

# Problem1

Yufei Yin

## Question 1a, (4 points):

Create a function in R named counts. This function should take as parameters a numeric vector  $x$  and also a number indicating a number of bins  $n$ . The function will consider the range  $[\min(x), \max(x)]$ , and then consider a partition of this interval into  $n$  non-overlapping equally sized half open intervals:  $I_1 = [\min(x), b_1)$ ;  $I_2 = [b_1, b_2)$ , ...,  $I_n = [b_{n-1}, \max(x))$ . Note that since the intervals are equally sized, the value of  $b_i$  is constrained. The function will then return a vector of length  $n$  such that the  $i$ -th element of this vector is the number of coordinates of  $x$  that lie in the interval  $I_i$ . Provide the code for this function: It should be of the following form:

```
counts = function(...) {  
  ...  
  return(...)  
}
```

```
counts = function (x, n) {  
  lowvalue = floor(min(x))  
  highvalue = ceiling(max(x))  
  width2use = (highvalue-lowvalue)/n  
  BinEdges = seq(from = lowvalue, to = highvalue, by = width2use)  
  a = BinEdges[-length(BinEdges)]  
  b = BinEdges[-1]  
  out = rep(NA, length(a))  
  for(index in seq_along(a)){  
    out[index]=sum(x>=a[index] & x<b[index])  
  }  
  return(out)  
}
```

## Question 1b, (3 points):

Create a function in R called `histo`. This function should take the same `x` and `n` parameters as the `count` function you wrote in the previous question, and it should plot a histogram of the vector `x` with `n` bins. The body of this function may make use of the `count` function you wrote in the previous question. The only plotting functions you may make use of are the `plot` function and the `lines` function ('just add an egg'). You may not make use of the `hist` function. Provide your code for this function. Hint: There are several ways to do this, one way would be to create a new and empty plot with:

```
plot(1,
     type="n",
     xlab="x",
     ylab="Counts",
     xlim=c(...),
     ylim=c(...))
```

Then make a for loop through the bins and call `lines` in the body of the for loop so that three lines (delinating the left, top, and right of the box) are drawn. As before (although this time without a return statement), your code should be in the following form:

```
histo = function( ... ){
  ...
}
```

```
histo = function(x, n){
  lowvalue = floor(min(x))
  highvalue = ceiling(max(x))
  width2use = (highvalue-lowvalue)/n
  BinEdges = seq(from = lowvalue, to = highvalue, by = width2use)
  a = BinEdges[-length(BinEdges)]
  b = BinEdges[-1]

  # counts function from question 1a
  out = counts(x,n)

  # corner case
  if (highvalue == max(x)){
    out[length(out)] = sum(x>=a[length(out)] & x<=b[length(out)])
  }

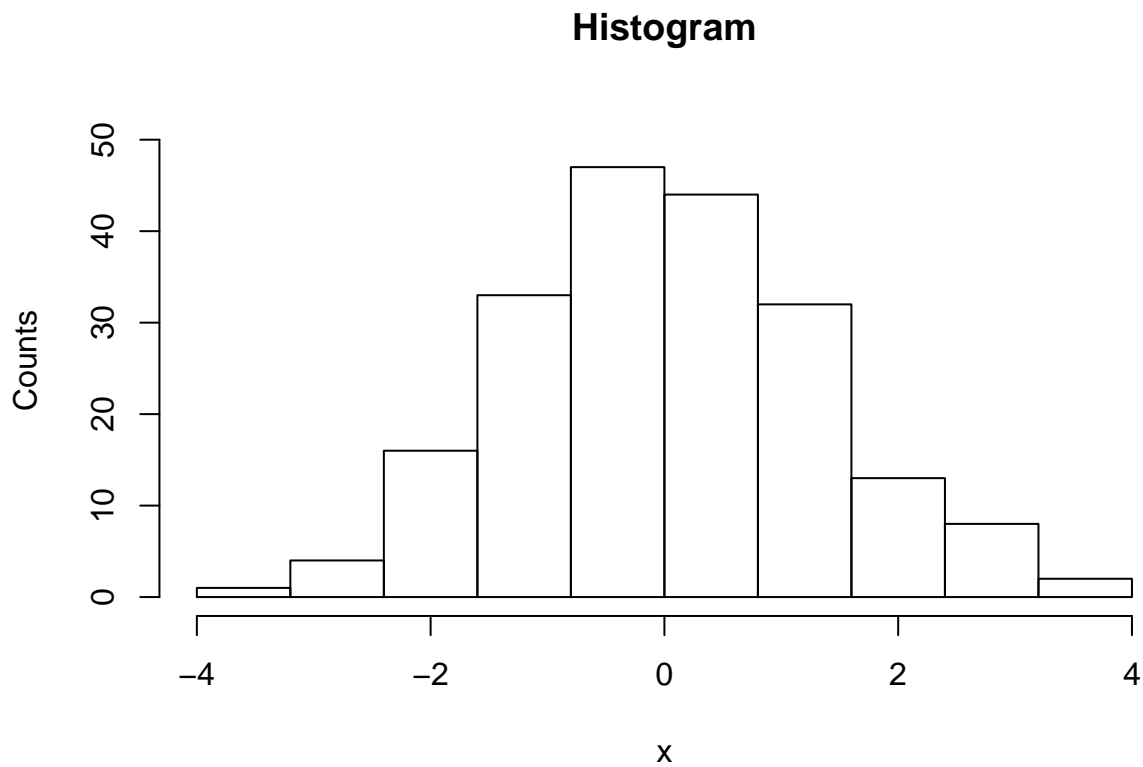
  # plot
  plot(1,
       type="n",
       xlab="x",
       ylab="Counts",
       main="Histogram",
       bty="n",
       xlim = c(lowvalue, highvalue),
       ylim = c(0,max(out)*1.1))
  for (i in seq_along(a)){
    lines(c(a[i],a[i],b[i],b[i],a[i]),c(0,out[i],out[i],0,0))
  }
}
```

### Question 1c, (2 points):

We will now test the `histo` function. Create a vector of length 200 such that the first 100 elements are independent draws from a normal distribution with mean -1 and variance 1 and the second 100 elements are independent draws from a normal distribution with mean 1 and variance 1. Call `histo` on this vector with 10 bins and provide a graphic including the resulting plot.

```
set.seed(1)
v1 = rnorm(n = 100, mean = -1, sd = 1)
v2 = rnorm(n = 100, mean = 1, sd = 1)
x = c(v1, v2)
```

```
histo(x = x, n = 10)
```



### Question 1d, (2 points):

We will now test a corner case of the `histo` function. Call `histo` on the vector  $x = (0, 0, 0, 1, 1, 2)$  with 3 bins, and provide a graphic including the resulting plot. Note that this is a corner case because the middle values lie at the boundary between two bins, and so their contribution to the histogram's height requires that the half-open nature of the intervals be respected.

```
x = c(0, 0, 0, 1, 1, 2)
n = 3
histo(x = x, n = n)
```

