

MT4608 Sampling Theory

Computer Practical 1

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In this practical we are going to use the R package **survey** to reproduce some of the results we have calculated manually in class and in tutorials, and then to ...

1 Getting started with the package **sampling**

The **sampling** package should already be installed in RStudio Cloud, but if you are working on another machine, you need to install it from CRAN.

We start by loading the **survey** package and the caribou dataset of Table 2.1 of the notes. The dataset is in the package **mt4608**, written specifically for this module¹.

```
library(survey) # load survey package
library(mt4608) # load this module package
```

The caribou survey data are contained in the package **mt4608**, and can be accessed as follows:

```
data(caribou) # get the dataset
caribou # look at it
help(caribou) # get a description of it
```

Calculate the total area and total number of strips in the survey region:

```
A = unique(caribou$area[caribou$stratum==1]) +
    unique(caribou$area[caribou$stratum==2])
N = unique(caribou$N[caribou$stratum==1]) +
    unique(caribou$N[caribou$stratum==2])
```

Estimate the mean number of animals per survey strip, together with a 95% confidence interval. To do this with the **survey** package, you first need to associate a survey design with the survey data, using the command **svydesign**. Look at the help for **svydesign** to understand more about the arguments; here **ids=~1** tells it that there are no clusters (you have to specify **ids**), and **fpc** if the “finite population correction” factor that we call *f* in the notes.

¹This package is not as robust as packages on CRAN, so please be a bit patient/tolerant - things may go a little wrong in situations that I did not anticipate when writing it.

```
n = dim(caribou)[1] # sample size
srs <- svydesign(ids=~1,data=caribou,fpc=rep(n/N,n))
```

Now using the survey data and the design, we can estimate the population mean and 95% confidence interval very easily using the commands `svymean` and `confint`. To do this you have to tell `svymean` which column of the data frame is the y -variable (the response). In our case it is the `count`:

```
ybar <- svymean(~count,srs)
ybar
confint(ybar)
```

1.1 Exercise 1

1. By doing appropriate calculations with the data, decide whether the function `confint` assumed that the sample mean has a normal distribution or a t-distribution.
2. Estimate the total number of caribou in the survey region, together with a 95% confidence interval