Class06

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General format for R script: ADD <- FUNCTION(x,y){x+y} add <- function(x,y=1){x+y} What would happen if we add x+yadd(1,1) [1] 2 add(c(100,1,100),1) [1] 101 2 101 add(c(100,1,100),c(100,1,100))[1] 200 2 200 add(10) [1] 11 add(1,1) [1] 2

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Make a function that generates a random nucleotide sequence of any length

```
Nucleotides <- c("A","T","G","C")</pre>
sequence <- sample(Nucleotides, 100, replace = TRUE)</pre>
sequence
  [1] "C" "A" "C" "C" "G" "C" "A" "G" "G" "C" "C" "T" "G" "C" "C" "C" "G"
 [19] "T" "T" "G" "T" "G" "T" "T" "A" "T" "C" "A" "T" "A" "T" "G" "A" "C" "A"
 [37] "G" "G" "T" "C" "C" "A" "G" "A" "C" "T" "C" "A" "C" "T" "T" "A" "G" "C"
 [55] "T" "G" "C" "T" "A" "G" "A" "C" "C" "G" "T" "C" "A" "A" "A" "A" "A" "G" "A"
 [73] "T" "C" "C" "T" "T" "C" "T" "G" "G" "C" "T" "A" "T" "T" "C" "G" "A" "G"
 [91] "G" "A" "A" "G" "C" "G" "G" "T" "T"
Generate_DNA <- function(length){</pre>
  Nucleotides <- c("A","T","G","C")</pre>
  sequence <- sample(Nucleotides, size = length, replace=TRUE)</pre>
  return(sequence)
Generate_DNA(10)
 [1] "G" "C" "G" "T" "C" "C" "T" "A" "G" "T"
This working snippet ROCKS! Now i can make it into an angelic function
library(bio3d)
unique(bio3d::aa.table$aa1)[1:20]
 [1] "A" "R" "N" "D" "C" "Q" "E" "G" "H" "I" "I" "K" "M" "F" "P" "S" "T" "W" "Y"
[20] "V"
amino_acids <- unique(bio3d::aa.table$aa1)[1:20]</pre>
sample(amino_acids, size = 30, replace = TRUE)
 [1] "N" "T" "O" "H" "C" "F" "P" "H" "T" "K" "H" "L" "H" "L" "T" "T" "K" "D" "H"
[20] "V" "Y" "Y" "E" "F" "L" "Y" "K" "T" "N" "F"
Generate_AA <- function(length){</pre>
  amino_acids <- unique(bio3d::aa.table$aa1)[1:20]</pre>
  string <- sample(amino_acids, size = length, replace = TRUE)</pre>
  string <- paste(string, collapse = "")</pre>
  return(string)
Generate_AA(100)
```

[1] "TIGGCTNWSTECGFHYISLVTIHQSGVTKRENQPRSVFTDERNNYNRCHGYNGYWSGTTNCVFNWFVVFEYYNWNNVWWFGSNVVIF.

I want to generate random sequences of length 6 - 12

AIAHNG
>id.7
FTQMKLL
>id.8
KKRQGAVH
>id.9
LDSTFLIRH
>id.10
CEPMIYRIRC
>id.11
FKMGYGSWKFY
>id.12
SCPFMDLHFDFF