Package compilation for saemix 3.4 and basic run

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Copy files

```
cmd<-paste("rm -r ",file.path(workDir,"*"),sep="")</pre>
system(cmd)
dir.create(workDir)
## Warning in dir.create(workDir): '/home/eco/work/saemix/versions/saemix3.4'
## existe déjà
dir.create(file.path(workDir, "saemix"))
cmd<-paste("cp -rp ",file.path(saemixDir,"documentation","docsaem.pdf")," ",file.path(saemixDir,"inst",</pre>
system(cmd)
cmd<-paste("cp -rp ",file.path(saemixDir,"R")," ", file.path(workDir,"saemix","/"),sep="")</pre>
system(cmd)
cmd<-paste("cp -rp ",file.path(saemixDir,"data")," ", file.path(workDir,"saemix","/"),sep="")</pre>
system(cmd)
cmd<-paste("cp -rp ",file.path(saemixDir,"inst")," ", file.path(workDir,"saemix","/"),sep="")</pre>
system(cmd)
for(ifile in c("CHANGES", "DESCRIPTION")) {
  cmd<-paste("cp ",file.path(saemixDir,ifile)," ", file.path(workDir, "saemix"),sep="")</pre>
  system(cmd)
cmd<-paste("cp ",file.path(saemixDir,"inst","CITATION")," ", file.path(workDir,"saemix","inst"),sep="")</pre>
system(cmd)
```

Compilation

- building on win-ftp
 - to specify when submitting Possibly mis-spelled words in DESCRIPTION: IAME (17:930) al (17:663) et (17:660) github (17:954) saemix (17:965)
 - removed: Found the following (possibly) invalid URLs: URL: http://group.monolix.org/ From: DESCRIPTION Status: 403 Message: Forbidden
 - to change: no commas between keywords in R (maybe for vignettes)
 - previous version, solved now:

```
```{r}
Found the following \keyword or \concept entries
which likely give several index terms:
```

```
File 'backward.procedure.Rd':
 \keyword{backward,}
 \keyword{selection,}
... (others)
```

- Examples with CPU or elapsed time > 5s user system elapsed rapi.saemix 47.006 0.096 47.115 cow.saemix 25.185 0.120 25.311 PD1.saemix 18.925 0.048 19.108 toenail.saemix 16.656 0.020 16.676 compare.saemix 9.152 0.020 9.171 theo.saemix 6.481 0.008 6.490 yield.saemix 5.419 0.016 5.435
- new NOTE (October 2023)
- checking HTML version of manual ... NOTE Skipping checking HTML validation: no command 'tidy' found
- compilation to pdf (as comments)

Current dependencies, versions downloaded on 29/07/25: - varTestnlme - nlive - mkin

```
setwd(workDir)
system("R CMD build saemix")
Test examples
if(testExamples)
 system("R CMD check --as-cran --run-donttest saemix_3.4.tar.gz") else
 system("R CMD check --as-cran saemix_3.4.tar.gz")
Reverse dependencies
cmd<-paste("cp ",file.path("/home/eco/work/saemix/versions/reverseDependencies/varTestnlme_1.3.5.tar.gz</pre>
system(cmd)
cmd<-paste("cp ",file.path("/home/eco/work/saemix/versions/reverseDependencies/nlive_0.8.0.tar.gz")," "</pre>
system(cmd)
cmd<-paste("cp ",file.path("/home/eco/work/saemix/versions/reverseDependencies/mkin_1.2.10.tar.gz")," "</pre>
system(cmd)
if(checkReverseDependencies) {
 # Check which packages depend on saemix
 package_dependencies(packages="saemix", reverse=TRUE)
 # TODO download these packages into workDir
}
$saemix
[1] "GrowthCurveME" "mkin"
 "nlive"
 "varTestnlme"
if(installPackage) {
 # install saemix current version and check packages
 install.packages(pkgs=file.path(workDir, "saemix_3.4.tar.gz"), repos=NULL)
 #result <- check_packages_in_dir(workDir, revdep = list())</pre>
 result <- check_packages_in_dir(workDir, revdep = list("varTestnlme"))</pre>
 summary(result)
}
```

**Examples on CRAN** Examples with CPU or elapsed time > 5s user system elapsed rapi.saemix 32.510 0.092 32.602 cow.saemix 17.020 0.072 17.093 toenail.saemix 13.531 0.000 13.532 PD1.saemix 12.789 0.035 12.910 compare.saemix 6.331 0.016 6.347 theo.saemix 5.171 0.000 5.172

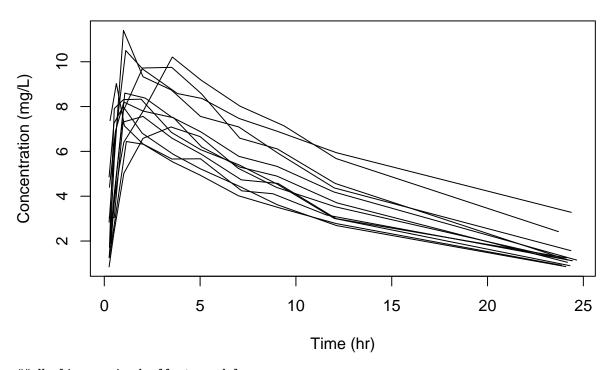
#### Warnings

## Check

Install package in development mode

```
dev_mode() # development mode
v Dev mode: ON
install.packages(pkgs=file.path(workDir, "saemix_3.4.tar.gz"), repos=NULL)
Installation du package dans '/home/eco/R-dev'
(car 'lib' n'est pas spécifié)
library(saemix)
library(testthat)
##
Attachement du package : 'testthat'
L'objet suivant est masqué depuis 'package:devtools':
##
##
 test_file
Running theopp example
?theo.saemix
Aucune documentation pour 'theo.saemix' n'a été trouvée dans les packages et les bibliothèques :
vous pourriez essayer '??theo.saemix'
Aucune documentation pour 'saemix' n'a été trouvée dans les packages et les bibliothèques :
vous pourriez essayer '??saemix'
Run on the theophylline example
Object of class SaemixData
##
 longitudinal data for use with the SAEM algorithm
Dataset theo.saemix
##
 Structured data: Concentration ~ Dose + Time | Id
##
 X variable for graphs: Time (hr)
##
 covariates: Weight (kg), Sex (-)
Dataset characteristics:
##
 number of subjects:
 12
##
 number of observations: 120
##
 average/min/max nb obs: 10.00 / 10 / 10
First 10 lines of data:
##
 Ιd
 Dose Time Concentration Weight Sex mdv cens occ ytype
 1 319.992 0.25
1
 2.84
 79.6
 1
 1
 0
 0
2
 1 319.992 0.57
 6.57
 79.6
 1
 0
 0
 1
3
 1 319.992 1.12
 10.50
 79.6
 1
 0
 0
 1
 1
4
 1 319.992 2.02
 9.66
 79.6
 1
 0
 1
 1 319.992 3.82
5
 8.58
 79.6
 0
 0 1
 1
 1
6
 1 319.992 5.10
 8.36
 79.6
 1
 0
 1
 1
7
 1 319.992 7.03
 7.47
 1 0
 0 1
 79.6
 1
8
 1 319.992 9.05
 6.89
 79.6
 1 0
```

```
9 1 319.992 12.12 5.94 79.6 1 0 0 1 1 ## 10 1 319.992 24.37 3.28 79.6 1 0 0 1 1
```



```
Nonlinear mixed-effects model
 Model function: One-compartment model with first-order absorption
##
 Model type: structural
<srcref: file "" chars 10:12 to 19:1>
##
 Nb of parameters: 3
##
 parameter names: ka V CL
##
 distribution:
 Parameter Distribution Estimated
##
[1,] ka
 log-normal
 Estimated
[2,] V
 log-normal
 Estimated
[3,] CL
 log-normal
##
 Variance-covariance matrix:
##
 ka V CL
ka 1 0 0
 0 1 0
CL 0 0 1
 Error model: constant , initial values: a.1=1 \,
##
 Covariate model:
 ka V CL
[1,] 0 1 0
[2,] 0 0 0
##
 Initial values
 ka V
Pop.CondInit 1.0 20 0.50
Cov.CondInit 0.1 0 -0.01
Nonlinear mixed-effects model fit by the SAEM algorithm
```

```
Data

Object of class SaemixData
 longitudinal data for use with the SAEM algorithm
Dataset theo.saemix
##
 Structured data: Concentration ~ Dose + Time | Id
 X variable for graphs: Time (hr)
 covariates: Weight (kg), Sex (-)
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 0 1
3
 1 319.992 1.12
 10.50
 79.6
 1 0
 0 1
4
 1 319.992 2.02
 9.66
 79.6 1 0
5 1 319.992 3.82
 8.58
 79.6 1 0
 0 1
 1 319.992 5.10
6
 8.36
 79.6
 1 0
 0 1
7.47
 79.6 1 0
 0 1
8 1 319.992 9.05
 6.89 79.6 1 0
 0 1
9 1 319.992 12.12
 5.94
 79.6 1 0
 1
10 1 319.992 24.37
 79.6 1 0
 3.28

 Model

Nonlinear mixed-effects model
Model function: One-compartment model with first-order absorption
Model type: structural
<srcref: file "" chars 10:12 to 19:1>
<bytecode: 0x604f6ab1c388>
##
 Nb of parameters: 3
##
 parameter names: ka V CL
##
 distribution:
 Parameter Distribution Estimated
[1,] ka log-normal Estimated
[2,] V
 log-normal
 Estimated
[3,] CL
 log-normal
 Estimated
##
 Variance-covariance matrix:
 ka V CL
ka 1 0 0
 0 1 0
CL 0 0 1
 Error model: constant, initial values: a.1=1
##
 Covariate model:
##
 [,1] [,2] [,3]
Weight 0 1
##
 Initial values
##
 ka V
Pop.CondInit 1.0 20 0.50
Cov.CondInit 0.1 0 -0.01

---- Key algorithm options ----
```

```
##
 Estimation of individual parameters (MAP)
##
 Estimation of standard errors and linearised log-likelihood
##
 Estimation of log-likelihood by importance sampling
##
 Number of iterations: K1=300, K2=100
##
 Number of chains: 5
##
 Seed: 632545
 Number of MCMC iterations for IS: 5000
##
##
 Simulations:
##
 nb of simulated datasets used for npde: 1000
##
 nb of simulated datasets used for VPC: 100
##
 Input/output
##
 save the results to a file: FALSE
##
 save the graphs to files: FALSE
 Results
----- Fixed effects -----

 Estimate SE
 Parameter
 CV(%) p-value
 1.5588 0.3071 19.7 -
[1,] ka
[2,] V
 18.8423 5.6328 29.9 -
[3,] beta_Weight(V) 0.0073 0.0042 58.0 0.085
 ## [4,] CL
[5,] a.1

----- Variance of random effects -----

 Parameter Estimate SE
 CV(%)
ka omega2.ka 0.414 0.1853 45
V omega2.V 0.012
 0.0078 64
CL omega2.CL 0.077 0.0368 48

----- Correlation matrix of random effects -----

 omega2.ka omega2.V omega2.CL
omega2.ka 1 0
omega2.V 0
 1
 0
omega2.CL 0
 0

----- Statistical criteria -----

Likelihood computed by linearisation
##
 -2LL= 341.3649
##
 AIC = 357.3649
 BIC = 361.2442
##
Likelihood computed by importance sampling
##
 -2LL= 342.6478
 AIC = 358.6478
##
 BIC = 362.5271
Nonlinear mixed-effects model fit by the SAEM algorithm
```

```
Data

Object of class SaemixData
 longitudinal data for use with the SAEM algorithm
Dataset theo.saemix
 Structured data: Concentration ~ Dose + Time | Id
 X variable for graphs: Time (hr)
##
##
 covariates: Weight (kg), Sex (-)
Dataset characteristics:
 number of subjects:
 12
##
 number of observations: 120
 average/min/max nb obs: 10.00 / 10 / 10
First 10 lines of data:
##
 Ιd
 Dose Time Concentration Weight Sex mdv cens occ ytype
1
 1 319.992 0.25
 2.84
 79.6
 1
 0
 0 1
2
 1 319.992 0.57
 6.57
 79.6
 1
 0
 0 1
3
 1 319.992 1.12
 10.50 79.6 1 0
4 1 319.992 2.02
 9.66 79.6 1 0
 0 1
 8.58
 1 319.992 3.82
5
 79.6 1 0
6 1 319.992 5.10
 8.36 79.6 1 0
 0 1
7 1 319.992 7.03
 7.47 79.6 1 0
8 1 319.992 9.05
 0 1
 6.89
 79.6 1 0
 1
 1 319.992 12.12
 5.94
 79.6 1 0
 0 1
 1
10 1 319.992 24.37
 3.28
 79.6 1 0
 0 1

 Model

Nonlinear mixed-effects model
 Model function: One-compartment model with first-order absorption
 Model type: structural
<srcref: file "" chars 10:12 to 19:1>
<bytecode: 0x604f6ab1c388>
Nb of parameters: 3
##
 parameter names: ka V CL
##
 distribution:
 Parameter Distribution Estimated
[1,] ka
 log-normal Estimated
[2,] V
 log-normal Estimated
[3,] CL
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 Estimated
 Variance-covariance matrix:
##
 ka V CL
ka 1 0 0
V 0 1 0
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 Error model: constant , initial values: a.1=1
##
 Covariate model:
##
 [,1] [,2] [,3]
Weight 0 1
##
 Initial values
 ka V
##
Pop.CondInit 1.0 20 0.50
Cov.CondInit 0.1 0 -0.01

```

```
---- Key algorithm options ----

##
 Estimation of individual parameters (MAP)
##
 Estimation of standard errors and linearised log-likelihood
##
 Estimation of log-likelihood by importance sampling
##
 Number of iterations: K1=300, K2=100
##
 Number of chains: 5
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##
 Number of MCMC iterations for IS: 5000
##
 Simulations:
##
 nb of simulated datasets used for npde: 1000
##
 nb of simulated datasets used for VPC: 100
##
 Input/output
 save the results to a file: FALSE
##
##
 save the graphs to files: FALSE

 Results

----- Fixed effects -----

Warning in .local(x, ...): NAs introduits lors de la conversion automatique
 Estimate SE
 CV(%) p-value
##
 Parameter
[1,] ka
 1.5588 0.3071 19.7
[2,] V
 18.8423 5.6328 29.9 -
[3,] beta_Weight(V) 0.0073 0.0042 58.0 0.085
[4,] CL
 2.7717 0.2431 8.8 -
[5,] a.1
 0.7389 0.0565 7.7 -

----- Variance of random effects -----

##
 Parameter Estimate SE
 CV(%)
ka omega2.ka 0.414 0.1853 45
V omega2.V 0.012
 0.0078 64
CL omega2.CL 0.077 0.0368 48
----- Correlation matrix of random effects -----

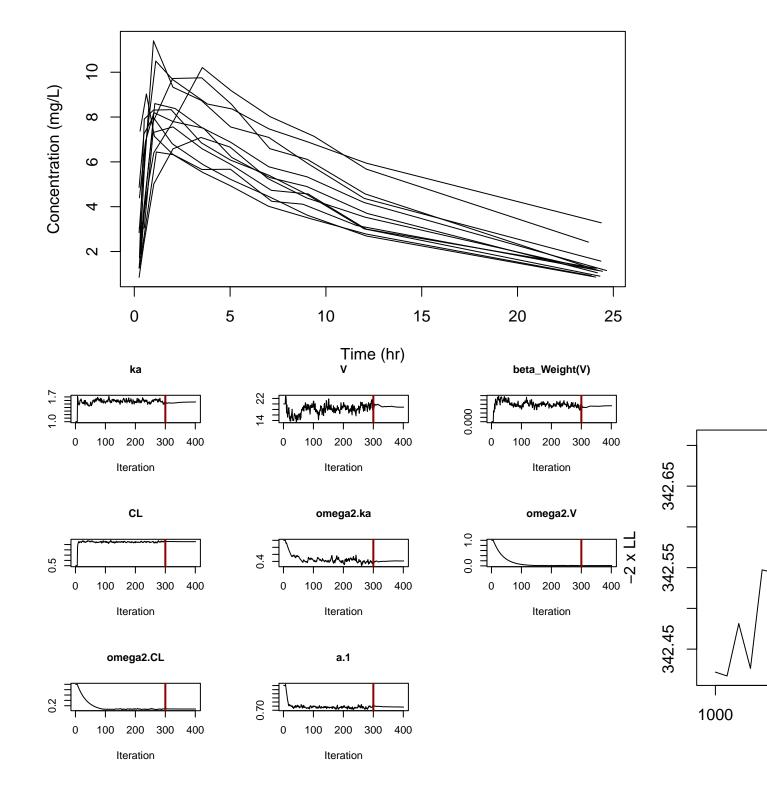
##
 omega2.ka omega2.V omega2.CL
 0
 0
omega2.ka 1
omega2.V 0
 1
 0
omega2.CL 0
 0
 1

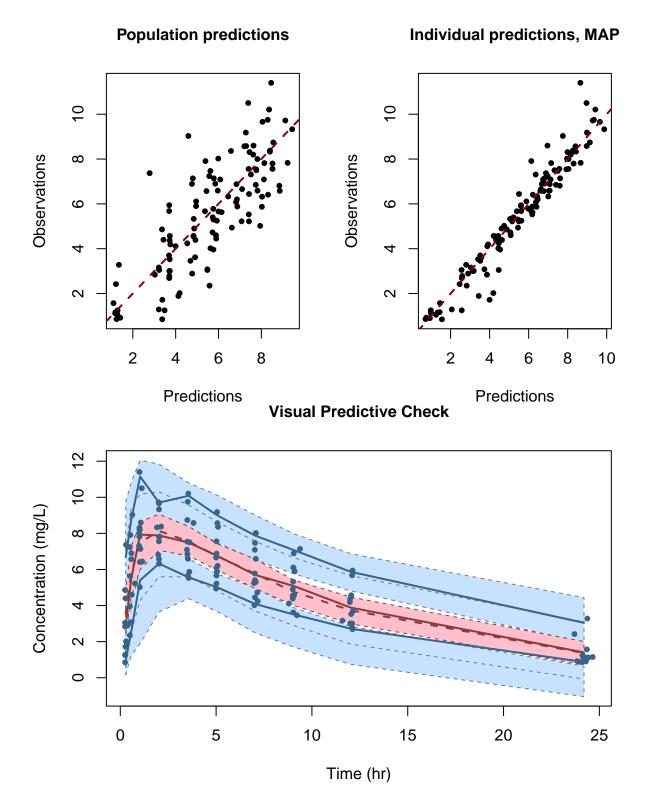
----- Statistical criteria -----

Likelihood computed by linearisation
##
 -2LL= 341.3649
##
 AIC = 357.3649
##
 BIC = 361.2442
##
Likelihood computed by importance sampling
##
 -2LL= 342.6478
##
 AIC = 358.6478
##
 BIC = 362.5271
```

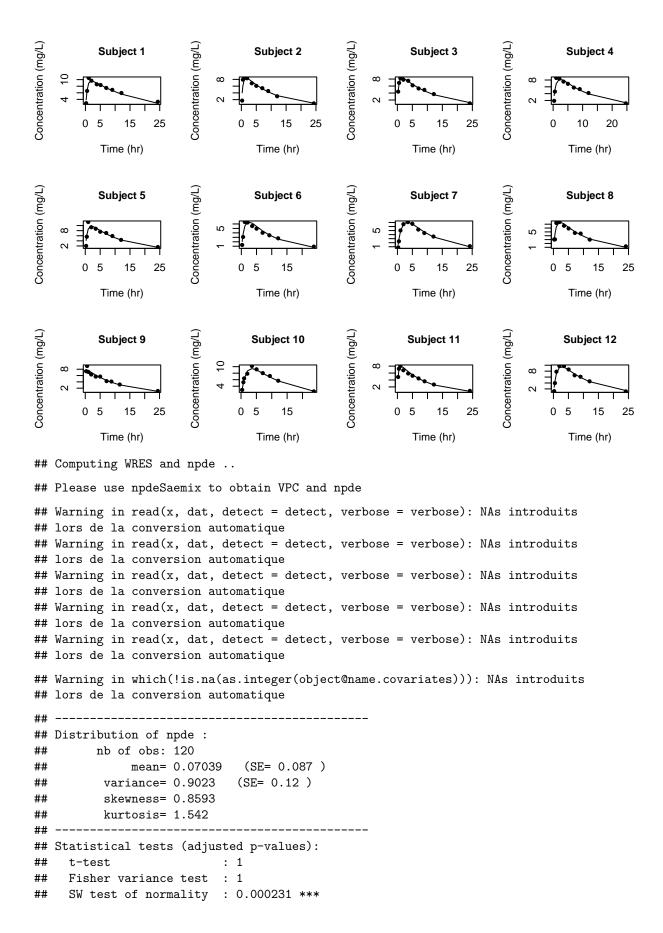
```
Object of class SaemixSimData
 data simulated according to a non-linear mixed effect model
Characteristics of original data
 number of subjects: 12
##
 summary of response:
##
##
 Min. 1st Qu. Median Mean 3rd Qu.
 Max.
 3.513 5.665 5.447 7.325 11.400
##
 0.850
Characteristics of simulated data
##
 number of simulated datasets: 1000
##
 summary of simulated response
 Min. 1st Qu. Median
##
 Mean 3rd Qu.
 Max.
-2.364 3.675 5.615 5.479
 7.359 14.679
##
 CL
 ka
1 1.7863333 29.59934 1.681678
2 1.9372081 31.98292 3.178891
3 2.2543380 33.18598 2.854727
4 1.2081689 31.53805 2.694492
5 1.4877219 27.10964 2.399152
6 1.0608154 38.09466 4.028817
7 0.6907825 32.17493 3.280839
8 1.3001684 34.17845 3.297077
9 6.3284102 32.50705 2.832542
10 0.7559841 26.58247 1.891913
11 3.1478359 35.17647 3.763632
12 0.9503826 26.06740 2.424328
Plot results:
```

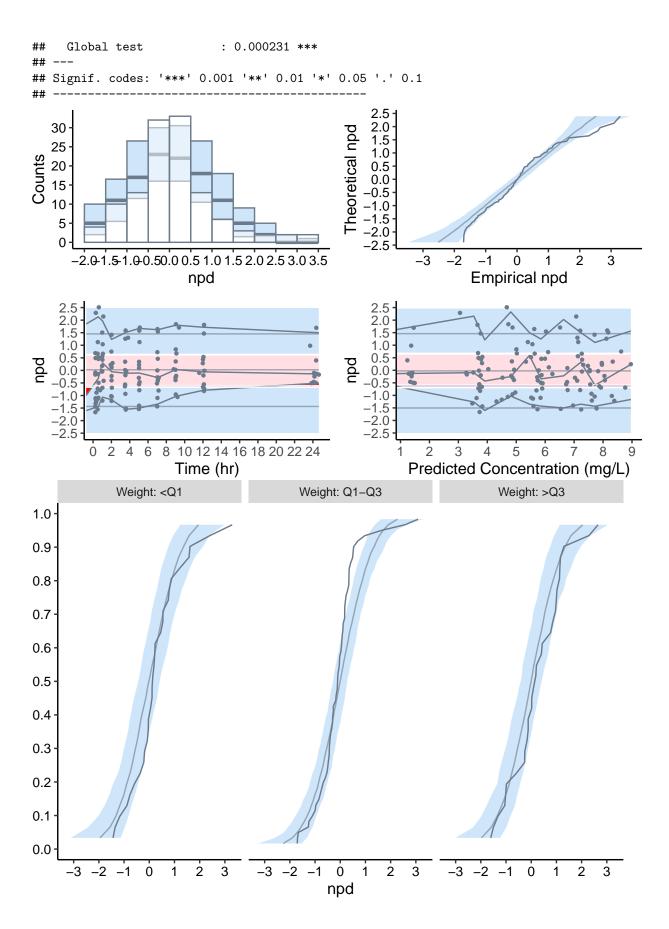
## Simulating data using nsim = 1000 simulated datasets ## Computing WRES and npde ...

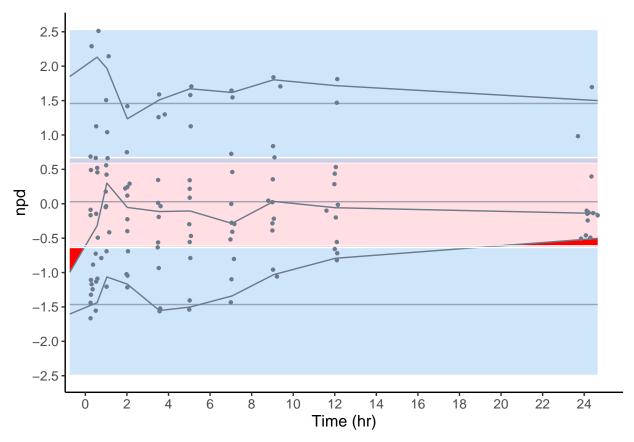




## Computing WRES and npde ...







Bootstrap

## v Dev mode: OFF