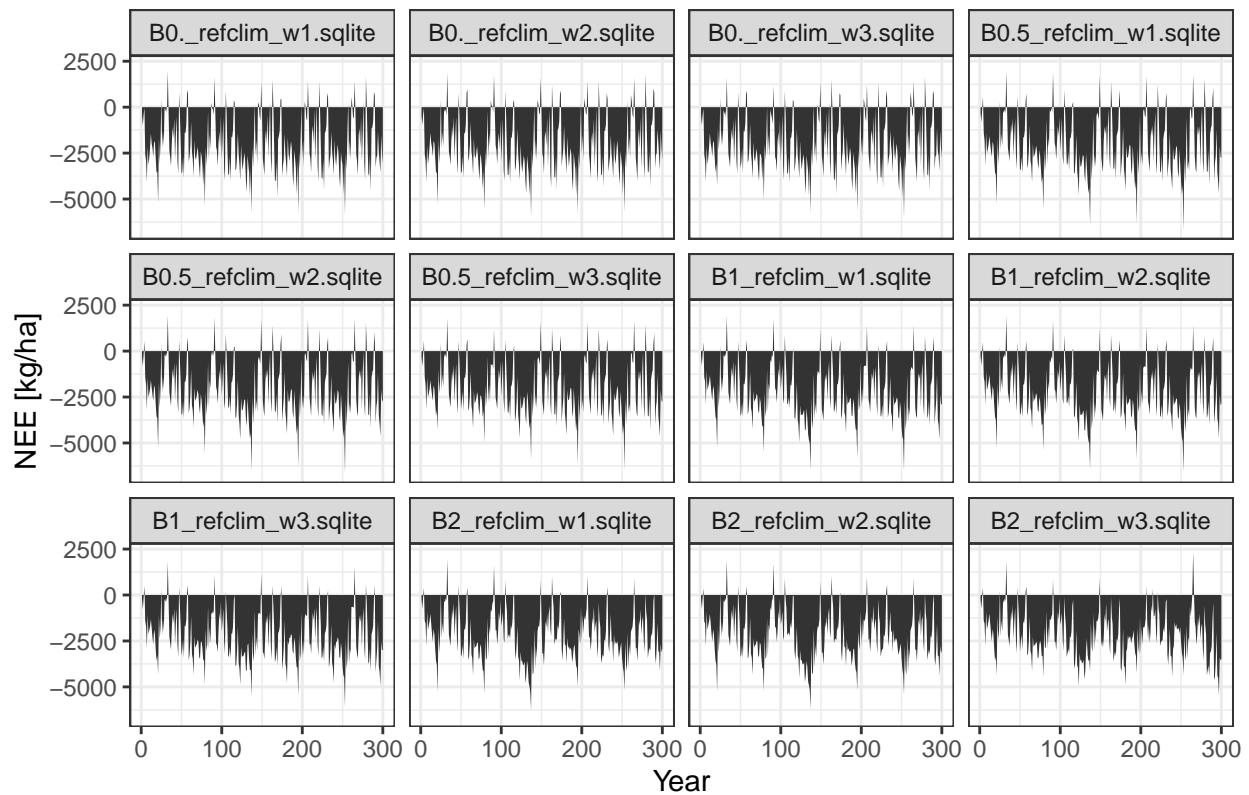
```
ggplot(lnd, aes(year, NPP_kg, fill=factor(species, levels=new_order_gg)))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
  ggtitle("Net Primary Productivity")+
  facet_wrap(~run, ncol=4)+
  labs(x = "Year", y="NPP [kg/ha]", fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+
  theme_bw()
```



```
# PLOT NEE AT LANDSCAPE LEVEL
```

```
ggplot(variables.all, aes(year, NEE))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE)) +
  ggtitle("Net Ecosystem Exchange") +
  facet_wrap(~case, ncol=4) +
  labs(x = "Year", y="NEE [kg/ha]") +
  theme(plot.title = element_text(hjust = 0.5)) +
  theme_bw()
```

## Net Ecosystem Exchange



```
# PLOT H count AT LANDSCAPE LEVEL
```

```
ggplot(variables.all, aes(year, H.count))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

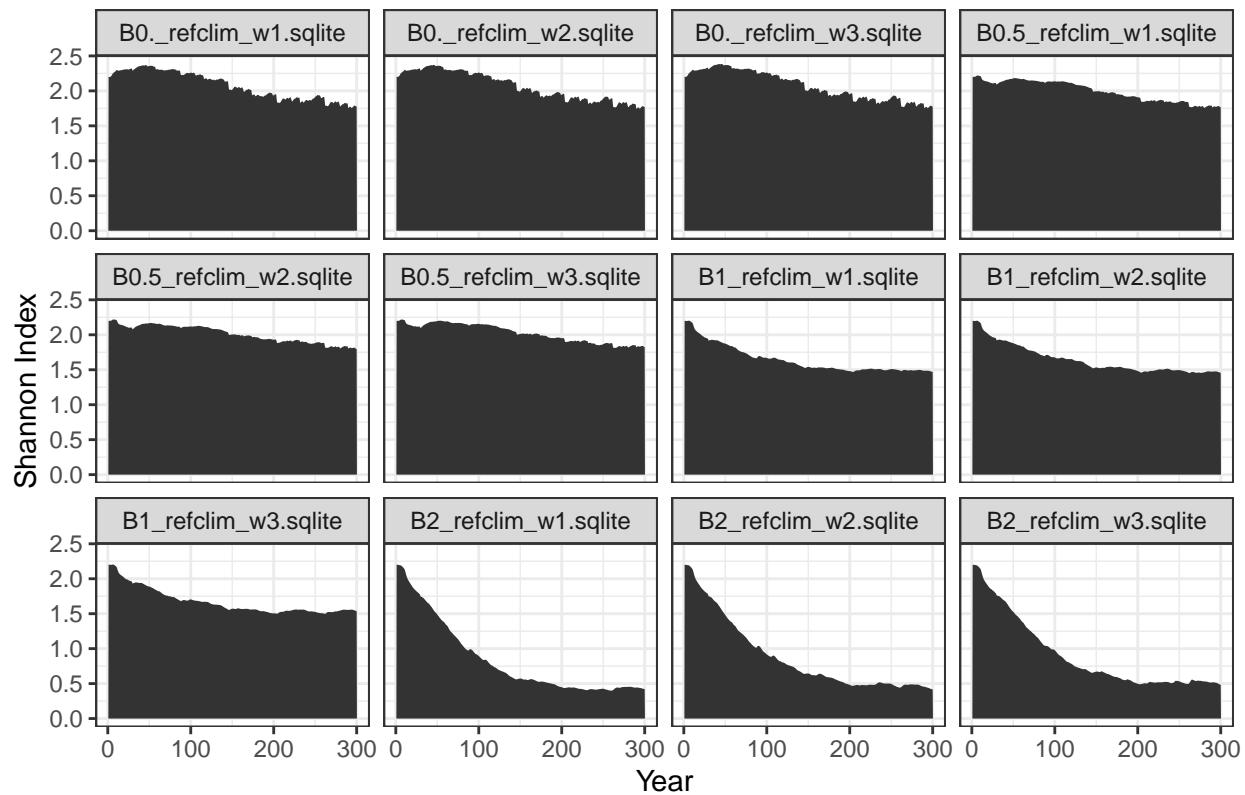
ggtitle("Shannon Biodiversity Index")+

```
facet_wrap(~case, ncol=4)+
```

```
labs(x = "Year", y="Shannon Index")+
  theme(plot.title = element_text(hjust = 0.5))+
```

```
theme_bw()
```

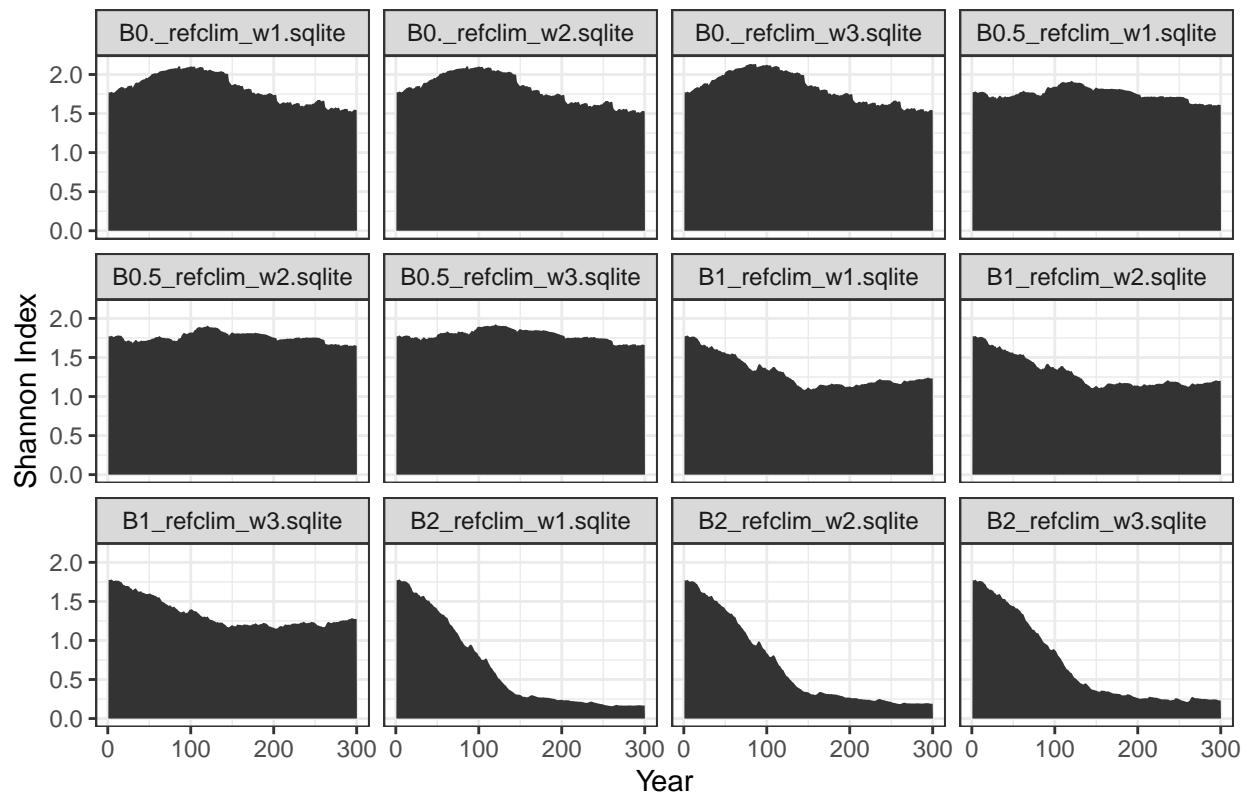
## Shannon Biodiversity Index



```
# PLOT H basal area AT LANDSCAPE LEVEL
```

```
ggplot(variables.all, aes(year, H.BA))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

## Shannon Biodiversity Index on Basal Area



```
# PLOT H basal area AT LANDSCAPE LEVEL
```

```
ggplot(variables.all, aes(year, NEP))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

ggtitle("Net Ecosystem Productivity")+

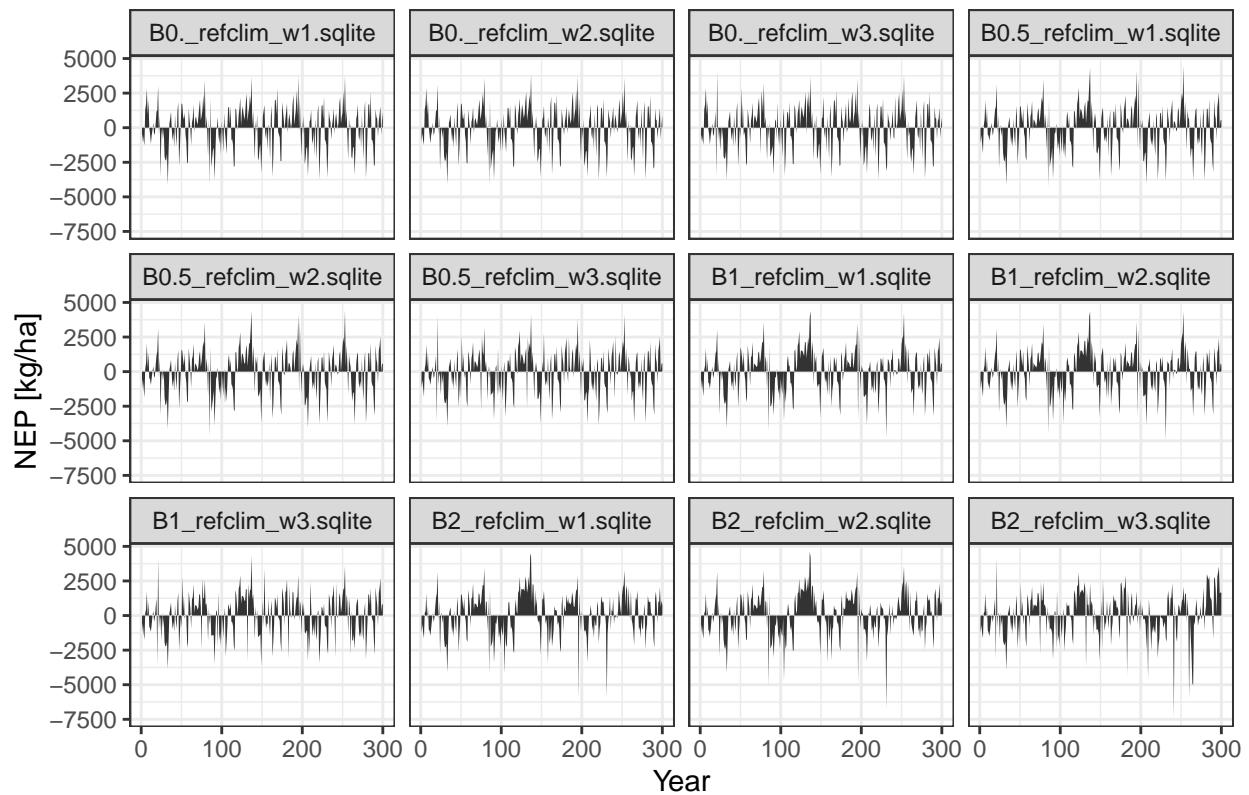
```
facet_wrap(~case, ncol=4)+
```

```
labs(x = "Year", y="NEP [kg/ha]")+
```

```
theme(plot.title = element_text(hjust = 0.5))+
```

```
theme_bw()
```

## Net Ecosystem Productivity



```
# PLOT GPP AT LANDSCAPE LEVEL
```

```
ggplot(variables.all, aes(year, GPP))+
  geom_area() +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

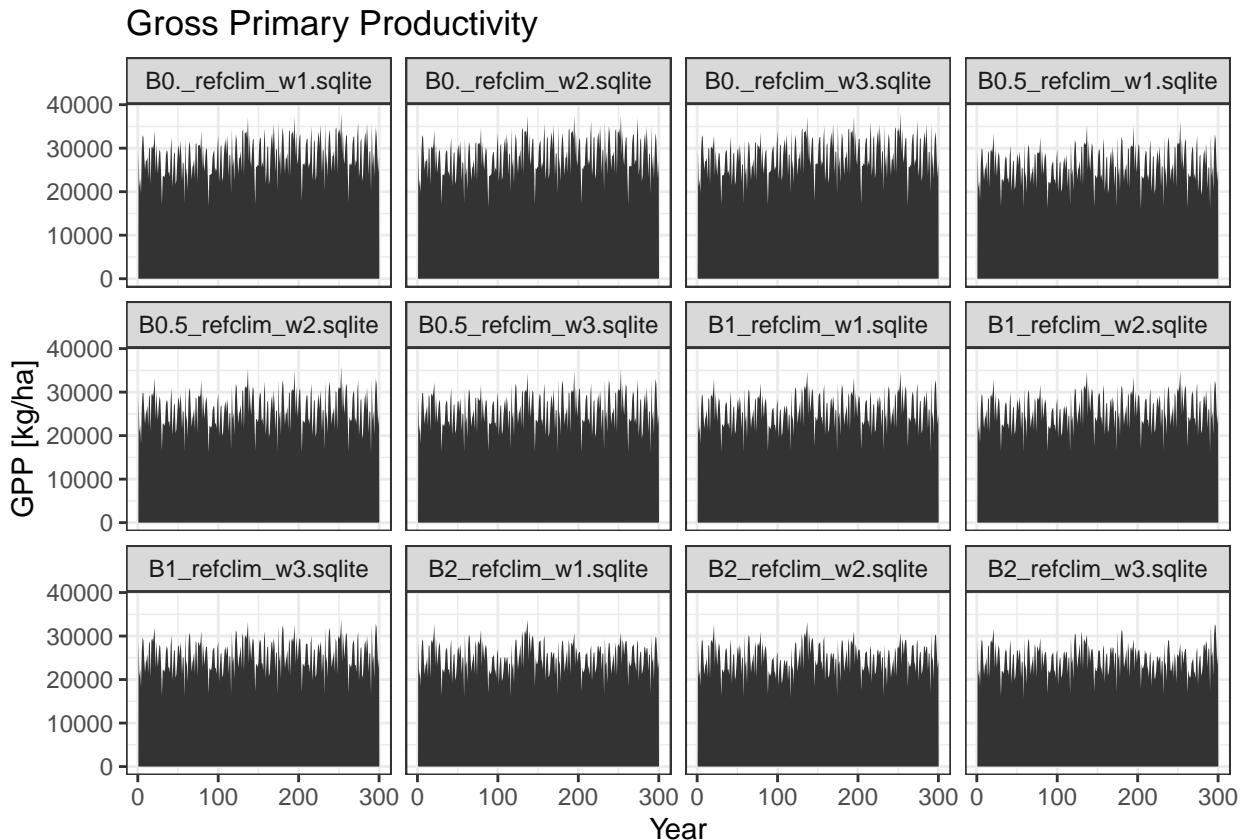
ggtitle("Gross Primary Productivity")+

```
facet_wrap(~case, ncol=4)+
```

```
labs(x = "Year", y="GPP [kg/ha]")+
```

```
theme(plot.title = element_text(hjust = 0.5))+
```

```
theme_bw()
```



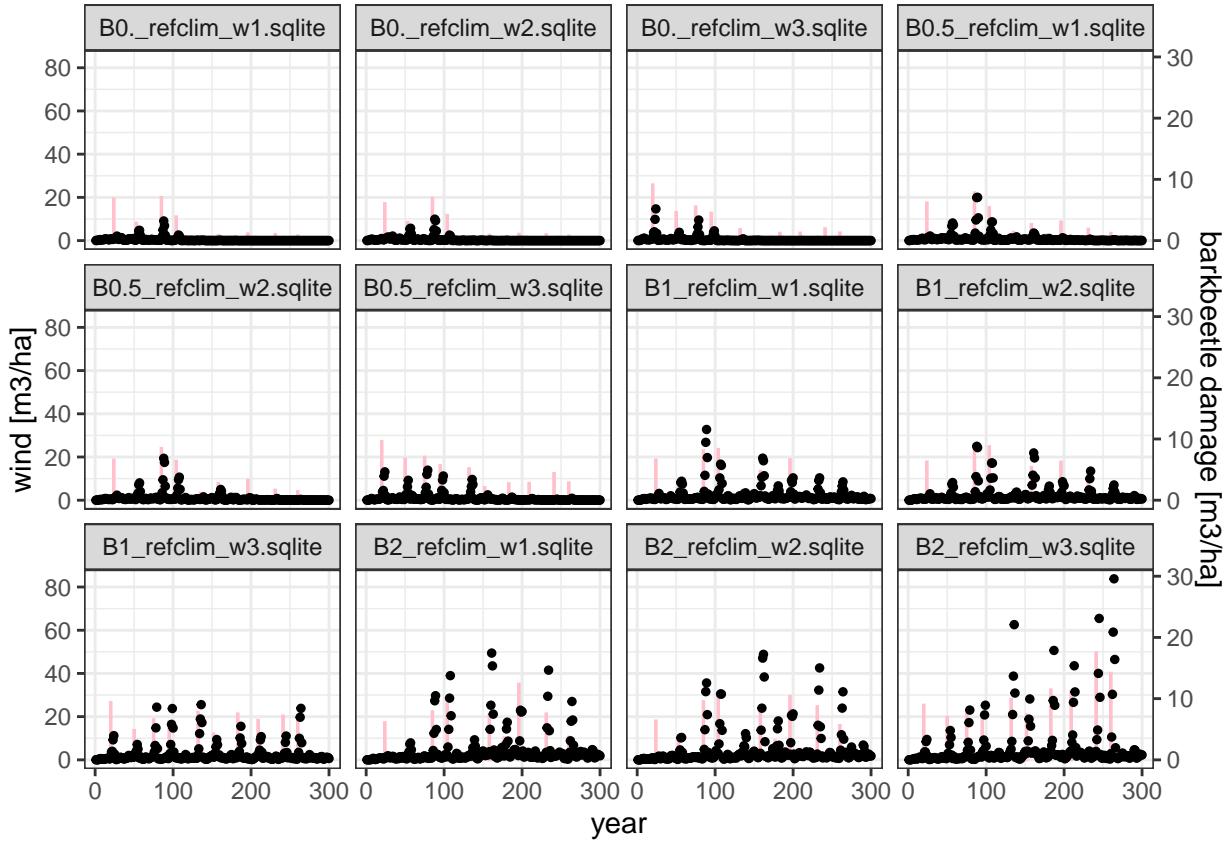
```
# PLOT SECOND Y AXIS FOR KILLED VOLUME BY DISTURBANCES IN LANDSCAPE AVG LINE 75

area<-lnd$area[1]
ylim.bb <- c(0, 600000)
ylim.w <- c(0, 1700000)

b <- diff(ylim.w)/diff(ylim.bb)
a <- ylim.w[1] - b*ylim.bb[1]

# TO MAKE 2 LINE AND 2 DIFFERENT SCALE PLOT "https://stackoverflow.com/questions/3099219/ggplot-with-2-
ggplot(damage.all,aes(year,killedVolume/area))+
  geom_col(fill="pink",col="pink")+
  geom_point(aes(y = a+ barkbeetle/area*b), data = damage.all,size=1) +
  scale_y_continuous(name="wind [m3/ha]", sec.axis = sec_axis(~ (. - a)/b, name = "barkbeetle damage [m3/ha]"),
  facet_wrap(~case, ncol=4)+
  theme_bw()
```

## Warning: Removed 3480 rows containing missing values ('position\_stack()').

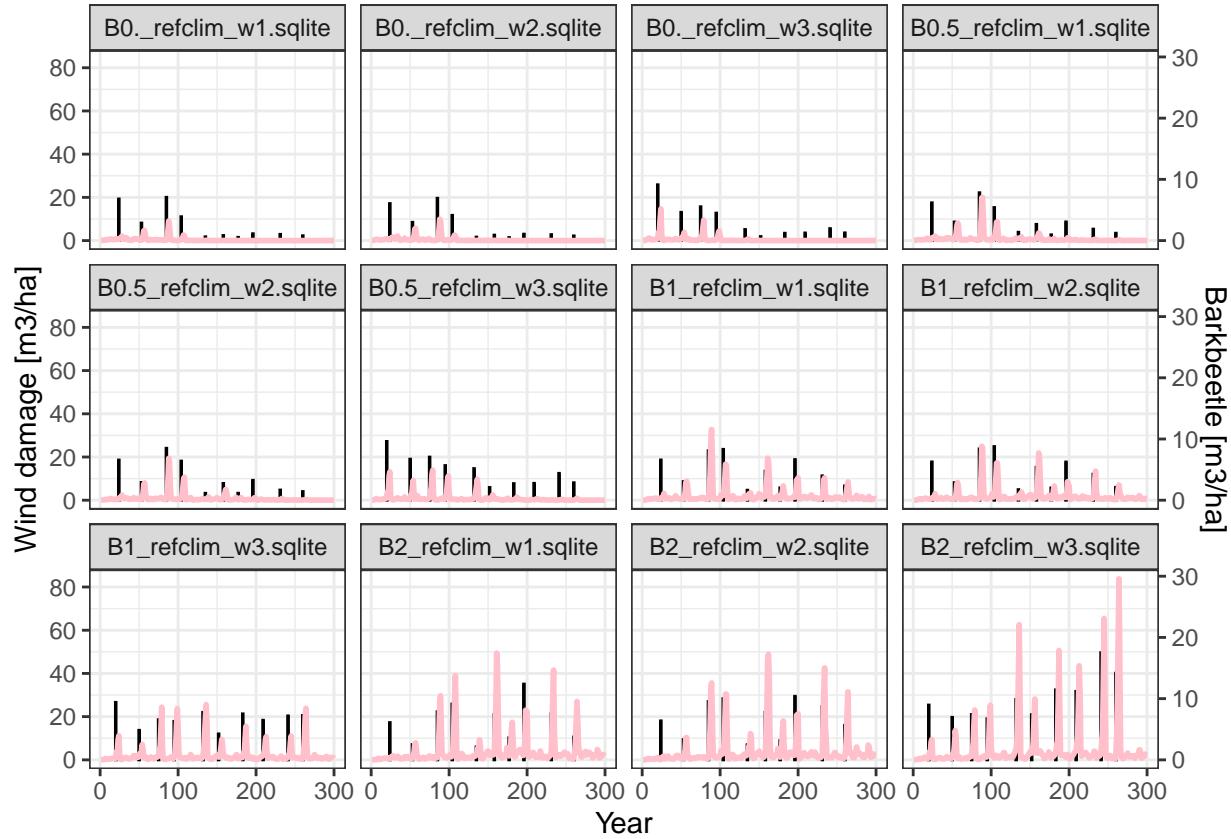


```
# variant 2

b <- diff(ylim.w)/diff(ylim.bb)
a <- ylim.w[1] - b*ylim.bb[1]

# TO MAKE 2 LINE AND 2 DIFFERENT SCALE PLOT "https://stackoverflow.com/questions/3099219/ggplot-with-2-
ggplot(damage.all,aes(year,killedVolume/area))+
  geom_col(fill="grey",col="black", size=0.5)+
  geom_line(aes(y = a+ barkbeetle/area*b), data = damage.all, size=1, col="pink") +
  scale_y_continuous(name="Wind damage [m³/ha]", sec.axis = sec_axis(~ (. - a)/b, name = "Barkbeetle [m³/ha]"),
  xlab("Year")+
  facet_wrap(~case, ncol=4)+
  theme_bw()

## Warning: Removed 3480 rows containing missing values ('position_stack()').
```



```
# Here gave me the right values
absolute_damage <- (damage.all %>% group_by(case) %>% summarise(mean(barkbeetle/area),mean(na.omit(killedVolume/area)))
absolute_damage
```

```
## # A tibble: 12 x 3
##   case           `mean(barkbeetle/area)` `mean(na.omit(killedVolume/area))` 
##   <chr>          <dbl>                   <dbl>
## 1 B0.5_refclim_w1.sqlite 0.289                  9.50
## 2 B0.5_refclim_w2.sqlite 0.290                 10.1 
## 3 B0.5_refclim_w3.sqlite 0.307                 13.9 
## 4 B0._refclim_w1.sqlite 0.112                  7.24
## 5 B0._refclim_w2.sqlite 0.126                  7.04
## 6 B0._refclim_w3.sqlite 0.143                  9.04
## 7 B1_refclim_w1.sqlite 0.727                 13.4 
## 8 B1_refclim_w2.sqlite 0.688                 13.6 
## 9 B1_refclim_w3.sqlite 0.795                 19.1 
## 10 B2_refclim_w1.sqlite 1.35                  17.7 
## 11 B2_refclim_w2.sqlite 1.40                  19.1 
## 12 B2_refclim_w3.sqlite 1.72                  28.8
```

```
## # ... with abbreviated variable name 1: `mean(na.omit(killedVolume/area))`
```

```
# PLOT SECOND Y AXIS IN RELATIVE KILLED VOLUME BY DISTURBANCES IN LANDSCAPE AVG
```

```
ylim.bb <- c(0, 600000)
ylim.w <- c(0, 1700000)
```

```

b <- diff(ylim.bb)/diff(ylim.w)
a <- ylim.bb[1] - b*ylim.w[1]

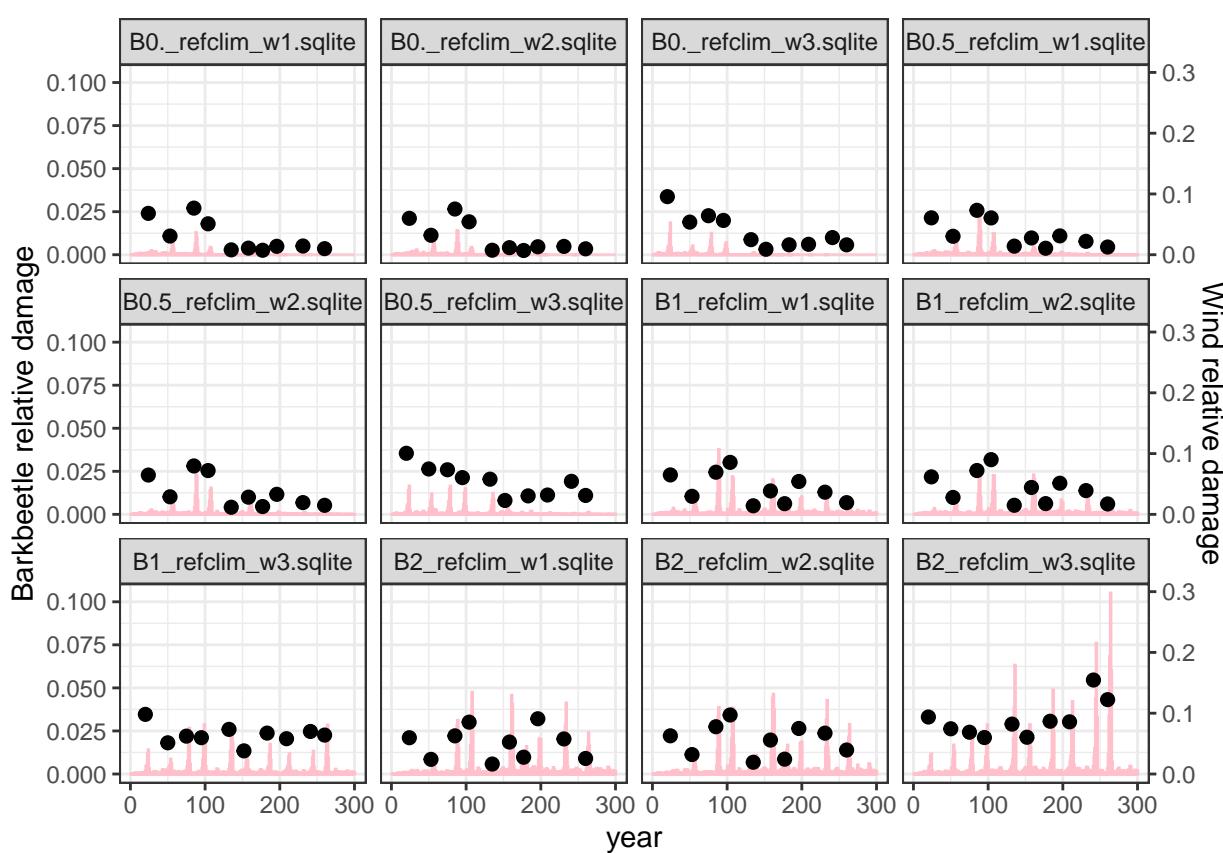
# TO MAKE 2 LINE AND 2 DIFFERENT SCALE PLOT "https://stackoverflow.com/questions/3099219/ggplot-with-2-
ggplot(damage.all,aes(year,barkbeetle/area/tot_vol))+  

  geom_col(fill="pink",col="pink")+
  geom_point(aes(y = a + killedVolume/area/tot_vol*b), data = damage.all,size=2) +
  scale_y_continuous(name="Barkbeetle relative damage", sec.axis = sec_axis(~ (. - a)/b, name = "Wind re-
  facet_wrap(~case, ncol=4)+  

  theme_bw()

```

## Warning: Removed 3480 rows containing missing values ('geom\_point()').



```

# SET THE SCALE OF THE Y AXIS. NAME IS THE NAME OF THE LABEL. FIRST Y AXIS SCALE REFERENCE ONE. SEC_AXI
rel_damage <- (damage.all %>%
  group_by(case) %>%
  summarise((barkbeetle/tot_vol),
            (killedVolume/tot_vol)))

```

## Warning: Returning more (or less) than 1 row per 'summarise()' group was deprecated in  
## dplyr 1.1.0.  
## i Please use 'reframe()' instead.  
## i When switching from 'summarise()' to 'reframe()', remember that 'reframe()'  
## always returns an ungrouped data frame and adjust accordingly.

```
## Call `lifecycle::last_lifecycle_warnings()` to see where this warning was
## generated.
```

```
## `summarise()` has grouped output by 'case'. You can override using the
## `.` argument.
```

```
rel_damage
```

```
## # A tibble: 3,600 x 3
## # Groups:   case [12]
##   case      `barkbeetle/tot_vol` `killedVolume/tot_vol` 
##   <chr>     <dbl>           <dbl>
## 1 B0.5_refclim_w1.sqlite 0.609        NA
## 2 B0.5_refclim_w1.sqlite 1.84         NA
## 3 B0.5_refclim_w1.sqlite 1.46         NA
## 4 B0.5_refclim_w1.sqlite 3.64         NA
## 5 B0.5_refclim_w1.sqlite 8.06         NA
## 6 B0.5_refclim_w1.sqlite 9.52         NA
## 7 B0.5_refclim_w1.sqlite 9.15         NA
## 8 B0.5_refclim_w1.sqlite 9.14         NA
## 9 B0.5_refclim_w1.sqlite 7.59         NA
## 10 B0.5_refclim_w1.sqlite 8.78         NA
## # ... with 3,590 more rows
```

```
relative_damage <- (damage.all %>%
  group_by(case) %>%
  summarise(mean(barkbeetle/tot_vol),
            mean(na.omit(killedVolume/tot_vol))))
```

# SUMMARISE THE DATA

```
## # A tibble: 12 x 3
##   case      `mean(barkbeetle/tot_vol)` `mean(na.omit(killedVolume~1)`
##   <chr>     <dbl>           <dbl>
## 1 B0.5_refclim_w1.sqlite 18.6          609.
## 2 B0.5_refclim_w2.sqlite 18.7          649.
## 3 B0.5_refclim_w3.sqlite 20.0          954.
## 4 B0._refclim_w1.sqlite  7.81          516.
## 5 B0._refclim_w2.sqlite  8.82          504.
## 6 B0._refclim_w3.sqlite  9.71          677.
## 7 B1_refclim_w1.sqlite   41.1          763.
## 8 B1_refclim_w2.sqlite   38.8          774.
## 9 B1_refclim_w3.sqlite   45.9          1140.
## 10 B2_refclim_w1.sqlite  66.6          892.
## 11 B2_refclim_w2.sqlite  70.2          978.
## 12 B2_refclim_w3.sqlite  92.3          1578.
## # ... with abbreviated variable name 1: `mean(na.omit(killedVolume/tot_vol))`
```

```
# variant 1
ylim.bb <- c(0, 600000)
ylim.w <- c(0, 1700000)

b <- diff(ylim.w)/diff(ylim.bb)
```

```

a <- ylim.w[1] - b*ylim.bb[1]

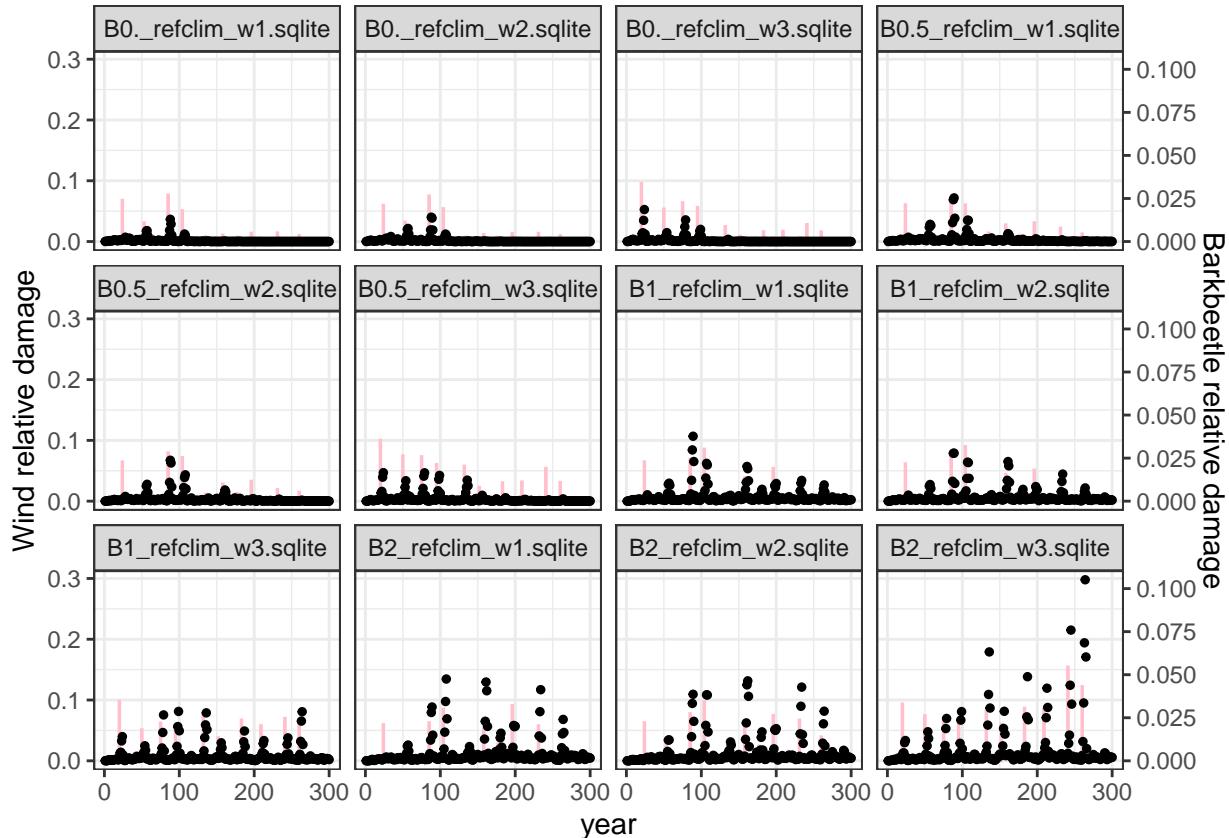
# TO MAKE 2 LINE AND 2 DIFFERENT SCALE PLOT "https://stackoverflow.com/questions/3099219/ggplot-with-2-
ggplot(damage.all,aes(year,killedVolume/area/tot_vol))+  

  geom_col(fill="pink",col="pink")+
  geom_point(aes(y = a+ barkbeetle/area/tot_vol*b), data = damage.all,size=1) +
  scale_y_continuous(name="Wind relative damage", sec.axis = sec_axis(~ (. - a)/b, name = "Barkbeetle re-
  facet_wrap(~case, ncol=4)+  

  theme_bw()

```

## Warning: Removed 3480 rows containing missing values ('position\_stack()'').



```

# variant 2

b <- diff(ylim.w)/diff(ylim.bb)
a <- ylim.w[1] - b*ylim.bb[1]

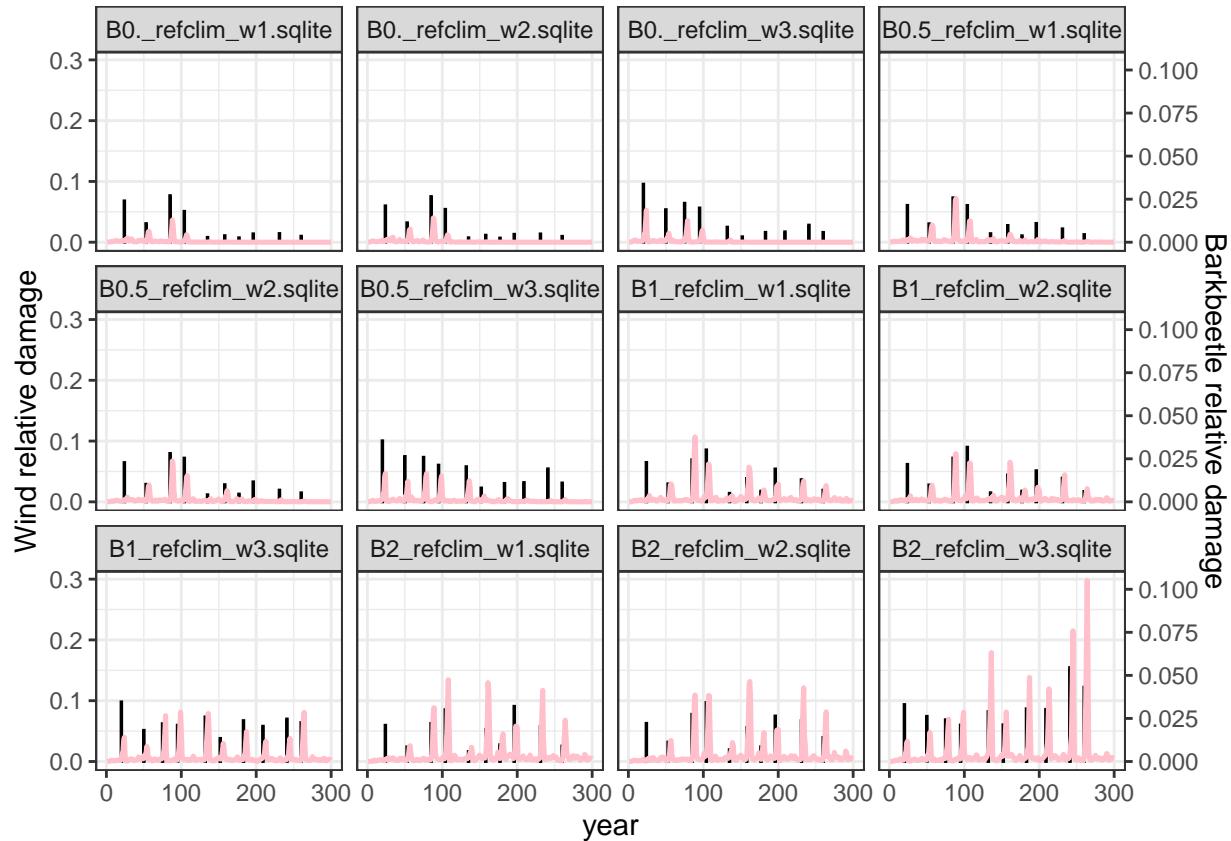
# TO MAKE 2 LINE AND 2 DIFFERENT SCALE PLOT "https://stackoverflow.com/questions/3099219/ggplot-with-2-
ggplot(damage.all,aes(year,killedVolume/area/tot_vol))+  

  geom_col(fill="grey",col="black")+
  geom_line(aes(y = a+ barkbeetle/area/tot_vol*b), data = damage.all, size=0.9, col="pink") +
  scale_y_continuous(name="Wind relative damage", sec.axis = sec_axis(~ (. - a)/b, name = "Barkbeetle re-
  facet_wrap(~case, ncol=4)+  

  theme_bw()

```

```
## Warning: Removed 3480 rows containing missing values ('position_stack()').
```



```
# Grid for the paper! #
```

```
# PLOT NUMBER OF STEMS GEOM_AREA AT LANDSCAPE LEVEL BY SPECIES
```

```
ggplot(lnd, aes(x=year, y=count_ha, fill=factor(species, levels=new_order_gg)))+
  geom_area(show.legend = F)+
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

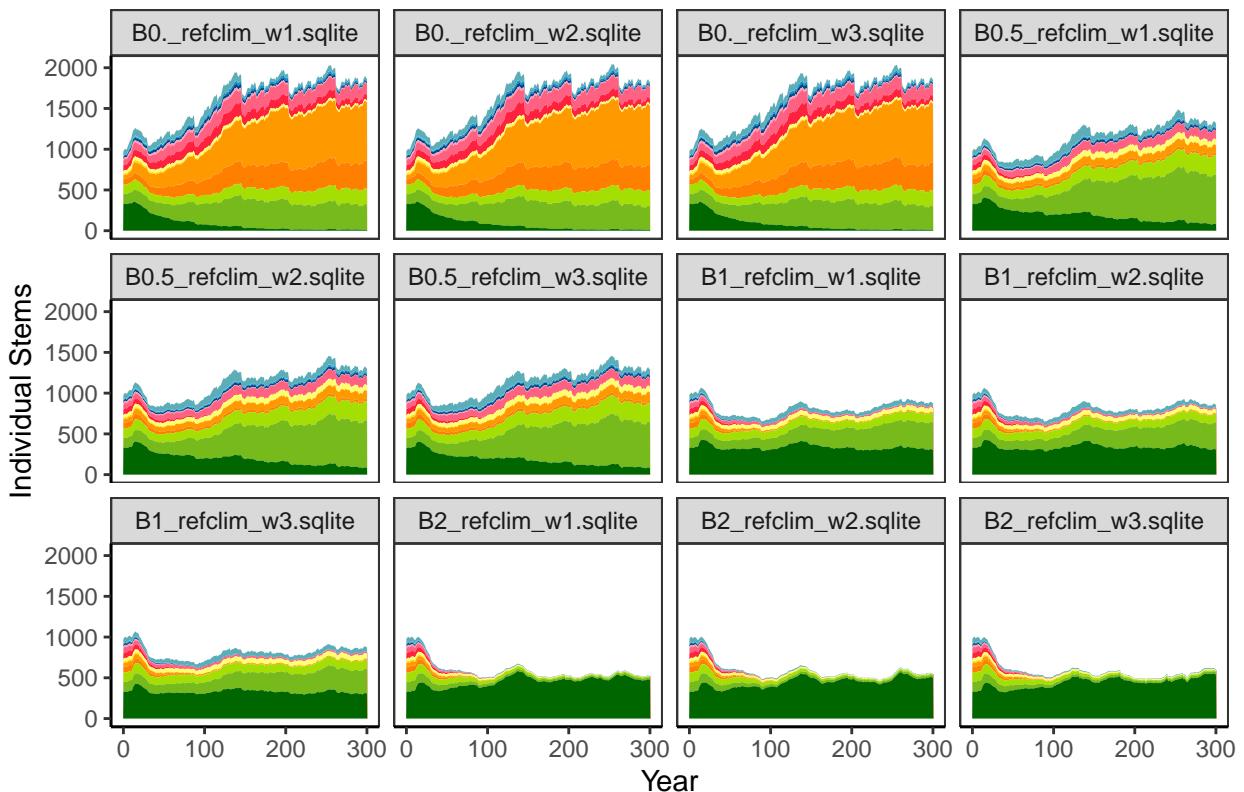
N. individual stems by species

```
+ facet_wrap(~run, ncol=4)+
```

```
  labs(x = "Year",y="Individual Stems", fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+
```

```
  theme_bw()+
  theme(axis.line = element_line(color='black'),
        plot.background = element_blank(),
        panel.grid.minor = element_blank(),
        panel.grid.major = element_blank())
```

## N. individual stems by species



```
#theme_bw() # this is to not have the gray - classic you will have a blank on -> in case for others l
```

```
# https://www.statology.org/ggplot-remove-gridlines/
```

```
# Total Carbon in Kg (total_carbon_kg    double    total carbon in living biomass (aboveground compartment)
```

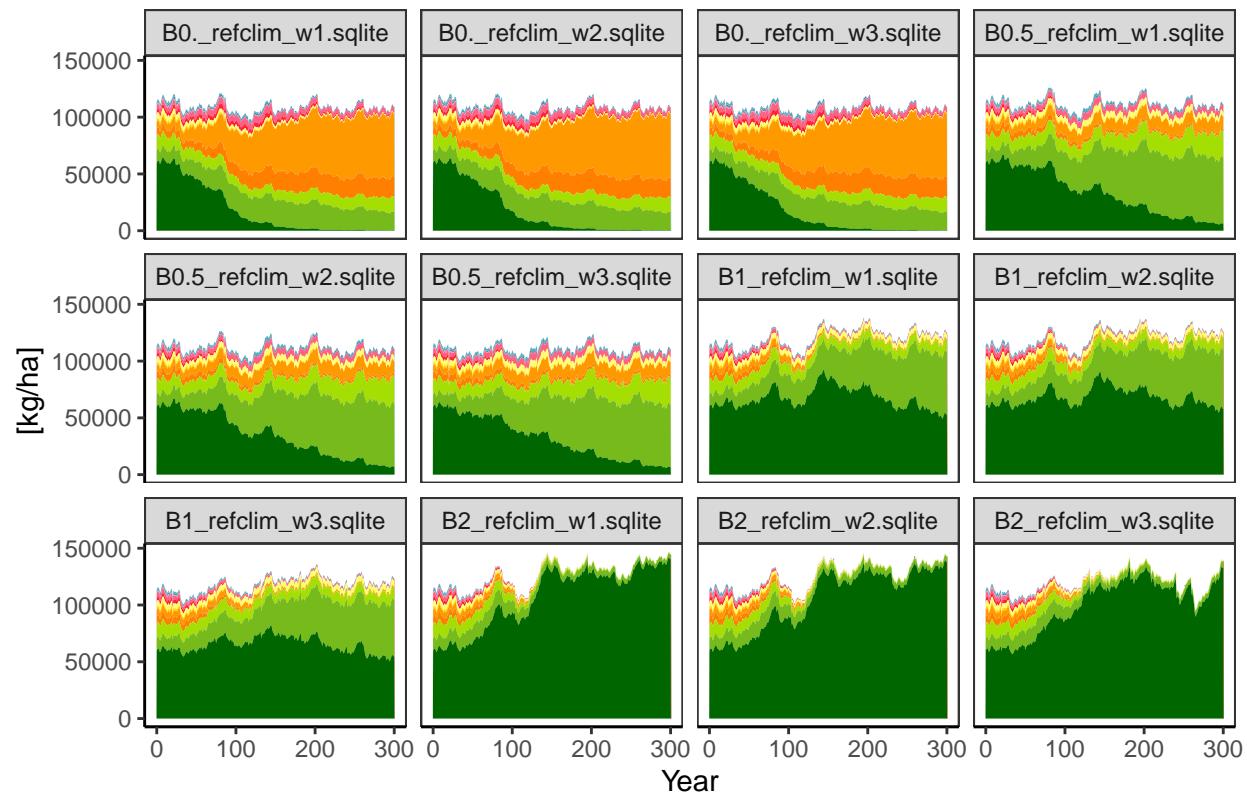
```
ggplot(lnd, aes(year, total_carbon_kg, fill=factor(species, levels=new_order_gg)))+
  geom_area(show.legend = F) +
  scale_fill_manual(values=cols[new_order_gg], guide=guide_legend(reverse=TRUE))+
```

```
  ggttitle("Total Carbon in Living Biomass")+
  facet_wrap(~run, ncol=4)+
```

```
  labs(x = "Year",y=" [kg/ha]",fill = "Species")+
  theme(plot.title = element_text(hjust = 0.5))+
```

```
  theme_bw()+
  theme(axis.line = element_line(color='black'),
        plot.background = element_blank(),
        panel.grid.minor = element_blank(),
        panel.grid.major = element_blank())
```

## Total Carbon in Living Biomass

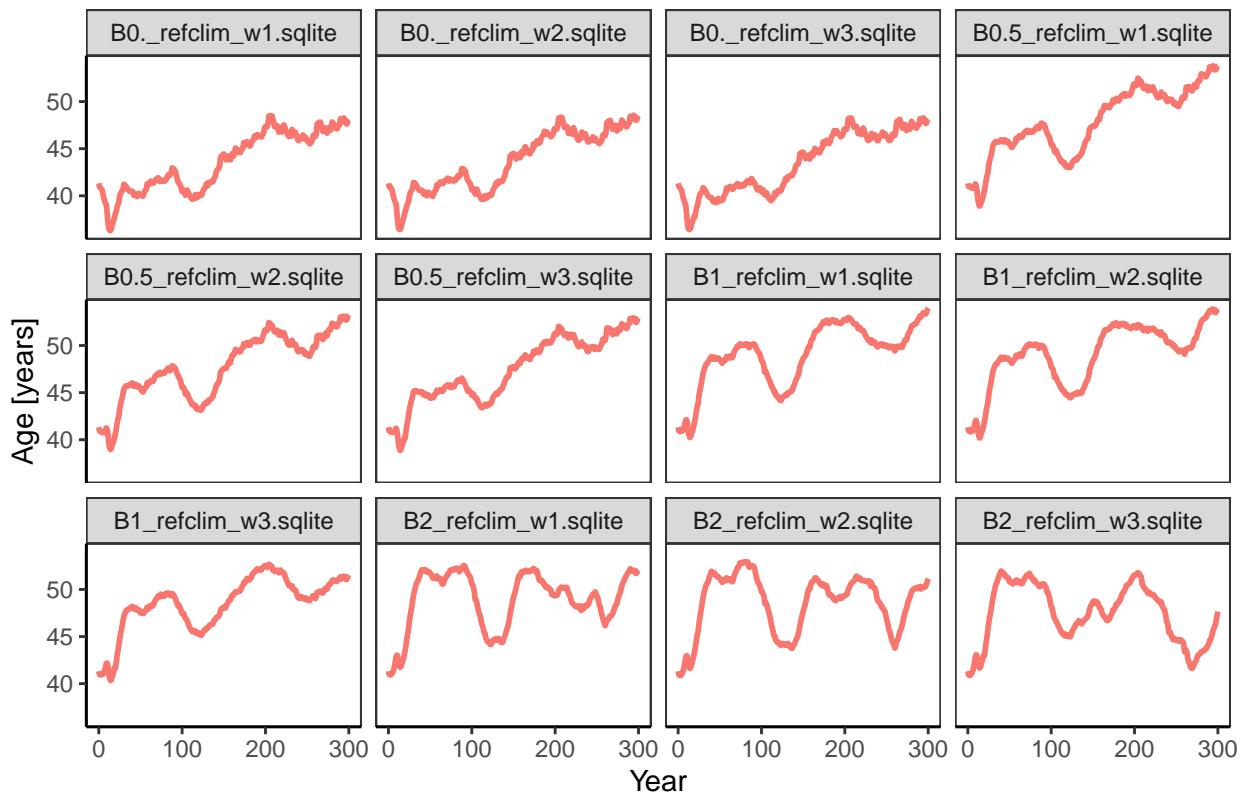


```
# theme_classic()

# AGE

ggplot(dys, aes(x=year, y=age_mean, color="red"))+
  geom_line(size=1, show.legend = F) +
  ggtitle("Avarage Trees Age")+
  facet_wrap(~run, ncol=4)+
  labs(x = "Year",y="Age [years]")+
  theme(plot.title = element_text(hjust = 0.5))+
  theme_bw()+
  theme(axis.line = element_line(color='black'),
        plot.background = element_blank(),
        panel.grid.minor = element_blank(),
        panel.grid.major = element_blank())
```

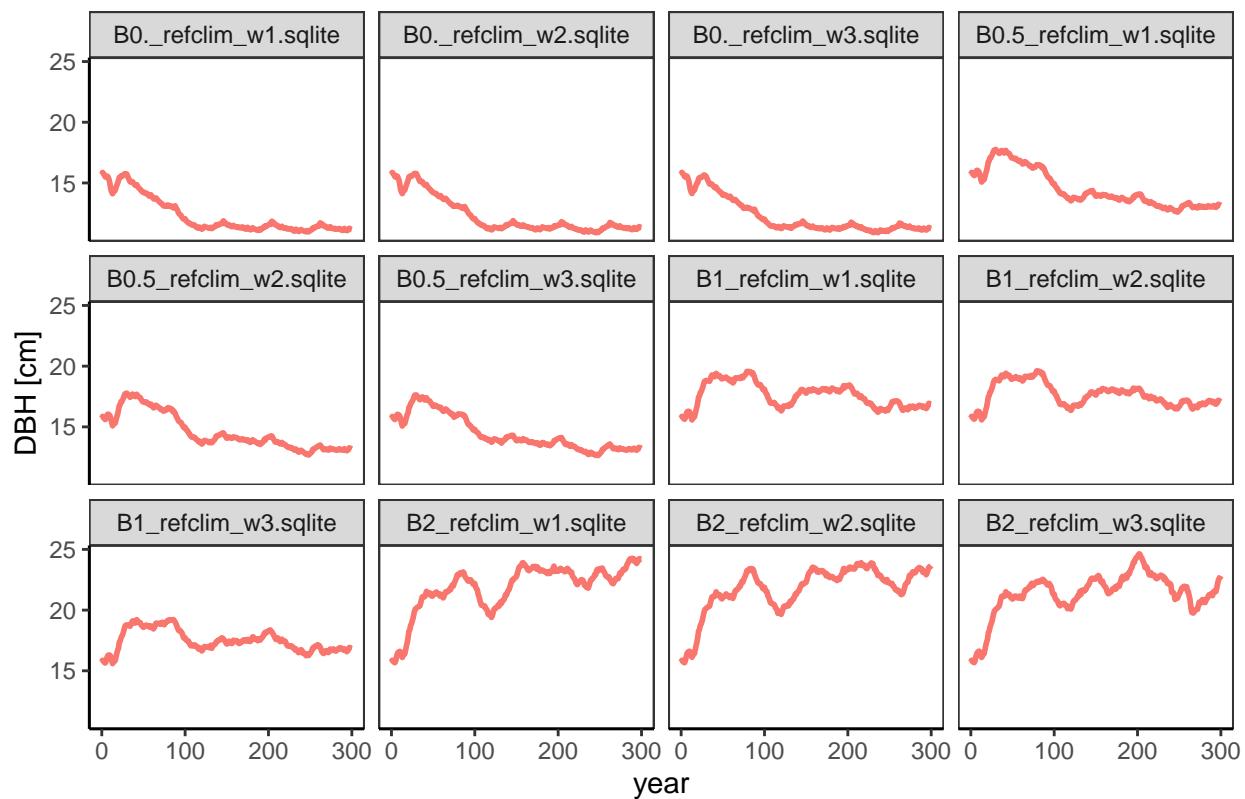
## Average Trees Age



```
# PLOT DBH GEOM_AREA AT LANDSCAPE LEVEL AVERAGE ALL SP TOGETHER
```

```
ggplot(data=dys, aes(x=year, y=dbh_mean, color=case)) +
  geom_line(size=1, show.legend = F) +
  ggtitle("Average DBH by species") +
  facet_wrap(~run, ncol=4) +
  theme(plot.title = element_text(hjust = 0.5)) +
  ylab("DBH [cm]") +
  theme_bw() +
  theme(axis.line = element_line(color='black'),
        plot.background = element_blank(),
        panel.grid.minor = element_blank(),
        panel.grid.major = element_blank())
```

## Average DBH by species



```

#_
# NEW KILLED VOLUME CALCULATION INCLUDING BARK BEETLE
# Make the disturbance impact:

# Wind AND bark beetle disturbance impact in the landscape volume in percentages
# wind and bark beetle regime

#_
killed_volume_w <- sum(w$killedVolume)
killed_volume_w

## [1] 29911115

killed_volume_bb <- sum(bb$killedVolume)
killed_volume_bb

## [1] 42348794

killed_volume_dist <- killed_volume_w + killed_volume_bb
killed_volume_dist

## [1] 72259909

```

```

killed_volume_per_year_dist <- killed_volume_dist/300
killed_volume_per_year_dist

## [1] 240866.4

killed_volume_per_year_dist_ha <- killed_volume_per_year_dist/17749.26
killed_volume_per_year_dist_ha

## [1] 13.5705

killed_volume_w <- sum(wind$killedVolume)
killed_volume_w

## [1] 5110448

killed_volume_bb <- sum(barkbeetle$killedVolume)
killed_volume_bb

## [1] 9154517

killed_volume_dist <- killed_volume_w + killed_volume_bb
killed_volume_dist

## [1] 14264965

killed_volume_per_year_dist <- killed_volume_dist/200
killed_volume_per_year_dist

## [1] 71324.83

killed_volume_per_year_dist_ha <- killed_volume_per_year_dist/17749.26
killed_volume_per_year_dist_ha

## [1] 4.018468

# FILTER AND GROUP FOR THE NEEDED COLUMNS AND GROUP BY YEAR TO CREATE NEW DATAFRAMES FOR THE % ANALYSIS

dfnew1 <- lnd[,c(1,8)]

df_vol = dfnew1 %>% group_by(year) %>%
  summarise(tot_vol = sum(volume_m3),
            .groups = 'drop')

prop_killed_vol_ha_year <- df_vol %>% mutate(perc.vol=100*killed_volume_per_year_dist_ha/tot_vol)
prop_killed_vol_ha_year

```

```

## # A tibble: 301 x 3
##   year tot_vol perc.vol
##   <int>   <dbl>    <dbl>
## 1     0   3495.  0.115
## 2     1   3589.  0.112
## 3     2   3551.  0.113
## 4     3   3531.  0.114
## 5     4   3408.  0.118
## 6     5   3454.  0.116
## 7     6   3563.  0.113
## 8     7   3652.  0.110
## 9     8   3663.  0.110
## 10    9   3616.  0.111
## # ... with 291 more rows

```

```
summary(prop_killed_vol_ha_year)
```

```

##      year      tot.vol      perc.vol
##  Min.   : 0   Min.   :3015   Min.   :0.1028
##  1st Qu.: 75  1st Qu.:3352  1st Qu.:0.1123
##  Median :150  Median :3469  Median :0.1158
##  Mean   :150  Mean   :3471  Mean   :0.1161
##  3rd Qu.:225  3rd Qu.:3578  3rd Qu.:0.1199
##  Max.   :300  Max.   :3911  Max.   :0.1333

```

```
dfnew1 <- landscape[,c(1,8)]
```

```

df.vol = dfnew1 %>% group_by(year) %>%
  summarise(tot.vol = sum(volume_m3),
            .groups = 'drop')

```

```

prop_killed_vol_ha_year <- df.vol %>% mutate(perc.vol=100*killed_volume_per_year_dist_ha/tot.vol)
prop_killed_vol_ha_year

```

```

## # A tibble: 301 x 3
##   year tot.vol perc.vol
##   <int>   <dbl>    <dbl>
## 1     0   291.   1.38
## 2     1   299.   1.34
## 3     2   296.   1.36
## 4     3   294.   1.37
## 5     4   284.   1.41
## 6     5   288.   1.40
## 7     6   297.   1.35
## 8     7   304.   1.32
## 9     8   305.   1.32
## 10    9   301.   1.34
## # ... with 291 more rows

```

```
summary(prop_killed_vol_ha_year)
```

```

##      year      tot.vol      perc.vol
##  Min.   : 0   Min.   :259.1   Min.   :0.9747
##  1st Qu.: 75  1st Qu.:304.8   1st Qu.:1.1002
##  Median :150  Median :336.3   Median :1.1950
##  Mean   :150  Mean   :334.9   Mean   :1.2161
##  3rd Qu.:225  3rd Qu.:365.3   3rd Qu.:1.3182
##  Max.   :300  Max.   :412.3   Max.   :1.5512

hist(prop_killed_vol_ha_year$perc.vol,
  main = "Landscape proportion of killed volume [m3/ha] by wind and bark beetles per year in CC",
  cex.main = 1, xlab = "[Killed volume / Total volume] = [%]",
  ylab = "Frequency [years]",
  cex.lab = 1,
  col="lightblue",
  breaks = "FD")

```

### Landscape proportion of killed volume [m3/ha] by wind and bark beetles per year in CC

