

# Models Fitted

Loïc Pages

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## Introduction

```
rm(list=ls())
library(knitr)
library(spamM)

## Registered S3 methods overwritten by 'registry':
##   method           from
##   print.registry_field proxy
##   print.registry_entry proxy

## spamM (Rousset & Ferdy, 2014, version 4.5.30) is loaded.
## Type 'help(spamM)' for a short introduction,
## 'news(package='spamM')' for news,
## and 'citation('spamM')' for proper citation.
## Further infos, slides, etc. at https://gitlab.mbb.univ-montp2.fr/francois/spamm-ref.

library(tidyverse)

## -- Attaching core tidyverse packages ----- tidyverse 2.0.0 --
## v dplyr     1.1.4    v readr     2.1.5
## vforcats   1.0.0    v stringr   1.5.1
## v ggplot2   3.5.1    v tibble    3.2.1
## v lubridate 1.9.4    v tidyverse 1.3.1
## v purrr    1.0.2

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

library(splines)
library(patchwork)

setwd("/media/loic/Commun/0Travail/Stage 2025 ISEM/Models")
centauree_data <- read.csv("donneesIPM_short.csv")
```

```

centauree_data_complet <- read.csv("donneesIPM.csv")

#Supprimer plantes dont l'age est inconnu
centauree_data <- centauree_data[!is.na(centauree_data$age0), ]
centauree_data$age1 <- ifelse(centauree_data$Stage1=="V",centauree_data$age0+1,NA)

#Forcer l'age maximal à 8
length(centauree_data$age0[centauree_data$age0 >= 8])

## [1] 93

centauree_data$age0[centauree_data$age0 > 8] <- 8

spaMM.options(separation_max=70)

annees <- 1995:2022
populations <- c("Po","Au","Pe","E1","E2","Cr")
taille_range <- seq(0.5, 20, by = 0.5)
age_range <- 1:8

fake_data <- expand.grid(
  year = annees,
  Pop = populations,
  Size0Mars = taille_range,
  age0 = age_range
)

fake_data <- fake_data %>%
  mutate(Nrw = row_number())

```

## Survival probability

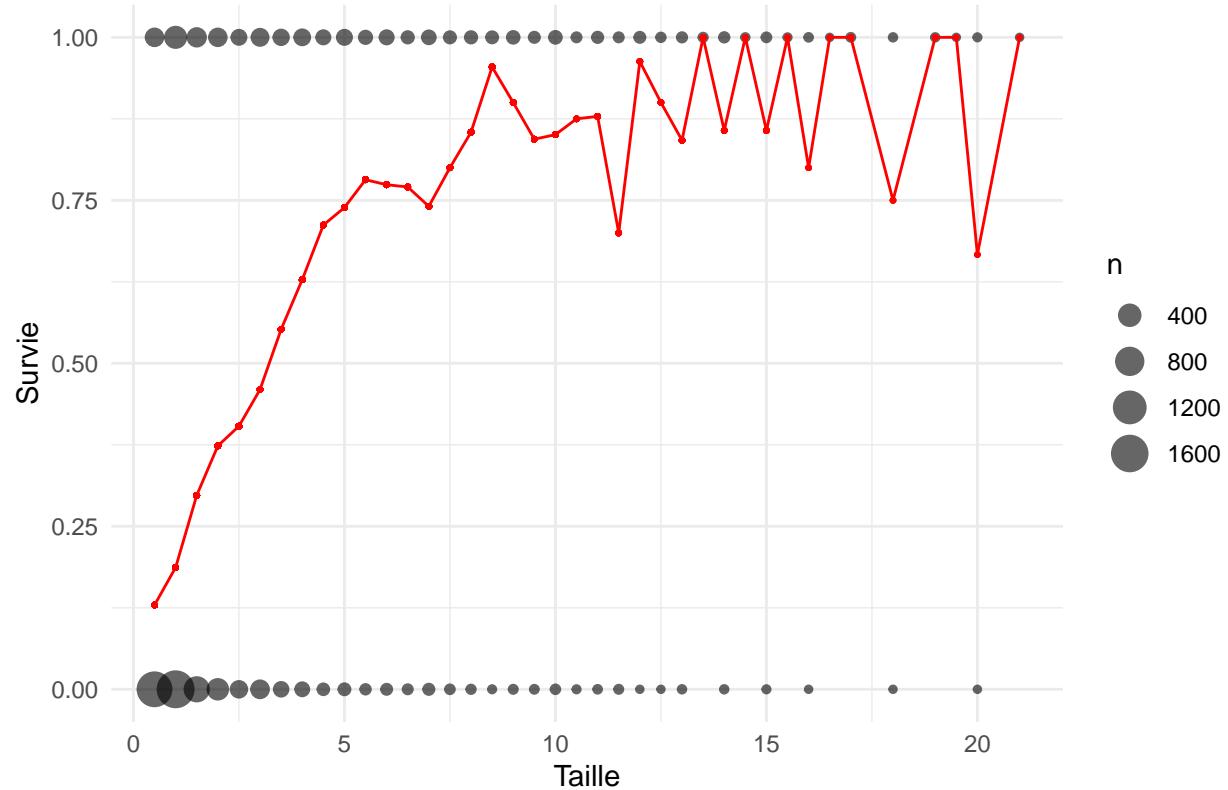
```

survdata <- centauree_data[centauree_data$Flowering0!=1,]
survdata <- survdata[!is.na(survdata$Size1Mars),]

survdata %>%
  group_by(Size0Mars) %>%
  mutate(survivalProba = sum(SurvieMars, na.rm = TRUE) / n()) %>%
  ggplot(aes(x = Size0Mars, y = SurvieMars)) +
  geom_count(alpha = 0.6) + # Points dimensionnés selon la fréquence
  geom_point(aes(y = survivalProba), color = "red", size = 0.5) +
  geom_line(aes(y = survivalProba), color = "red") +
  labs(title = "Relation entre la taille et la survie",
       x = "Taille",
       y = "Survie") +
  ylim(0, 1) +
  theme_minimal()

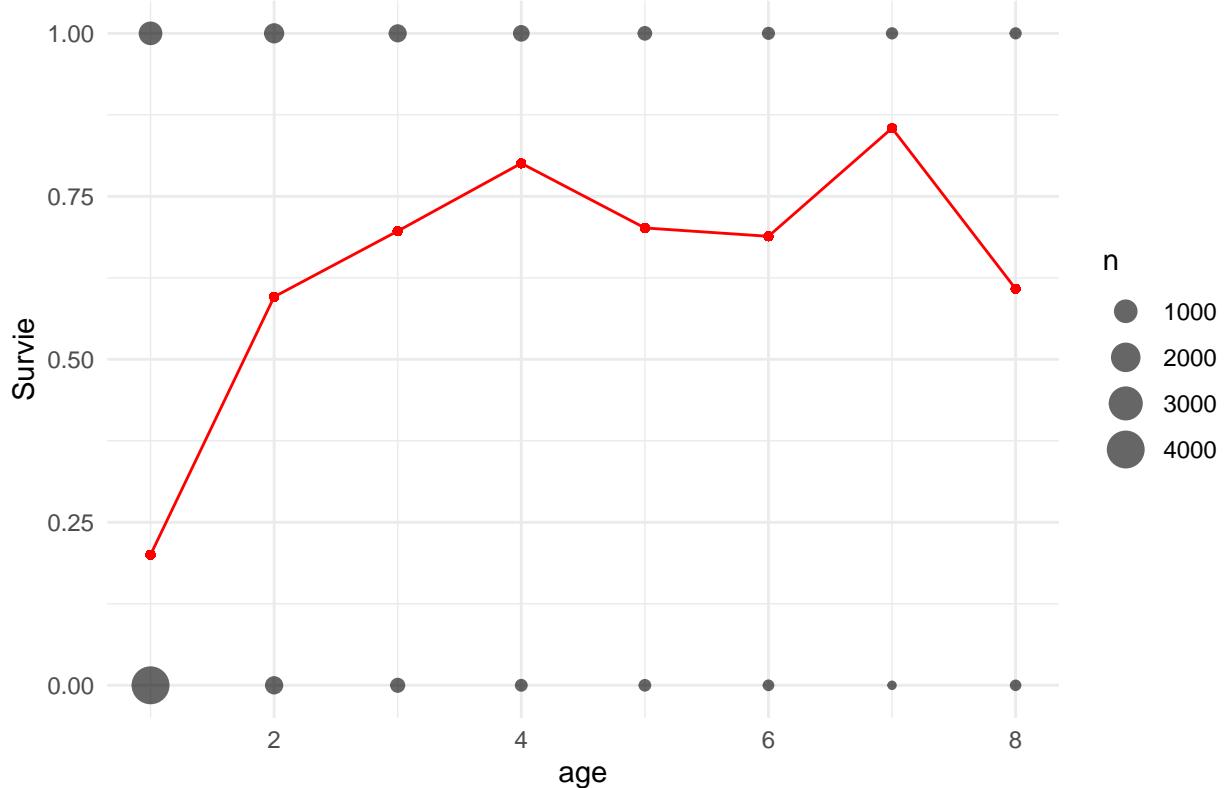
```

## Relation entre la taille et la survie



```
survdata %>%
  group_by(age0) %>%
  mutate(survivalProba = sum(SurvieMars, na.rm = TRUE) / n()) %>%
  ggplot(aes(x = age0, y = SurvieMars)) +
  geom_count(alpha = 0.6) + # Points dimensionnés selon la fréquence
  geom_point(aes(x = age0, y = survivalProba), color = "red", size = 1) +
  geom_line(aes(x = age0, y = survivalProba), color = "red") +
  labs(title = "Relation entre l'age et la survie",
       x = "age",
       y = "Survie") +
  ylim(0, 1) +
  theme_minimal()
```

## Relation entre l'age et la survie



```
Survglm1 <- fitme(SurviveMars ~ 1 + bs(Size0Mars, df=5, degree=3) + bs(age0, degree=3, knots = 6.5) + (Size0Mars + age0 | year) + (age0 | Pop),
                     family=binomial,
                     data=survdata,
                     method="PQL/L")

Survglm2 <- fitme(SurviveMars ~ 1 + bs(Size0Mars, df=5, degree=3) + bs(age0, degree=3, knots = 6.5) + (Size0Mars + age0 | year) + (age0 | Pop),
                     family=binomial,
                     data=survdata,
                     method="PQL/L")

Survglm3 <- fitme(SurviveMars ~ 1 + bs(Size0Mars, df=5, degree=3) + bs(age0, degree=3, knots = c(1.5, 6.5)) +
                     family=binomial,
                     data=survdata,
                     method="PQL/L")

## Warning in (function (formula, resid.formula = NULL, data, prior.weights, :
## 'c()' detected in formula: did you mean cbind() for binomial response or for
## poly()?

Survglm4 <- fitme(SurviveMars ~ 1 + bs(Size0Mars, df=5, degree=3) + bs(age0, degree=3, knots = c(1.5, 6.5)) +
                     family=binomial,
                     data=survdata,
                     method="PQL/L")
```

```

## Warning in (function (formula, resid.formula = NULL, data, prior.weights, :
## 'c(' detected in formula: did you mean cbind() for binomial response or for
## poly()?

Survglm5 <- fitme(SurviveMars ~ 1 + bs(Size0Mars, df=5, degree=3) + bs(age0, degree=3, knots = 6.5)
+ (Size0Mars|year) + (Size0Mars + age0|Pop) ,
family=binomial,
data=survdata,
method="PQL/L")

```

```
summary(Survglm1)
```

```

## formula: SurviveMars ~ 1 + bs(Size0Mars, df = 5, degree = 3) + bs(age0,
##      degree = 3, knots = 6.5) + (Size0Mars | year) + (age0 | Pop)
## Estimation of ranCoefs by ML (p_v approximation of logL).
## Estimation of fixed effects by h-likelihood approximation.
## family: binomial( link = logit )
## ----- Fixed effects (beta) -----
##                                     Estimate Cond. SE t-value
## (Intercept)                 -2.2406  0.2462 -9.1006
## bs(Size0Mars, df = 5, degree = 3)1  0.1103  0.1999  0.5517
## bs(Size0Mars, df = 5, degree = 3)2  0.8657  0.1272  6.8056
## bs(Size0Mars, df = 5, degree = 3)3  4.2121  0.4778  8.8158
## bs(Size0Mars, df = 5, degree = 3)4  3.8150  0.9696  3.9348
## bs(Size0Mars, df = 5, degree = 3)5  3.5933  1.2790  2.8095
## bs(age0, degree = 3, knots = 6.5)1  2.5146  0.2908  8.6480
## bs(age0, degree = 3, knots = 6.5)2 -0.8001  0.5510 -1.4520
## bs(age0, degree = 3, knots = 6.5)3  2.1587  0.6255  3.4513
## bs(age0, degree = 3, knots = 6.5)4  0.1423  0.4088  0.3480
## ----- Random effects -----
## Family: gaussian( link = identity )
##     --- Random-coefficients Cov matrices:
##   Group      Term    Var. Corr.
##   year (Intercept)  1.02
##   year  Size0Mars  0.01061 -0.672
##   Pop   (Intercept)  0.1431
##   Pop     age0  0.009096 -0.76
## # of obs: 7293; # of groups: year, 27; Pop, 6
## ----- Likelihood values -----
##          logLik
## h-likelihood: -3345.794
## logL      (p_v(h)): -3347.972

```

```
summary(Survglm2)
```

```

## formula: SurviveMars ~ 1 + bs(Size0Mars, df = 5, degree = 3) + bs(age0,
##      degree = 3, knots = 6.5) + (Size0Mars + age0 | year) + (age0 |
##      Pop)
## Estimation of ranCoefs by ML (p_v approximation of logL).
## Estimation of fixed effects by h-likelihood approximation.
## family: binomial( link = logit )
## ----- Fixed effects (beta) -----

```

```

##                               Estimate Cond. SE t-value
## (Intercept)                  -2.24874  0.2508 -8.9673
## bs(Size0Mars, df = 5, degree = 3)1  0.09729  0.2001  0.4862
## bs(Size0Mars, df = 5, degree = 3)2  0.87965  0.1261  6.9774
## bs(Size0Mars, df = 5, degree = 3)3  4.33060  0.4547  9.5240
## bs(Size0Mars, df = 5, degree = 3)4  3.76265  0.9328  4.0336
## bs(Size0Mars, df = 5, degree = 3)5  3.44384  1.2172  2.8294
## bs(age0, degree = 3, knots = 6.5)1  2.48418  0.3082  8.0608
## bs(age0, degree = 3, knots = 6.5)2 -0.84350  0.5967 -1.4135
## bs(age0, degree = 3, knots = 6.5)3  2.18330  0.7078  3.0848
## bs(age0, degree = 3, knots = 6.5)4  0.06919  0.5440  0.1272
## ----- Random effects -----
## Family: gaussian( link = identity )
##     --- Random-coefficients Cov matrices:
##   Group      Term      Var.  Corr. Corr..1
##   year (Intercept) 1.304
##   year   Size0Mars 0.0005913      -1
##   year      age0  0.04756 -0.663  0.663
##   Pop (Intercept) 0.1543
##   Pop      age0  0.01036 -0.7826
## # of obs: 7293; # of groups: year, 27; Pop, 6
## ----- Likelihood values -----
##           logLik
## h-likelihood: -3359.315
## logL      (p_v(h)): -3341.086

```

```
summary(Survglm3)
```

```

## formula: SurvieMars ~ 1 + bs(Size0Mars, df = 5, degree = 3) + bs(age0,
##   degree = 3, knots = c(1.5, 6.5)) + (Size0Mars | year) + (age0 |
##   Pop)
## Estimation of ranCoefs by ML (p_v approximation of logL).
## Estimation of fixed effects by h-likelihood approximation.
## family: binomial( link = logit )
## ----- Fixed effects (beta) -----
##                               Estimate Cond. SE t-value
## (Intercept)                  -2.23985  0.2463 -9.09280
## bs(Size0Mars, df = 5, degree = 3)1  0.10932  0.1999  0.54682
## bs(Size0Mars, df = 5, degree = 3)2  0.86720  0.1272  6.81565
## bs(Size0Mars, df = 5, degree = 3)3  4.22462  0.4787  8.82432
## bs(Size0Mars, df = 5, degree = 3)4  3.80246  0.9711  3.91554
## bs(Size0Mars, df = 5, degree = 3)5  3.59587  1.2796  2.81021
## bs(age0, degree = 3, knots = c(1.5, 6.5))1  0.01095  0.3491  0.03138
## bs(age0, degree = 3, knots = c(1.5, 6.5))2  2.61115  0.5908  4.41987
## bs(age0, degree = 3, knots = c(1.5, 6.5))3 -0.95676  0.7692 -1.24392
## bs(age0, degree = 3, knots = c(1.5, 6.5))4  2.37011  0.7184  3.29903
## bs(age0, degree = 3, knots = c(1.5, 6.5))5  0.13872  0.4095  0.33875
## ----- Random effects -----
## Family: gaussian( link = identity )
##     --- Random-coefficients Cov matrices:
##   Group      Term      Var.  Corr.
##   year (Intercept) 1.021
##   year   Size0Mars 0.01077 -0.6711
##   Pop (Intercept) 0.1436

```

```

##      Pop          age0  0.009156 -0.7602
## # of obs: 7293; # of groups: year, 27; Pop, 6
## ----- Likelihood values -----
##           logLik
##      h-likelihood: -3345.466
## logL      (p_v(h)): -3347.771

```

```
summary(Survglm4)
```

```

## formula: SurvieMars ~ 1 + bs(Size0Mars, df = 5, degree = 3) + bs(age0,
##      degree = 3, knots = c(1.5, 6.5)) + (Size0Mars + age0 | year) +
##      (age0 | Pop)
## Estimation of ranCoefs by ML (p_v approximation of logL).
## Estimation of fixed effects by h-likelihood approximation.
## family: binomial( link = logit )
## ----- Fixed effects (beta) -----
##                                         Estimate Cond. SE t-value
## (Intercept)                         -2.2482  0.2508 -8.9626
## bs(Size0Mars, df = 5, degree = 3)1    0.0967  0.2001  0.4833
## bs(Size0Mars, df = 5, degree = 3)2    0.8804  0.1261  6.9830
## bs(Size0Mars, df = 5, degree = 3)3    4.3379  0.4554  9.5250
## bs(Size0Mars, df = 5, degree = 3)4    3.7544  0.9336  4.0213
## bs(Size0Mars, df = 5, degree = 3)5    3.4392  1.2172  2.8256
## bs(age0, degree = 3, knots = c(1.5, 6.5))1  0.1132  0.3500  0.3234
## bs(age0, degree = 3, knots = c(1.5, 6.5))2  2.4172  0.5974  4.0460
## bs(age0, degree = 3, knots = c(1.5, 6.5))3  -0.8176  0.8049 -1.0158
## bs(age0, degree = 3, knots = c(1.5, 6.5))4  2.2926  0.7865  2.9149
## bs(age0, degree = 3, knots = c(1.5, 6.5))5  0.0687  0.5444  0.1262
## ----- Random effects -----
## Family: gaussian( link = identity )
##     --- Random-coefficients Cov matrices:
##   Group      Term      Var.  Corr. Corr..1
##   year (Intercept)  1.303
##   year  Size0Mars  0.0006054      -1
##   year      age0   0.04765 -0.6614  0.6614
##   Pop (Intercept)  0.1546
##   Pop      age0   0.01038 -0.7827
## # of obs: 7293; # of groups: year, 27; Pop, 6
## ----- Likelihood values -----
##           logLik
##      h-likelihood: -3359.220
## logL      (p_v(h)): -3341.027

```

```
summary(Survglm5)
```

```

## formula: SurvieMars ~ 1 + bs(Size0Mars, df = 5, degree = 3) + bs(age0,
##      degree = 3, knots = 6.5) + (Size0Mars | year) + (Size0Mars +
##      age0 | Pop)
## Estimation of ranCoefs by ML (p_v approximation of logL).
## Estimation of fixed effects by h-likelihood approximation.
## family: binomial( link = logit )
## ----- Fixed effects (beta) -----
##                                         Estimate Cond. SE t-value

```

```

## (Intercept) -2.2250 0.2397 -9.2840
## bs(Size0Mars, df = 5, degree = 3)1 0.1091 0.1999 0.5459
## bs(Size0Mars, df = 5, degree = 3)2 0.8570 0.1278 6.7070
## bs(Size0Mars, df = 5, degree = 3)3 4.1960 0.4967 8.4476
## bs(Size0Mars, df = 5, degree = 3)4 3.9689 1.0148 3.9109
## bs(Size0Mars, df = 5, degree = 3)5 3.6233 1.3663 2.6519
## bs(age0, degree = 3, knots = 6.5)1 2.5437 0.2918 8.7182
## bs(age0, degree = 3, knots = 6.5)2 -0.8567 0.5554 -1.5426
## bs(age0, degree = 3, knots = 6.5)3 2.2147 0.6319 3.5047
## bs(age0, degree = 3, knots = 6.5)4 0.1441 0.4172 0.3455
## ----- Random effects -----
## Family: gaussian( link = identity )
## --- Random-coefficients Cov matrices:
## Group Term Var. Corr. Corr..1
## year (Intercept) 1.011
## year Size0Mars 0.01131 -0.6471
## Pop (Intercept) 0.1314
## Pop Size0Mars 0.001472 0.353
## Pop age0 0.009911 -0.9029 -0.3149
## # of obs: 7293; # of groups: year, 27; Pop, 6
## ----- Likelihood values -----
## logLik
## h-likelihood: -3348.440
## logL (p_v(h)): -3346.683

```

```

Survpredict1 <- predict(Survglm1, newdata = fake_data)[,1]
Survpredict2 <- predict(Survglm2, newdata = fake_data)[,1]
Survpredict3 <- predict(Survglm3, newdata = fake_data)[,1]
Survpredict4 <- predict(Survglm4, newdata = fake_data)[,1]
Survpredict5 <- predict(Survglm5, newdata = fake_data)[,1]

```

```

plot_survie <- function(data = fake_data, prediction, var, c1, valc1 = 1, c2, valc2 = "Au", fact) {
  data %>%
    mutate(surv_predi = prediction) %>%
    filter(!!sym(c1) == valc1, !!sym(c2) == valc2) %>%
    ggplot(aes(x = .data[[var]], y = surv_predi)) +
    geom_line(aes(color = as.factor(.data[[fact]]))) +
    theme_minimal() +
    ylim(0, 1)
}

```

## Survie en fonction de la taille

En fixant la population : voir l'effet année

```

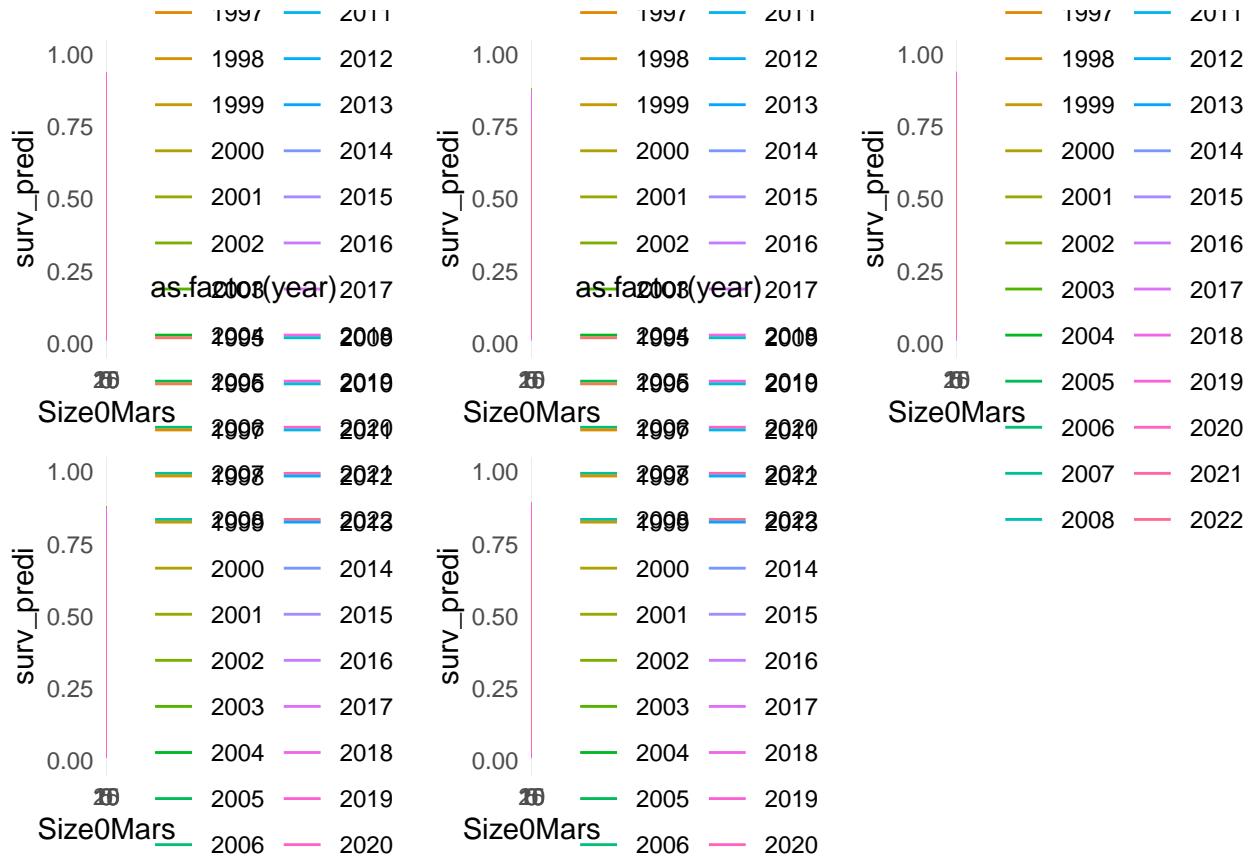
var <- "Size0Mars"
c1 <- "age0"
valc1 <- 1
c2 <- "Pop"
valc2 <- "Au"
fact <- "year"

```

```

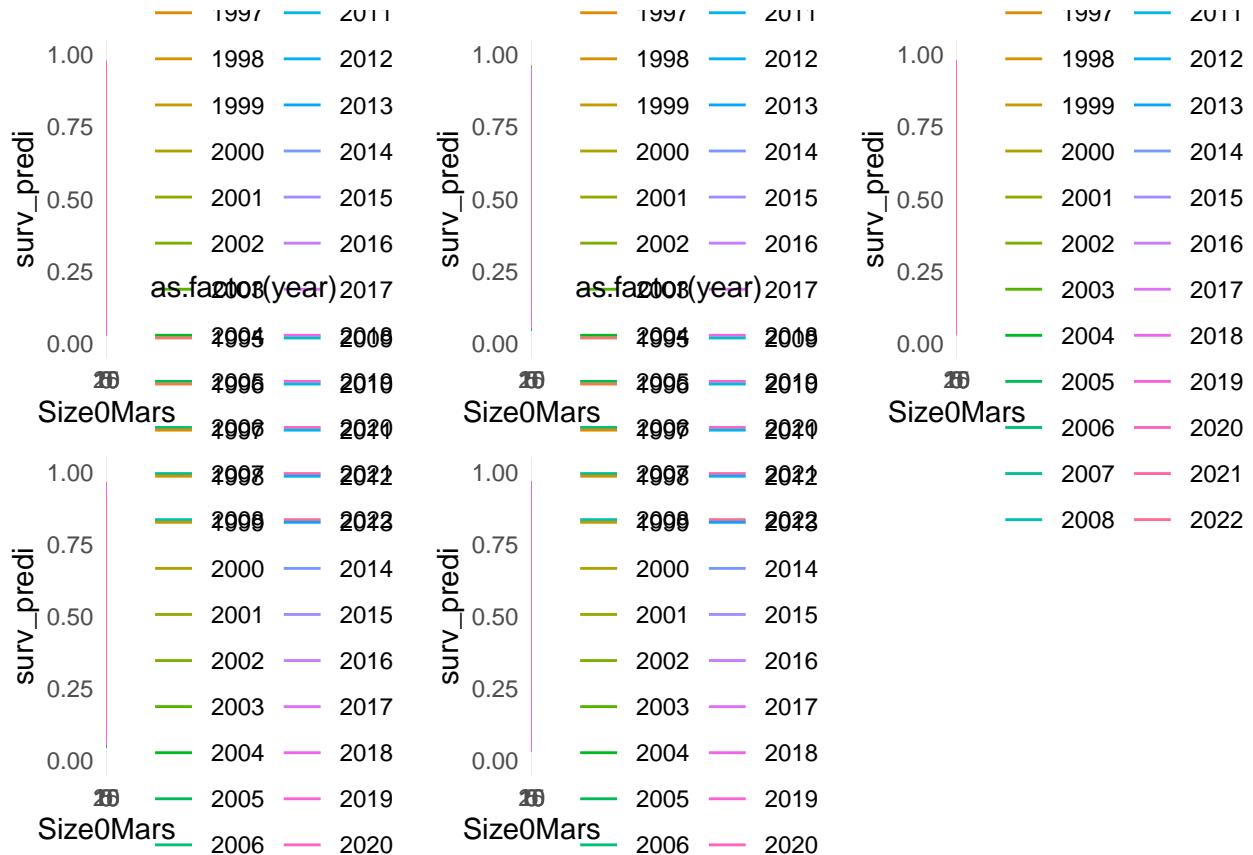
wrap_plots(
plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

```



```

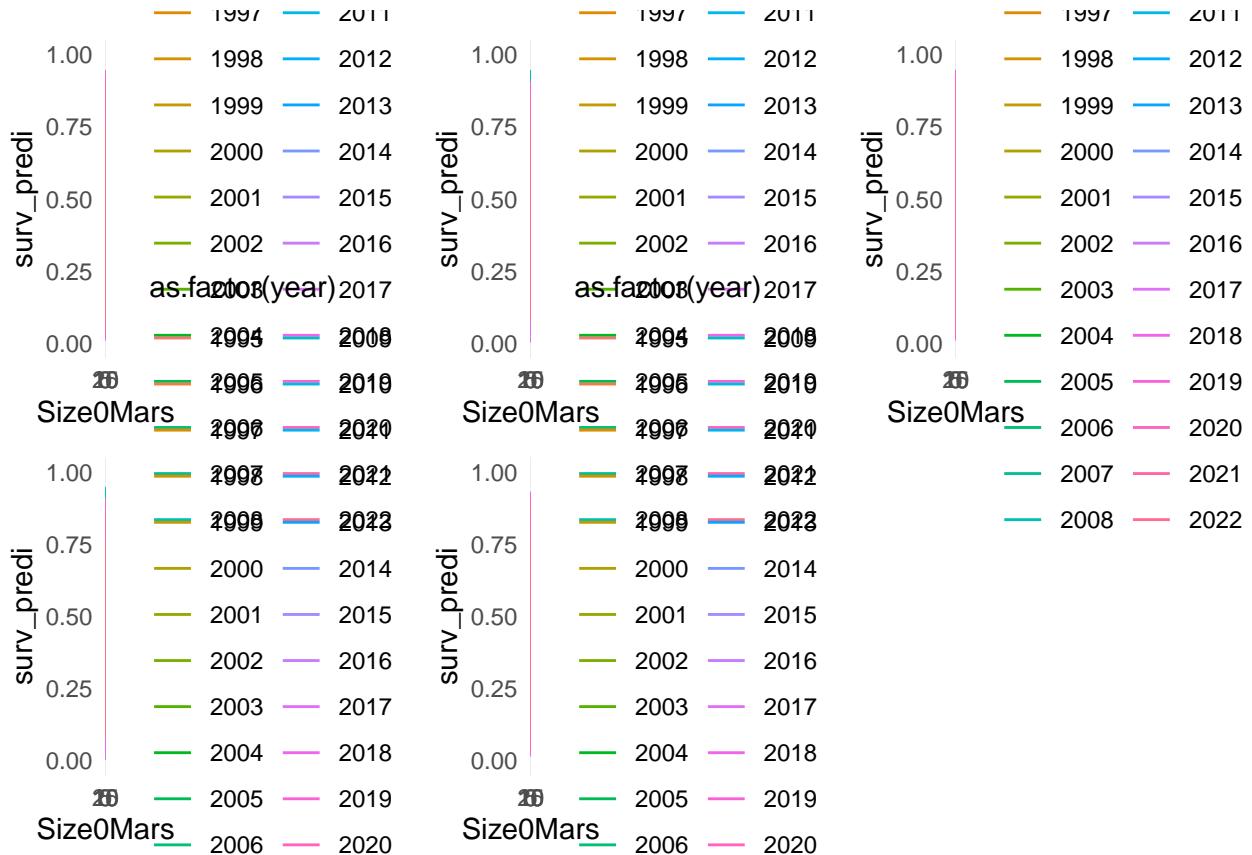
valc1 <- 3
wrap_plots(
plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)
```



```

valc1 <- 8
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

```

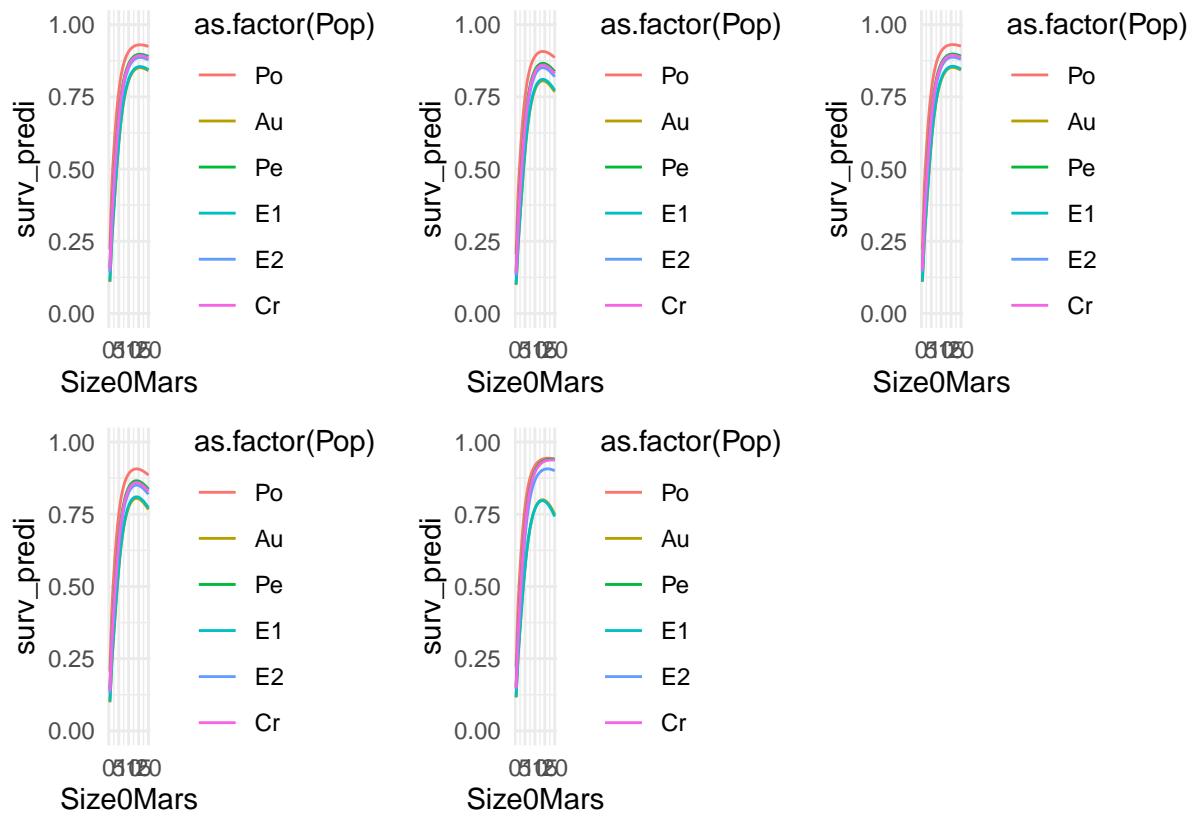


En fixant l'année : voir l'effet population

```

var <- "Size0Mars"
c1 <- "age0"
valc1 <- 1
c2 <- "year"
valc2 <- 2000
fact <- "Pop"

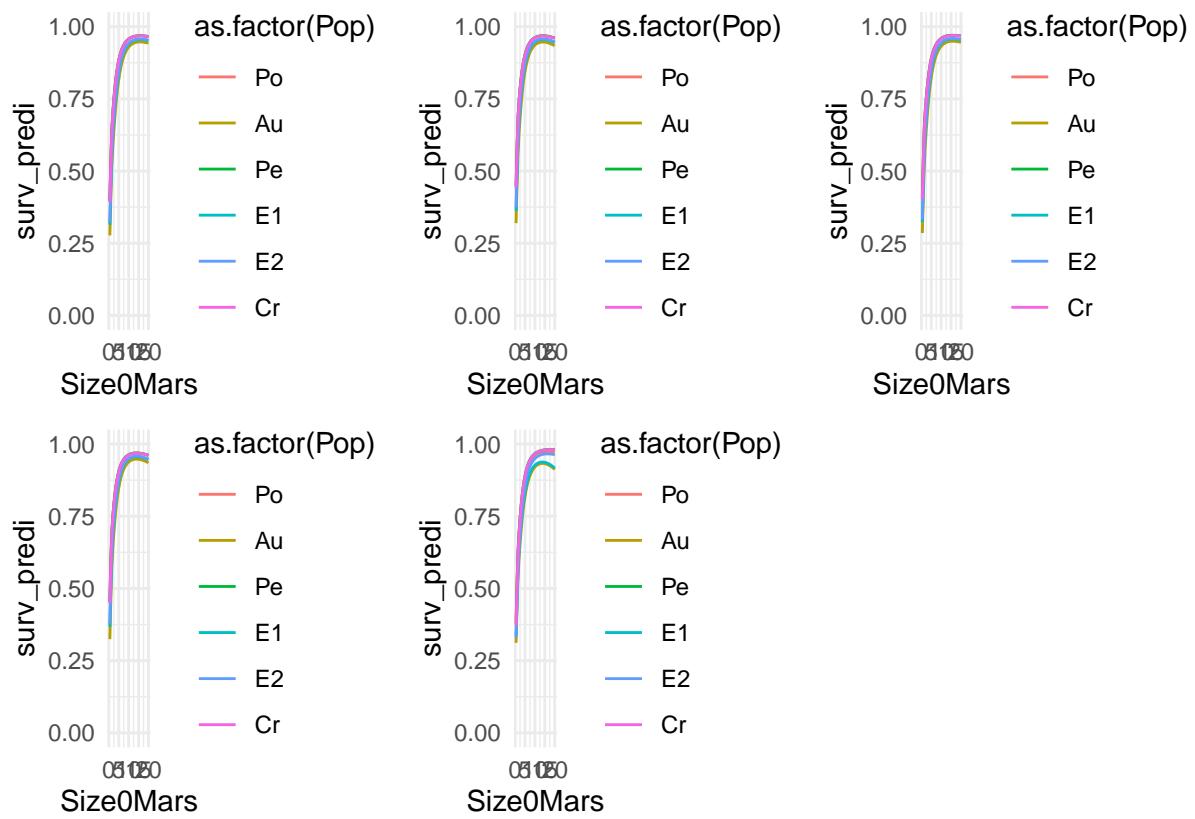
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)
  
```



```

valc1 <- 3
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

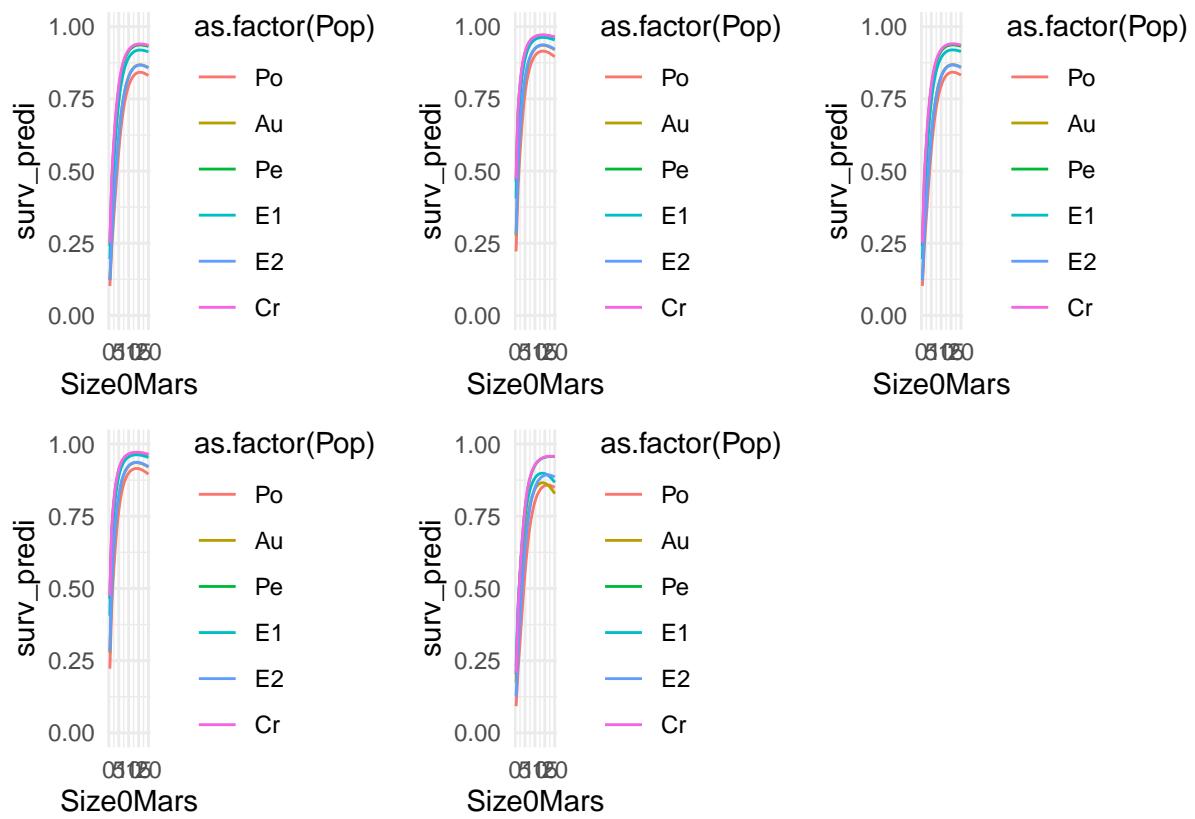
```



```

valc1 <- 8
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

```



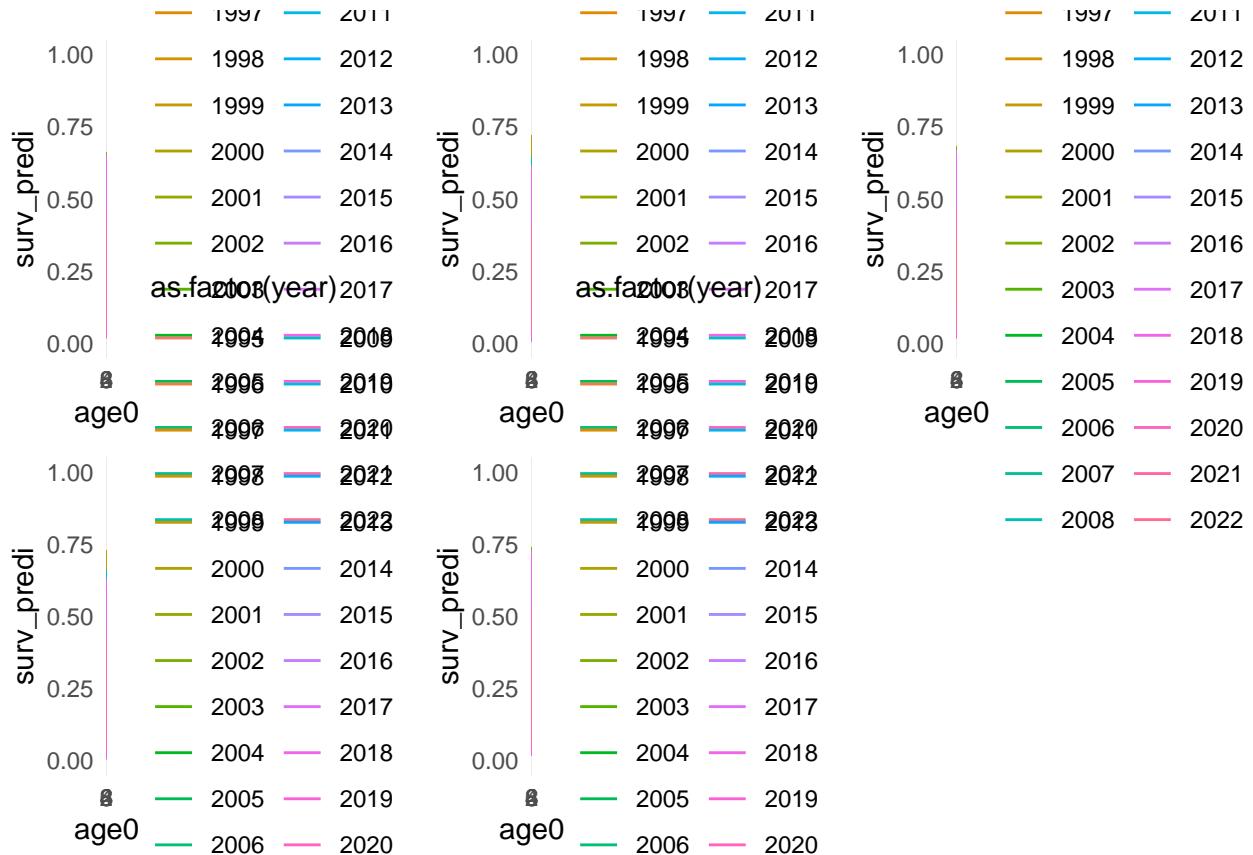
## Survie en fonction de l'âge

En fixant la population : voir l'effet année

```

var <- "age0"
c1 <- "Size0Mars"
valc1 <- 1
c2 <- "Pop"
valc2 <- "Au"
fact <- "year"

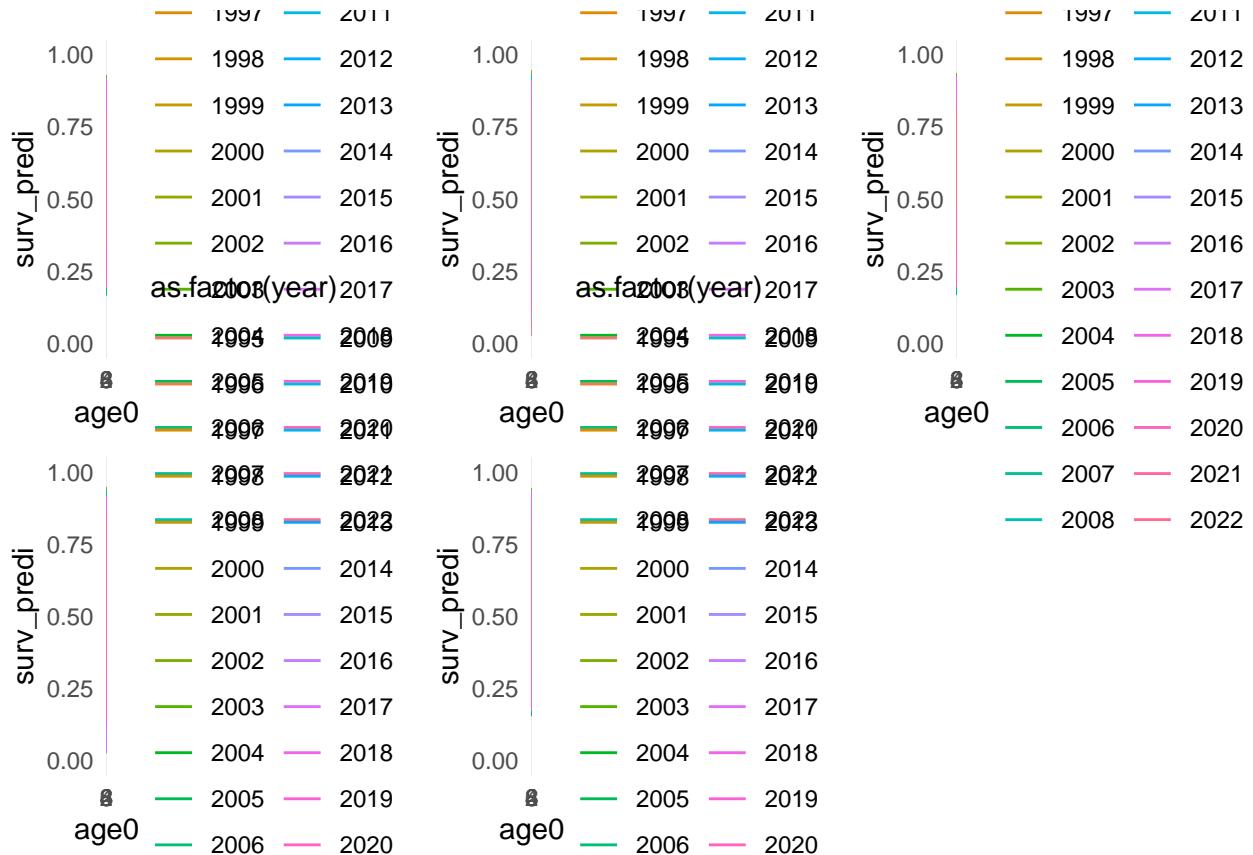
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)
  
```



```

valc1 <- 5
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

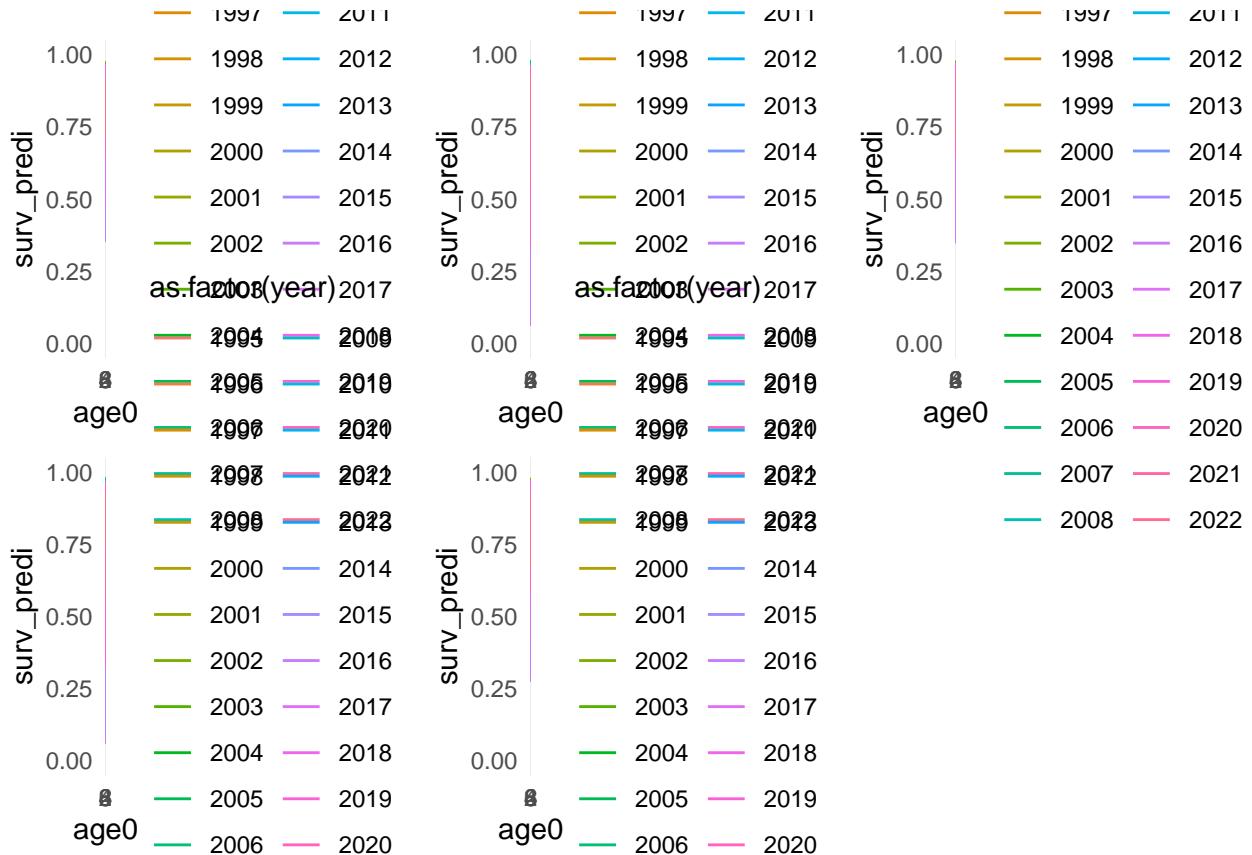
```



```

valc1 <- 10
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

```

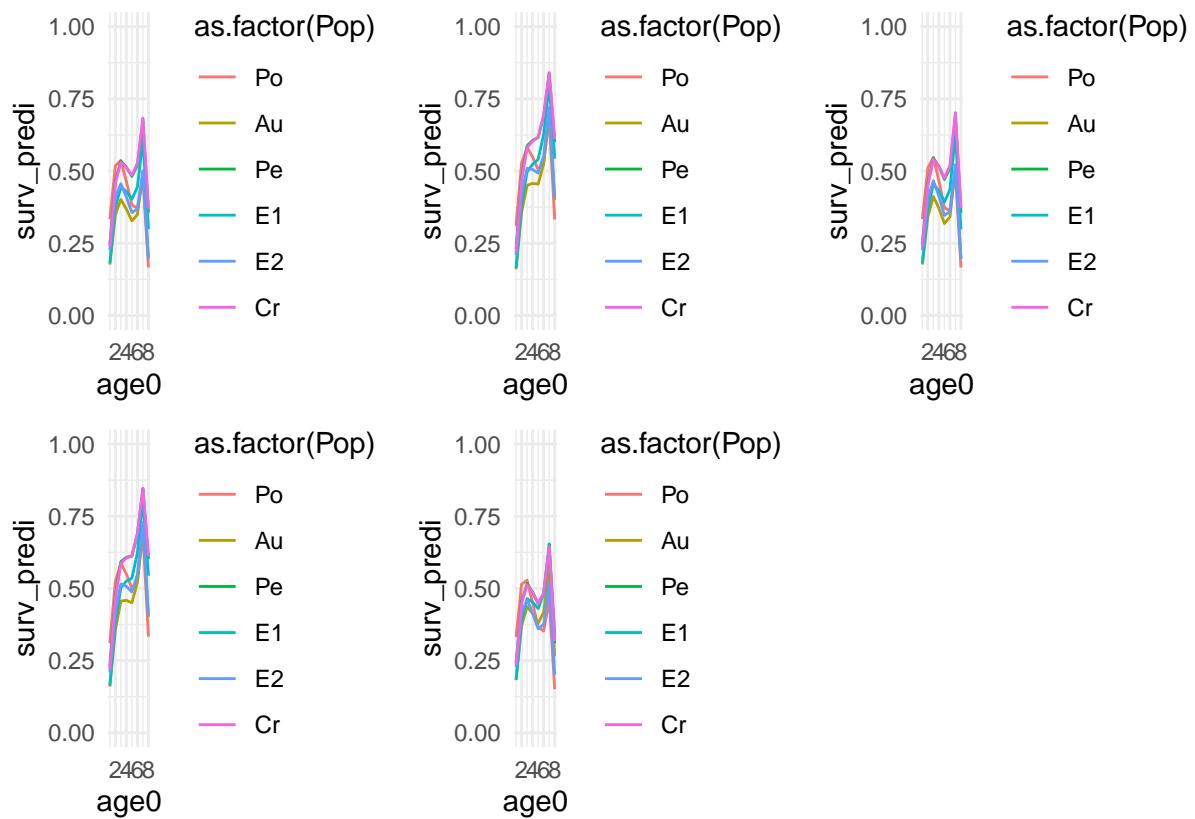


En fixant l'année : voir l'effet population

```

var <- "age0"
c1 <- "SizeOMars"
valc1 <- 1
c2 <- "year"
valc2 <- 2000
fact <- "Pop"

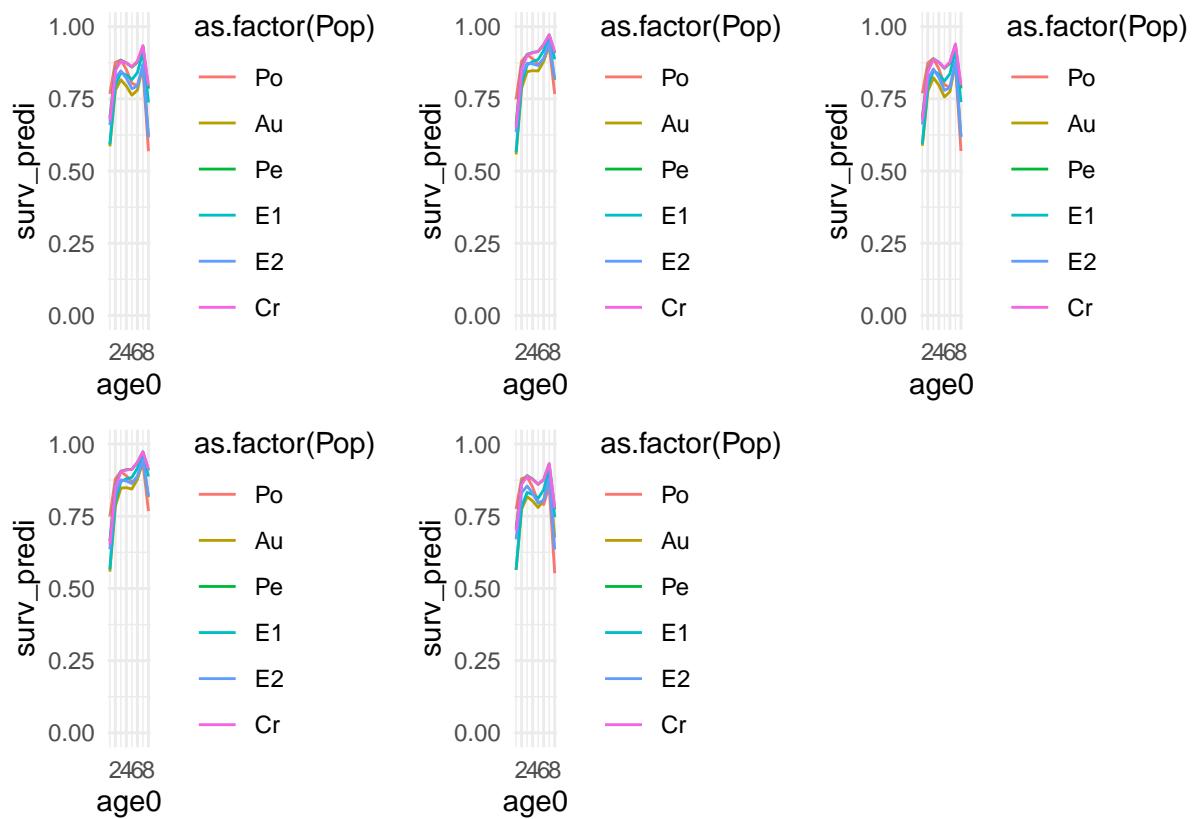
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)
  
```



```

valc1 <- 5
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

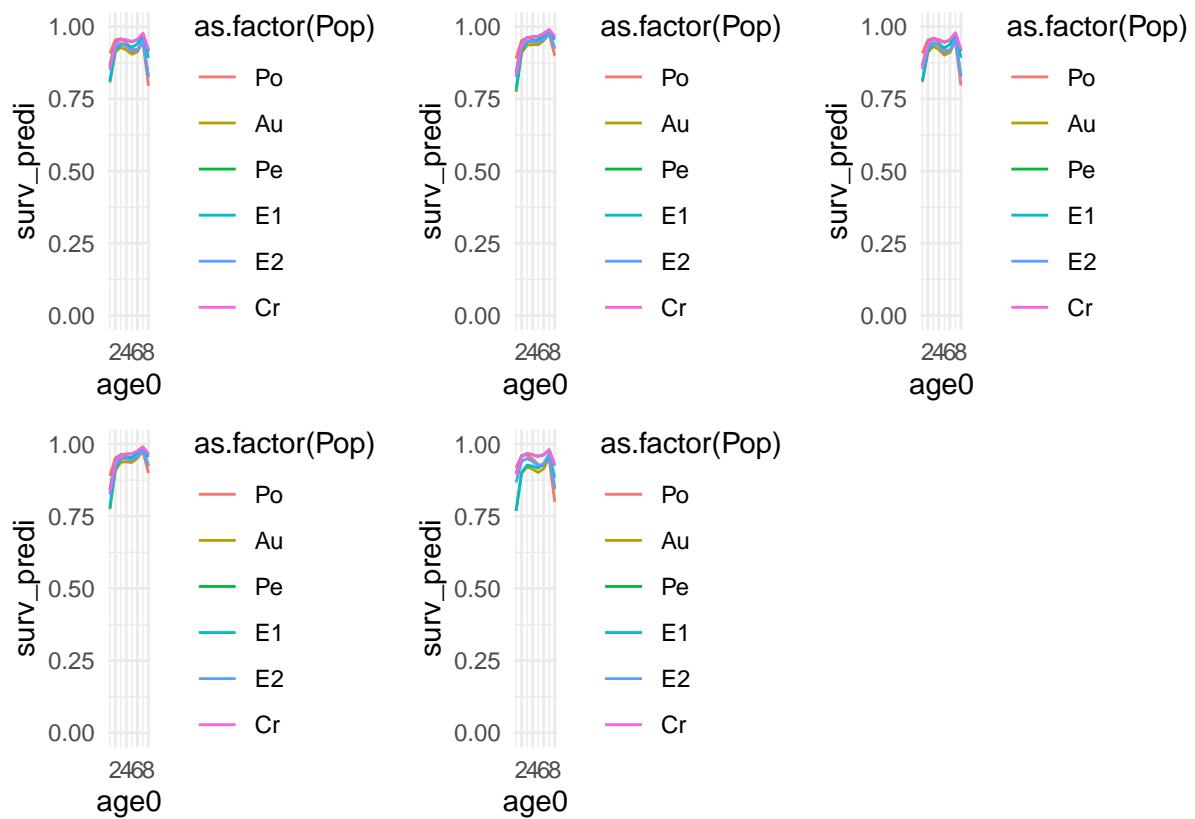
```



```

valc1 <- 10
wrap_plots(
  plot_survie(prediction = Survpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_survie(prediction = Survpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

```



## Nombre de capitules

```

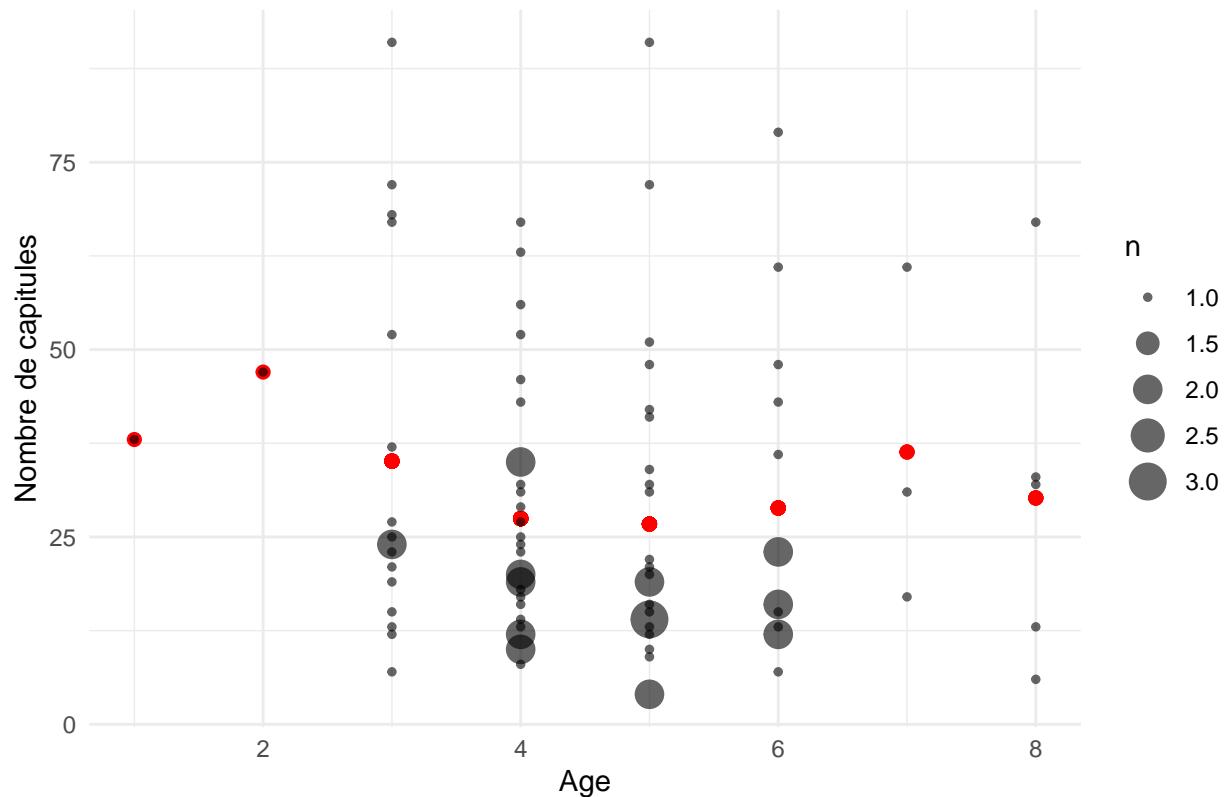
cptldata <- centauree_data[!is.na(centauree_data$Cpt10),]
cptldata <- cptldata[!cptldata$Flowering0==0,]

# Nombre de capitules moyen / age
cptldata %>%
  group_by(age0) %>%
  mutate(meancptl=mean(Cpt10)) %>%

ggplot(aes(x = age0, y = meancptl)) +
  geom_point(color = "red", size = 2) +
  geom_count(aes(y=Cpt10), alpha=0.6) +
  labs(title = "Relation entre l'age et le nombre de capitules",
       x = "Age",
       y = "Nombre de capitules") +
  theme_minimal()

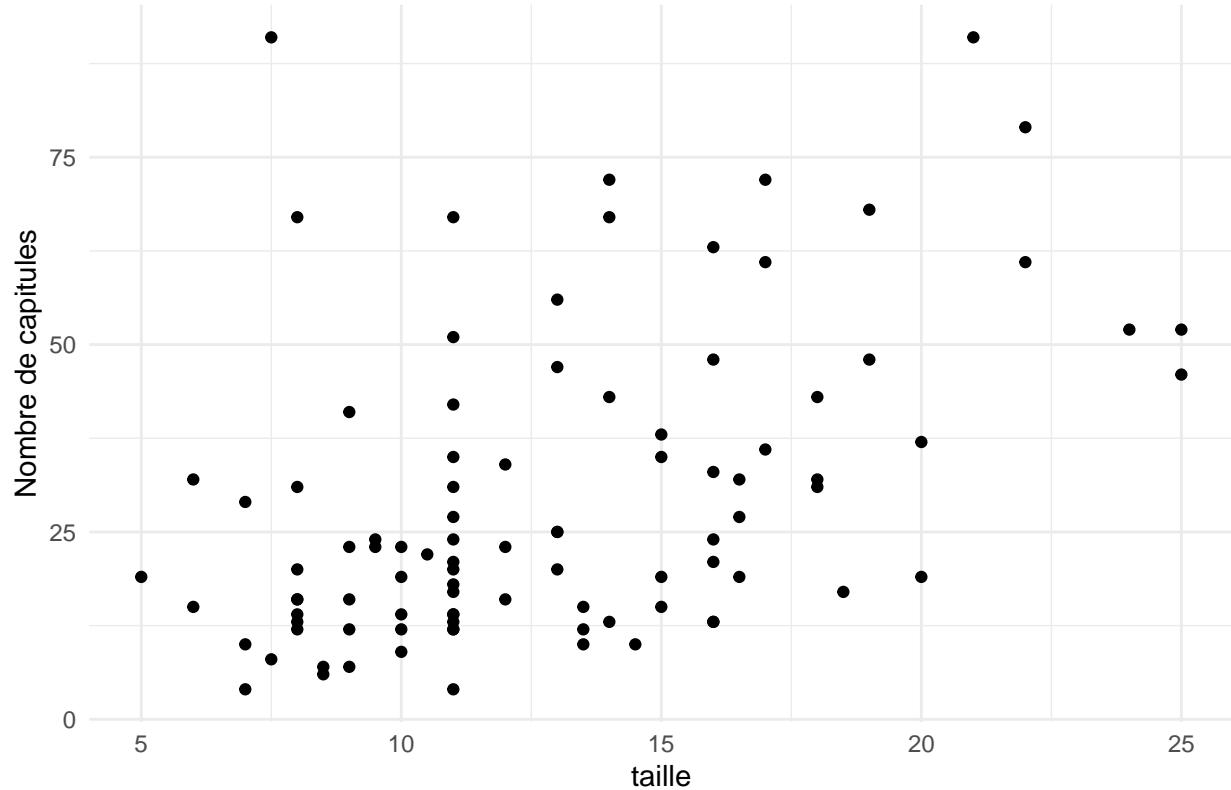
```

## Relation entre l'age et le nombre de chapitres



```
# Nombre de chapitre / année
cptldata %>%
  ggplot(aes(x=Size0Mars,y=Cptl0)) +
  geom_point() +
  labs(title = "Relation entre la taille et le nombre de chapitres",
       x = "taille",
       y = "Nombre de chapitres") +
  theme_minimal()
```

## Relation entre la taille et le nombre de chapitres



```
Cptlglm1 <- fitme(Cptl0 ~ 1 + Size0Mars,
                     data=cptldata)

Cptlglm2 <- fitme(Cptl0 ~ 1 + poly(Size0Mars,2),
                     data=cptldata)

Cptlglm3 <- fitme(Cptl0 ~ 1 + poly(Size0Mars,3),
                     data=cptldata)

Cptlglm4 <- fitme(Cptl0 ~ 1 + Size0Mars +(1|year),
                     data=cptldata)

Cptlglm5 <- fitme(Cptl0 ~ 1 + Size0Mars + age0,
                     data=cptldata)
```

```
summary(Cptlglm1)

## formula: Cptl0 ~ 1 + Size0Mars
## ML: Estimation of phi by ML.
##      Estimation of fixed effects by ML.
## family: gaussian( link = identity )
## -----
##             Estimate Cond. SE t-value
## (Intercept) 2.951   5.6590  0.5214
## Size0Mars    2.074   0.4166  4.9795
```

```

## ----- Residual variance -----
## Coefficients for log(phi) ~ 1 :
##           Estimate Cond. SE
## (Intercept) 5.786  0.1451
## Estimate of phi=residual var: 325.6
## ----- Likelihood values -----
##           logLik
## logL       : -409.6181

```

```
summary(Cptlglm2)
```

```

## formula: Cptl0 ~ 1 + poly(Size0Mars, 2)
## ML: Estimation of phi by ML.
##   Estimation of fixed effects by ML.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept) 29.58    1.836 16.108
## poly(Size0Mars, 2)1 89.85   17.898  5.020
## poly(Size0Mars, 2)2 22.36   17.898  1.249
## ----- Residual variance -----
## Coefficients for log(phi) ~ 1 :
##           Estimate Cond. SE
## (Intercept) 5.769  0.1451
## Estimate of phi=residual var: 320.3
## ----- Likelihood values -----
##           logLik
## logL       : -408.8444

```

```
summary(Cptlglm3)
```

```

## formula: Cptl0 ~ 1 + poly(Size0Mars, 3)
## ML: Estimation of phi by ML.
##   Estimation of fixed effects by ML.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept) 29.58    1.825 16.210
## poly(Size0Mars, 3)1 89.85   17.785  5.052
## poly(Size0Mars, 3)2 22.36   17.785  1.257
## poly(Size0Mars, 3)3 -19.56   17.785 -1.100
## ----- Residual variance -----
## Coefficients for log(phi) ~ 1 :
##           Estimate Cond. SE
## (Intercept) 5.757  0.1451
## Estimate of phi=residual var: 316.3
## ----- Likelihood values -----
##           logLik
## logL       : -408.2434

```

```
summary(Cptlglm4)
```

```

## formula: Cptl0 ~ 1 + Size0Mars + (1 | year)
## Estimation of fixed effects by ML.
## Estimation of lambda and phi by 'outer' ML, maximizing logL.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept)    4.198   5.9466  0.7059
## Size0Mars      2.010   0.4253  4.7263
## ----- Random effects -----
## Family: gaussian( link = identity )
##     --- Variance parameters ('lambda'):
## lambda = var(u) for u ~ Gaussian;
## year : 19.26
## # of obs: 95; # of groups: year, 16
## ----- Residual variance -----
## phi estimate was 308.566
## ----- Likelihood values -----
##           logLik
## logL      (p_v(h)): -409.4104

```

```
summary(Cptlglm5)
```

```

## formula: Cptl0 ~ 1 + Size0Mars + age0
## ML: Estimation of phi by ML.
##     Estimation of fixed effects by ML.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept)  0.9787   9.3256  0.1049
## Size0Mars    2.0965   0.4246  4.9372
## age0        0.3645   1.3705  0.2660
## ----- Residual variance -----
## Coefficients for log(phi) ~ 1 :
##           Estimate Cond. SE
## (Intercept)  5.785   0.1451
## Estimate of phi=residual var: 325.4
## ----- Likelihood values -----
##           logLik
## logL      : -409.5828

```

```

Cptlpredict1 <- predict(Cptlglm1, newdata = fake_data)[,1]
Cptlpredict2 <- predict(Cptlglm2, newdata = fake_data)[,1]
Cptlpredict3 <- predict(Cptlglm3, newdata = fake_data)[,1]
Cptlpredict4 <- predict(Cptlglm4, newdata = fake_data)[,1]
Cptlpredict5 <- predict(Cptlglm5, newdata = fake_data)[,1]

```

```

plot_capitule <- function(data = fake_data, prediction, var, c1, valc1 = 1, c2, valc2 = "Au", fact) {
  data %>%
    mutate(cptl_predi = prediction) %>%
    filter(!sym(c1) == valc1, !sym(c2) == valc2) %>%
    ggplot(aes(x = .data[[var]], y = cptl_predi)) +
    geom_line(aes(color = as.factor(.data[[fact]]))) +
    theme_minimal() +

```

```

    ylim(0,50)
}

```

## Nombre de chapitres en fonction de la taille

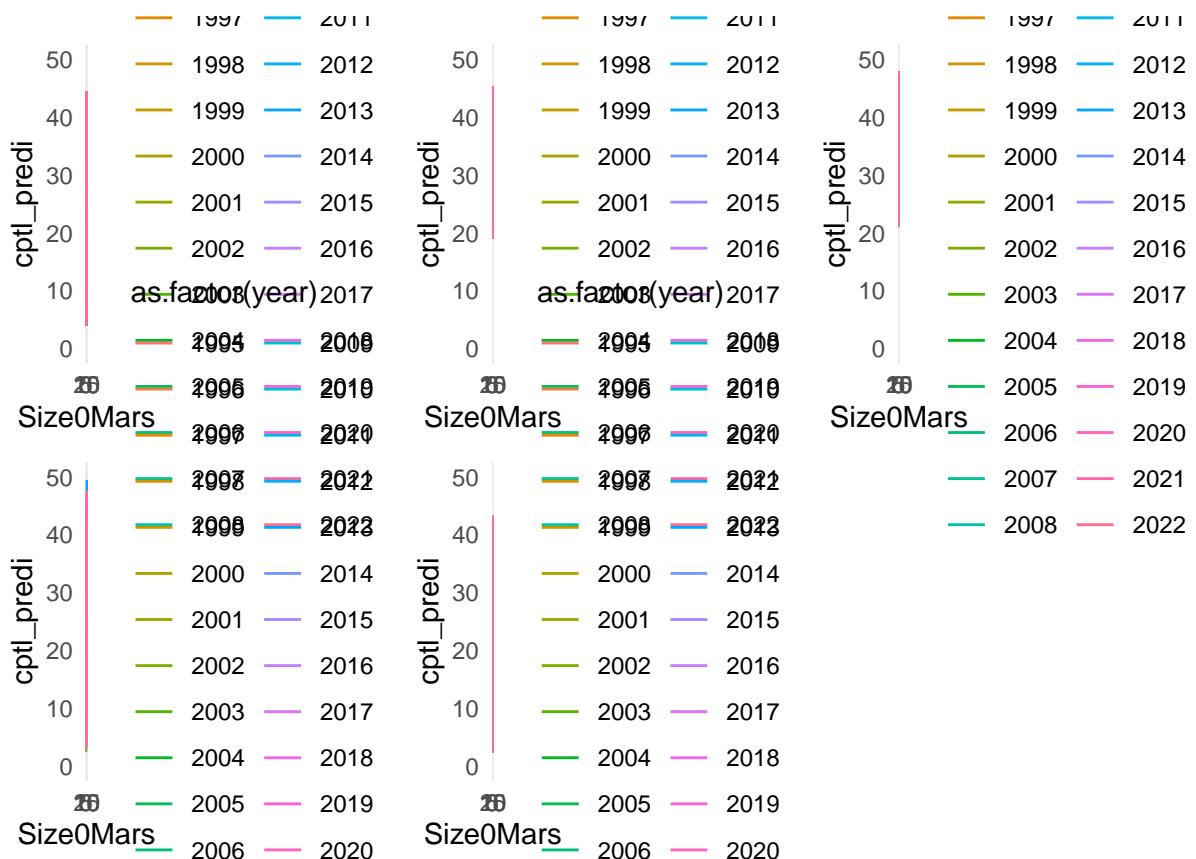
En fixant la population : voir l'effet année

```

var <- "Size0Mars"
c1 <- "age0"
valc1 <- 1
c2 <- "Pop"
valc2 <- "Au"
fact <- "year"

wrap_plots(
  plot_capitule(prediction = Cptlpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

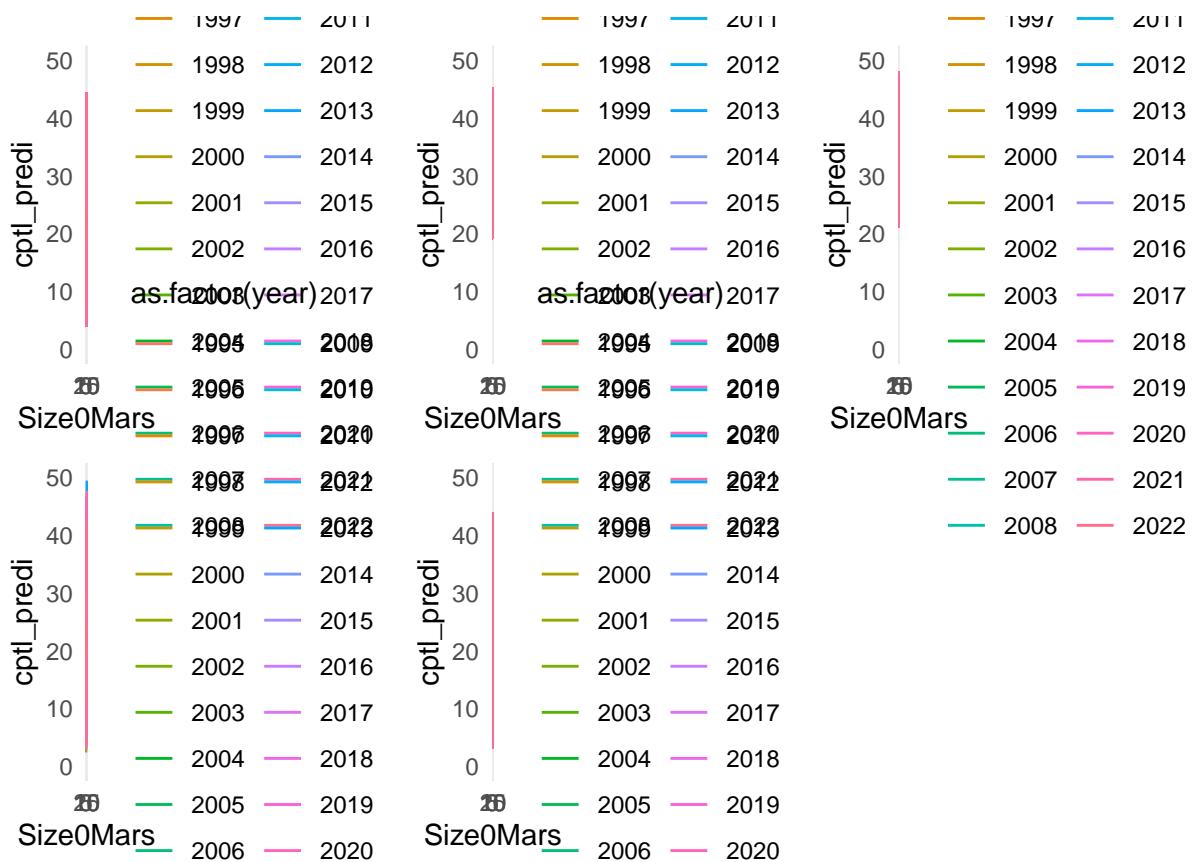
```



```

valc1 <- 3
wrap_plots(
  plot_capitule(prediction = Cptlpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

```



## Nombre de chapitres en fonction de l'âge

En fixant la population : voir l'effet année

```

var <- "age0"
c1 <- "Size0Mars"
valc1 <- 1
c2 <- "Pop"
valc2 <- "Au"
fact <- "year"

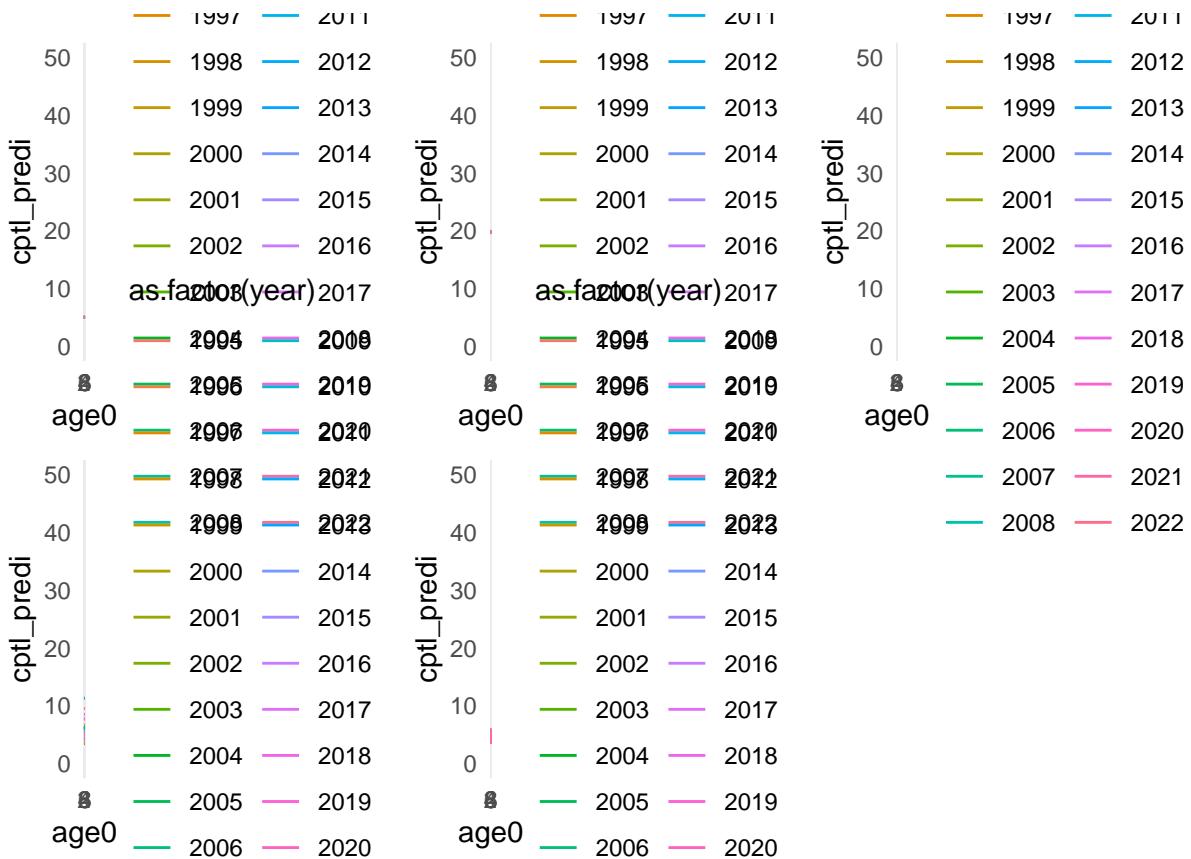
wrap_plots(
  plot_capitule(prediction = Cptlpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),

```

```

plot_capitule(prediction = Cptlpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_capitule(prediction = Cptlpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_capitule(prediction = Cptlpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

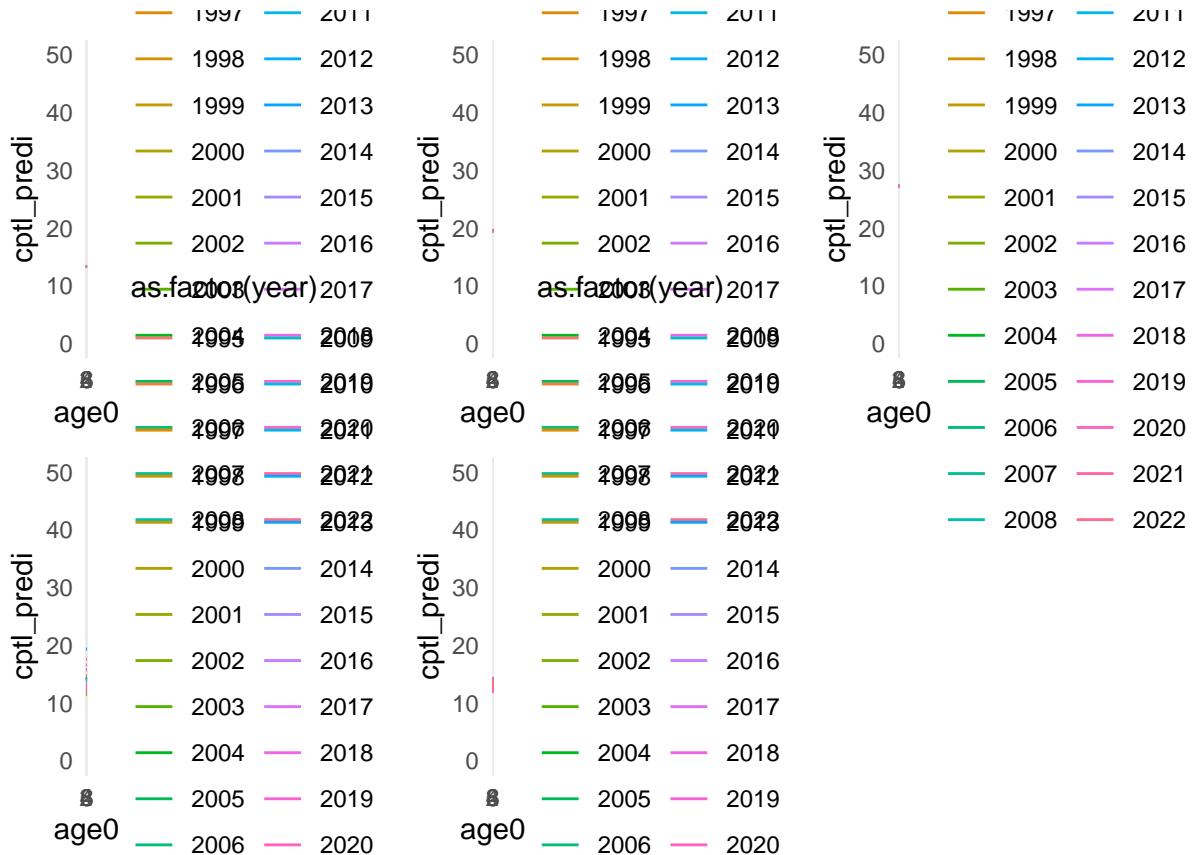
```



```

valc1 <- 5
wrap_plots(
plot_capitule(prediction = Cptlpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_capitule(prediction = Cptlpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_capitule(prediction = Cptlpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_capitule(prediction = Cptlpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
plot_capitule(prediction = Cptlpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

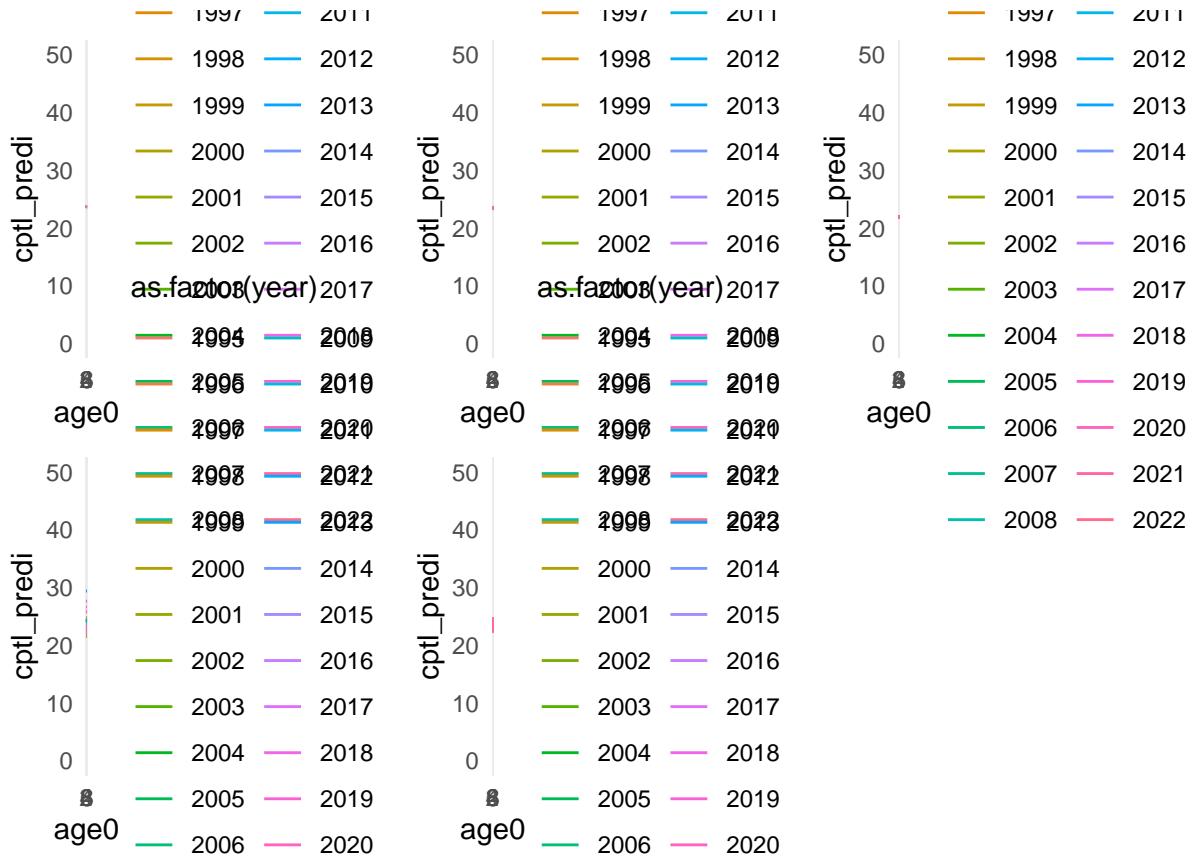
```



```

valc1 <- 10
wrap_plots(
  plot_capitule(prediction = Cptlpredict1, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict2, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict3, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict4, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact),
  plot_capitule(prediction = Cptlpredict5, var=var, c1=c1, valc1=valc1, c2=c2, valc2=valc2, fact=fact)
)

```



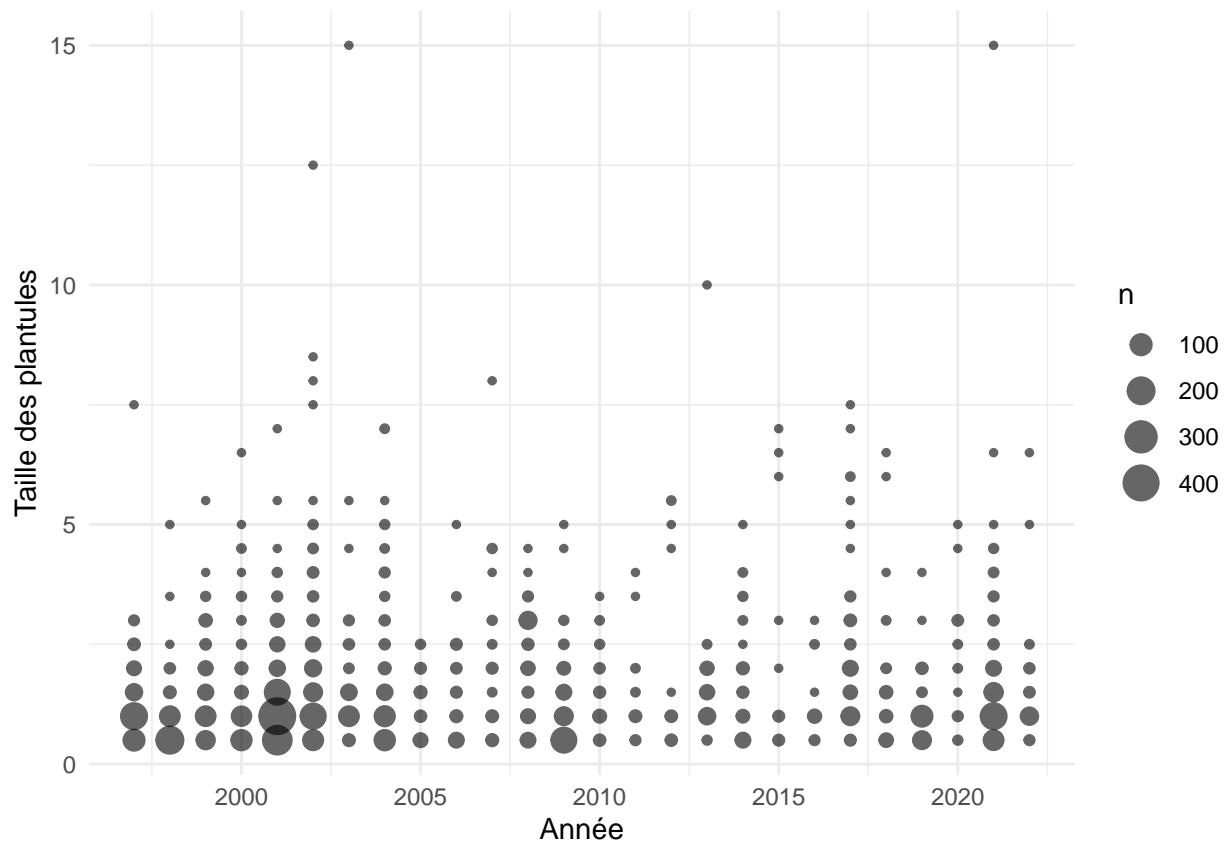
## Taille des plantules

```
plantule_data <- centauree_data_complet[centauree_data_complet$age0==1,]
```

```
# Taille des plantules / année

plantule_data %>%
  ggplot(aes(x = year, y = Size0Mars)) +
  geom_count(alpha=0.6) +
  labs(x = "Année",
       y = "Taille des plantules") +
  theme_minimal()
```

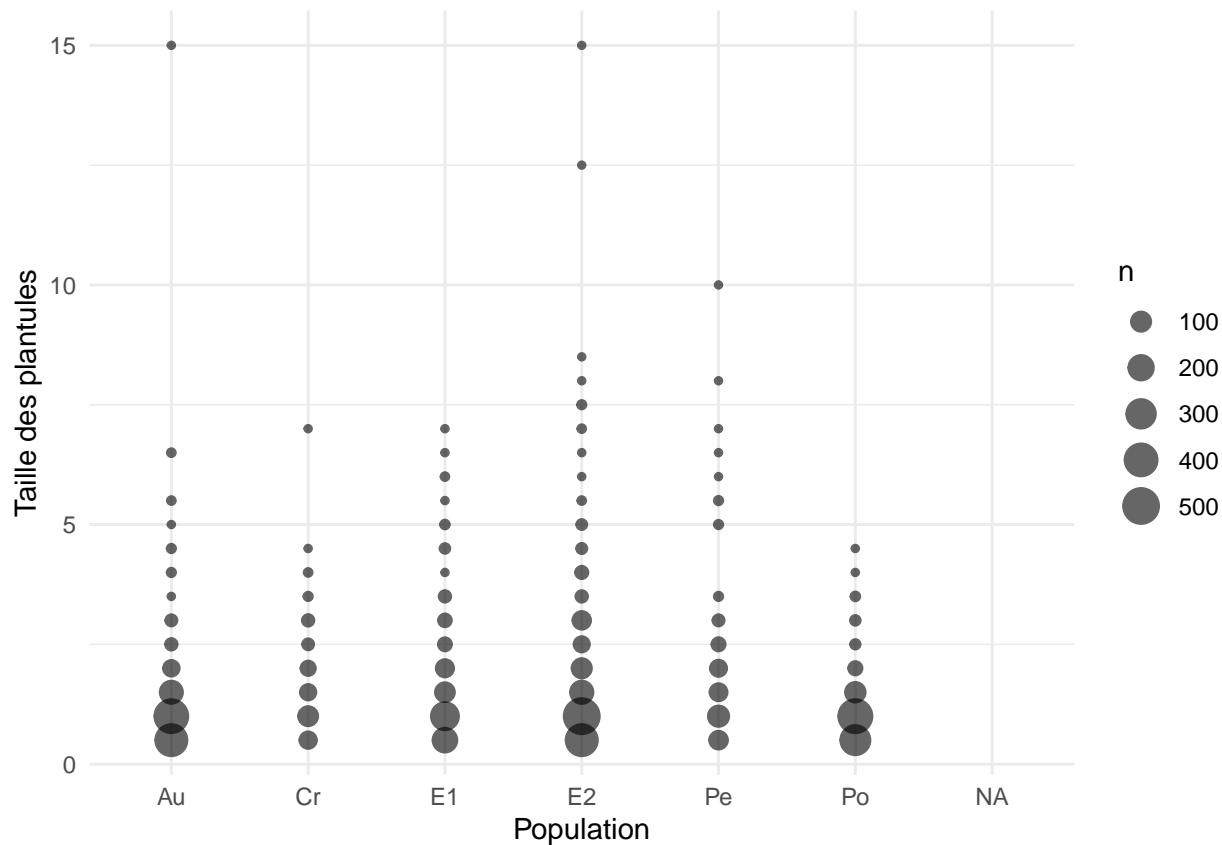
```
## Warning: Removed 2360 rows containing non-finite outside the scale range
## ('stat_sum()'').
```



```
# Taille des plantules / population

plantule_data %>%
  ggplot(aes(x = Pop, y = Size0Mars)) +
  geom_count(alpha=0.6) +
  labs(x = "Population",
       y = "Taille des plantules") +
  theme_minimal()
```

```
## Warning: Removed 2360 rows containing non-finite outside the scale range
## ('stat_sum()').
```



```
Pltglm1 <- fitme(Size0Mars ~ 1 + (1|year) + (1|Pop) + (1|Pop:year),
                   data=plantule_data)
```

```
Pltglm2 <- fitme(Size0Mars ~ 1 + (1|Pop) + (1|Pop:year),
                   data=plantule_data)
```

```
Pltglm3 <- fitme(Size0Mars ~ 1 + (1|Pop:year),
                   data=plantule_data)
```

```
Pltglm4 <- fitme(Size0Mars ~ 1 + (1|year) + (1|Pop:year),
                   data=plantule_data)
```

```
Pltglm5 <- fitme(Size0Mars ~ 1 + (1|year) + (1|Pop),
                   data=plantule_data)
```

```
summary(Pltglm1)
```

```
## formula: Size0Mars ~ 1 + (1 | year) + (1 | Pop) + (1 | Pop:year)
## Estimation of fixed effects by ML.
## Estimation of lambda and phi by 'outer' ML, maximizing logL.
## family: gaussian( link = identity )
##   ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept) 1.382  0.1267 10.91
##   ----- Random effects -----
```

```

## Family: gaussian( link = identity )
## --- Variance parameters ('lambda'):
## lambda = var(u) for u ~ Gaussian;
##   year : 0.03363
##   Pop : 0.06829
##   Pop:year : 0.3535
## # of obs: 4648; # of groups: year, 26; Pop, 6; Pop:year, 136
## ----- Residual variance -----
## phi estimate was 0.688727
## ----- Likelihood values -----
##           logLik
## logL      (p_v(h)): -5891.028

```

```
summary(Pltglm2)
```

```

## formula: Size0Mars ~ 1 + (1 | Pop) + (1 | Pop:year)
## Estimation of fixed effects by ML.
## Estimation of lambda and phi by 'outer' ML, maximizing logL.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept) 1.374  0.1213  11.32
## ----- Random effects -----
## Family: gaussian( link = identity )
## --- Variance parameters ('lambda'):
## lambda = var(u) for u ~ Gaussian;
##   Pop : 0.06649
##   Pop:year : 0.3966
## # of obs: 4648; # of groups: Pop, 6; Pop:year, 136
## ----- Residual variance -----
## phi estimate was 0.688347
## ----- Likelihood values -----
##           logLik
## logL      (p_v(h)): -5891.66

```

```
summary(Pltglm3)
```

```

## formula: Size0Mars ~ 1 + (1 | Pop:year)
## Estimation of fixed effects by ML.
## Estimation of lambda and phi by 'outer' ML, maximizing logL.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept) 1.354  0.06375  21.24
## ----- Random effects -----
## Family: gaussian( link = identity )
## --- Variance parameters ('lambda'):
## lambda = var(u) for u ~ Gaussian;
##   Pop:year : 0.4668
## # of obs: 4648; # of groups: Pop:year, 136
## ----- Residual variance -----
## phi estimate was 0.688088
## ----- Likelihood values -----

```

```

##          logLik
## logL      (p_v(h)): -5895.849

summary(Pltglm4)

## formula: Size0Mars ~ 1 + (1 | year) + (1 | Pop:year)
## Estimation of fixed effects by ML.
## Estimation of lambda and phi by 'outer' ML, maximizing logL.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept) 1.357  0.06779 20.01
## ----- Random effects -----
## Family: gaussian( link = identity )
##     --- Variance parameters ('lambda'):
## lambda = var(u) for u ~ Gaussian;
##   year : 0.01713
##   Pop:year : 0.4463
## # of obs: 4648; # of groups: year, 26; Pop:year, 136
## ----- Residual variance -----
## phi estimate was 0.688217
## ----- Likelihood values -----
##          logLik
## logL      (p_v(h)): -5895.717

summary(Pltglm5)

## formula: Size0Mars ~ 1 + (1 | year) + (1 | Pop)
## Estimation of fixed effects by ML.
## Estimation of lambda and phi by 'outer' ML, maximizing logL.
## family: gaussian( link = identity )
## ----- Fixed effects (beta) -----
##           Estimate Cond. SE t-value
## (Intercept) 1.357  0.1032 13.14
## ----- Random effects -----
## Family: gaussian( link = identity )
##     --- Variance parameters ('lambda'):
## lambda = var(u) for u ~ Gaussian;
##   year : 0.08524
##   Pop : 0.04151
## # of obs: 4648; # of groups: year, 26; Pop, 6
## ----- Residual variance -----
## phi estimate was 0.77653
## ----- Likelihood values -----
##          logLik
## logL      (p_v(h)): -6050.509

Pltpredict1 <- predict(Pltglm1, newdata = fake_data)[,1]
Pltpredict2 <- predict(Pltglm2, newdata = fake_data)[,1]
Pltpredict3 <- predict(Pltglm3, newdata = fake_data)[,1]
Pltpredict4 <- predict(Pltglm4, newdata = fake_data)[,1]
Pltpredict5 <- predict(Pltglm5, newdata = fake_data)[,1]

```

```

plot_plantule <- function(data = fake_data, prediction, var, fact) {
  data %>%
    mutate(plt_predi = prediction) %>%
    ggplot(aes(x = .data[[var]], y = plt_predi)) +
    geom_line(aes(color = as.factor(.data[[fact]]))) +
    theme_minimal()
}

plot_plantule2 <- function(data = fake_data, prediction, var, fact) {
  data %>%
    mutate(plt_predi = prediction) %>%
    ggplot(aes(x = .data[[var]], y = plt_predi)) +
    geom_point(aes(color = as.factor(.data[[fact]]))) +
    theme_minimal()
}

```

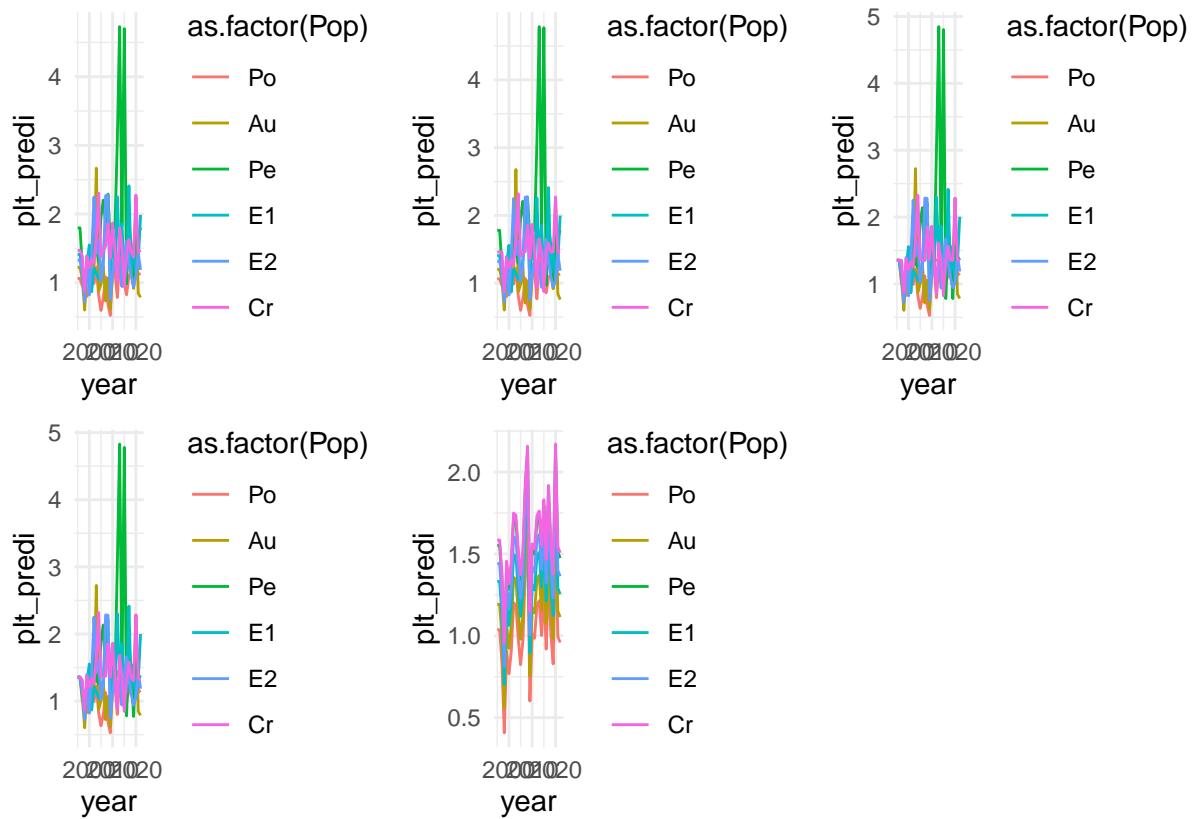
## Taille des plantules en fonction de l'année

```

var <- "year"
fact <- "Pop"

wrap_plots(
  plot_plantule(prediction = Pltpredict1, var=var, fact=fact),
  plot_plantule(prediction = Pltpredict2, var=var, fact=fact),
  plot_plantule(prediction = Pltpredict3, var=var, fact=fact),
  plot_plantule(prediction = Pltpredict4, var=var, fact=fact),
  plot_plantule(prediction = Pltpredict5, var=var, fact=fact)
)

```



## Taille des plantules en fonction de la population

```

var <- "Pop"
fact <- "year"

wrap_plots(
  plot_plantule2(prediction = Pltpredict1, var=var, fact=fact),
  plot_plantule2(prediction = Pltpredict2, var=var, fact=fact),
  plot_plantule2(prediction = Pltpredict3, var=var, fact=fact),
  plot_plantule2(prediction = Pltpredict4, var=var, fact=fact),
  plot_plantule2(prediction = Pltpredict5, var=var, fact=fact)
)
  
```

