Getting started with RLumCarlo

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Scope

RLumCarlo is collection of energy-band models to simulate luminescence signals using Monte-Carlo (MC) methods. This document aims at providing an overview and a brief introduction to RLumCarlo.

The models in RLumCarlo

The following tables lists the models implemented in RLumCarlo along with the R function call and the corresponding R and C++ files. The modelling takes place in the C++ functions which are wrapped by the R functions with a similar name. If you, however, want to cross-check the code, you should inspect files with the ending '.cpp'.

MODEL.NAME	R.CALL	FILES
MC_CW_IRSL_DELOC	run_MC_CW_IRSL_DELOC()	R/run_MC_CW_IRSL_DELOC.R src/MC_C_MC_CW_IRSL_DELOC.cpp
MC_CW_IRSL_LOC	run_MC_CW_IRSL_LOC()	R/run_MC_CW_IRSL_LOC.R src/MC_C_MC_CW_IRSL_LOC.cpp
MC_CW_IRSL_TUN	run_MC_CW_IRSL_TUN()	R/run_MC_CW_IRSL_TUN.R src/MC_C_MC_CW_IRSL_TUN.cpp
MC_ISO_DELOC	run_MC_ISO_DELOC()	R/run_MC_ISO_DELOC.R src/MC_C_MC_ISO_DELOC.cpp
MC_ISO_LOC	run_MC_ISO_LOC()	R/run_MC_ISO_LOC.R src/MC_C_MC_ISO_LOC.cpp
MC_ISO_TUN	run_MC_ISO_TUN()	R/run_MC_ISO_TUN.R src/MC_C_MC_ISO_TUN.cpp
MC_LM_OSL_DELOC	run_MC_LM_OSL_DELOC()	R/run_MC_LM_OSL_DELOC.R src/MC_C_MC_LM_OSL_DELOC.cpp
MC_LM_OSL_LOC	$run_MC_LM_OSL_LOC()$	R/run_MC_LM_OSL_LOC.R src/MC_C_MC_LM_OSL_LOC.cpp
MC_LM_OSL_TUN	run_MC_LM_OSL_TUN()	R/run_MC_LM_OSL_TUN.R src/MC_C_MC_LM_OSL_TUN.cpp
MC_TL_DELOC	$run_MC_TL_DELOC()$	R/run_MC_TL_DELOC.R src/MC_C_MC_TL_DELOC.cpp
MC_TL_LOC	$run_MC_TL_LOC()$	R/run_MC_TL_LOC.R src/MC_C_MC_TL_LOC.cpp
MC_TL_TUN	run_MC_TL_TUN()	R/run_MC_TL_TUN.R src/MC_C_MC_TL_TUN.cpp

Each model can be run by calling one of the R functions starting with run_. Currently three different model

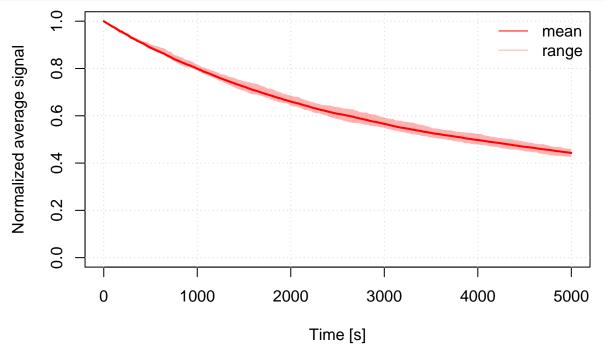
types (TUN: tunneling, LOC: localised transition, DELOC: delocalised transition) are implemented for the stimulation types TL, IRSL, LM-OSL, and ISO (isothermal). Please note that each model has different parameters and requirements.

Examples

Example 1: A first example

The first examples simulates an iso-thermal curve using the tunneling model.

```
results <- run_MC_ISO_TUN(
    E = 1.2,
    s = 1e10,
    T = 200,
    rho = 0.007,
    times = seq(0, 5000)
) %T>%
    plot_RLumCarlo(norm = TRUE, legend = TRUE)
```



The modelling output is an object of class RLumCarlo_Model_Output, which is basically a list consisting of an array and a vector.

```
str(results)
```

```
## List of 2
## $ signal: num [1:5001, 1:21, 1:10] 0 0 0 0 0 0 0 0 0 0 0 ...
## ..- attr(*, "dimnames")=List of 3
## ...$ : NULL
## ...$ : NULL
## ...$ : NULL
## $ time : int [1:5001] 0 1 2 3 4 5 6 7 8 9 ...
## - attr(*, "class")= chr "RLumCarlo_Model_Output"
## - attr(*, "model")= chr "run_MC_ISO_TUN"
```

While this represents the full modelling output results, the interpretation might be less straight forward and the user may want to condense the information via summary(). The function summary() is also used internally by the function plot_RLumCarlo().

```
df <- summary(results)</pre>
```

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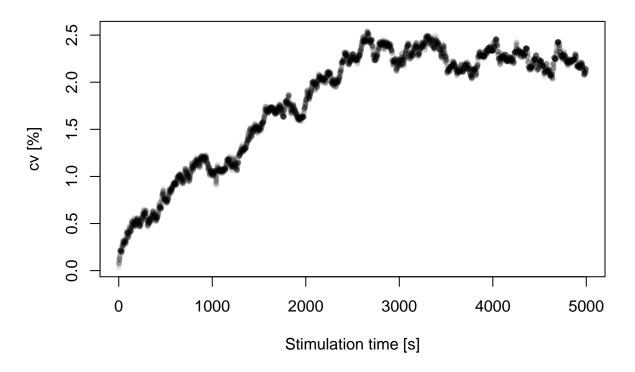
```
##
         time
                         mean
                                            y_min
                                                               y_max
##
    Min.
                0
                    Min.
                            :0.04671
                                       Min.
                                               :0.04484
                                                           Min.
                                                                  :0.04847
##
    1st Qu.:1250
                    1st Qu.:0.05409
                                       1st Qu.:0.05195
                                                           1st Qu.:0.05651
    Median:2500
                    Median: 0.06421
                                       Median :0.06174
                                                           Median : 0.06730
##
##
    Mean
            :2500
                    Mean
                            :0.06812
                                       Mean
                                               :0.06638
                                                           Mean
                                                                  :0.07044
##
    3rd Qu.:3750
                    3rd Qu.:0.08005
                                       3rd Qu.:0.07872
                                                           3rd Qu.:0.08199
##
    Max.
            :5000
                            :0.10554
                                               :0.10547
                                                                  :0.10557
                    Max.
                                       Max.
                                                           Max.
##
          sd
                               var
##
    Min.
            :3.764e-05
                                 :1.417e-09
                         Min.
    1st Qu.:1.002e-03
                         1st Qu.:1.003e-06
##
    Median :1.167e-03
                         Median :1.361e-06
##
    Mean
            :1.120e-03
                                 :1.331e-06
                         Mean
                         3rd Qu.:1.760e-06
##
    3rd Qu.:1.327e-03
##
    Max.
            :1.602e-03
                                 :2.567e-06
                         Max.
head(df)
##
     time
                mean
                         y_min
                                    y_{max}
## 1
        0 0.1055450 0.1054750 0.1055660 3.763821e-05 1.416635e-09
##
        1 0.1055209 0.1053236 0.1055660 7.840228e-05 6.146917e-09
##
  3
        2 0.1054894 0.1053236 0.1055660 8.256933e-05 6.817694e-09
## 4
        3 0.1054635 0.1053236 0.1055660 7.801698e-05 6.086649e-09
```

The call summarises the modelling results and returns a terminal output and a data.frame with, e.g., the mean or the standard deviation, which can be used to create plots for further insight. For instance, the stimulation time agains the relative standard deviation:

4 0.1054556 0.1053236 0.1055660 8.705057e-05 7.577802e-09

5 0.1054042 0.1052167 0.1055369 1.211621e-04 1.468026e-08

```
plot(
    x = df$time,
    y = (df$sd / df$mean) * 100,
    pch = 20,
    col = rgb(0,0,0,.1),
    xlab = "Stimulation time [s]",
    ylab = "cv [%]"
)
```



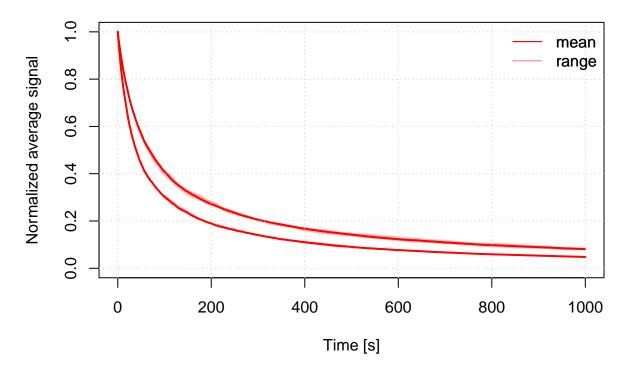
Example 2: Combining two plots

The following example uses continuous wave (CW) infrared light stimulation (IRSL), and combines two plots in one single plot window.

```
times <- seq(0, 1000)

## Run MC simulation
run_MC_CW_IRSL_TUN(A = 0.12, rho = 0.003, times = times) %>%
   plot_RLumCarlo(norm = TRUE, legend = TRUE)

run_MC_CW_IRSL_TUN(A = 0.21, rho = 0.003, times = times) %>%
   plot_RLumCarlo(norm = TRUE, add = TRUE)
```



Example 3: Testing parameters

The example above can be further extended to test the effect of different parameters. Contrary to the example above, here the results are stored in a list and plot_RLumCarlo() is called only one time.

```
s <- 3.5e12
rho <- 0.015
E <- 1.45
r_c <- c(0,0.7,0.77,0.86, 0.97)
times <- seq(100, 450) # time = temperature
results <- lapply(r_c, function(x) {
   run_MC_TL_TUN(
        s = s,
        E = E,
        rho = rho,
        r_c = x,
        times = times
)</pre>
```

NULL

