Package ‘RLumCarlo’

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BugReports https://github.com/R-Lum/RLumCarlo/issues

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**RLumCarlo-package**  

*Monte-Carlo Methods for Simulating Luminescence Phenomena.*  

---  

**Description**  

A collection of functions to simulate luminescence production in dosimetric materials using Monte-Carlo methods. Implemented are models for delocalised, localised and tunnelling transitions. Supported stimulation modes are TL, CW-OSL, LM-OSL, LM-IRSL, and ITL (ISO-TL).
Details

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References


create_ClusterSystem

Create dosimetric cluster system

Description

In order to allow interaction of an spatial a correlation clusters in RLumCarlo, first a dosimetric system needs to be created in a three-dimensional space, which is the purpose of this function.
Usage

create_ClusterSystem(n = 100, h = 0.5, plot = FALSE, ...)

Arguments

- **n**: numeric (with default): number of clusters to be created in an arbitrary 3-dimensional cube. x, y, z distances range between 0 and 1.
- **h**: numeric (with default): numeric scalar the cut the cluster tree using stats::cutree. The number must range between 0 and 1.
- **plot**: logical (with default): enables/disables plot output
- **...**: further arguments to be passed to the plot output

Value

The function returns a list of class RLumCarlo_clusters consisting of numeric vector of cluster groups and a matrix of the cluster positions in the arbitrary space. If plot = TRUE the system is displayed using scatterplot3d::scatterplot3d

Function version

0.1.0

How to cite


Author(s)

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

See Also

stats::dist, stats::hclust, stats::cutree

Examples

create_ClusterSystem(n = 10, plot = TRUE)
plot_RLumCarlo

Plot RLumCarlo Monte-Carlo Simulation Results

Description

Visualise 'RLumCarlo' modelling results without extracting the values manually. Typically visualised are the averaged signal or the number of remaining electrons, with a polygon indicating modelling uncertainties.

Usage

plot_RLumCarlo(
  object,
  plot_value = "mean",
  plot_uncertainty = "range",
  FUN = NULL,
  norm = FALSE,
  add = FALSE,
  ...
)

Arguments

object list of class RLumCarlo_Model_Output (required): input object to be plotted, usually the required input object is generated by one of the functions starting with run_. Alternatively a list of such objects can be provided.

plot_value character (with default): type of curve value to be displayed. Allowed are mean (the default) and sum (meaningful if different systems are combined). NULL disables the value visualisation.

plot_uncertainty character (with default): type of the displayed uncertainty. Allowed values are range, sd (standard deviation) and var (variance). NULL disables the uncertainty visualisation.

FUN function (optional): own function that can be applied to the y-values before normalisation and plotting

norm logical (with default): normalise curve to the highest intensity value

add logical (with default): allows overplotting of results by adding curves to an existing plot. This argument is handled automatically if object is of type list

... further argument, that can be passed to control the plot output largely following the argument names in graphics::plot.default. Currently supported are: xlab, ylab, xlim, ylim, main, lwd, type, pch, lty, col, grid, legend. The arguments lwd, type, pch, lty, col can be provided as a vector if object is a list

Details

For colouring the curves, the package khroma::khroma-package is used to provide colours that can be best distinguished, in particular by colour-blind users.
Value

This function returns a graphical output which is the visualisation of the modelling output.

Function version

0.1.0

How to cite


Author(s)

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)
Johannes Friedrich, University of Bayreuth (Germany)

Examples

```r
## plain plot
DELOC <- run_MC_TL_DELOC(
  s = 3.5e12,
  E = 1.45,
  R = 0.1,
  method = 'seq',
  clusters = 100,
  times = 150:350)
plot_RLumCarlo(legend = TRUE)

## TL with FUN to correct for thermal quenching
f <- function(x) x * 1/(1 + (2e+6 * exp(-0.55/(8.617e-5 * (DELOC$time + 273))))))
plot_RLumCarlo(object = DELOC,
               FUN = f)
```

Description

Runs a Monte-Carlo (MC) simulation of continuous wave infrared stimulated luminescence (CW-IRSL) using the generalized one trap (GOT) model. Localized transitions refer to transitions which do not involve the conduction or valence band. These transitions take place between the ground state of the trapped charge, and also involve an energy state of the recombination centre.


Usage

```
run_MC_CW_IRSL_LOC(
  A,
  times,
  clusters = 10,
  n_filled = 100,
  r,
  method = "par",
  output = "signal",
  ...
)
```

Arguments

- **A** numeric **(required)**: The optical excitation rate from the ground state of the trap to the excited state \((s^{-1})\)
- **times** numeric **(required)**: The sequence of time steps within the simulation \((s)\)
- **clusters** numeric **(with default)**: The number of created clusters for the MC runs. The input can be the output of `create_ClusterSystem`. In that case `n_filled` indicate absolute numbers of a system.
- **n_filled** integer **(with default)**: The number of filled electron traps at the beginning of the simulation \((\text{dimensionless})\). Can be a vector of \(\text{length(clusters)}\), shorter values are recycled.
- **r** numeric **(required)**: The retrapping ratio for localized transitions
- **method** character **(with default)**: Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.
- **output** character **(with default)**: output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)
- ... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

Details

The model

\[ I_{LOC}(t) = -\frac{dn}{dt} = A \times \frac{n^2}{(r + n)} \]

where in the function:
- \(A\) := optical excitation rate from the ground state into the excited state \((s^{-1})\)
- \(r\) := retrapping ratio for localized transitions
- \(t\) := time \((s)\)
- \(n\) := number of filled electron traps

Value

This function returns an object of class `RLumCarlo_Model_Output` which is a list consisting of an array with dimension `length(times) \times clusters` and a numeric time vector.
Function version

0.1.0

How to cite


Author(s)

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References


Further reading


Examples

run_MC_CW_IRSL_LOC(
  A = 0.12,
  times = 0:100,
  clusters = 50,
  n_filled = 100,
  r = 1e-7,
  method = "seq",
  output = "signal"
) %>%
plot_RLumCarlo(legend = TRUE)

---

run_MC_CW_IRSL_TUN

Run Monte-Carlo Simulation for CW-IRSL (tunnelling transitions)

Description

Runs a Monte-Carlo (MC) simulation of continuous wave infrared stimulated luminescence (CW-IRSL) using the model for tunnelling transitions. Tunnelling refers to quantum mechanical tunnelling processes from the excited state of the trap, into a recombination centre.
run_MC_CW_IRSL_TUN

Usage

run_MC_CW_IRSL_TUN(
    A,
    rho,
    times,
    clusters = 10,
    r_c = 0,
    delta.r = 0.1,
    N_e = 200,
    method = "seq",
    output = "signal",
    ...
)

Arguments

A numeric (required): The effective optical excitation rate for the tunnelling process (s^-1).

rho numeric (required): The density of recombination centres (defined as ρ’ in Huntley 2006) (dimensionless).

times numeric (required): The sequence of time steps within the simulation (s).

clusters numeric (with default): The number of created clusters for the MC runs. The input can be the output of create_ClusterSystem. In that case n_filled indicate absolute numbers of a system.

r_c numeric (with default): Critical distance (>0) that must be provided if the sample has been thermally and/or optically pretreated. This parameter expresses the fact that electron-hole pairs within a critical radius r_c have already recombined.

delta.r numeric (with default): Increments of the dimensionless distance parameter r’

N_e numeric (width default): The total number of electron traps available (dimensionless). Can be a vector of length(clusters), shorter values are recycled.

method character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.

output character (with default): output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)

... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

Details

The model

\[ I_{TU}N(r', t) = -dn/dt = A \ast \exp(-\rho'^{-1/3} \ast r') \ast n(r', t) \]

Where in the function:
A := effective optical excitation rate for the tunnelling process (s^-1)
run_MC_CW_IRSL_TUN

\[ r' := \text{the dimensionless tunnelling radius} \]
\[ \rho' := r \rho \text{ the dimensionless density of recombination centres (see Huntley (2006))} \]
\[ t := \text{time (s)} \]
\[ n := \text{the instantaneous number of electrons corresponding to the radius } r' \text{ at time } t \]

Value

This function returns an object of class RLumCarlo_Model_Output which is a list consisting of an array with dimension length(times) x length(r) x clusters and a numeric time vector.

Function version

0.2.0

How to cite


Author(s)

Johannes Friedrich, University of Bayreuth (Germany), Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References


Further reading


Examples

```r
run_MC_CW_IRSL_TUN(
  A = 0.8,
  rho = 1e-4,
  times = 0:50,
  r_c = 0.05,
  delta.r = 0.1,
)```
run_MC_CW_OSL_DELOC

```r
method = "seq",
clusters = 10,
output = "signal") %>%
plot_RLumCarlo(norm = TRUE, legend = TRUE)
```

---

**Description**

Runs a Monte-Carlo (MC) simulation of continuous wave optically stimulated luminescence (CW-OSL) using the one trap one recombination centre (OTOR) model. The term delocalized here refers to the involvement of the conduction band.

**Usage**

```r
run_MC_CW_OSL_DELOC(
  A,
  times,
  clusters = 10,
  N_e = 200,
  n_filled = N_e,
  R,
  method = "par",
  output = "signal",
  ...
)
```

**Arguments**

- **A** numeric (required): The optical excitation rate from trap to conduction band (s^-1)
- **times** numeric (required): The sequence of temperature steps within the simulation (s)
- **clusters** numeric (with default): The number of created clusters for the MC runs. The input can be the output of `create_ClusterSystem`. In that case `n_filled` indicate absolute numbers of a system.
- **N_e** integer (with default): The total number of electron traps available (dimensionless). Can be a vector of `length(clusters)`, shorter values are recycled.
- **n_filled** integer (with default): The number of filled electron traps at the beginning of the simulation (dimensionless). Can be a vector of `length(clusters)`, shorter values are recycled.
- **R** numeric (required): The retrapping ratio for delocalized transitions (dimensionless)
method character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.

output character (with default): Output is either the 'signal' (the default) or 'remaining_e' (the remaining charges, electrons, in the trap)

... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

Details

The model

\[ I_{DELOC}(t) = -dn/dt = A \times \left( n^2 / (N \times R + n(1 - R)) \right) \]

Where in the function:
\( t := \text{time (s)} \)
\( A := \text{the optical excitation rate from trap to conduction band (1/s)} \)
\( n := n_{\text{filled}}, \text{the instantaneous number of electrons} \)
\( N := N_{e} \text{ the available number of electron traps available} \)
\( R := \text{retrapping ratio for delocalized transitions} \)

Value

This function returns an object of class RLumCarlo_Model_Output which is a list consisting of an array with dimension length(times) x clusters and a numeric time vector.

Function version

0.1.0

How to cite


Author(s)

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References


Further reading

Examples

## brief example
run_MC_CW_OSL_DELOC(
    A = 0.12,
    R = 0.1,
    times = 0:10,
    clusters = 10,
    method = "seq")
plot_RLumCarlo(legend = TRUE)

## A long example
## Not run:
A <- c(0.1, 0.3, 0.5, 1)
times <- seq(0, 60, 1)
s <- 1e12
E <- 1
R <- c(1e-7, 1e-6, 0.01, 0.1) # sequence of different R values
clusters <- 1000 # number of Monte Carlo simulations
N_e <- c(200, 500, 700, 400) # number of free electrons
n_filled <- c(200, 500, 100, 70) # number of filled traps
method <- "par"
output <- "signal"
col <- c(1, 2, 3, 4) # different colours for the individual curves
plot_uncertainty <- c(TRUE, FALSE, TRUE, FALSE) # do you want to see the uncertainty?
add_TF <- c(FALSE, rep(TRUE, (length(R)-1)))

## loop to plot different curves into one plot
for (u in 1:length(R)){
    results <- run_MC_CW_OSL_DELOC(
        A = A[u],
        times,
        clusters = clusters,
        N_e = N_e[u],
        n_filled = n_filled[u],
        R = R[u],
        method = method,
        output = output)
    plot_RLumCarlo(
        results,
        add = add_TF[u],
        legend = FALSE,
        col = col[u],
        main = "Delocalised Transition")
}
# add your legend with your parameters
legend("topright",
    ncol = 4,
    cex = 0.55,
    title = "parameters",
    legend=c(
        paste0("A = ", A),
        ...)
Description

Runs a Monte-Carlo (MC) simulation of isothermally stimulated luminescence (ISO-TL or ITL) using the one trap one recombination centre (OTOR) model. Delocalised refers to involvement of the conduction band.

Usage

```r
run_MC_ISO_DELOC(
  s,
  E,
  T = 20,
  times,
  clusters = 10,
  N_e = 200,
  n_filled = N_e,
  R,
  method = "par",
  output = "signal",
  ...
)
```

Arguments

- `s` numeric (required): The frequency factor of the trap (s^-1)
- `E` numeric (required): Thermal activation energy of the trap (eV)
- `T` numeric (with default): Constant stimulation temperature (°C)
- `times` numeric (with default): The sequence of time steps within the simulation (s)
- `clusters` numeric (with default): The number of created clusters for the MC runs. The input can be the output of `create_ClusterSystem`. In that case `n_filled` indicate absolute numbers of a system.
- `N_e` integer (with default): The total number of electron traps available (dimensionless). Can be a vector of `length(clusters)`, shorter values are recycled.
n_filled \hspace{10pt} \textbf{integer} (with default): The number of filled electron traps at the beginning of the simulation (dimensionless). Can be a vector of length(clusters), shorter values are recycled.

\( R \hspace{10pt} \textbf{numeric (required)}: \) The delocalized retrapping ratio (dimensionless)

\textbf{method} \hspace{10pt} \textbf{character} (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.

\textbf{output} \hspace{10pt} \textbf{character} (with default): Output is either the 'signal' (the default) or 'remaining_e' (the remaining charges, electrons, in the trap)

... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

\textbf{Details}

\textbf{The model}

\[ I_{DELOC}(t) = -\frac{dn}{dt} = (s \times \exp(-E/(k_B \times T_{ISO}))) \times \left(\frac{n^2}{N \times R + n(1 - R)}\right) \]

Where in the function:

\( t := \text{time} \)

\( k_B := \text{Boltzmann constant} \ (8.617 \times 10^{-5} \text{ eV K}^{-1}) \)

\( T_{ISO} := \text{temperature of the isothermal experiment (°C)} \)

\( n := n_{\text{filled}}, \text{the number of filled electron traps at the beginning of the simulation} \)

\( E := \text{the trap depth (eV)} \)

\( s := \text{the frequency factor in (s}^{-1})\)

\( N := N_e, \text{the total number of electron traps available (dimensionless)} \)

\( R := \text{the retrapping ratio for delocalized transitions} \)

\textbf{Value}

This function returns an object of class \texttt{RLumCarlo_Model_Output} which is a list consisting of an array with dimension length(times) x clusters and a \textbf{numeric} time vector.

\textbf{Function version}

0.1.0

\textbf{How to cite}


\textbf{Author(s)}

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)
References


Further reading


Examples

```r
run_MC_ISO_DELOC(
  s = 3.5e12,
  E = 1.45,
  T = 200,
  R = 1,
  method = 'seq',
  times = 0:100) %>%
plot_RLumCarlo(legend = TRUE)
```

---

**run_MC_ISO_LOC**

*Run Monte-Carlo simulation for ISO-TL (localized transitions)*

**Description**

Runs a Monte-Carlo (MC) simulation of isothermally stimulated luminescence (ISO-TL or ITL) using the generalized one trap (GOT) model. Localized transitions refer to transitions which do no involve the conduction or valence band. These transitions take place between the ground state and an excited state of the trapped charge, and also involve an energy state of the recombination centre.

**Usage**

```r
run_MC_ISO_LOC(
  s,
  E,
  T = 20,
  times,
  clusters = 10,
  n_filled = 100,
  r,
  method = "par",
  output = "signal",
  ...
)
```
Arguments

$s$ numeric (required): The frequency factor of the trap (s^{-1})
$E$ numeric (required): Thermal activation energy of the trap (eV)
$T$ numeric (with default): Constant stimulation temperature (°C)
$\text{times}$ numeric (with default): The sequence of time steps within the simulation (s)
$\text{clusters}$ numeric (with default): The number of created clusters for the MC runs. The input can be the output of create_ClusterSystem. In that case n_filled indicate absolute numbers of a system.
$n\_\text{filled}$ integer (with default): The number of filled electron traps at the beginning of the simulation (dimensionless). Can be a vector of length(clusters), shorter values are recycled.
$r$ numeric (required): The retrapping ratio for localized transitions.
$\text{method}$ character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.
$output$ character (with default): output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)
...

Details

The model

\[ I_{\text{LOC}}(t) = -\frac{dn}{dt} = (s \cdot \exp(-E/(k_B \cdot T_{\text{ISO}}))) \cdot (n^2/(r + n)) \]

Where in the function:
$t :=$ time (s)
$k_B :=$ Boltzmann constant (8.617 x 10^{-5} eV K^{-1})
$T_{\text{ISO}} :=$ isothermal temperature (°C)
$n := n\_\text{filled}$
$s :=$ frequency factor of the trap (1/s)
$E :=$ activation energy of the trap (eV)
$r :=$ retrapping ratio for localized transitions

Value

This function returns an object of class RLumCarlo_Model_Output which is a list consisting of an array with dimension length(times) x clusters and a numeric time vector.

Function version

0.1.0
**How to cite**


**Author(s)**

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

**References**


**Examples**

```r
run_MC_ISO_LOC(
  E = 1.45,
  s = 3.5e12,
  T = 200,
  times = 0:100,
  method = 'seq',
  r = 1)
plot_RLumCarlo(legend = TRUE)
```

**Description**

Runs a Monte-Carlo (MC) simulation of isothermally stimulated luminescence (ISO-TL or ITL) using the tunnelling (TUN) model. Tunnelling refers to quantum mechanical tunnelling processes from the excited state of the trapped charge, into the recombination centre.

**Usage**

```r
run_MC_ISO_TUN(  
  E,  
  s,  
  T = 200,  
  rho,  
  times,  
  clusters = 10,  
  r_c = 0,  
)```
delta.r = 0.1,
N_e = 200,
method = "par",
output = "signal",
...)

Arguments

- **E** numeric (required): Thermal activation energy of the trap (eV).
- **s** numeric (required): The effective frequency factor for the tunnelling process (s^-1).
- **T** numeric (with default): Constant stimulation temperature (°C).
- **rho** numeric (required): The dimensionless density of recombination centres (defined as \( \rho' \) in Huntley 2006) (dimensionless).
- **times** numeric (required): The sequence of time steps within the simulation (s).
- **clusters** numeric (with default): The number of created clusters for the MC runs. The input can be the output of `create_ClusterSystem`. In that case `n_filled` indicate absolute numbers of a system.
- **r_c** numeric (with default): Critical distance (>0) that must be provided if the sample has been thermally and/or optically pretreated. This parameter expresses the fact that electron-hole pairs within a critical radius \( r_c \) have already recombined.
- **delta.r** numeric (with default): Fractional change of the dimensionless distance of nearest recombination centres (\( r' \))
- **N_e** numeric (with default): The total number of electron traps available (dimensionless). Can be a vector of length(\( \text{clusters} \)), shorter values are recycled.
- **method** character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.
- **output** character (with default): output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)

... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

Details

The model

\[
I_{TUN}(r', t) = -\frac{dn}{dt} = (s \cdot \exp\left(-\frac{E}{(k_B \cdot T_{ISO})}\right)) \cdot \exp\left(-\left(\rho'\right)^{-1/3} \cdot r'\right) \cdot n(r', t)
\]

Where in the function:

- **E** := thermal activation energy (eV)
- **s** := the effective frequency factor for the tunnelling process (s^-1)
- **T_{ISO}** := the temperature of the isothermal experiment (°C)
- **k_B** := Boltzmann constant (8.617 x 10^-5 eV K^-1)
\( r' := \) the dimensionless tunnelling radius \\
\( \rho' := r \rho_0 \) the dimensionless density of recombination centres see Huntley (2006) \\
\( t := \) time (s) \\
\( n := \) the instantaneous number of electrons corresponding to the radius \( r' \)

**Value**

This function returns an object of class \texttt{RLumCarlo_Model_Output} which is a list consisting of an array with dimension length(times) x length(r) x clusters and a numeric time vector.

**Function version**

0.1.0

**How to cite**


**Author(s)**

Johannes Friedrich, University of Bayreuth (Germany), Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

**References**


**Examples**

```r
## short example
run_MC_ISO_TUN(
  E = .8,
  s = 1e16,
  T = 50,
```

---

**run_MC_ISO_TUN**
run_MC_LM_OSL_DELOC

Run Monte-Carlo Simulation for LM-OSL (delocalized transitions)

Description

Runs a Monte-Carlo (MC) simulation of linearly modulated optically stimulated luminescence (LM-OSL) using the one trap one recombination centre (OTOR) model. Delocalised refers to involvement of the conduction band.

Usage

run_MC_LM_OSL_DELOC(
  A,
  times,
  clusters = 10,
  N.e = 200,
  n_filled = N.e,
  R,
  method = "par",
  output = "signal",
  ...)

## Not run:
## long (meaningful) example
results <- run_MC_ISO_TUN(
  E = 0.8,
  s = 1e16,
  T = 50,
  rho = 1e-4,
  times = 0:100,
  clusters = 1000,
  N.e = 200,
  r_c = 0.1,
  delta.r = 0.05,
  method = "par")
plot_RLumCarlo(results, legend = TRUE)

## End(Not run)
**Arguments**

- **A**  
  *numeric (required)*: The optical excitation rate from trap to conduction band (s^-1)

- **times**  
  *numeric (required)*: The sequence of time steps within the simulation (s)

- **clusters**  
  *numeric (with default)*: The number of created clusters for the MC runs. The input can be the output of `create_ClusterSystem`. In that case `n_filled` indicate absolute numbers of a system.

- **N_e**  
  *integer (with default)*: The total number of electron traps available (dimensionless). Can be a vector of length(clusters), shorter values are recycled.

- **n_filled**  
  *integer (with default)*: The number of filled electron traps at the beginning of the simulation (dimensionless). Can be a vector of length(clusters), shorter values are recycled.

- **R**  
  *numeric (required)*: The retrapping ratio for delocalized transitions

- **method**  
  *character (with default)*: Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.

- **output**  
  *character (with default)*: output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)

- ...  
  further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

**Details**

**The model**

\[
I_{DELOC}(t) = -\frac{dn}{dt} = A \cdot t / P \cdot \frac{n \cdot R + n(1 - R)}{N}
\]

Where in the function:
- \( t := \text{time (s)} \)
- \( A := \text{the optical excitation rate from trap to conduction band (1/s)} \)
- \( n := \text{n_filled, the instantaneous number of electrons} \)
- \( R := \text{the retrapping ratio for delocalized transitions} \)
- \( N := \text{N_e, the total number of electron traps available (dimensionless)} \)
- \( P := \text{total stimulation time (s)} \)

**Value**

This function returns an object of class `RLumCarlo_Model_Output` which is a list consisting of an array with dimension length(times) x clusters and a numeric time vector.

**Function version**

0.1.0
run_MC_LM_OSL_LOC

How to cite


Author(s)

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References


Further reading


Examples

```r
run_MC_LM_OSL_DELOC(
  A = 0.12,
  R = 0.1,
  times = 0:50,
  method = "seq",
  clusters = 10) %>%
plot_RLumCarlo(legend = TRUE)
```

---

run_MC_LM_OSL_LOC

*Run Monte-Carlo Simulation for LM-OSL (localized transitions)*

Description

Runs a Monte-Carlo (MC) simulation of linearly modulated optically stimulated luminescence (LM-OSL) using the generalized one trap (GOT) model. Localized transitions refer to transitions which do not involve the conduction or valence band. These transitions take place between the ground state and an excited state of the trap, and also involve a an energy state of the recombination centre.
run_MC_LM_OSL_LOC

Usage

run_MC_LM_OSL_LOC(
    A,
    times,
    clusters = 10,
    n_filled = 100,
    r,
    method = "par",
    output = "signal",
    ...
)

Arguments

A numeric (required): The optical excitation rate from the ground state into the excited state of the trap (s\(^{-1}\))

times numeric (required): The sequence of time steps within the simulation (s)

clusters numeric (with default): The number of created clusters for the MC runs. The input can be the output of create_ClusterSystem. In that case n_filled indicate absolute numbers of a system.

n_filled integer (with default): The number of filled electron traps at the beginning of the simulation (dimensionless). Can be a vector of length(clusters), shorter values are recycled.

r numeric (required): The retrapping ratio for localized transitions

method character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.

output character (with default): output is either the 'signal' (the default) or 'remaining_e' (the remaining charges, electrons, in the trap)

... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

Details

The model

\[ I_{LOC}(t) = -\frac{dn}{dt} = (A \times t/P) \times (n^2/(r + n)) \]

Where in the function:

A := optical excitation rate from the ground state into the excited state of the trap (1/s)

P := total excitation time (s)

t := time (s)

n := n_filled, the instantaneous number of electrons

r := the retrapping ratio for localized transitions
run_MC_LM_OSL_LOC

Value

This function returns an object of class RLumCarlo_Model_Output which is a list consisting of an array with dimension length(times) x clusters and a numeric time vector.

Function version

0.1.0

How to cite


Author(s)

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References


Examples

## short example
run_MC_LM_OSL_LOC(
  A = 1,
  times = 0:40,
  clusters = 10,
  n_filled = 100,
  r = 1e-7,
  method = "seq",
  output = "signal") %>%
plot_RLumCarlo(legend = TRUE)

## Not run:
## the long (meaningful) example
results <- run_MC_LM_OSL_LOC(
  A = 1,
  times = 0:100,
  clusters = 100,
  n_filled = 100,
  r = 1e-7,
  method = "par",
  output = "signal")

## plot
run_MC_LM_OSL_TUN Run Monte-Carlo Simulation for LM-OSL (tunnelling transitions)

Description

Runs a Monte-Carlo (MC) simulation of linearly modulated optically stimulated luminescence (LM-OSL) using the tunnelling (TUN) model. Tunnelling refers to quantum mechanical tunnelling processes from the excited state of the trapped charge, into a recombination centre.

Usage

run_MC_LM_OSL_TUN(
  A,
  rho,
  times,
  clusters = 10,
  r_c = 0,
  delta.r = 0.1,
  N_e = 200,
  method = "par",
  output = "signal",
  ...
)

Arguments

A numeric (required): The effective optical excitation rate for the tunnelling process
rho numeric (required): The dimensionless density of recombination centres (defined as $\rho'$ in Huntley 2006) (dimensionless)
times numeric (required): The sequence of time steps within the simulation (s)
clusters numeric (with default): The number of MC runs
r_c numeric (with default): Critical distance (>0) that is to be used if the sample has been thermally and/or optically pretreated. This parameter expresses the fact that electron-hole pairs within a critical radius $r_c$ have already been recombined.
delta.r numeric (with default): Increments of dimensionless distance $r'$
N_e numeric (width default): The total number of electron traps available (dimensionless). Can be a vector of length(clusters), shorter values are recycled.
method character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.

output character (with default): output is either the 'signal' (the default) or 'remaining_e' (the remaining charges, electrons, in the trap)

... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

Details

The model

$$I_{TUN}(r', t) = -\frac{dn}{dt} = (A \ast t / P) \ast exp\left(-\left(\rho'\right)^{-1/3} * r'\right) \ast n(r', t)$$

Where in the function:
A := the optical excitation rate for the tunnelling process (s^-1)
t := time (s)
P := maximum stimulation time (s)
r' := the dimensionless tunnelling radius
\(\rho' := \rho_0\) the dimensionless density of recombination centres see Huntley (2006)
n := the instantaneous number of electrons corresponding to the radius r'

Value

This function returns an object of class RLumCarlo_Model_Output which is a list consisting of an array with dimension length(times) x length(r) x clusters and a numeric time vector.

Function version

0.1.0

How to cite


Author(s)

Johannes Friedrich, University of Bayreuth (Germany), Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References


Examples

```r
## the short example
run_MC_LM_OSL_TUN(
  A = 1,
  rho = 1e-3,
  times = 0:100,
  clusters = 10,
  N_e = 100,
  r_c = 0.1,
  delta.r = 1e-1,
  method = "seq",
  output = "signal") %>%
  plot_RLumCarlo(norm = TRUE)

## Not run:
## the long (meaningful) example
results <- run_MC_LM_OSL_TUN(
  A = 1,
  rho = 1e-3,
  times = 0:1000,
  clusters = 30,
  N_e = 100,
  r_c = 0.1,
  delta.r = 1e-1,
  method = "par",
  output = "signal")

plot_RLumCarlo(results, norm = TRUE)
## End(Not run)
```

Run Monte-Carlo Simulation for TL (delocalized transitions)
**Description**

Runs a Monte-Carlo (MC) simulation of thermoluminescence (TL) using the one trap one recombination centre (OTOR) model. Delocalised refers to involvement of the conduction band. The heating rate in this function is assumed to be 1 K/s.

**Usage**

```r
run_MC_TL_DELOC(
  s,
  E,
  times,
  b = 1,
  clusters = 10,
  N_e = 200,
  n_filled = N_e,
  R = 1,
  method = "par",
  output = "signal",
  ...
)
```

**Arguments**

- `s` **numeric (required)**: The frequency factor of the trap (\(s^{-1}\))
- `E` **numeric (required)**: Thermal activation energy of the trap (eV)
- `times` **numeric (required)**: The sequence of temperature steps within the simulation (s). The default heating rate is set to 1 K/s. The final temperature is \(\max(times) \times b\)
- `b` **numeric (with default)**: the heating rate in K/s
- `clusters` **numeric (with default)**: The number of created clusters for the MC runs. The input can be the output of `create_ClusterSystem`. In that case `n_filled` indicate absolute numbers of a system.
- `N_e` **integer (with default)**: The total number of electron traps available (dimensionless). Can be a vector of `length(clusters)`, shorter values are recycled.
- `n_filled` **integer (with default)**: The number of filled electron traps at the beginning of the simulation (dimensionless). Can be a vector of `length(clusters)`, shorter values are recycled.
- `R` **numeric (with default)**: Re-trapping ratio for delocalized transitions
- `method` **character (with default)**: Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.
- `output` **character (with default)**: output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)
- `...` further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal
Details

The model

\[ \dot{I}_{DELOC}(t) = -\frac{dn}{dt} = \left(s \times \exp\left(-\frac{E}{(k_B \times T)}\right)\right) \times \left(\frac{n^2}{(N \times R + n(1 - R))}\right) \]

Where in the function:
- \( E \) := the thermal activation energy (eV)
- \( s \) := the frequency factor in \((s^{-1})\)
- \( t \) := time (s)
- \( k_B \) := Boltzmann constant \((8.617 \times 10^{-5} \text{ eV K}^{-1})\)
- \( T \) := temperature \((°C)\)
- \( R \) := Delocalised retrapping ratio
- \( n := n_{\text{filled}} \), the instantaneous number of electrons
- \( N := N_{\text{e}} \), the total number of electron traps available (dimensionless)

Why times and \( b \) instead of temperature?

The parameter to control the temperature is a function of the stimulation times (the parameter \( times \)) and the heating rate (\( b \)). Thus, the final temperature is \( \text{max(times)} \times b \). For a heating rate (\( b = 1 \)) the final temperature is \( \text{max(times)} \). While this might be a little bit confusing, it also allows you to control the time resolution of the simulation, i.e. you can simulate more points per second.

Value

This function returns an object of class \texttt{RLumCarlo_Model_Output} which is a list consisting of an array with dimension \( \text{length(times)} \times \text{clusters} \) and a numeric time vector.

Function version

0.1.0

How to cite


Author(s)

Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References

Further reading

Examples

```r
## the short example
run_MC_TL_DELOC(
  s = 3.5e12,
  E = 1.45,
  R = 0.1,
  method = 'seq',
  clusters = 100,
  times = 150:350)
plot_RLumCarlo(legend = TRUE)

## Not run:
## the long (meaningful) example
# define your parameters
times <- seq(100, 450, 1)
s <- rep(3.5e12, 4)
E <- rep(1.45, 4)
R <- c(0.7e-6, 1e-6, 0.01, 0.1)
clusters <- 300
N_e <- c(400, 500, 700, 400)
n_filled <- c(400, 500, 300, 70)
method <- "par"
output <- "signal"
col <- c(1, 2, 3, 4) # different colours for the individual curves
plot_uncertainty <- c(TRUE, TRUE, TRUE, TRUE) # do you want to see the uncertainty?
add_TF <- c(FALSE, rep(TRUE, (length(R) - 1)))

# loop to plot different curves into one plot
for (u in 1:length(R)){
  results <- run_MC_TL_DELOC(
    times=times,
    s = s[u],
    E = E[u],
    clusters = clusters,
    N_e = N_e[u],
    n_filled = n_filled[u],
    R = R[u],
    method = method,
    output = output)
  plot_RLumCarlo(results,
  add = add_TF[u],
  legend = FALSE,
  col=col[u],
  main = " your plot",
  ylim=c(0,20))
```

run_MC_TL_LOC

Run Monte-Carlo Simulation for TL (localized transitions)

Description

Runs a Monte-Carlo (MC) simulation of thermoluminescence (TL) using the generalized one trap (GOT) model. Localized transitions refer to transitions which do not involve the conduction or valence band. These transitions take place between the ground state and an excited state of the trapped charge, and also involve an energy state of the recombination centre. The heating rate in this function is assumed to be 1 K/s.

Usage

run_MC_TL_LOC(s, E, times, b = 1, clusters = 10, n_filled = 100, r, method = "par", output = "signal", ...)

Arguments

s numeric (required): The frequency factor of the trap (s^-1)
E numeric (required): Thermal activation energy of the trap (eV)
times numeric (required): The sequence of temperature steps within the simulation (s). The default heating rate is set to 1 K/s. The final temperature is max(times) * b

b numeric (with default): The heating rate in K/s

clusters numeric (with default): The number of created clusters for the MC runs. The input can be the output of create_ClusterSystem. In that case n_filled indicate absolute numbers of a system.

n_filled integer (with default): The number of filled electron traps at the beginning of the simulation (dimensionless). Can be a vector of length(clusters), shorter values are recycled.

r numeric (required): The localized retrapping ratio (dimensionless)

method character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.

output character (with default): output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)

... further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal

Details

The model

$$I_{LOC}(t) = -\frac{dn}{dt} = (s \exp(-E/(k_B * T))) * (\frac{n^2}{r+n})$$

Where in the function:
E := the thermal activation energy (eV)
s := the frequency factor for the trap (s^-1)
t := time (s)
k_B := Boltzmann constant (8.617 x 10^-5 eV K^-1)
T := temperature (°C)
n := the instantaneous number of electrons
r := the retrapping ratio for localized transitions

Value

This function returns an object of class RLumCarlo_Model_Output which is a list consisting of an array with dimension length(times) x clusters and a numeric time vector.

Function version

0.1.0

How to cite

run_MC_TL_TUN

Author(s)
Sebastian Kreutzer, Institute of Geography, Heidelberg University (Germany)

References

Examples
## the short example
run_MC_TL_LOC(
  s = 1e14,
  E = 0.9,
  times = 50:100,
  b = 1,
  method = "seq",
  clusters = 30,
  r = 1)
plot_RLumCarlo()

## Not run:
## the long (meaningful) example
results <- run_MC_TL_LOC(
  s = 1e14,
  E = 0.9,
  times = 50:100,
  method = "par",
  clusters = 100,
  r = 1)

## plot
plot_RLumCarlo(results)

## End(Not run)

run_MC_TL_TUN

Run Monte-Carlo Simulation for TL (tunnelling transitions)

Description
Runs a Monte-Carlo (MC) simulation of thermoluminescence (TL) caused by tunnelling (TUN) transitions. Tunnelling refers to quantum mechanical tunnelling processes from the excited state of the trap into a recombination centre. The heating rate in this function is assumed to be 1 K/s.
run_MC_TL_TUN

Usage

run_MC_TL_TUN(s, E, rho, r_c = 0, times, b = 1, clusters = 10, N_e = 200, delta.r = 0.1, method = "par", output = "signal", ...
)

Arguments

- `s` list (required): The effective frequency factor for the tunnelling process (s^-1)
- `E` numeric (required): Thermal activation energy of the trap (eV)
- `rho` numeric (required): The dimensionless density of recombination centres (defined as $\rho$' in Huntley 2006)
- `r_c` numeric (with default): Critical distance (>0) that is to be used if the sample has been thermally and/or optically pretreated. This parameter expresses the fact that electron-hole pairs within a critical radius $r_c$ have already recombined.
- `times` numeric (required): The sequence of temperature steps within the simulation (s). The default heating rate is set to 1 K/s. The final temperature is $\max(times) \times b$
- `b` numeric (with default): the heating rate in K/s
- `clusters` numeric (with default): The number of created clusters for the MC runs. The input can be the output of create_ClusterSystem. In that case $n_{filled}$ indicate absolute numbers of a system.
- `N_e` numeric (with default): The total number of electron traps available (dimensionless). Can be a vector of length(clusters), shorter values are recycled.
- `delta.r` numeric (with default): The increments of the dimensionless distance $r'$
- `method` character (with default): Sequential 'seq' or parallel 'par' processing. In the parallel mode the function tries to run the simulation on multiple CPU cores (if available) with a positive effect on the computation time.
- `output` character (with default): output is either the 'signal' (the default) or 'remaining_e' (the remaining charges/electrons in the trap)
- `...` further arguments, such as cores to control the number of used CPU cores or verbose to silence the terminal
Details

The model

\[ I_{TU}(r', t) = -\frac{dn}{dt} = (s * \exp(-E/(k_B * T))) * \exp(-(\rho')^{-1/3} * r') * n(r', t) \]

Where in the function:
- \( s := \) frequency for the tunnelling process (s^-1)
- \( E := \) thermal activation energy (eV)
- \( k_B := \) Boltzmann constant (8.617 x 10^-5 eV K^-1)
- \( T := \) temperature (°C)
- \( r' := \) the dimensionless tunnelling radius
- \( \rho' := \rho \text{'}, the dimensionless density of recombination centres (see Huntley (2006))
- \( t := \) time (s)
- \( n := \) the instantaneous number of electrons at distance \( r' \)

Value

This function returns an object of class `RLumCarlo_Model_Output` which is a list consisting of an array with dimension length(times) x length(r) x clusters and a numeric time vector.

Function version

0.1.0

How to cite


Author(s)

Johannes Friedrich, University of Bayreuth (Germany), Sebastian Kreutzer, Geography & Earth Sciences, Aberystwyth University (United Kingdom)

References


Further reading

Examples

```r
## the short example
run_MC_TL_TUN(
  s = 1e12,
  E = 0.9,
  rho = 1,
  r_c = 0.1,
  times = 80:120,
  b = 1,
  clusters = 50,
  method = 'seq',
  delta.r = 1e-1) %>%
plot_RLumCarlo()

## Not run:
## the long (meaningful example)
results <- run_MC_TL_TUN(
  s = 1e12,
  E = 0.9,
  rho = 0.01,
  r_c = 0.1,
  times = 80:220,
  clusters = 100,
  method = 'par',
  delta.r = 1e-1)

## plot
plot_RLumCarlo(results)

## End(Not run)
```
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