

# class 6

Anqi Feng

my first function

```
add<-function(x,y=1){  
  x+y  
}
```

can i just use this?

```
add(10,10)
```

```
[1] 20
```

```
add(10)
```

```
[1] 11
```

```
add(1,1)
```

```
[1] 2
```

```
add(x=1,y=100)
```

```
[1] 101
```

```
add(c(100,1,100),1)
```

```
[1] 101    2 101
```

```
add(10)
```

```
[1] 11
```

Q;make a function: “generate\_DNA()” that makes a random nucleotide sequence of any length

```
bases <- c("A", "C", "G", "T")
sequence <- sample(bases, size=5, replace=TRUE)
```

That is my working lines, now i can make it into a function

```
generate_dna<-function(length){
  bases <- c("A", "C", "G", "T")
  sequence <- sample(bases, size=length, replace=TRUE)
  return(sequence)
}
```

```
generate_dna(10)
```

```
[1] "C" "G" "T" "G" "G" "G" "T" "G" "G" "T"
```

```
bio3d::aa.table
```

|     | aa3 | aa1 | mass    | formula      | name          |
|-----|-----|-----|---------|--------------|---------------|
| ALA | ALA | A   | 71.078  | C3 H5 N 01   | Alanine       |
| ARG | ARG | R   | 157.194 | C6 H13 N4 01 | Arginine      |
| ASN | ASN | N   | 114.103 | C4 H6 N2 02  | Asparagine    |
| ASP | ASP | D   | 114.079 | C4 H4 N 03   | Aspartic Acid |
| CYS | CYS | C   | 103.143 | C3 H5 N 01 S | Cystein       |
| GLN | GLN | Q   | 117.126 | C4 H9 N2 02  | Glutamine     |
| GLU | GLU | E   | 128.106 | C5 H6 N 03   | Glutamic Acid |
| GLY | GLY | G   | 57.051  | C2 H3 N 01   | Glycine       |
| HIS | HIS | H   | 137.139 | C6 H7 N3 01  | Histidine     |
| ILE | ILE | I   | 113.158 | C6 H11 N 01  | Isoleucine    |
| LEU | LEU | L   | 113.158 | C6 H11 N 01  | Leucine       |
| LYS | LYS | K   | 129.180 | C6 H13 N2 01 | Lysine        |
| MET | MET | M   | 131.196 | C5 H9 N 01 S | Methionine    |
| PHE | PHE | F   | 147.174 | C9 H9 N 01   | Phenylalanine |

|     |     |   |         |               |                                  |
|-----|-----|---|---------|---------------|----------------------------------|
| PRO | PRO | P | 97.115  | C5 H7 N 01    | Proline                          |
| SER | SER | S | 87.077  | C3 H5 N 02    | Serine                           |
| THR | THR | T | 101.104 | C4 H7 N 02    | Threonine                        |
| TRP | TRP | W | 186.210 | C11 H10 N2 01 | Tryptophan                       |
| TYR | TYR | Y | 163.173 | C9 H9 N 02    | Tyrosine                         |
| VAL | VAL | V | 99.131  | C5 H9 N 01    | Valine                           |
| ABA | ABA | X | 85.104  | C4 H7 N1 01   | alpha-aminobutyric acid          |
| ASH | ASH | D | 115.087 | C4 H5 N 03    | Aspartic acid Neutral            |
| CIR | CIR | R | 157.170 | C6 H11 N3 02  | citrulline                       |
| CME | CME | C | 179.260 | C5 H9 N 02 S2 | s,s-(2-hydroxyethyl)thiocysteine |
| CMT | CMT | C | 115.154 | C4 H5 N 01 S  | o-methylcysteine                 |
| CSD | CSD | C | 134.134 | C3 H4 N 03 S  | s-cysteinesulfinic acid          |
| CSO | CSO | C | 119.142 | C3 H5 N 02 S  | s-hydroxycysteine                |
| CSW | CSW | C | 135.142 | C3 H5 N 03 S  | cysteine-s-dioxide               |
| CSX | CSX | C | 119.142 | C3 H5 N 02 S  | s-oxy cysteine                   |
| CYM | CYM | C | 102.135 | C3 H4 N 01 S  | Cystein Negative                 |
| CYX | CYX | C | 102.135 | C3 H4 N 01 S  | Cystein SSbond                   |
| DDE | DDE | H | 280.346 | C13 H22 N5 02 | diphthamide                      |
| GLH | GLH | E | 129.114 | C5 H7 N 03    | Glutatmic acid Neutral           |
| HID | HID | H | 137.139 | C6 H7 N3 01   | Histidine                        |
| HIE | HIE | H | 137.139 | C6 H7 N3 01   | Histidine                        |
| HIP | HIP | H | 138.147 | C6 H8 N3 01   | Histidine Positive               |
| HSD | HSD | H | 137.139 | C6 H7 N3 01   | Histidine                        |
| HSE | HSE | H | 137.139 | C6 H7 N3 01   | Histidine                        |
| HSP | HSP | H | 138.147 | C6 H8 N3 01   | Histidine Positive               |
| IAS | IAS | D | 115.087 | C4 H5 N 03    | beta-aspartyl                    |
| KCX | KCX | K | 172.182 | C7 H12 N2 03  | lysine nz-carboxylic acid        |
| LYN | LYN | K | 129.180 | C6 H13 N2 01  | Lysine Neutral                   |
| MHO | MHO | M | 147.195 | C5 H9 N 02 S  | s-oxymethionine                  |
| MLY | MLY | K | 156.225 | C8 H16 N2 01  | n-dimethyl-lysine                |
| MSE | MSE | M | 178.091 | C5 H9 N 01 SE | selenomethionine                 |
| OCS | OCS | C | 151.141 | C3 H5 N 04 S  | cysteinesulfonic acid            |
| PFF | PFF | F | 165.164 | C9 H8 F N 01  | 4-fluoro-l-phenylalanine         |
| PTR | PTR | Y | 243.153 | C9 H10 N 05 P | o-phosphotyrosine                |
| SEP | SEP | S | 167.057 | C3 H6 N 05 P  | phosphoserine                    |
| TPO | TPO | T | 181.084 | C4 H8 N 05 P  | phosphothreonine                 |

```
unique(bio3d::aa.table$aa1)[1:20]
```

```
[1] "A" "R" "N" "D" "C" "Q" "E" "G" "H" "I" "L" "K" "M" "F" "P" "S" "T" "W" "Y"
[20] "V"
```

```
generate_protein<-function(l){
  aas<-unique(bio3d::aa.table$aa1)[1:20]
  poly <- sample(aas, size=l, replace=TRUE)
  poly<-paste(poly, collapse = "")
  return(poly)
}
```

```
generate_protein(10)
```

```
[1] "CGDQVMISGA"
```

Generate protein sequences of length 6 to 12

```
answer<-sapply(6:12, generate_protein)
```

```
cat(paste( ">id.", 6:12, "\n", answer, sep="", "\n"))
```

```
>id.6
PRGSSC
>id.7
PYFDTES
>id.8
FIAYDKKH
>id.9
EARHTDTRS
>id.10
PMKHDTISIE
>id.11
LAYDKVTPQGK
>id.12
RVNCEECWAEAG
```