

# Data processing for volumetric BMP measurements

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## 1 Introduction

Volumetric methods are commonly used for measuring cumulative biogas and methane production in order to determine methane potential and anaerobic biodegradability of a given substrate. In volumetric methods, biogas volumes are measured under constant pressure using different methods such as liquid displacement and syringe methods [1]. In the liquid displacement method, the biogas is transferred to an external system measuring the volume. As biogas is produced it is collected in the external liquid displacement system displacing an amount of the barrier liquid equivalent to the biogas volume. In the syringe method, lubricated syringes are manually inserted into the bottle of interest. Here, the pressure increase resulting from biogas production forces the piston inwards the syringe until the pressure equilibrates the surrounding pressure (most commonly atmospheric). Produced biogas volume is equivalent to the displacement of the piston and can be read directly of the syringe [2].

Determination of  $\text{CH}_4$  volume from volumetric measurements requires data processing. The `calcBgVol()` function was developed to process volumetric data. A newly developed biogas package (available for the R environment since 2015) address issues with time-consuming calculations and lack of reproducibility among laboratories for obtaining BMP [3]. The biogas package consists of ten functions including `cumBg()`, which are used for processing volumetric, manometric, gravimetric, and gas density measurements. The resulting biogas and methane production and production rates can be further used to calculate BMP using the `summBg()` function.

This document describes how to process volumetric biogas measurements using the `calcBgVol()` function. We assumed that readers are familiar with biogas data collection and R.

## 2 Overview of the Function

`calcBgVol()` is a “high-level” function within the biogas package. The purpose of `calcBgVol()` is to convert volume data collected in the laboratory to cumu-

lative biogas and CH<sub>4</sub> production and production rates. Typically, these values will then be used for calculation of BMP, e.g. using the `summBg()` function. `calcBgVol()` can handle data from multiple bottles and accepts three different data structures. For simple operations (e.g. interpolation and standardization of biogas volume) `calcBgVol()` is supported by calls to external “low-level” functions (refer to Section 5, Table 1).

Two methods are commonly used to calculate methane production from volume and composition measurements. Method 1 is based on normalized CH<sub>4</sub> concentrations, whereas method 2 accounts for the actual CH<sub>4</sub> in the bottle headspace. Both methods are available through `calcBgVol()` and results are expected to be virtual identical. The examples below describe cumulative biogas calculation from three datasets of different structures (‘‘wide’’, ‘‘long’’, and ‘‘longcombo’’) using volumetric method 1. For method 2 examples, refer to vignette for manometric `cumBg` function (`cumBgMan_function.Rnw`).

## 2.1 Function Arguments

The arguments for the function are:

```
## function (dat, comp = NULL, temp = NULL, pres = NULL,
##         interval = TRUE, data.struct = "longcombo",
##         id.name = "id", time.name = "time",
##         vol.name = "vol", comp.name = "xCH4",
##         headspace = NULL, vol.hs.name = "vol.hs",
##         cmethod = "removed", imethod = "linear",
##         extrap = FALSE, addt0 = TRUE, showt0 = TRUE,
##         dry = FALSE, empty.name = NULL,
##         std.message = !quiet, check = TRUE,
##         temp.std = getOption("temp.std", as.numeric(NA)),
##         pres.std = getOption("pres.std", as.numeric(NA)),
##         unit.temp = getOption("unit.temp", "C"),
##         unit.pres = getOption("unit.pres", "atm"),
##         quiet = FALSE)
## NULL
```

Most of the arguments have default values, but to calculate CH<sub>4</sub> production we must provide values for at least `dat` (data frame with response variable - here volume measurements), `comp` (data frame with methane concentration), `pres` (biogas pressure), and `temp` (biogas temperature)<sup>1</sup> along with the names of a few columns in the input data frames. If `temp` and/or `pres` arguments are not provided, biogas volumes will not be standardized.

<sup>1</sup>. By default, temperature is in °C and pressure in atm, but these can be changed in the function call with the `temp.unit` and `pres.unit` arguments, or globally with `options`. The same default values apply for temperature and pressure for presentation of biogas and methane, but these can be changed in the function call with the `temp.std` and `pres.std` arguments.

By default `interval = TRUE` and `data.struct` is set as `‘‘longcombo’’`. `‘‘wide’’` and `‘‘long’’` structured data will be restructured to `‘‘longcombo’’` internally by `cumBgDataPrep()`, when specified by the `data.struct` argument (refer to Section 2.2). When data are cumulative, the interval argument should be set to `FALSE`. When data are mixed interval and cumulative response variables (`empty.name != TRUE`), the interval argument is ignored.

Similarly, there is an `id.name` argument for the bottle identification code (ID) column (default is `‘‘id’’`). The default value is `‘‘id’’`. For `data.struct = ‘‘wide’’`, there is no ID column. Instead data for each bottle, have individual columns and column names, which are used as ID codes. Here, the name of the column containing the response variables (`vol.name`), is set as the name of the first column with response variables. All following columns are assumed to have measurement data.

Furthermore, we need to specify the name of the time column containing time data using the `time.name` argument (default is `‘‘time’’`). If separate data frames are used for `dat` and `comp`, the name must match. Time data may be POSIXct objects, but then `t0` will not be added to rows by the `calcBgVol()` function. In addition the `addt0` argument is used to add row with “time zero” (`time.name = 0`) for each bottle in order to calculate production rates for the first observation (default is `TRUE`). `showt0` determines if the “time zero” should be returned in the output (default is `TRUE` if `time.name` is numeric and contains 0 and otherwise `FALSE`). The use of `t0` output is for plotting.

The `comp.name` argument is used to indicate which column within the `comp` data frame contains the  $\text{CH}_4$  content. Default is `‘‘xCH4’’`. The `comp` argument may be a single value instead of a data frame or column. When providing a single value for `comp`, this value is applied to all observations, and the `comp.name` argument is not needed. Note that if the `comp` argument is not provided, `calcBgVol()` will return results for biogas only and not  $\text{CH}_4$ . The definition of `‘‘xCH4’’` depends on the `cmethod` argument. By default (`cmethod = ‘‘removed’’`) the function calculates volumes following [4] as the product of standardized volume of biogas removed and normalized  $\text{CH}_4$  content (as mole fraction in dry biogas, normalized so the sum of mole fractions of  $\text{CH}_4$  and  $\text{CO}_2$  sum to unity). If results should be based on the sum of methane removed and methane remaining in the bottle headspace, `cmethod` should be set to `‘‘total’’`. When `cmethod = ‘‘total’’`,  $\text{CH}_4$  concentration is expressed relative to all biogas components except water ( $\text{CH}_4$ ,  $\text{CO}_2$ ,  $\text{N}_2$ ,  $\text{H}_2\text{S}$ , etc.) instead of  $\text{CH}_4$  and  $\text{CO}_2$  only.

If any  $\text{CH}_4$  measurements are missing, `‘‘xCH4’’` is interpolated by the external “low-level” function `interp()`. Here, the `imethod` argument can be used to define interpolation method (default is `‘‘linear’’`), which is passed to `interp()`. Similar, an `extrap` argument is passed to `interp()` (default is `FALSE`). The `extrap` argument is used to indicate if composition data (`comp.name`) should be extrapolated (e.g. in the case of missing initial composition values). Extrapolation is constant (the value of the nearest observation is taken). In general, extrapolation and extensive interpolation should be avoided.

Additionally, a data frame containing headspace volumes is required if `cmethod`

= `‘total’` and should contain at least a headspace volume column (`vol.hs.name`) and a bottle identification column, with the same column name as in `dat` and `comp` data frames. The headspace volume column can be set using the `vol.hs.name` argument (default is `‘vol.hs’`).

By default biogas is assumed to be saturated with water vapor. If biogas volumes are already standardized to dry conditions (as with AMPTS II data) we need to set `dry = TRUE`.

## 2.2 Data Structures

Input data may be structured in one of three ways: `‘longcombo’`, `‘long’`, and `‘wide’`. Default is `‘longcombo’`, where the composition column is in the `dat` data frame and no separate composition data frame is required. In the following examples all three data structures will be addressed. For more details about each of the three data structure see `biogas_quick_start` vignette.

### 3 Examples: Calculation of Cumulative Production of Biogas and CH<sub>4</sub> and Production Rates from Volumetric Measurements

Calculation of cumulative biogas and CH<sub>4</sub> production and production rates, typically requires two data frames: Biogas volume measurements and biogas composition (CH<sub>4</sub> concentration).

#### 3.1 ‘‘longcombo’’ Data Structure

In this example, we will use a ‘‘longcombo’’ example dataset included in the biogas package: `s3lcombo` for both biogas volumes and composition. These data are from an experiment carried out with municipal wastewater sludge. Only three bottles, all with inoculum and substrate, were included: D, E, and F. 300 mL glass serum bottles with butyl rubber septa and screw caps were used. Volume was measured using syringes. Methane and carbon dioxide contents were determined by gas chromatography using a thermal conductivity detector and normalized so methane and carbon dioxide sum to 1.0. This is a small dataset, with versions available for all three different data structures. It cannot be used to calculate BMP, because no inoculum-only bottles were included.

```
data("s3lcombo")

dim(s3lcombo)

## [1] 21 4

s3lcombo

##   id  time.d vol.ml  xCH4
## 1  D  0.9438   103    NA
## 2  E  0.9451   106    NA
## 3  F  0.9472   107    NA
## 4  D  2.9060   192 0.6983
## 5  E  2.9090   181 0.6817
## 6  F  2.9100   203 0.6869
## 7  D  5.8860   141 0.6800
## 8  E  5.8880   133 0.6800
## 9  F  5.8900   140 0.6800
## 10 D 10.0000   112 0.6646
## 11 E 10.0000   111 0.6644
## 12 F 10.0100   110 0.6632
## 13 D 23.1000   200 0.6946
## 14 E 23.1000   190 0.6871
## 15 F 23.1000   200 0.6829
## 16 D 34.0100   109 0.6626
```

```
## 17 E 34.0100 110 0.6556
## 18 F 34.0100 112 0.6527
## 19 D 57.8400 146 0.6651
## 20 E 57.8400 136 0.6600
## 21 F 57.8400 138 NA
```

```
summary(s3lcombo)
```

```
## id          time.d          vol.ml          xCH4
## D:7  Min.    : 0.9438  Min.    :103.0  Min.    :0.6527
## E:7  1st Qu.: 2.9100  1st Qu.:110.0  1st Qu.:0.6632
## F:7  Median :10.0000  Median :136.0  Median :0.6800
##      Mean   :19.2421  Mean    :141.9  Mean    :0.6741
##      3rd Qu.:34.0100  3rd Qu.:181.0  3rd Qu.:0.6829
##      Max.   :57.8400  Max.    :203.0  Max.    :0.6983
##                                     NA's    :4
```

Calculating cumulative biogas and CH<sub>4</sub> production and production rates, is a common first step in processing data from a BMP test. Subsequently, BMP can be calculated by the “high-level” function `summBg()` included in the biogas package. Cumulative biogas and CH<sub>4</sub> production and production rates from volumetric data with `s3lcombo` data frame as the only input, can be calculated from `calcBgVol()`.

To calculate CH<sub>4</sub> production from these “`longcombo`” data, we must provide values for at least `dat` and `comp`, which is in a combined data frame (we will use `s3lcombo`), `temp` (biogas temperature), and `pres` (biogas pressure) along with the names of a few columns in our input data frame.

We can use the default values “`longcombo`”, “`id`”, and “`xCH4`” for the `data.struct`, `id.name`, and `comp.name` arguments, respectively. Whereas, the `vol.name` argument needs to be specified as “`vol.ml`”. Similar, default values can be used for `cmethod = “removed”`, evaluating CH<sub>4</sub> concentration based on normalized CH<sub>4</sub> and CO<sub>2</sub> values and for `imethod = “linear”`, resulting in internal linear interpolation of `xCH4` by calling the `interp` function. Furthermore, initial composition (`xCH4`) values are missing in the `s3lcombo` data frame. Therefore, we set `extrap = TRUE` for extrapolation of these initial CH<sub>4</sub> values

By default biogas is assumed to be saturated with water vapor (`dry = FALSE`). Biogas volumes have not been standardized to dry conditions and hence, default value for the `dry` argument matches requirements for making an proper evaluation of the dataset.

```
cum.prod.lc <- calcBgVol(s3lcombo, temp = 25, pres = 1,
                        time.name = 'time.d', vol.name = 'vol.ml',
                        comp.name = 'xCH4',
                        extrap = TRUE)
## Biogas composition is interpolated.
```

```
## Using a standard pressure of 1 atm and standard temperature of 0
C for standardizing volume.
```

The output becomes:

```
head(cum.prod.lc)

##   id  time.d  vol.ml  xCH4  temperature  pressure  vBg
## 1  D  0.0000    NA    NA            NA        NA  0.00000
## 2  D  0.9438   103  0.6983            25         1  91.40334
## 3  D  2.9060   192  0.6983            25         1 170.38293
## 4  D  5.8860   141  0.6800            25         1 125.12497
## 5  D 10.0000   112  0.6646            25         1  99.39004
## 6  D 23.1000   200  0.6946            25         1 177.48222
##          vCH4      cvBg      cvCH4      rvBg      rvCH4
## 1  0.00000  0.00000  0.00000      NA      NA
## 2  63.82696  91.40334  63.82696  96.84609  67.627628
## 3 118.97840 261.78628 182.80536  86.83260  60.635206
## 4  85.08498 386.91124 267.89033  41.98824  28.552006
## 5  66.05462 486.30129 333.94496  24.15898  16.056058
## 6 123.27915 663.78351 457.22411  13.54826   9.410622

dim(cum.prod.lc)

## [1] 24 12
```

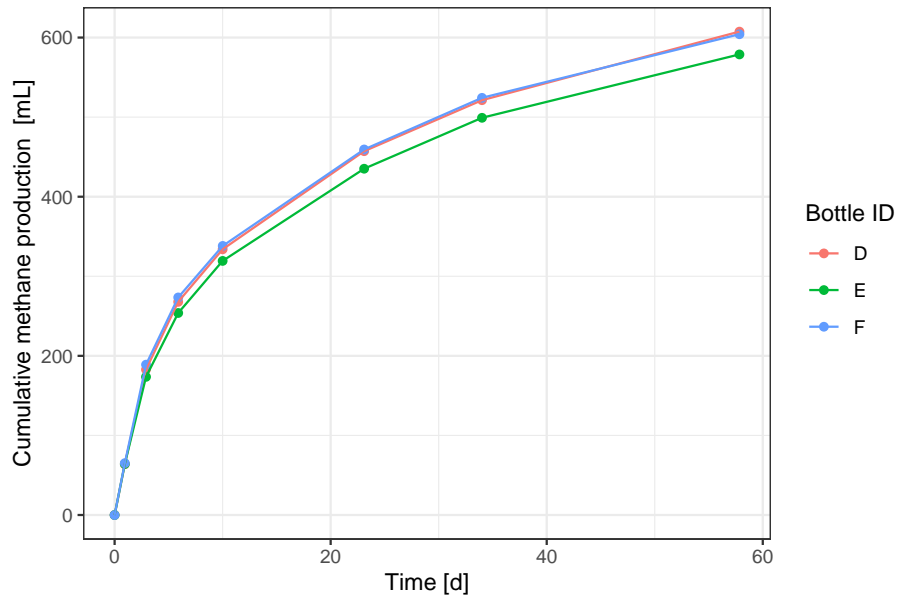
The returned data frame has maintained the ‘‘longcombo’’ structure with all the original columns in `s3lcombo`, plus additional columns from volumetric biogas calculations.

In these columns, `v` stands for (standardized) volume, `cv` (standardized) cumulative volume, `rv` stands for (standardized) volume production rate, and `Bg` and `CH4` for biogas and methane. `cvBg` contains standardized cumulative biogas production and `cvCH4` contains standardized cumulative  $\text{CH}_4$  production.

Plotting results is recommended for easier understanding, and it can be used to catch strange responses. Here we will use the `ggplot` function from the `ggplot2` package to plot it.

```
library(ggplot2)

ggplot(cum.prod.lc, aes(time.d, cvCH4, colour = factor(id))) +
  geom_point() +
  geom_line(aes(group = id)) +
  labs(x = "Time [d]", y = "Cumulative methane production [mL]", colour = "Bottle ID") +
  theme_bw()
```



### 3.2 ‘‘wide’’ Data Structure

In this example, we will use a ‘‘wide’’ structured example dataset included in the `biogas` package, having the data frame `feedVol` for biogas volumes and `feedSetup` for bottle identification and substrate and inoculum masses. The `feedSetup` data frame is not relevant for `calcBgVol()`, but is required subsequently for `summBg()` to calculate BMP. BMP measurement data are from a batch test carried out on animal feed ingredients along with cellulose as a control. The experiment included 12 batch bottles:

- Three bottles with inoculum only (BK)
- Three bottles with cellulose and inoculum (CEL)
- Three bottles with animal feed ingredient 1 and inoculum (SC)
- Three bottles with animal feed ingredient 2 and inoculum (SD)

A typical automated volumetric method called AMPTS II was used to measure biogas production: an online, standardized lab-measurement platform for BMP tests. Applying AMPTS II, the measured volumes are standardized, and the composition is 100% methane. Therefore, the `comp` argument is set to 1 when calling the `calcBgVol()` function. Furthermore, pressure is set to a fixed value (1 atm or 101.325 kPa) and temperature to 0°C.

```
data("feedVol")
dim(feedVol)
```



```

## [1] 44 13

head(feedVol)

##   time.d      1      2      3      4      5      6      7      8
## 1      0  0.0  0.0  0.0  0.0  0.0  0.0  0.0  0.0
## 2      1 70.6 68.5 69.7 78.6 78.1 85.9 383.8 390.7
## 3      2 100.5 97.6 100.3 357.0 369.0 384.4 902.4 927.4
## 4      3 125.0 120.8 125.9 846.0 895.2 902.4 1517.3 1563.0
## 5      4 144.0 138.3 144.5 1291.5 1323.7 1373.0 1911.5 1940.0
## 6      5 158.6 152.1 158.9 1513.2 1542.1 1600.1 2044.9 2080.7
##      9     10     11     12
## 1      0.0  0.0  0.0  0.0
## 2     399.3 249.5 277.8 280.6
## 3     927.0 521.1 561.8 562.9
## 4    1557.6 830.6 876.9 876.0
## 5    1955.3 1112.7 1160.3 1150.0
## 6    2082.0 1348.9 1413.0 1388.9

summary(feedVol)

##      time.d      1      2      3
## Min.   : 0.00  Min.   : 0.0  Min.   : 0.0  Min.   : 0.0
## 1st Qu.:10.75  1st Qu.:214.2  1st Qu.:200.7  1st Qu.:212.2
## Median :21.50  Median :256.4  Median :239.7  Median :251.3
## Mean   :21.50  Mean   :228.7  Mean   :213.5  Mean   :225.1
## 3rd Qu.:32.25  3rd Qu.:269.3  3rd Qu.:249.8  3rd Qu.:264.5
## Max.   :43.00  Max.   :269.3  Max.   :249.8  Max.   :264.5
##      4      5      6      7
## Min.   : 0  Min.   : 0  Min.   : 0  Min.   : 0
## 1st Qu.:1877  1st Qu.:1892  1st Qu.:1915  1st Qu.:2328
## Median :1975  Median :1995  Median :1995  Median :2558
## Mean   :1784  Mean   :1807  Mean   :1807  Mean   :2339
## 3rd Qu.:2011  3rd Qu.:2047  3rd Qu.:2022  3rd Qu.:2645
## Max.   :2025  Max.   :2047  Max.   :2022  Max.   :2675
##      8      9      10      11
## Min.   : 0  Min.   : 0  Min.   : 0  Min.   : 0
## 1st Qu.:2376  1st Qu.:2367  1st Qu.:2247  1st Qu.:2335
## Median :2603  Median :2603  Median :2818  Median :2866
## Mean   :2373  Mean   :2377  Mean   :2455  Mean   :2504
## 3rd Qu.:2674  3rd Qu.:2685  3rd Qu.:3015  3rd Qu.:3055
## Max.   :2681  Max.   :2699  Max.   :3124  Max.   :3141
##      12
## Min.   : 0
## 1st Qu.:2298
## Median :2854

```

```
## Mean      :2487
## 3rd Qu.   :3044
## Max.      :3120
```

As with the ‘‘longcombo’’ data, cumulative production of CH<sub>4</sub> is needed in order to calculate BMP. Again, we can calculate these with the `calcBgVol()` function, using the `feedVol` data frame and the `comp = 1` argument as inputs. For this particular example, standardized, cumulative CH<sub>4</sub> volume is present in the input data frame (as with all AMPTS II datasets) and the function only restructures the data and calculates interval volumes and rates. In other cases (see `s3volw` for example), volume standardization and other steps (e.g. interpolation or summing interval values to get cumulative) are needed.

To calculate CH<sub>4</sub> production from these ‘‘wide’’ structured data, we must provide values for at least `dat`, `comp`, `temp`, and `pres` along with the names of a few columns in our input data frame. The `dat` argument is set as the `feedVol` data frame, whereas `comp`, `temp`, and `pres` are set as single values of 1 (mole fraction), 0 (degree C), and 1 (atm), respectively. Since a single value is provided for `comp`, the `comp.name` argument is not required.

For `data.struct != ‘‘longcombo’’` the data structure needs to be specified. Here we set `data.struct = ‘‘wide’’`. Furthermore, we need to specify the name of the time column in `feedVol` using the `time.name` argument.

As mentioned, there are no ID columns for `data.struct = ‘‘wide’’`. Instead column name of the first column with response variables (volume measurements) is used as ID code. In this example observations are numbered 1 to 12 and hence, the `vol.name` argument is set to 1. All following columns are assumed to also contain response variables. By default (`cmethod = ‘‘removed’’`) CH<sub>4</sub> concentrations are evaluated based on normalized CH<sub>4</sub> and CO<sub>2</sub> values. Similar, default values can be used for `imethod = ‘‘linear’’`, resulting in internal linear interpolation of `xCH4` by calling the `interp()` function, and `extrap = FALSE` as all composition data are provided (set to 1 for all observations).

By default biogas is assumed to be saturated with water vapor. For AMPTS II data, biogas volume are already standardized to dry conditions. Therefore, we set `dry = TRUE`. Additionally, the response variables are cumulative data and hence, we set `interval = FALSE`.

```
cum.prod.w <- calcBgVol(feedVol, comp = 1, temp = 0, pres = 1,
                        data.struct = "wide",
                        time.name = "time.d", vol.name = "1",
                        dry = TRUE,
                        interval = FALSE)

## Only a single value was provided for biogas composition (1), so
## applying it to all observations.
## Using a standard pressure of 1 atm and standard temperature of 0
## C for standardizing volume.
```

Note the message about standard temperature and pressure—it is important to make sure these values are correct, therefore users are reminded by a message<sup>2</sup>. Also, note the message about applying single composition value to all observations. When defining pressure as a single value, pressure is assumed to be constant and the same for all observations.

The output becomes:

```
head(cum.prod.w)

##   id time.d   vol xCH4 temperature pressure  vBg vCH4  cvBg
## 1  1     0   0.0   1      0           1  0.0  0.0  0.0
## 2  1     1  70.6   1      0           1 70.6 70.6 70.6
## 3  1     2 100.5   1      0           1 29.9 29.9 100.5
## 4  1     3 125.0   1      0           1 24.5 24.5 125.0
## 5  1     4 144.0   1      0           1 19.0 19.0 144.0
## 6  1     5 158.6   1      0           1 14.6 14.6 158.6
##   cvCH4 rvBg rvCH4
## 1   0.0  NA   NA
## 2  70.6 70.6 70.6
## 3 100.5 29.9 29.9
## 4 125.0 24.5 24.5
## 5 144.0 19.0 19.0
## 6 158.6 14.6 14.6

dim(cum.prod.w)

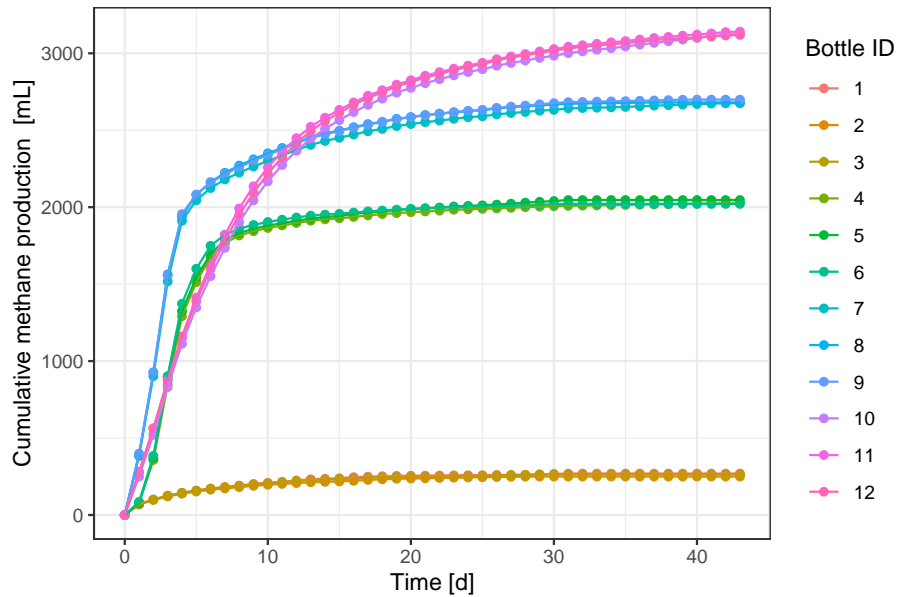
## [1] 528 12
```

The data frame that is returned has been sorted and restructured to ‘‘longcombo’’ structure and has all the original columns in `feedVol`, plus additional columns from the volumetric calculation method (refer to Section 3.1).

As with the ‘‘longcombo’’ data example, the results may be easier to understand graphically using the `ggplot` function from the `ggplot2` package.

```
ggplot(cum.prod.w, aes(time.d, cvCH4, colour = factor(id))) +
  geom_point() +
  geom_line(aes(group = id)) +
  labs(x = "Time [d]", y = "Cumulative methane production [mL]", colour = "Bottle ID") +
  theme_bw()
```

<sup>2</sup> Remember that standard conditions can be set in the function call with `temp.std` and `pres.std`, or globally with `options()`.



### 3.3 ‘‘long’’ Data Structure

In this example, we will use long structured example datasets included with the biogas package: `vol` for biogas volumes and `comp` for composition. These data are from a BMP test carried out on two different substrates A and B, and cellulose included as a ‘‘control’’. The experiment included 12 batch bottles:

- 3 bottles with substrate A and inoculum
- 3 bottles with substrate B and inoculum
- 3 bottles with cellulose and inoculum
- 3 bottles with inoculum only

Bottles consisted of 500 mL or 1.0 L glass bottles, and were sealed with a butyl rubber septum and a screw cap. Initial substrate and inoculum masses were determined. A typical volumetric method was used to measure biogas production: accumulated biogas was measured and removed intermittently using syringes, and composition was measured for some of these samples.

```
data("vol")
dim(vol)
## [1] 288 4
head(vol)
```

```
##      id      date.time days vol
## 1 2_1 2014-06-07 07:00:00 1.98 393
## 2 2_1 2014-06-08 07:00:00 2.98 260
## 3 2_1 2014-06-09 07:00:00 3.98 245
## 4 2_1 2014-06-10 07:00:00 4.98 225
## 5 2_1 2014-06-11 07:00:00 5.98 200
## 6 2_1 2014-06-12 08:00:00 7.02 175

summary(vol)

##      id      date.time      days
## 2_1 : 24  Min. :2014-06-07 07:00:00  Min. : 1.98
## 2_2 : 24  1st Qu.:2014-06-13 20:00:00  1st Qu.: 8.52
## 2_3 : 24  Median :2014-06-28 06:00:00  Median :22.94
## 2_4 : 24  Mean   :2014-07-16 15:29:22  Mean   :41.33
## 2_5 : 24  3rd Qu.:2014-07-25 22:45:00  3rd Qu.:50.63
## 2_6 : 24  Max.   :2014-12-19 04:30:00  Max.   :196.92
## (Other):144
##      vol
## Min.   : 98.0
## 1st Qu.:171.5
## Median :225.0
## Mean   :271.7
## 3rd Qu.:300.0
## Max.   :840.0
##
```

```
data("comp")

dim(comp)

## [1] 132  4

head(comp)

##      id      date.time days      xCH4
## 516 2_1 2014-06-12 08:00:00  7.02 0.7104731
## 519 2_1 2014-06-19 08:00:00 14.02 0.7024937
## 522 2_1 2014-06-26 05:00:00 20.90 0.6659919
## 524 2_1 2014-07-03 04:00:00 27.85 0.6789466
## 525 2_1 2014-07-10 03:00:00 34.81 0.6951429
## 528 2_1 2014-07-24 04:00:00 48.85 0.6693053

summary(comp)

##      id      date.time      days
```

```

## 2_1 :11 Min. :2014-06-12 08:00:00 Min. : 7.02
## 2_2 :11 1st Qu.:2014-06-26 05:00:00 1st Qu.: 20.90
## 2_3 :11 Median :2014-07-24 04:00:00 Median : 48.85
## 2_4 :11 Mean :2014-07-31 07:47:43 Mean : 56.01
## 2_5 :11 3rd Qu.:2014-08-28 04:00:00 3rd Qu.: 83.85
## 2_6 :11 Max. :2014-10-13 07:00:00 Max. :129.98
## (Other):66
## xCH4
## Min. :0.5647
## 1st Qu.:0.6393
## Median :0.6598
## Mean :0.6587
## 3rd Qu.:0.6786
## Max. :0.7115
##

```

To calculate CH<sub>4</sub> production from these ‘‘long’’ structured data, we must provide values for at least `dat` (we will use `vol`), `comp` (we will use `comp`), `temp`, and `pres`. For `data.struct != ‘‘longcombo’’` the data structure needs to be specified. Here we set `data.struct = ‘‘long’’`. Furthermore, we need to specify the name of the time column in `feedVol` using the `time.name` argument.

We can use default values ‘‘id’’, ‘‘vol’’, and ‘‘xCH4’’ for the `id.name`, `vol.name` and `comp.name` arguments, respectively. Similar, default values can be used for `cmethod = ‘‘removed’’`, evaluating CH<sub>4</sub> concentration based on normalized CH<sub>4</sub> and CO<sub>2</sub> values, for `imethod = ‘‘linear’’`, resulting in internal linear interpolation of `xCH4` by calling the `interp()` function, and for `dry = TRUE` assuming biogas to be saturated with water vapor. Additionally, the response variables are interval data only and hence, we can use the default `interval = TRUE`.

In addition to interpolation for later observations, an extrapolation argument (`extrap`) can be provided if required. We do not have initial biogas composition (compare the heads of the `vol` and `comp` data frames) so extrapolation is needed, in addition to interpolation. Therefore, we set `extrap = TRUE`.

```

cum.prod.l <- calcBgVol(vol, comp = comp, temp = 35, pres = 1,
                        data.struct = "long",
                        time.name = "days", vol.name = "vol", comp.name = "xCH4",
                        extrap = TRUE)

## Biogas composition is interpolated.
## Using a standard pressure of 1 atm and standard temperature of 0
## C for standardizing volume.

```

The output becomes:

```

head(cum.prod.l)

##   id          date.time days vol      xCH4 temperature
## 1 2_1          <NA> 0.00  NA        NA           NA
## 2 2_1 2014-06-07 07:00:00 1.98 393 0.7104731      35
## 3 2_1 2014-06-08 07:00:00 2.98 260 0.7104731      35
## 4 2_1 2014-06-09 07:00:00 3.98 245 0.7104731      35
## 5 2_1 2014-06-10 07:00:00 4.98 225 0.7104731      35
## 6 2_1 2014-06-11 07:00:00 5.98 200 0.7104731      35
##   pressure      vBg      vCH4      cvBg      cvCH4      rvBg
## 1         NA    0.0000    0.0000    0.0000    0.0000        NA
## 2          1 328.9470 233.7080 328.9470 233.7080 166.1348
## 3          1 217.6240 154.6160 546.5710 388.3240 217.6240
## 4          1 205.0687 145.6958 751.6397 534.0198 205.0687
## 5          1 188.3284 133.8023 939.9681 667.8221 188.3284
## 6          1 167.4031 118.9354 1107.3712 786.7574 167.4031
##      rvCH4
## 1         NA
## 2 118.0343
## 3 154.6160
## 4 145.6958
## 5 133.8023
## 6 118.9354

dim(cum.prod.l)

## [1] 300 13

```

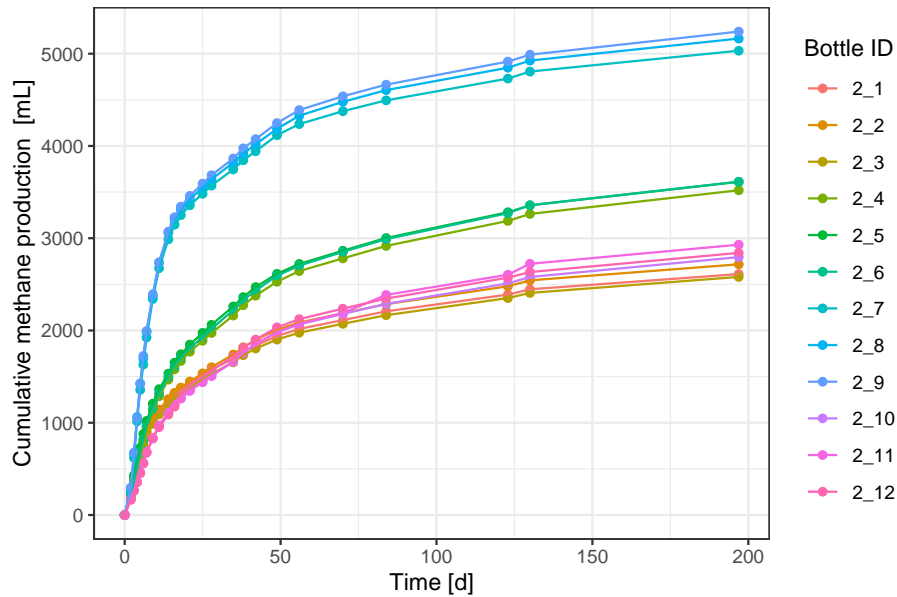
The returned data frame has been restructured to ‘‘longcombo’’ structure and contains all the original columns in `vol`, plus additional columns from volumetric biogas calculations (refer to Section 3.1).

As with the ‘‘longcombo’’ data example, the results may be easier to understand graphically using the `ggplot` function from the `ggplot2` package.

```

ggplot(cum.prod.l, aes(days, cvCH4, colour = factor(id))) +
  geom_point() +
  geom_line(aes(group = id)) +
  labs(x = "Time [d]", y = "Cumulative methane production [mL]",
       colour = "Bottle ID") +
  theme_bw()

```



### 3.4 Other Arguments

Instead of setting fixed single values for temperature and pressure arguments, these can be taken from columns within `dat`. Fixed data were created by adding temperature and pressure data to the `vol` data frame used in the ‘‘long’’ data structure example. Temperature is set to a fixed value for all observations, whereas pressure is set to vary between observations using `rnorm()` having a mean value of 1 atm and a standard deviation of 0.001. It is possible to have both pressure and temperature values that varies between measurements. Though, the approach (function call) would be similar to the shown below.

```
data("vol")

dim(vol)

## [1] 288 4

head(vol)

##   id      date.time days vol
## 1 2_1 2014-06-07 07:00:00 1.98 393
## 2 2_1 2014-06-08 07:00:00 2.98 260
## 3 2_1 2014-06-09 07:00:00 3.98 245
## 4 2_1 2014-06-10 07:00:00 4.98 225
## 5 2_1 2014-06-11 07:00:00 5.98 200
## 6 2_1 2014-06-12 08:00:00 7.02 175
```



```
summary(vol)
```

```
##      id      date.time      days
## 2_1   : 24   Min.    :2014-06-07 07:00:00   Min.    : 1.98
## 2_2   : 24   1st Qu.:2014-06-13 20:00:00   1st Qu.: 8.52
## 2_3   : 24   Median  :2014-06-28 06:00:00   Median  :22.94
## 2_4   : 24   Mean    :2014-07-16 15:29:22   Mean    :41.33
## 2_5   : 24   3rd Qu.:2014-07-25 22:45:00   3rd Qu.:50.63
## 2_6   : 24   Max.    :2014-12-19 04:30:00   Max.    :196.92
## (Other):144
##      vol
## Min.    : 98.0
## 1st Qu.:171.5
## Median  :225.0
## Mean    :271.7
## 3rd Qu.:300.0
## Max.    :840.0
##
```

```
data("comp")
```

```
dim(comp)
```

```
## [1] 132  4
```

```
head(comp)
```

```
##      id      date.time  days      xCH4
## 516 2_1 2014-06-12 08:00:00  7.02 0.7104731
## 519 2_1 2014-06-19 08:00:00 14.02 0.7024937
## 522 2_1 2014-06-26 05:00:00 20.90 0.6659919
## 524 2_1 2014-07-03 04:00:00 27.85 0.6789466
## 525 2_1 2014-07-10 03:00:00 34.81 0.6951429
## 528 2_1 2014-07-24 04:00:00 48.85 0.6693053
```

```
summary(comp)
```

```
##      id      date.time      days
## 2_1   :11   Min.    :2014-06-12 08:00:00   Min.    : 7.02
## 2_2   :11   1st Qu.:2014-06-26 05:00:00   1st Qu.:20.90
## 2_3   :11   Median  :2014-07-24 04:00:00   Median  :48.85
## 2_4   :11   Mean    :2014-07-31 07:47:43   Mean    :56.01
## 2_5   :11   3rd Qu.:2014-08-28 04:00:00   3rd Qu.:83.85
## 2_6   :11   Max.    :2014-10-13 07:00:00   Max.    :129.98
## (Other):66
##      xCH4
```

```
## Min. :0.5647
## 1st Qu.:0.6393
## Median :0.6598
## Mean :0.6587
## 3rd Qu.:0.6786
## Max. :0.7115
##
```

Adding temperature and pressure to the `dat` data frame.

```
vol$temp <- 35
vol$pres <- NA
vol$pres <- rnorm(vol$pres, mean = 1, sd = 0.001)
```

```
head(vol)
```

```
##   id      date.time days vol temp      pres
## 1 2_1 2014-06-07 07:00:00 1.98 393    35 1.0004518
## 2 2_1 2014-06-08 07:00:00 2.98 260    35 0.9991467
## 3 2_1 2014-06-09 07:00:00 3.98 245    35 0.9981821
## 4 2_1 2014-06-10 07:00:00 4.98 225    35 1.0008298
## 5 2_1 2014-06-11 07:00:00 5.98 200    35 0.9999292
## 6 2_1 2014-06-12 08:00:00 7.02 175    35 1.0005784
```

The arguments required to process these data using `calcBgVol()` are the same as in the ‘‘long’’ data structure example above, except from the `temp` and `pres` arguments. Instead of fixed values, we now need to call a column name from `vol`. Therefore, we set `temp = ‘temp’` and `pres = ‘pres’`

```
cum.prod <- calcBgVol(vol, comp = comp, temp = "temp", pres = "pres",
                      data.struct = "long",
                      time.name = "days", vol.name = "vol", comp.name = "xCH4",
                      extrap = TRUE)
```

```
## Biogas composition is interpolated.
## Using a standard pressure of 1 atm and standard temperature of 0
## C for standardizing volume.
```

The returned data frame has been restructured to ‘‘longcombo’’ structure and contains all the original columns in `vol`, plus additional columns from volumetric biogas calculations. In a similar matter, the `comp` argument can be set as a single value instead of a column or data frame.

By default, time zero rows are added from the `addt0` argument. Time zero rows can be omitted by setting `showt0 = FALSE`.

```

cum.prod <- calcBgVol(vol, comp = comp, temp = "temp", pres = "pres",
  data.struct = "long",
  time.name = "days", vol.name = "vol", comp.name = "xCH4",
  extrap = TRUE, showt0 = FALSE)

## Biogas composition is interpolated.
## Using a standard pressure of 1 atm and standard temperature of 0
## C for standardizing volume.

head(cum.prod)

##   id          date.time days vol temp      pres      xCH4
## 1 2_1 2014-06-07 07:00:00 1.98 393   35 1.0004518 0.7104731
## 2 2_1 2014-06-08 07:00:00 2.98 260   35 0.9991467 0.7104731
## 3 2_1 2014-06-09 07:00:00 3.98 245   35 0.9981821 0.7104731
## 4 2_1 2014-06-10 07:00:00 4.98 225   35 1.0008298 0.7104731
## 5 2_1 2014-06-11 07:00:00 5.98 200   35 0.9999292 0.7104731
## 6 2_1 2014-06-12 08:00:00 7.02 175   35 1.0005784 0.7104731
##      vBg      vCH4      cvBg      cvCH4      rvBg      rvCH4
## 1 329.1043 233.8198 329.1043 233.8198 166.2143 118.0908
## 2 217.4274 154.4763 546.5317 388.2961 217.4274 154.4763
## 3 204.6740 145.4154 751.2057 533.7115 204.6740 145.4154
## 4 188.4939 133.9198 939.6996 667.6313 188.4939 133.9198
## 5 167.3905 118.9265 1107.0901 786.5578 167.3905 118.9265
## 6 146.5674 104.1322 1253.6575 890.6899 140.9302 100.1271

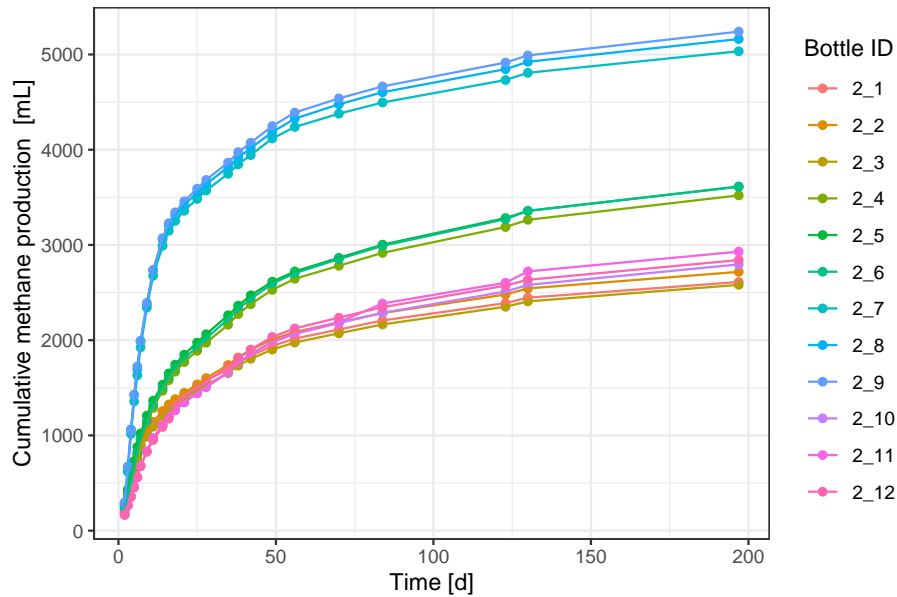
```

The results are shown graphically using the `ggplot` function from the `ggplot2` package.

```

ggplot(cum.prod, aes(days, cvCH4, colour = factor(id))) +
  geom_point() +
  geom_line(aes(group = id)) +
  labs(x = "Time [d]", y = "Cumulative methane production [mL]",
    colour = "Bottle ID") +
  theme_bw()

```



Note the lack of observations in time zero in the plot.

Furthermore, we can avoid adding the time zero values in the first place by setting `addt0 = FALSE`.

```
cum.prod <- calcBgVol(vol, comp = comp, temp = "temp", pres = "pres",
  data.struct = "long",
  time.name = "days", vol.name = "vol", comp.name = "xCH4",
  extrap = TRUE, addt0 = FALSE)
```

```
## Biogas composition is interpolated.
## Using a standard pressure of 1 atm and standard temperature of 0
## C for standardizing volume.
```

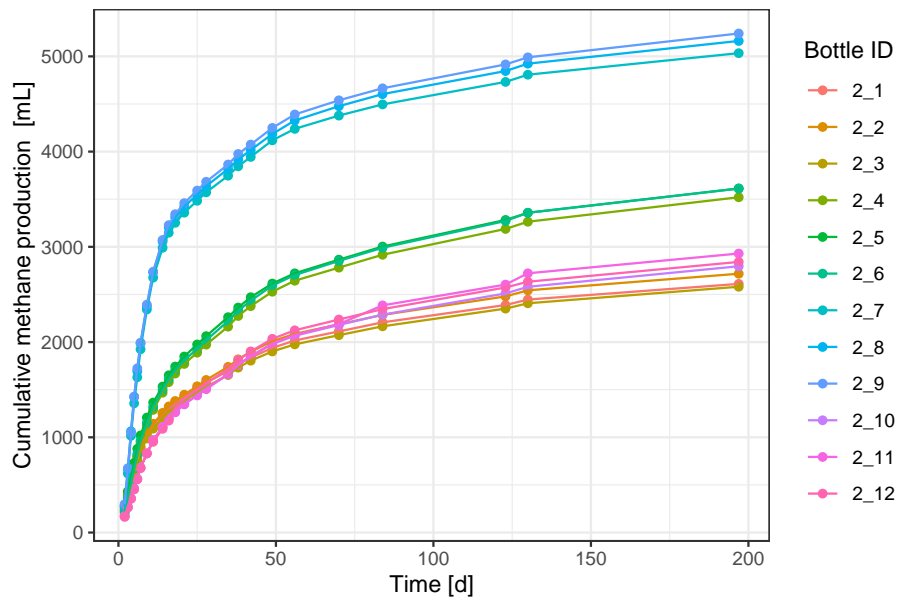
```
head(cum.prod)
```

```
##   id      date.time days  vol  temp    pres    xCH4
## 1 2_1 2014-06-07 07:00:00 1.98 393    35 1.0004518 0.7104731
## 2 2_1 2014-06-08 07:00:00 2.98 260    35 0.9991467 0.7104731
## 3 2_1 2014-06-09 07:00:00 3.98 245    35 0.9981821 0.7104731
## 4 2_1 2014-06-10 07:00:00 4.98 225    35 1.0008298 0.7104731
## 5 2_1 2014-06-11 07:00:00 5.98 200    35 0.9999292 0.7104731
## 6 2_1 2014-06-12 08:00:00 7.02 175    35 1.0005784 0.7104731
##      vBg    vCH4    cvBg    cvCH4    rvBg    rvCH4
## 1 329.1043 233.8198 329.1043 233.8198      NA      NA
## 2 217.4274 154.4763 546.5317 388.2961 217.4274 154.4763
## 3 204.6740 145.4154 751.2057 533.7115 204.6740 145.4154
## 4 188.4939 133.9198 939.6996 667.6313 188.4939 133.9198
```

```
## 5 167.3905 118.9265 1107.0901 786.5578 167.3905 118.9265
## 6 146.5674 104.1322 1253.6575 890.6899 140.9302 100.1271
```

Note how rates are not calculated for the first observations when `addt0 = FALSE`, in contrary to `showt0 = FALSE` where initial rates are calculated. In both cases, cumulative production values for the first observations are omitted from the plot.

```
ggplot(cum.prod, aes(days, cvCH4, colour = factor(id))) +
  geom_point() +
  geom_line(aes(group = id)) +
  labs(x = "Time [d]", y = "Cumulative methane production [mL]",
        colour = "Bottle ID") +
  theme_bw()
```



By default, input data are checked for unreasonable values using the `check` argument. If `check = TRUE`, the input values of `xCH4` in biogas are checked. If data seems unreasonable ( $0 > xCH4 > 1$ ), a warning is returned. In the example below, a single methane value in the composition data frame `comp` has been set to 1.5 for the purpose of demonstrating the function of the `check` argument.

```
data("vol")
data("comp")

comp[10,"xCH4"] <- 1.5
cum.prod <- calcBgVol(vol, comp = comp, temp = 35, pres = 1,
```

```

      data.struct = "long",
      time.name = "days", vol.name = "vol", comp.name = "xCH4",
      extrap = TRUE)

## Biogas composition is interpolated.
## Warning in cumBgDataPrep(dat = dat, dat.type = "vol", dat.name =
vol.name, : Methane concentration was > 1.0 mol/mol for at least one
observation, so is assumed to be a percentage, and was corrected by
dividing by 100. Range of new values: 0.00564685459645082-0.015
## Using a standard pressure of 1 atm and standard temperature of 0
C for standardizing volume.

```

Note the warning about methane concentration being outside the range 0 - 1 (exceeding 1 mol/mol) for at least one observation. By default it is assumed to be a percentage and corrected. If this warning is not provided (`check= TRUE`), we are not made aware of this default correction of the observation.

```

cum.prod <- calcBgVol(vol, comp = comp, temp = 35, pres = 1,
  data.struct = "long",
  time.name = "days", vol.name = "vol", comp.name = "xCH4",
  extrap = TRUE, check = FALSE)

## Biogas composition is interpolated.
## Using a standard pressure of 1 atm and standard temperature of 0
C for standardizing volume.

```

By default, the unit of temperature and pressure values are set as degree celsius and atm, respectively. As stated above, these can be set using the `unit.temp` and `unit.pres` arguments. For the purpose of demonstrating the function of the unit arguments, the already fixed pressure and temperature values in `vol` have been manipulated to be K and kPa, respectively.

```

vol$temp <- 35 + 273.15
vol$pres <- NA
vol$pres <- rnorm(vol$pres, mean = 101.325, sd = 0.101325)

head(vol)

##   id      date.time days vol  temp  pres
## 1 2_1 2014-06-07 07:00:00 1.98 393 308.15 101.5705
## 2 2_1 2014-06-08 07:00:00 2.98 260 308.15 101.2465
## 3 2_1 2014-06-09 07:00:00 3.98 245 308.15 101.3776
## 4 2_1 2014-06-10 07:00:00 4.98 225 308.15 101.2777
## 5 2_1 2014-06-11 07:00:00 5.98 200 308.15 101.4288
## 6 2_1 2014-06-12 08:00:00 7.02 175 308.15 101.2345

```



in question. Though, this operation often requires additional setup information (e.g. inoculum and substrate mass), which is most commonly provided in an external data frame (`setup`).

## 5 Function Internals

In general, `cumBg*`(`)` functions consist of four sections: argument checks, restructure and sorting data, interpolation if needed, and biogas standardization and calculations. Restructuring and sorting data and interpolation are handled by the external functions `cumBgDataPrep()` and `interp()`, respectively. From `interp()` gas composition, cumulative biogas production, and other variables can be interpolated to a specified time if required. From `cumBgDataPrep()` ‘‘wide’’ and ‘‘long’’ data structure are restructured to ‘‘longcombo’’ data, which is required for `calcBgVol()` to process data. Additionally, data are sorted, headspace is added if provided, and composition data is corrected if it seems to be a percentage. If data of interest are mixed (interval and cumulative, `empty.name = TRUE`), these will be sorted and biogas volume standardized within `cumBgDataPrep()` to obtain interval data only. Subsequently, the restructured and sorted data is standardized in `calcBgVol()` by the external function `stdVol()`, if not already standardized.

`calcBgVol()` accepts all data structures, but the volumetric calculation methods within `calcBgVol()` only process ‘‘longcombo’’ data. ‘‘wide’’ and ‘‘long’’ data are restructured internally by the ‘‘low-level’’ function `cumBgDataPrep()`. `cumBgDataPrep()` sorts and restructures biogas data for further calculation of cumulative biogas and methane production and production rates. The `data.struct` argument is internally passed to `cumBgDataPrep()` and restructured to ‘‘longcombo’’ structure prior to being processed by volumetric calculation methods.

Table 1: Operations done with the ‘‘low-level’’ functions in `calcBgVol()`. All functions are vectorized. See help files for more details.

Operation	Function
Standardize gas volume	<code>stdVol()</code>
Interpolate composition etc.	<code>interp()</code>
Structurize and sort data	<code>cumBgDataPrep()</code>

## References

- [1] B.K. Richards, R.J. Cummings, T.E. White, and W.J. Jewell. Methods of assessing microbial activity and inhibition under anaerobic conditions: a literature review. *Rev Environ Sci Biotechnol*, 3: 93-115, 2004.



- [2] J. Filer, H. H. Ding and S. Chang Biochemical Methane Potential (BMP) Assay Method for Anaerobic Digestion Research. *Water*, 11, 921, 2019.
- [3] Hafner, S.D., Koch, K., Carrere, H., Astals, S., Weinrich, S., Rennuit, C. 2018 Software for biogas research: Tools for measurement and prediction of methane production. *SoftwareX* 7: 205-210
- [4] B.K. Richards, R.J. Cummings, T.E. White, and W.J. Jewell. Methods for kinetic-analysis of methane fermentation in high solids biomass digesters. *Biomass & Bioenergy*, 1(2):65–73, 1991.