# Moving to saemix 3.0

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# TODO list

- kompare files with 2.4 to take into account changes made for the CRAN compiler
- error models
  - combined 2 versus combined 1
- bugfixes
  - simulated annealing
- documentation
  - error models
  - new models from Belhal for responses defined by their likelihood
  - interface to mkin
  - new references
  - new defaults: no plot

# Comparing 2.4 and 3.0

## **Function list**

#### 2.4

aaa\_generics.R func\_diagnostics.R func\_FIM.R main\_estep.R main.R SaemixObject.R compute\_LL.R func\_distcond.R func\_plots.R main\_initialiseMainAlgo.R SaemixData.R SaemixRes.R func\_aux.R func\_estimParam.R func\_simulations.R main\_mstep.R SaemixModel.R zzz.R

#### 3.0

aaa\_generics.R func\_aux.R func\_estimParam.R func\_stepwise.R main.R SaemixRes.R backward.R func\_compare.R func\_FIM.R main\_estep.R SaemixData.R stepwise.R compute\_LL.R func\_distcond\_noplot.R func\_plots.R main\_initialiseMainAlgo.R SaemixModel.R zzz.R forward.R func\_distcond.R func\_simulations.R main\_mstep.R SaemixObject.R

#### functions in 3.0 not in 2.4

- covariate selection algorithm (Maud)
  - forward.R, backward.R, stepwise.R, func\_stepwise.R, func\_compare.R
  - func\_diagnostics.R (? Maud or something else ?)
- version without plots of func distcond.R

# Function comparison

See **changes 2.4 to 3.0.ods** for side-by-side comparison of functions tracing the changes between 2.4 and 3.0, as well as the changes carried over from 2.4 immediately.

#### Loading

#### aaa\_generics.R TODO

- check generic definition of read
- check aliases for some internal functions

#### zzz.R

• check if the current version works or if we need to add date

#### Classes

#### SaemixData

- check object validation
  - create testthat functions
- harmonise output messages across the package
  - no messages to be output to stdin by default
- new slot automatic in 3.0
  - normally allows automatic recognition and filling in arguments not given
  - $\square$  TODO: testthat function
  - also test validate.names()
- use is(x, "data.frame") instead of testing class(x)=="data.frame" (class may have more than 1 value)

#### SaemixModel

- main change in 3.0: added discrete response with a modelType argument
  - for joint models, will need some tweak (maybe as a vector of response types?)
- $\bullet \ \ new \ function \ validate\_covariance.model$ 
  - $\square$  TODO: change name to validate.covariance.model
  - check function (add testthat) and add documentation

## SaemixRes

- check definition of ypred and ppred
  - ypred should be f(theta\_MAP) and ppred should be E(f(theta))

#### SaemixObject

- maybe change name saemix.simul to saemix.simulate
- $\square$  TODO: logLik.saemix, AIC.SaemixObject, BIC.SaemixObject
  - Maud made changes in the 2.4
  - but Johannes also made some changes in 3.0
  - $-\Box$  TODO: add Johannes's changes to the 2.4 version
- options
  - — □ TODO: see CRAN for the rules (like for 2.4, in compute\_LL.R)

## Computational functions

#### compute\_LL.R

• check if alias ggq.mlx exists

#### func aux.R

- modified combined error model
  - □ TODO: add to documentation
- conditional distribution function split into 2 functions according to nature of the model
  - □ TODO: add to documentation
- check computation of compute.LLy

#### func FIM.R

- $\Box$  TODO: secure the code for discrete data models (only compute LL by linearisation ? do not use altogether ?)
- $\square$  TODO: check cat and replace with message
  - check name of option to print out messages (maybe name of option changed)

#### Main algorithm

#### main\_initialiseMainAlgo.R

- $\hfill\Box$  TODO: check cat and replace with message
  - check name of option to print out messages (maybe name of option changed)

## main\_estep.R

- check computation of compute.LLy now completely replacing the computation of U.y
- 4th kernel added (Laplacian kernel)
  - $-\Box$  TODO: add default option (0 iterations)
  - $\square$  TODO: add to documentation and recommendations

#### main mstep.R

- check computation of sigma and influence of SA (nbiter.saemix changed to nbiter.sa)
  - − □ TODO: check
- $\square$  TODO: add Lucie's changes +++
  - ask Lucie to check afterwards

#### main.R

- $\Box$  TODO: check cat and replace with message
  - check name of option to print out messages (maybe name of option changed)

## Parameters, Simulations

#### func\_distcond.R

- commented out the plots for the moment
  - $-\Box$  TODO: add option to output the graphs

# $func\_estimParam.R$

- renamed the function to predict newdata as in 2.4 (also has an Roxygen documentation now)
  - copied the old file in newCode
  - $\square$  TODO: check that the code within is similar

## $func\_simulations.R$

- renamed simul.saemix in saemix.simul in 3.0 for consistency (all functions start with saemix and not end with it)
  - $\square$  TODO: add to documentation and CHANGELOG

## Plots

- $\square$  TODO: check cat and replace with message
  - check name of option to print out messages (maybe name of option changed)

# New functions by Maud

#### BIC criterion

- definition and computation included in SaemixObject (see above)
  - $\square$  TODO: add to documentation and CHANGELOG

# Algorithm

- create a notebook to test the algorithm and integrate it
- add test example
  - $\square$  TODO: add to documentation

#### $\mathbf{H}\mathbf{M}\mathbf{M}$

 $\Box$  TODO: check status with Maud