## **Problem 5**

(a) R code for generating simple randomization and statified block randomization is given below.

For block randomization, we apply a more technique. We introduce three more arrays: one to keep track of the current block size for each stratum, one to keep track of the current number of patients in a block for each stratum, and one to keep track of the number of patients assigned to group 1 in the current block for each stratum. We assign patients within each stratum to treatments on a rolling basis. Once a block is full, we randomly choose the next block size to be either 2 or 4.

```
blockSize = rep(0,4)
currCapl = rep(0,4)

for(i in 1:20){
    # Interpret the 0's and 1's as TRUE and FALSE, respectively
    if(dat[i,1] & dat[i,2]){ #(V1, V2) = (1,1)
        j = 1
    }else if(dat[i,1] & !dat[i,2]){ #(V1, V2) = (1,0)
        j = 2
    }else if(!dat[i,1] & dat[i,2]){ #(V1, V2) = (0,1)
        j = 3
```

```
}else{ #(V1, V2) = (0,0)
    j = 4
}

if(runif(1) < (blockSize[j]/2-currAssign[j])/(blockSize[j]-currCap[j])){
    dat[i,4] = 1
        currAssign[j] = currAssign[j]+1
}else{
    dat[i,4] = 0
}

currCap[j] = currCap[j]+1

if(currCap[j]==blockSize[j]){
    currCap[j] = currAssign[j] = 0
    blockSize[j] = ifelse(runif(1)<0.5,2,4)
}</pre>
```

The results of both randomization schemes for this particular seed are given below. Different seeds may give different results.

```
v1 v2 simple block
1
   1
      0
              1
2
      1
              1
                    0
    1
3
    0
      0
              1
                    0
4
   1
      0
              0
                    1
5
      0
              0
   1
                    1
6
   0
      0
              0
                    1
7
   1
      0
              1
                    0
      1
8
   1
              1
                    1
9
   0
      1
              1
                    0
10
      1
              1
                    1
   0
11
   0 1
              0
                    0
12
   1 0
              1
                    1
13
   1 1
                    0
              1
14
              1
                    1
   0 0
15
              0
   1
      1
                    1
16
   0 1
              0
                    1
17
     0
              0
                    0
   1
                    0
18
   0 0
              1
19
   1
     0
              1
                    1
20
              0
                    0
   0
      1
```

(b) For the above randomization, we check the balance in terms of absolute difference between patients assigned treatment A and patients assigned treatment B. R code is given below:

```
# (i) Overall balance
print(sprintf("Overall imbalance from simple rand: %d",
```

```
abs (sum(dat[,3]==1)-sum(dat[,3]==0)))
print(sprintf("Overall imbalance from stratified block rand: %d",
              abs (sum (dat [, 4] == 1) -sum (dat [, 4] == 0))))
"Overall imbalance from simple rand: 4"
"Overall imbalance from stratified block rand: 0"
# (ii) Across strata
stratum = rep(0,20)
for(i in 1:20) {
  if(dat[i,1] \& dat[i,2]) \{ \#(V1, V2) = (1,1) \}
    stratum[i] = 1
  else if(dat[i,1] & !dat[i,2]) { #(V1, V2) = (1,0)}
    stratum[i] = 2
  else if(!dat[i,1] & dat[i,2]) { #(V1, V2) = (0,1)}
    stratum[i] = 3
  else{ #(V1, V2) = (0,0)}
    stratum[i] = 4
 }
}
for(i in 1:4) {
 print(sprintf("For stratum %d:",i))
  st = which(stratum==i)
  print(sprintf("Imbalance from simple randomization: %d",
                abs (sum(dat[st, 3] == 1) - sum(dat[st, 3] == 0)))
 print(sprintf("Imbalance from stratified block randomization: %d",
                abs (sum (dat [st, 4] == 1) - sum (dat [st, 4] == 0))))
}
"For stratum 1:"
"Imbalance from simple randomization: 2"
"Imbalance from stratified block randomization: 0"
"For stratum 2:"
"Imbalance from simple randomization: 1"
"Imbalance from stratified block randomization: 1"
"For stratum 3:"
"Imbalance from simple randomization: 1"
"Imbalance from stratified block randomization: 1"
"For stratum 4:"
"Imbalance from simple randomization: 2"
"Imbalance from stratified block randomization: 0"
# (iii) Marginally across prognostic factors
for(i in 0:1){
 print(sprintf("For V1 = %d:",i))
  st = which(dat[,1]==i)
  print(sprintf("Imbalance from simple rand: %d",
```

```
abs(sum(dat[st,3]==1)-sum(dat[st,3]==0))))
 print(sprintf("Imbalance from stratified block randomization: %d",
                abs(sum(dat[st, 4] == 1) - sum(dat[st, 4] == 0))))
}
"For V1 = 0:"
"Imbalance from simple rand: 1"
"Imbalance from stratified block randomization: 1"
"For V1 = 1:"
"Imbalance from simple rand: 3"
"Imbalance from stratified block randomization: 1"
for(i in 0:1) {
 print(sprintf("For V2 = %d:",i))
  st = which(dat[,2]==i)
 print(sprintf("Imbalance from simple rand: %d",
                abs(sum(dat[st,3]==1)-sum(dat[st,3]==0))))
 print(sprintf("Imbalance from stratified block randomization: %d",
                abs(sum(dat[st, 4] == 1) - sum(dat[st, 4] == 0))))
}
"For V2 = 0:"
"Imbalance from simple rand: 3"
"Imbalance from stratified block randomization: 1"
"For V2 = 1:"
"Imbalance from simple rand: 1"
"Imbalance from stratified block randomization: 1"
```

Thus, at least for this seed, it seems that the stratified block randomization produced better balance overall, across the strata (V1, V2) = (1, 1) and (V1, V2) = (0, 0), and marginally across V1 = 1 and V2 = 0. In general, one should see better balance in all three categories, though individual results will vary.