

# pIRIR analysis B19-LU 6

## Table of contents

|          |  |           |
|----------|--|-----------|
| <b>1</b> | <b>Data structure</b>                              | <b>2</b>  |
| 1.1      | Sample set . . . . .                               | 2         |
| 1.2      | Measurement sequence . . . . .                     | 3         |
| 1.3      | Measurement plots . . . . .                        | 6         |
| <b>2</b> | <b>Data preparation</b>                            | <b>7</b>  |
| 2.1      | Data cleaning . . . . .                            | 7         |
| 2.2      | Dose rate calculation . . . . .                    | 8         |
| 2.3      | Corrected data set . . . . .                       | 8         |
| <b>3</b> | <b>Dose calculation</b>                            | <b>10</b> |
| 3.1      | De(t) plots . . . . .                              | 10        |
| 3.2      | Dose-response curve fitting optimization . . . . . | 11        |
| 3.3      | De calculation result table . . . . .              | 12        |
| 3.4      | Rejection criteria . . . . .                       | 13        |
| <b>4</b> | <b>Dose distribution</b>                           | <b>15</b> |
| 4.1      | Distribution plots . . . . .                       | 15        |
| 4.2      | Central age model . . . . .                        | 17        |
| <b>5</b> | <b>References</b>                                  | <b>18</b> |

**Analysed file:** 3-Permafrost-1mm-63-100-Fsp-pIR290-sample\_B19-LU-6-24al.binx

**Date of analysis:** 01 Jan 2023

This report describes the data analysis work flow for the pIRIR measurement of the sample *B19-LU 6*, measured by Steffi Hesse and Dr. Tobias Lauer. The data analysis was performed by [Dirk Mittelstraß](#) on behalf of Dr. Margret Fuchs from HZDR Innovation GmbH and paid for by the HZDR Innovation GmbH.

The data was processed using the open-source statistical programming language [R](#) and multiple open-source function libraries (called ‘packages’), especially the R package [Luminescence](#) by Kreutzer et al. (2012, 2022). This PDF was formatted and created with [Quarto](#).

# 1 Data structure

## 1.1 Sample set

First we prove if all aliquots are of the same sample and the same run by looking at the Risö Reader record parameters.

Table 1: Structure of the BIN file content

| index | SAMPLE   | DATE   | RUN | SET | POSITION | records |
|-------|----------|--------|-----|-----|----------|---------|
| 1     | Sample1  | 141022 | 2   | 1   | 1        | 48      |
| 2     | Sample3  | 141022 | 2   | 1   | 3        | 48      |
| 3     | Sample5  | 141022 | 2   | 1   | 5        | 48      |
| 4     | Sample7  | 141022 | 2   | 1   | 7        | 48      |
| 5     | Sample9  | 141022 | 2   | 1   | 9        | 48      |
| 6     | Sample11 | 141022 | 2   | 1   | 11       | 48      |
| 7     | Sample13 | 141022 | 2   | 1   | 13       | 48      |
| 8     | Sample15 | 141022 | 2   | 1   | 15       | 48      |
| 9     | Sample17 | 141022 | 2   | 1   | 17       | 48      |
| 10    | Sample19 | 141022 | 2   | 1   | 19       | 48      |
| 11    | Sample21 | 141022 | 2   | 1   | 21       | 48      |
| 12    | Sample23 | 141022 | 2   | 1   | 23       | 48      |
| 13    | Sample25 | 141022 | 2   | 1   | 25       | 48      |
| 14    | Sample27 | 141022 | 2   | 1   | 27       | 48      |
| 15    | Sample29 | 141022 | 2   | 1   | 29       | 48      |
| 16    | Sample31 | 141022 | 2   | 1   | 31       | 48      |
| 17    | Sample33 | 141022 | 2   | 1   | 33       | 48      |
| 18    | Sample35 | 141022 | 2   | 1   | 35       | 48      |
| 19    | Sample37 | 141022 | 2   | 1   | 37       | 48      |
| 20    | Sample39 | 151022 | 2   | 1   | 39       | 48      |
| 21    | Sample41 | 151022 | 2   | 1   | 41       | 48      |
| 22    | Sample43 | 151022 | 2   | 1   | 43       | 48      |
| 23    | Sample45 | 151022 | 2   | 1   | 45       | 48      |
| 24    | Sample47 | 151022 | 2   | 1   | 47       | 48      |

24 aliquots were processed with the same sequence and were of the same measurement run.

## 1.2 Measurement sequence

Table 2: Measurement sequence structure for aliquot 1

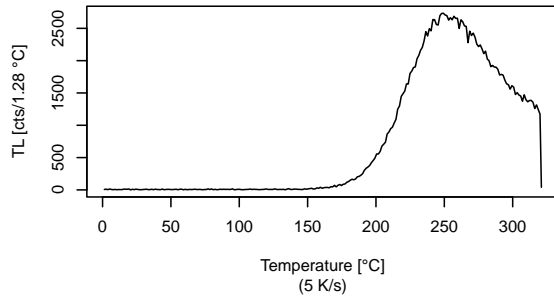
| #  | Type | max. Temp. [°C] | Irr. time [s] | Channels | Channel width [s] |
|----|------|-----------------|---------------|----------|-------------------|
| 1  | TL   | 321             | 0             | 250      | 1.285             |
| 2  | IRSL | 50              | 0             | 220      | 0.500             |
| 3  | IRSL | 290             | 0             | 420      | 0.500             |
| 4  | TL   | 0               | 1000          | 250      | 1.285             |
| 5  | IRSL | 50              | 1000          | 220      | 0.500             |
| 6  | IRSL | 290             | 1000          | 420      | 0.500             |
| 7  | IRSL | 325             | 1000          | 420      | 0.238             |
| 8  | TL   | 321             | 0             | 250      | 1.285             |
| 9  | IRSL | 50              | 0             | 220      | 0.500             |
| 10 | IRSL | 290             | 0             | 420      | 0.500             |
| 11 | TL   | 0               | 1000          | 250      | 1.285             |
| 12 | IRSL | 50              | 1000          | 220      | 0.500             |
| 13 | IRSL | 290             | 1000          | 420      | 0.500             |
| 14 | IRSL | 325             | 1000          | 420      | 0.238             |
| 15 | TL   | 0               | 1000          | 250      | 1.285             |
| 16 | IRSL | 50              | 1000          | 220      | 0.500             |
| 17 | IRSL | 290             | 1000          | 420      | 0.500             |
| 18 | TL   | 0               | 1000          | 250      | 1.285             |
| 19 | IRSL | 50              | 1000          | 220      | 0.500             |
| 20 | IRSL | 290             | 1000          | 420      | 0.500             |
| 21 | IRSL | 325             | 1000          | 420      | 0.238             |
| 22 | TL   | 0               | 2000          | 250      | 1.285             |
| 23 | IRSL | 50              | 2000          | 220      | 0.500             |
| 24 | IRSL | 290             | 2000          | 420      | 0.500             |
| 25 | TL   | 0               | 1000          | 250      | 1.285             |
| 26 | IRSL | 50              | 1000          | 220      | 0.500             |
| 27 | IRSL | 290             | 1000          | 420      | 0.500             |
| 28 | IRSL | 325             | 1000          | 420      | 0.238             |
| 29 | TL   | 0               | 4000          | 250      | 1.285             |
| 30 | IRSL | 50              | 4000          | 220      | 0.500             |
| 31 | IRSL | 290             | 4000          | 420      | 0.500             |
| 32 | TL   | 0               | 1000          | 250      | 1.285             |
| 33 | IRSL | 50              | 1000          | 220      | 0.500             |
| 34 | IRSL | 290             | 1000          | 420      | 0.500             |
| 35 | IRSL | 325             | 1000          | 420      | 0.238             |
| 36 | TL   | 0               | 6000          | 250      | 1.285             |
| 37 | IRSL | 50              | 6000          | 220      | 0.500             |

| #  | Type | max. Temp. [°C] | Irr. time [s] | Channels | Channel width [s] |
|----|------|-----------------|---------------|----------|-------------------|
| 38 | IRSL | 290             | 6000          | 420      | 0.500             |
| 39 | TL   | 0               | 1000          | 250      | 1.285             |
| 40 | IRSL | 50              | 1000          | 220      | 0.500             |
| 41 | IRSL | 290             | 1000          | 420      | 0.500             |
| 42 | IRSL | 325             | 1000          | 420      | 0.238             |
| 43 | TL   | 0               | 1000          | 250      | 1.285             |
| 44 | IRSL | 50              | 1000          | 220      | 0.500             |
| 45 | IRSL | 290             | 1000          | 420      | 0.500             |
| 46 | TL   | 0               | 1000          | 250      | 1.285             |
| 47 | IRSL | 50              | 1000          | 220      | 0.500             |
| 48 | IRSL | 290             | 1000          | 420      | 0.500             |

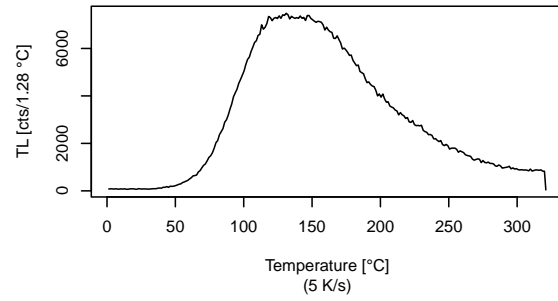


## 1.3 Measurement plots

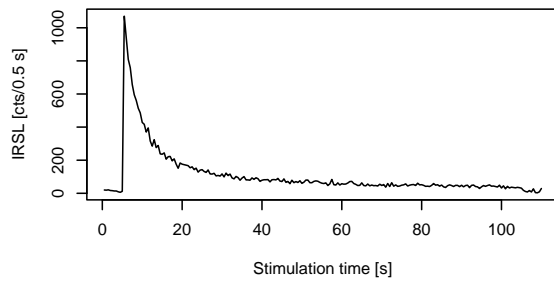
(step 1) natural dose TL



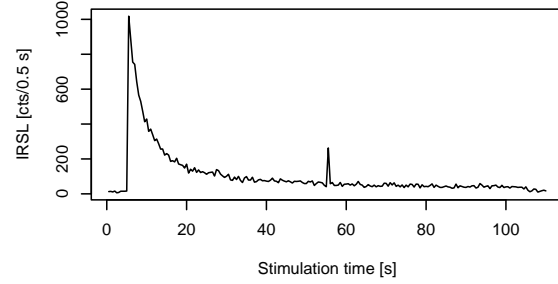
(step 4) test dose TL



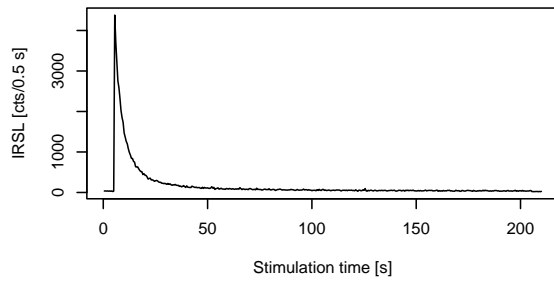
(step 2) natural dose IRSL at 50 °C



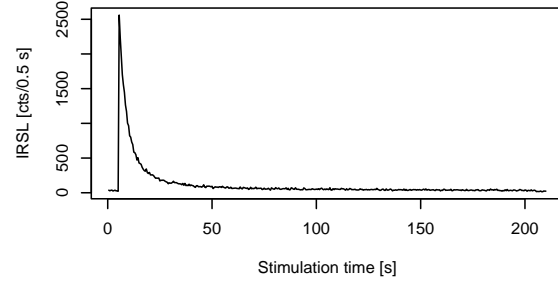
(step 5) test dose IRSL at 50 °C



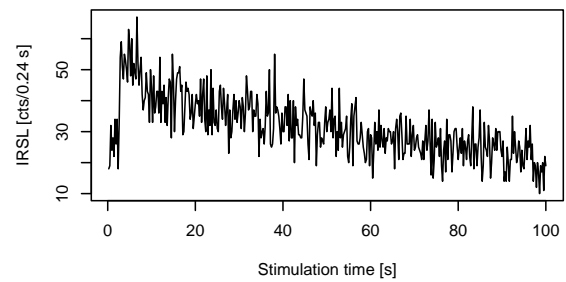
(step 3) natural dose pIRSL at 290 °C



(step 6) test dose pIRSL at 290 °C



(step 7) IRSL bleaching at 325 °C



## 2 Data preparation

### 2.1 Data cleaning

The sequence structure and the record plots reveal two issues which need to be solved before the data can be further analysed with the function `analyse_SAR.CWOSL()` of the `Luminescence` package:

- The data contains three different types of IRSL steps
- At the beginning and the end of the measurement records, the stimulation light was turned off.

We use the function `RLum.OSL_correction()` of the package `OSLdecomposition` (Mittelstraß et al. 2022) to separate IRSL records and to remove the zero-stimulation parts of the records. The correction procedure log is shown below.

```
CORRECTION STEP 1 ----- Check records for consistency in the detection settings -----
Frequency table of different sets of detection settings (Channels, Channel width):
      settings frequency record_type
1          220, 0.5          336      IRSL
3          420, 0.5          336      IRSL2
2 420, 0.238095238095238          144      IRSL3
RLum.Data.Curve@RecordType changed to IRSL2 or IRSL3 in sequence: 1, 2, 3, 4, 5, 6, 7, 8, 9,
Further data manipulations are performed just on IRSL records
(time needed: 0.65 s)
```

```
CORRECTION STEP 2 ----- Remove not stimulated measurement parts -----
Measurement parts with stimulation light turned off detected and removed:
  5 s at the beginning and 5 s at the end.
-> Length of 336 IRSL records reduced from 110 s to 100 s
(time needed: 1.7 s)
```

We perform the code again but only for IRSL2 records to clean also 290°C IRSL records.

```
CORRECTION STEP 1 ----- Check records for consistency in the detection settings -----
All IRSL2 records have the same detection settings
(time needed: 0.28 s)
```

```
CORRECTION STEP 2 ----- Remove not stimulated measurement parts -----
Measurement parts with stimulation light turned off detected and removed:
  5 s at the beginning and 0 s at the end.
-> Length of 336 IRSL2 records reduced from 210 s to 205 s
(time needed: 3.12 s)
```

## 2.2 Dose rate calculation

The beta-source dose rate is not given in the data set. However, Dr. Tobias Lauer provided a calibrated dose rate value of  $0.1053 \pm 0.001$  Gy/s measured in the same week as these measurements were performed. Thus it is not necessary to recalculate the dose rate.

We apply the dose rate on the IRR\_TIME property of the records to get the applied doses in each step. The dose rate error will be added to the equivalent dose error in a later step.

## 2.3 Corrected data set

The above data preparation steps accumulate to a more reasonable sequence structure, where we also translate the record length into measurement time, add measurement cycle and step names and write proper column names:

Table 3: Measurement sequence structure for aliquot 1

| Index | Record Type | Temperature (°C) | Dose (Gy) | Meas. time (sec) | Ch. width (sec) | Meas. cycle | Step      |
|-------|-------------|------------------|-----------|------------------|-----------------|-------------|-----------|
| 1     | TL          | 321              | 0.0       | 321.25           | 1.285           | 1           | preheat   |
| 2     | IRSL        | 50               | 0.0       | 100.00           | 0.500           | 1           | Lx        |
| 3     | IRSL2       | 290              | 0.0       | 205.00           | 0.500           | 1           | Lx        |
| 4     | TL          | 0                | 105.3     | 321.25           | 1.285           | 1           | preheat   |
| 5     | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 1           | Tx        |
| 6     | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 1           | Tx        |
| 7     | IRSL3       | 325              | 105.3     | 99.96            | 0.238           | 1           | bleaching |
| 8     | TL          | 321              | 0.0       | 321.25           | 1.285           | 2           | preheat   |
| 9     | IRSL        | 50               | 0.0       | 100.00           | 0.500           | 2           | Lx        |
| 10    | IRSL2       | 290              | 0.0       | 205.00           | 0.500           | 2           | Lx        |
| 11    | TL          | 0                | 105.3     | 321.25           | 1.285           | 2           | preheat   |
| 12    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 2           | Tx        |
| 13    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 2           | Tx        |
| 14    | IRSL3       | 325              | 105.3     | 99.96            | 0.238           | 2           | bleaching |
| 15    | TL          | 0                | 105.3     | 321.25           | 1.285           | 3           | preheat   |
| 16    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 3           | Lx        |
| 17    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 3           | Lx        |
| 18    | TL          | 0                | 105.3     | 321.25           | 1.285           | 3           | preheat   |
| 19    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 3           | Tx        |
| 20    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 3           | Tx        |
| 21    | IRSL3       | 325              | 105.3     | 99.96            | 0.238           | 3           | bleaching |
| 22    | TL          | 0                | 210.6     | 321.25           | 1.285           | 4           | preheat   |
| 23    | IRSL        | 50               | 210.6     | 100.00           | 0.500           | 4           | Lx        |

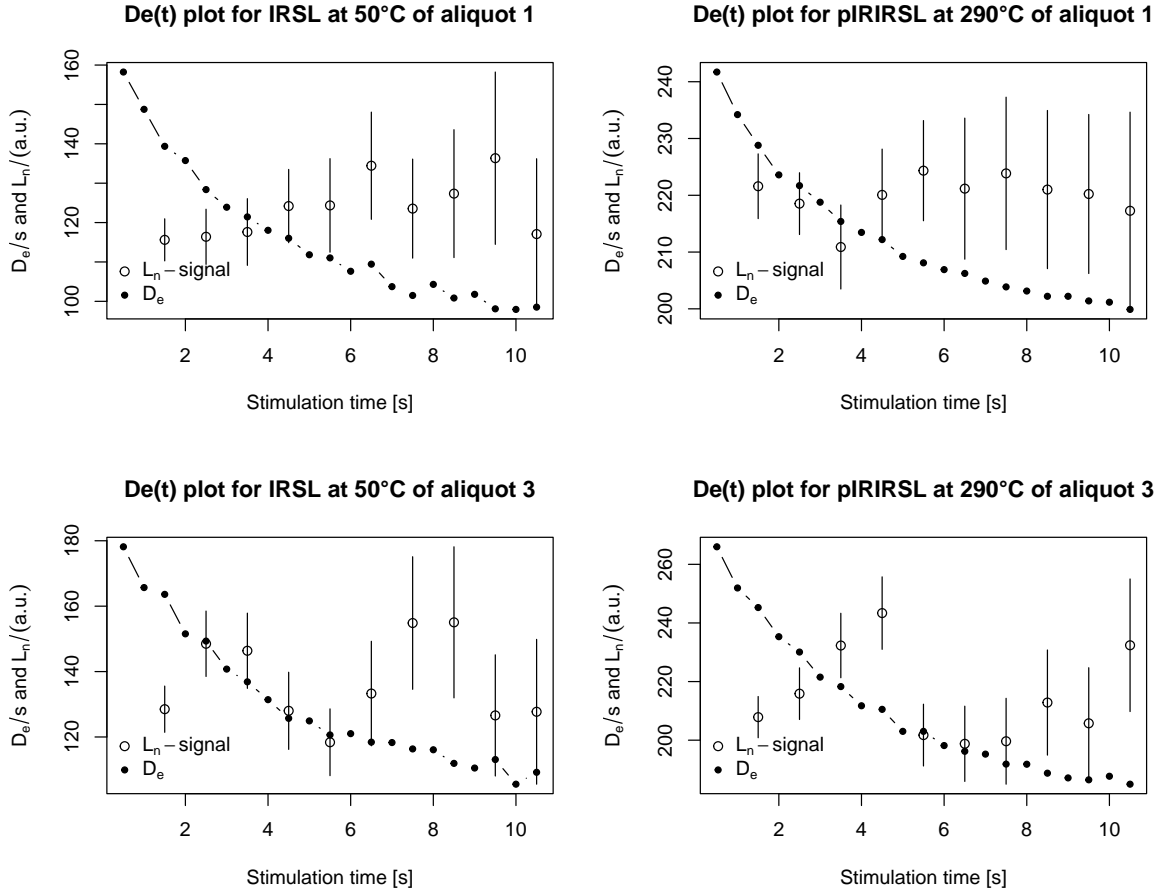


| Index | Record Type | Temperature (°C) | Dose (Gy) | Meas. time (sec) | Ch. width (sec) | Meas. cycle | Step      |
|-------|-------------|------------------|-----------|------------------|-----------------|-------------|-----------|
| 24    | IRSL2       | 290              | 210.6     | 205.00           | 0.500           | 4           | Lx        |
| 25    | TL          | 0                | 105.3     | 321.25           | 1.285           | 4           | preheat   |
| 26    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 4           | Tx        |
| 27    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 4           | Tx        |
| 28    | IRSL3       | 325              | 105.3     | 99.96            | 0.238           | 4           | bleaching |
| 29    | TL          | 0                | 421.2     | 321.25           | 1.285           | 5           | preheat   |
| 30    | IRSL        | 50               | 421.2     | 100.00           | 0.500           | 5           | Lx        |
| 31    | IRSL2       | 290              | 421.2     | 205.00           | 0.500           | 5           | Lx        |
| 32    | TL          | 0                | 105.3     | 321.25           | 1.285           | 5           | preheat   |
| 33    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 5           | Tx        |
| 34    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 5           | Tx        |
| 35    | IRSL3       | 325              | 105.3     | 99.96            | 0.238           | 5           | bleaching |
| 36    | TL          | 0                | 631.8     | 321.25           | 1.285           | 6           | preheat   |
| 37    | IRSL        | 50               | 631.8     | 100.00           | 0.500           | 6           | Lx        |
| 38    | IRSL2       | 290              | 631.8     | 205.00           | 0.500           | 6           | Lx        |
| 39    | TL          | 0                | 105.3     | 321.25           | 1.285           | 6           | preheat   |
| 40    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 6           | Tx        |
| 41    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 6           | Tx        |
| 42    | IRSL3       | 325              | 105.3     | 99.96            | 0.238           | 6           | bleaching |
| 43    | TL          | 0                | 105.3     | 321.25           | 1.285           | 7           | preheat   |
| 44    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 7           | Lx        |
| 45    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 7           | Lx        |
| 46    | TL          | 0                | 105.3     | 321.25           | 1.285           | 7           | preheat   |
| 47    | IRSL        | 50               | 105.3     | 100.00           | 0.500           | 7           | Tx        |
| 48    | IRSL2       | 290              | 105.3     | 205.00           | 0.500           | 7           | Tx        |

### 3 Dose calculation

#### 3.1 De(t) plots

De(t) plotting helps to identify potential age over- or underestimation due to partial signal resetting, unstable signal components or other signal related issues (Bailey et al. 2003). Thus, we evaluated the De(t) plots for the first 10 seconds of stimulation for all measurements using the `plot_DetPlot()` function of the `Luminescence` package. Below are shown the plots for aliquot 1 and aliquot 3 as examples:



Like for other B19-LU samples the IRSL at 50°C measured equivalent doses are lower than the pIR-IRSL measured, which is a indication of thermal fading of the luminescence signal in the samples and thus also a indication of De underestimation in the IRSL at 50°C signals (Thomsen et al. 2008).

However, the significant increase in the De values when shifting the signal integration window towards higher which can be observed for at least 50 % of the aliquots of B19-LU 3, 4 and 5, are less frequent in B19-LU 6.

### 3.2 Dose-response curve fitting optimization

To calculate the De values relevant for dating, we discard the IRSL at 50°C signals and use just the pIR IRSL measurements. The De values are calculated using the `analyse_SAR.CWOSL()` function of the `Luminescence` package.

As the dose calculation of sample **B19 LU-1** revealed a dependence of the De results and the rejection test results from the growth-curve parameters, see details in the B19 LU-1 report), the same parameters were also tested for B19 LU-6. The tables below compare the number of rejected aliquots as well as the Central Age Model (CAM) results in dependence to the chosen parameters:

Table 4: Analysis results setting `fit.method = EXP` (default model).

| Signal integration window | Aliquots that passed RC (of 24) | CAM paleodose [Gy] | Overdispersion [%] |
|---------------------------|---------------------------------|--------------------|--------------------|
| 1.5 sec                   | 24                              | 240.48 +- 4.26     | 8.10               |
| 3.5 sec                   | 24                              | 241.50 +- 4.02     | 7.79               |

Table 5: Analysis results setting `fit.method = GOK` (model by Guralnik et al. (2015)).

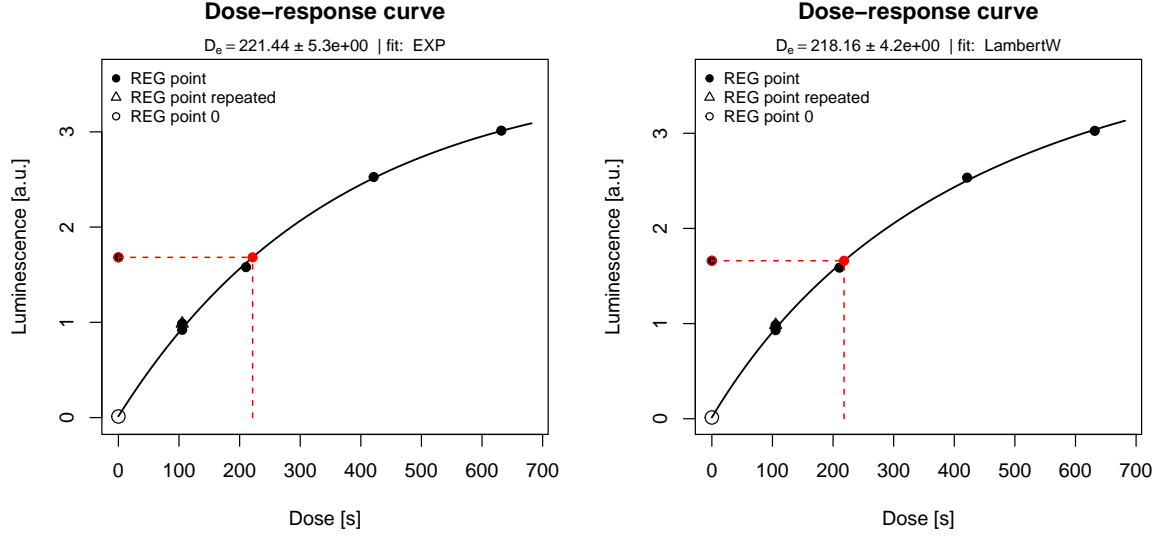
| Signal integration window | Aliquots that passed RC (of 24) | CAM paleodose [Gy] | Overdispersion [%] |
|---------------------------|---------------------------------|--------------------|--------------------|
| 1.5 sec                   | 24                              | 244.30 +- 4.17     | 7.67               |
| 3.5 sec                   | 24                              | 245.74 +- 3.93     | 7.37               |

Table 6: Analysis results setting `fit.method = LambertW` (model by Pagonis et al. (2020)).

| Signal integration window | Aliquots that passed RC (of 24) | CAM paleodose [Gy]    | Overdispersion [%] |
|---------------------------|---------------------------------|-----------------------|--------------------|
| 1.5 sec                   | 24                              | 244.87 +- 4.11        | 7.39               |
| <b>3.5 sec</b>            | <b>24</b>                       | <b>246.12 +- 3.90</b> | <b>7.25</b>        |

Changing the growth curve parameters has no significant impact on the result. Thus, the same parameters as used for B19 LU-1 were used for this data set (**bold** table row) to ensure

parameter uniformity. Please be aware that the results displayed in the tables above are not exactly reproducible as the used calculation procedures deploy Monte Carlo methods.



### 3.3 De calculation result table

Table 7: Equivalent doses for pIRIR290 measurements

| #  | De [Gy] | De error [Gy] | D01 [Gy] | D01 error [Gy] | Rejection criteria |
|----|---------|---------------|----------|----------------|--------------------|
| 1  | 218.16  | 4.72          | NA       | NA             | OK                 |
| 2  | 243.66  | 6.13          | NA       | NA             | OK                 |
| 3  | 225.72  | 7.92          | NA       | NA             | OK                 |
| 4  | 287.60  | 12.27         | NA       | NA             | OK                 |
| 5  | 211.27  | 4.10          | NA       | NA             | OK                 |
| 6  | 268.94  | 7.84          | NA       | NA             | OK                 |
| 7  | 247.31  | 7.05          | NA       | NA             | OK                 |
| 8  | 257.46  | 8.61          | NA       | NA             | OK                 |
| 9  | 243.85  | 5.19          | NA       | NA             | OK                 |
| 10 | 268.34  | 7.15          | NA       | NA             | OK                 |
| 11 | 222.19  | 4.77          | NA       | NA             | OK                 |
| 12 | 251.82  | 5.45          | NA       | NA             | OK                 |
| 13 | 248.43  | 8.31          | NA       | NA             | OK                 |
| 14 | 258.56  | 4.27          | NA       | NA             | OK                 |
| 15 | 252.35  | 5.82          | NA       | NA             | OK                 |
| 16 | 223.90  | 6.91          | NA       | NA             | OK                 |

| #  | De [Gy] | De error [Gy] | D01 [Gy] | D01 error [Gy] | Rejection criteria |
|----|---------|---------------|----------|----------------|--------------------|
| 17 | 267.54  | 8.26          | NA       | NA             | OK                 |
| 18 | 251.75  | 8.32          | NA       | NA             | OK                 |
| 19 | 248.89  | 8.56          | NA       | NA             | OK                 |
| 20 | 217.38  | 5.99          | NA       | NA             | OK                 |
| 21 | 260.50  | 7.23          | NA       | NA             | OK                 |
| 22 | 268.36  | 5.05          | NA       | NA             | OK                 |
| 23 | 248.82  | 7.93          | NA       | NA             | OK                 |
| 24 | 238.39  | 8.17          | NA       | NA             | OK                 |

24 of all aliquots passed the rejection criteria. The results of all aliquots in the table above include the dose rate errors.

### 3.4 Rejection criteria

Table 8: Rejection criteria thresholds (left) and results (right)

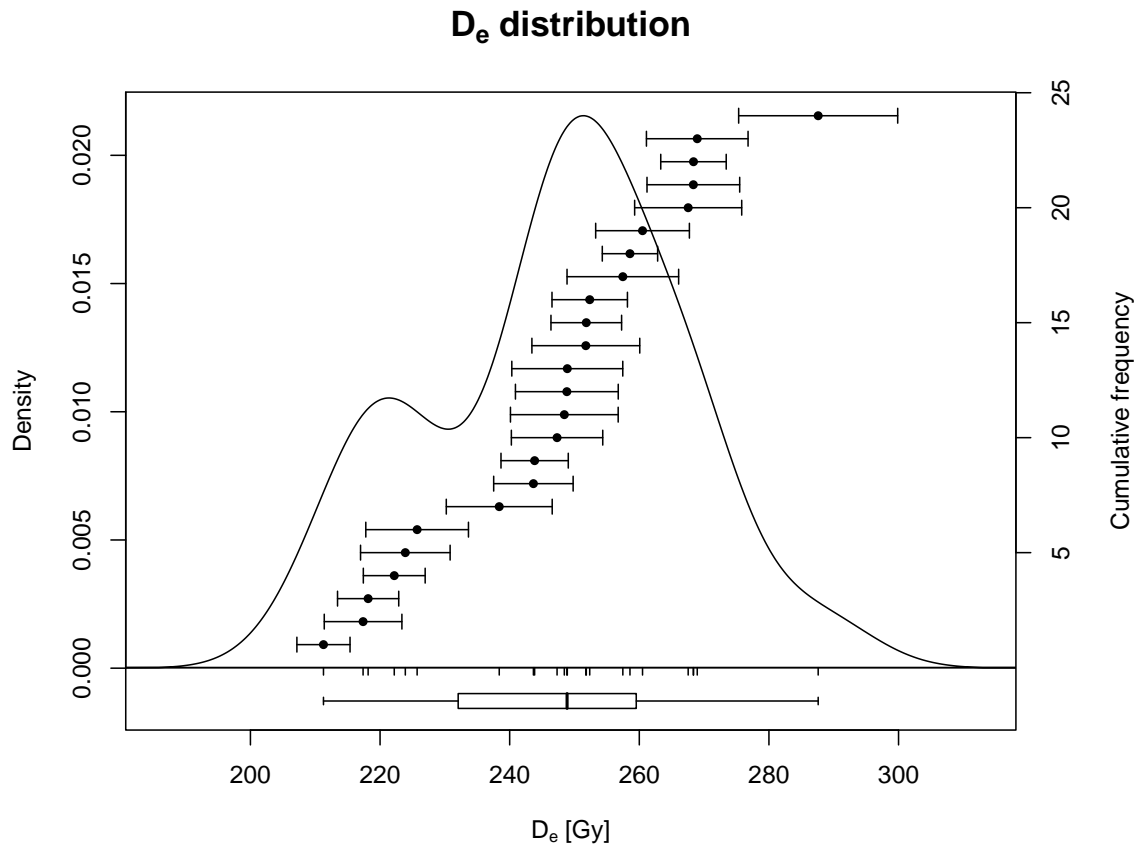
| # | Criterium               | Threshold | #  | A     | B     | C     | D     | E       |
|---|-------------------------|-----------|----|-------|-------|-------|-------|---------|
| A | Recycling ratio (R5/R1) | 0.1       | 1  | 1.058 | 0.008 | 0.009 | 0.019 | 218.161 |
| B | Recuperation rate 1     | 0.1       | 2  | 1.004 | 0.010 | 0.010 | 0.023 | 243.660 |
| C | Testdose error          | 0.1       | 3  | 1.012 | 0.006 | 0.012 | 0.034 | 225.722 |
| D | Palaeodose error        | 0.1       | 4  | 1.019 | 0.009 | 0.018 | 0.042 | 287.597 |
| E | De > max. dose point    | 631.8     | 5  | 1.028 | 0.007 | 0.007 | 0.017 | 211.271 |
|   |                         |           | 6  | 1.026 | 0.012 | 0.012 | 0.028 | 268.944 |
|   |                         |           | 7  | 1.023 | 0.011 | 0.011 | 0.027 | 247.310 |
|   |                         |           | 8  | 1.048 | 0.023 | 0.012 | 0.032 | 257.464 |
|   |                         |           | 9  | 1.026 | 0.005 | 0.010 | 0.019 | 243.847 |
|   |                         |           | 10 | 1.052 | 0.010 | 0.009 | 0.025 | 268.337 |
|   |                         |           | 11 | 1.073 | 0.009 | 0.008 | 0.019 | 222.194 |
|   |                         |           | 12 | 0.990 | 0.012 | 0.007 | 0.019 | 251.820 |
|   |                         |           | 13 | 1.024 | 0.005 | 0.012 | 0.032 | 248.427 |
|   |                         |           | 14 | 1.040 | 0.007 | 0.005 | 0.014 | 258.560 |
|   |                         |           | 15 | 1.046 | 0.009 | 0.011 | 0.021 | 252.350 |
|   |                         |           | 16 | 1.015 | 0.007 | 0.013 | 0.029 | 223.900 |
|   |                         |           | 17 | 1.058 | 0.009 | 0.011 | 0.029 | 267.541 |
|   |                         |           | 18 | 1.024 | 0.007 | 0.013 | 0.032 | 251.755 |
|   |                         |           | 19 | 0.998 | 0.013 | 0.013 | 0.033 | 248.887 |
|   |                         |           | 20 | 1.024 | 0.008 | 0.009 | 0.026 | 217.384 |
|   |                         |           | 21 | 1.042 | 0.011 | 0.011 | 0.026 | 260.501 |
|   |                         |           | 22 | 1.008 | 0.009 | 0.008 | 0.016 | 268.360 |
|   |                         |           | 23 | 1.028 | 0.007 | 0.014 | 0.030 | 248.818 |
|   |                         |           | 24 | 1.010 | 0.016 | 0.014 | 0.033 | 238.386 |

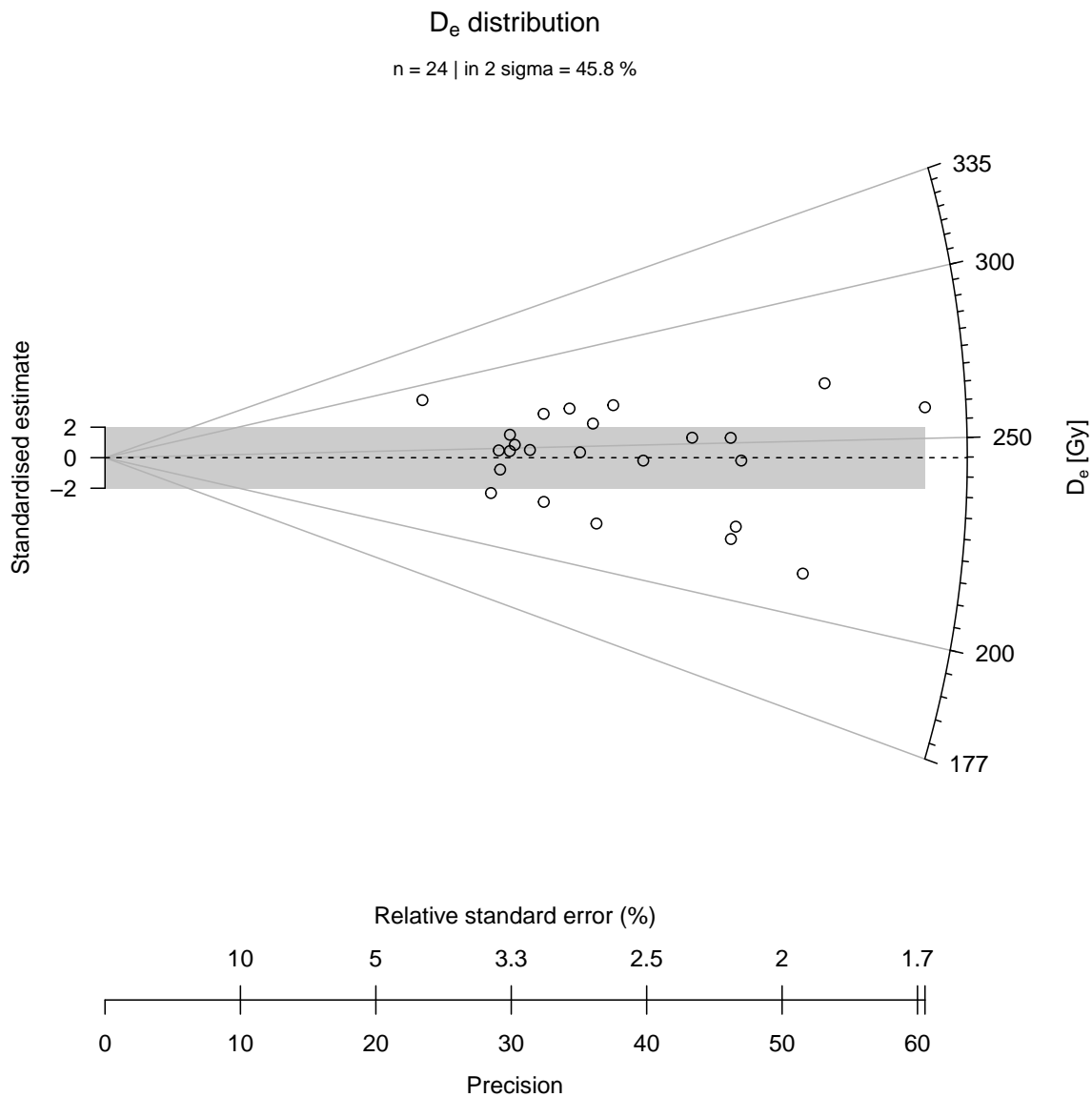
## 4 Dose distribution

**Important:** Those aliquots which did not passed the rejection criteria, where not included in any of the dose distribution calculations.

### 4.1 Distribution plots

The dose distribution is plotted below with the functions `plot_KDE()` and `plot_RadialPlot()` of the `Luminescence` package.







## 4.2 Central age model

Below is output of the function `calc_CentralDose()` of the `Luminescence` package shown, which calculates the central dose and the over-dispersion of the De distribution in accordance to the model given by Galbraith et al. (1999) .

```
[calc_CentralDose]

----- meta data -----
n:                24
log:              TRUE
----- dose estimate -----
abs. central dose: 246.08
abs. SE:           3.91
rel. SE [%]:       1.59
----- overdispersion -----
abs. OD:           17.83
abs. SE:           2.96
OD [%]:            7.25
SE [%]:            1.20
-----
```

*SE = standard error, OD = over-dispersion*

## 5 References

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