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Exercise 07: Control structures

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Introductory exercises

for loops

Write a for loop to compute

$$\sum_{i=1}^{10} 0.5^i.$$

```
q <- 0.5
n <- 10

res <- 0
for(i in 1:n){
  res <- res + q^i
}
res
```

```
## [1] 0.9990234
```

Now move this for loop structure inside a function named `geomSeries` with the arguments `q` and `n` so that the function returns

$$f(q, n) = \sum_{i=1}^n q^i.$$

```
geomSeries <- function(q, n) {
  res <- 0
  for(i in 1:n){
    res <- res + q^i
  }
  res
}
```

Your function should return

```
geomSeries(0.5, 10)
```

```
## [1] 0.9990234
```

while loops

Suppose S_n is defined as

$$S_n = \sum_{i=1}^n 0.5^i.$$

Use a while loop to calculate S_n until

$$|S_n - S_{n-1}| < 10^{-6}.$$

```
q <- 0.5
tol <- 1e-6

res <- 0
i <- 1
term <- q^i
```



```
while(abs(term) >= tol){  
  # summation  
  res <- res + term  
  
  # increase of loop variable  
  i <- i + 1  
  # new additive term  
  term <- q^i  
}  
res
```

```
## [1] 0.9999981
```

Functions, default values and if-else-structures

Now alter the function `geomSeries` that it accepts the arguments `q`, `n` and `tol`. Use the following default values:

```
geomSeries <- function(q, n = NULL, tol = 1e-6){  
  # your code here  
}
```

The wanted functionality of the modified function `geomSeries` is as follows:

- if the argument `n` is used then `geomSeries` should return

$$f(q, n) = \sum_{i=1}^n q^i.$$

- if the argument `n` is not used (i.e. `is.null(n) == TRUE`), than `geomSeries` should return an approximation of the infinite sum

$$f(q) = \sum_{i=1}^{\infty} q^i$$

until $|S_n - S_{n-1}| < 'tol'$.

```
geomSeries <- function(q, n = NULL, tol = 1.e-6) {  
  
  # Check if argument `n` is used or not  
  if(!is.null(n)){  
    # calculate for loop  
    res <- 0  
    for(i in 1:n){  
      res <- res + q^i  
    }  
  } else {  
    # calculate while loop  
    res <- 0  
    i <- 1  
    term <- q^i  
    while(abs(term) >= tol){
```



```
# summation
res <- res + term

# increase of loop variable
i <- i + 1
# new additive term
term <- q^i
}
res
}
return(res)
}
```

Have fun with programming in R! Your modified function `geomSeries` should work as follows:

```
geomSeries(0.5, n = 3)
```

```
## [1] 0.875
```

```
geomSeries(0.5, n = 5, tol = 1e-4)
```

```
## [1] 0.96875
```

```
geomSeries(0.5, tol = 1e-4)
```

```
## [1] 0.9998779
```



The \sqrt{N} law

The core function

Write a function `meanVarSdSe` that takes a numeric vector `x` as argument. The function should return a named vector that contains the mean, the variance, the standard deviation `sd` and the standard error `se` of `x`. The standard error is defined as

$$se(x) = \frac{sd(x)}{\sqrt{\#x}}, \quad (1)$$

where $\#x$ denotes the cardinality, i.e. the number of elements contained in `x`.

The code should have the following structure

and return a named vector according to

```
meanVarSdSe <- function(x){  
  n <- length(x)  
  c(mean = mean(x),  
    var = var(x),  
    sd = sd(x),  
    se = sd(x)/sqrt(n))  
}
```

```
x <- 1:100  
meanVarSdSe(x)
```

```
##      mean      var      sd      se  
## 50.500000 841.666667 29.011492 2.901149
```

You can use the functions `mean`, `var`, `sd` and `length`. Check the help files for these functions for further arguments that can be used optionally.

Look at the following code sequence. What result do you expect?

```
x <- c(NA, 1:100)  
meanVarSdSe(x)
```

Now run the code. Explain the result. Extend the function definition of `meanVarSdSe` with the argument `...`, as is illustrated as follows:

```
meanVarSdSe <- function(x, ...){  
  c(mean = mean(x, ...),  
    var = var(x, ...),  
    sd = sd(x, ...),  
    se = sd(x, ...)/sqrt(sum(!is.na(x))))  
}
```

so that the `na.rm = TRUE` argument can be passed optionally to the functions `mean`, `var` and `sd`. What is the correct value for $\#x$ in the case of missing values? Use `sum(!is.na(x))` as denominator in eq. (1). Read the help page for the function `is.na()`. The optimized function should return



```
meanVarSdSe( c(x, NA), na.rm = TRUE)
```

```
##      mean      var      sd      se
## 50.500000 841.666667 29.011492 2.901149
```

Convergence and Control Structures

The \sqrt{N} Law: the precision of the sample average improves with the square root of the sample size N . Simulate 10^6 Poisson distributed random numbers with expectation value $\lambda = 100$ via

```
set.seed(1) # why?
x <- rpois(n = 1e6, lambda = 100)
```

Write a loop (repeat or while) which calculates the mean, variance, standard deviation and standard error of the first N elements of x until ' $se \leq 0.05$ '. Start with $N = 2$ and multiply N after each iteration by a factor of 2. Store N and the calculated values for each iteration as rows in a matrix named `result`. Use the function `meanVarSdSe` developed in exercise,. Make sure that the column names of your result matrix match the following output. Your matrix should look like

```
tol <- 0.05
factor <- 2
```

```
result <- NULL
n <- 2
repeat{
  result <- rbind(result, c(N=n, meanVarSdSe(x[1:n])))
  if(result[nrow(result), "se"] <= tol) break
  n <- n * factor
}
```

```
tail(result) # see ?tail for details
```

```
##      N      mean      var      sd      se
## [11,] 2048  99.90137 104.63414 10.22908 0.22603293
## [12,] 4096  99.97803 101.65813 10.08257 0.15754008
## [13,] 8192 100.02466  99.13747  9.95678 0.11000792
## [14,] 16384 100.00885 100.58669 10.02929 0.07835384
## [15,] 32768 100.04257 101.13827 10.05675 0.05555623
## [16,] 65536  99.97209 100.32065 10.01602 0.03912508
```

Question: Why is a `for` loop not optimal for the specific task?

Check: Are these values plausible?

- Extract the columns `se` and `N` from the `result` matrix and store in individual vectors named `se` and `N`.

```
se <- result[, "se"]
N <- result[, "N"]
```

- Plot the standard error versus the sample size in a scatter plot. Make use of the additional argument `log = 'xy'`. Discuss the result.



```
plot(N, se, log = "xy")
```

- Now perform a linear regression and model the logarithm of the standard error se as a linear function of $\log(N)$. What coefficient \hat{b}_1 do you expect according equ. (1)?

```
lm01 <- lm(log(se) ~ log(N))
summary(lm01)
```

```
##
## Call:
## lm(formula = log(se) ~ log(N))
##
## Residuals:
##      Min       1Q   Median       3Q      Max
## -0.17541 -0.05660  0.01691  0.02669  0.37975
##
## Coefficients:
##              Estimate Std. Error t value Pr(>|t|)
## (Intercept)  2.26831     0.06804   33.34 9.71e-15 ***
## log(N)       -0.49841     0.01015  -49.10 < 2e-16 ***
## ---
## Signif. codes:
## 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
##
## Residual standard error: 0.1297 on 14 degrees of freedom
## Multiple R-squared:  0.9942, Adjusted R-squared:  0.9938
## F-statistic: 2410 on 1 and 14 DF, p-value: < 2.2e-16
```

Something's not working - why?

Lower the break condition to ' $se \leq 0.01$ '. Run the changed code. Make sure that you reinitialize N and $result$ before running the loop. What happens? Compare N and $\text{length}(x)$. Which expression causes the error?