



SAMPLING OF PARAMETER UNCERTAINTY IN CAMPSIS

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INTRODUCTION

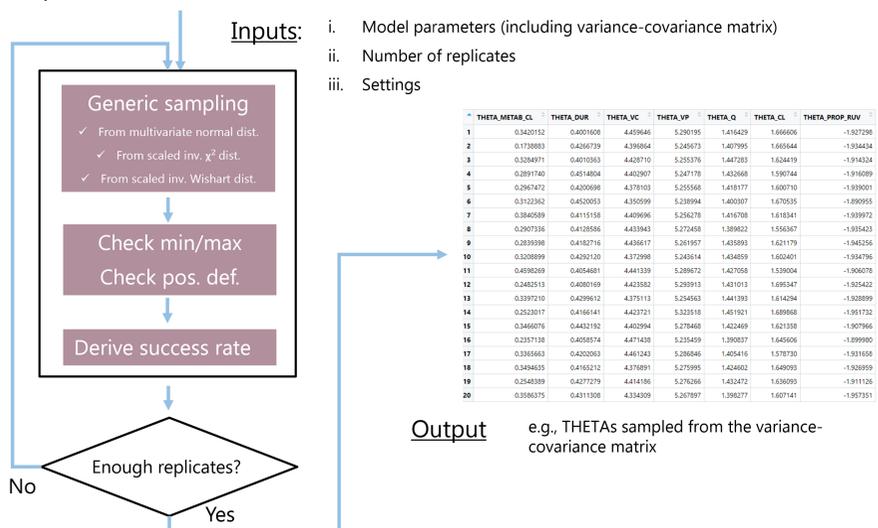
Campsis (www.campsis.org) is an open-source PK/PD simulation suite running R, freely available on CRAN [1] and GitHub [2], designed to facilitate the simulation of population PK/PD models, and by extension, the simulation of clinical trials. This poster presents a revised way of simulating with parameter uncertainty, including random sampling from the scaled inverse Chi-squared or Wishart distributions for the OMEGA and SIGMA matrix.

Objectives

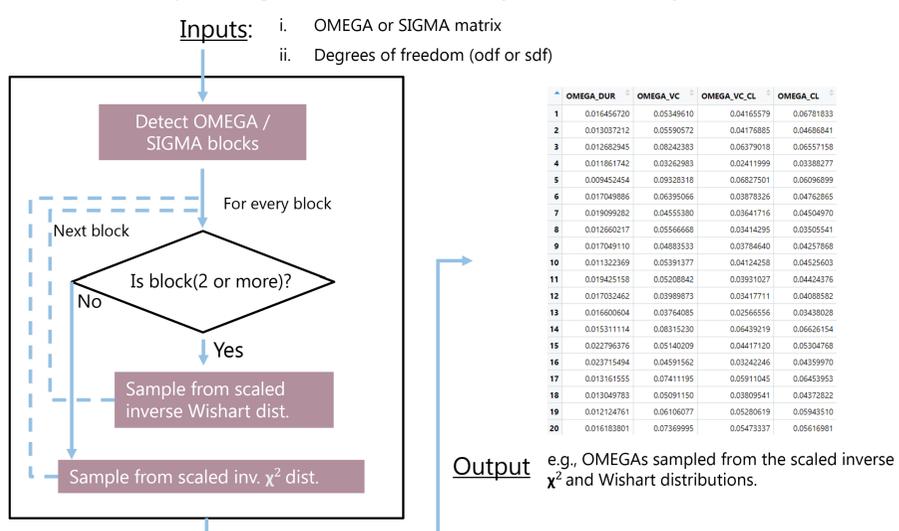
- Revisit the previous process of simulating with parameter uncertainty in Campsis.
- Sample parameters from various sources based on user choice:
 - **Variance-covariance matrix** (THETAs, OMEGAs and SIGMAs)
 - **Scaled inverse Chi-squared distribution** (for single OMEGA or SIGMA)
 - **Scaled inverse Wishart distribution** (for block(s) of OMEGAs or SIGMAs)
 - **External source** (e.g., manual data frame, CIR file, etc.)
- Allow the users to enter min and max values for each parameter and check these limits during parameter sampling
- When parameters are sampled from the variance-covariance matrix, check the OMEGA or SIGMA matrix is positive definite

DESIGN

The sampling of parameters with uncertainty was implemented in the package *Campsismod* [3] [4] as a generic iterative process (see diagram). First, the model parameters are sampled for each replicate. Second, the model parameters are checked based on the minimal and maximal values provided by the user and based on the positive definiteness of the OMEGA or SIGMA matrices. If the parameters are not valid, the process is repeated until valid parameters are found.



This iterative process is called once to sample the parameters from the variance-covariance matrix (THETAs and, on user's request, OMEGAs and SIGMAs) and is called a second time to sample the OMEGAs and SIGMAs from the scaled inverse Chi-squared or Wishart distributions (see diagram below). Before sampling the OMEGAs/SIGMAs, the structural blocks of the OMEGA/SIGMA matrix are automatically detected. Different degrees of freedom corresponding to these blocks may be entered by the user.



INPUTS / OUTPUTS

Inputs:

- Campsis model: it includes all model parameters and the variance-covariance matrix. All parameters include min and max values.
- Settings for random sampling:
 - By default, parameters are randomly sampled from the variance-covariance matrix
 - **Wishart** argument: OMEGAs/SIGMAs are sampled from the scaled inverse Chi-squared or Wishart distribution. In that case, degrees of freedom are required.
 - **odf** argument: degrees of freedom for the OMEGAs, single value (same for all) or a vector (one specific value per OMEGA block)
 - **sdf** argument: degrees of freedom for the SIGMAs, single value (same for all) or a vector (one specific value per SIGMA block)
 - **Check min/max** values: TRUE/FALSE
 - **Check positive definiteness** of OMEGA/SIGMA matrix: TRUE/FALSE
- Import from data frame: alternatively, model parameters can be provided in the form of an external data frame from a different source (e.g., from a bootstrap or SIR run).

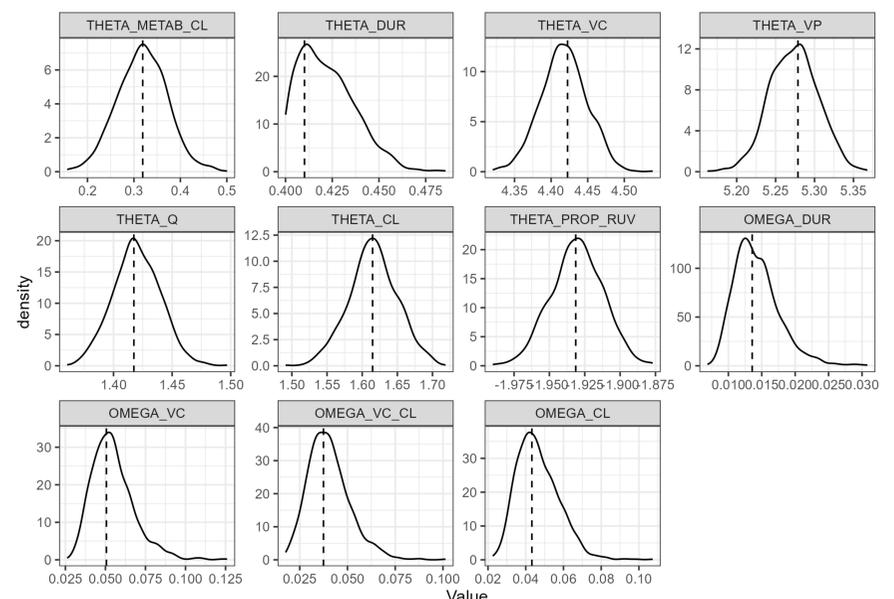
Outputs:

The output of the procedure is an object containing the original Campsis model and the set of sampled parameters, in the form of a data frame, ready for simulations with uncertainty.

EXAMPLES

```

1 library(campsis)
2
3 # Re-use an existing model from the library
4 model <- model_suite$testing$other$`2cpt_zo_all0_metab_effect_on_cl` %>%
5   add(Omega(name="VC_CL", index=2, index2=3, value=0.8, type="cor")) %>%
6   setMinMax(Theta("DUR"), min=0.40, max=Inf) # To illustrate min/max limits
7
8 # A) Sample OMEGAs/SIGMAs from the variance-covariance matrix
9 repModelA <- model %>%
10  replicate(1000, settings=AutoReplicationSettings(wishart=FALSE, checkPosDef=TRUE, checkMinMax=TRUE))
11
12 # B) Sample OMEGAs/SIGMAs from the scaled inverse Wishart distribution
13 repModelB <- model %>%
14  replicate(1000, settings=AutoReplicationSettings(wishart=TRUE, odf=40, sdf=1000, checkMinMax=TRUE))
15
16 # C) Sample parameters from an external source (e.g. CIR)
17 data <- read.csv("data.csv")
18 repModelC <- model %>%
19  replicate(1000, settings=ManualReplicationSettings(data=data))
20
21 # Export the first replicate from repModelA to a Campsis model (as an example)
22 model <- repModelA %>%
23   export(dest=CampsisModel(), index=1)
24
25 # Easy integration with Campsis to run trial simulations with replicates
26 settings <- Settings(Hardware(cpu=8, replicate_parallel=TRUE))
27 results <- progress::with_progress({
28   simulate(model=repModelA, dataset=dataset, seed=1, dest="mrgsolve", settings=settings)
29 })
30 # Show the sampled parameters using density plots (see plot below)
31 print(repModelB)
  
```



REFERENCES

- [1] <https://cran.r-project.org/web/packages/campsis/index.html>
- [2] <https://github.com/Calvagone/campsis> or www.campsis.org
- [3] <https://cran.r-project.org/web/packages/campsismod/index.html>
- [4] <https://github.com/Calvagone/campsismod>

