STAT542 Coding3 Bonus

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Generate Samples from an HMM

```
T = 200
A0 = rbind(c(0.8, 0.2),
           c(0.2, 0.8))
B0 = rbind(c(0.1, 0.2, 0.7),
           c(0.4, 0.3, 0.3))
w0 = c(0.5, 0.5)
para0 = list(mz = 2, mx = 3, w = w0,
           A = AO, B = BO)
genHMM = function(para, n){
  # return n samples from HMM with parameter = para
 z = para mz
 mx = para$mx
  w = para$w
  A = para A
  B = para$B
  Z = rep(0, n)
  X = rep(0, n)
  ## MY CODE: generate Z[1]
  Z[1] = sample(1:2, size = 1, prob = w, replace = TRUE)
  ## MY CODE: generate Z[i]
  for(i in 2:n){
    Z[i] = sample(1:2, size = 1, prob = A[Z[i-1],], replace = TRUE)
  ## MY CODE: generate X[i]
  for(i in 1:n){
    X[i] = sample(1:3, size = 1, prob = B[Z[i],], replace = TRUE)
  }
  return(X)
```

```
data = genHMM(para0, T)
```

The Baum-Welch (i.e., EM) Algorihtm

Define forward and backward functions

```
forward.prob = function(x, para){
  # Output the forward probability matrix alp
  # alp: T by mz, (t, i) entry = P(x_{1:t}, Z_t = i)
 T = length(x)
 mz = para$mz
 A = para A
 B = para$B
  w = para$w
 alp = matrix(0, T, mz)
  # fill in the first row of alp
 alp[1,] = w * B[, x[1]]
  # Recursively compute the remaining rows of alp
 for(t in 2:T){
   tmp = alp[t-1, ] %*% A
   alp[t,] = tmp * B[, x[t]]
 return(alp)
}
backward.prob = function(x, para){
 \# Output the backward probability matrix beta
  # beta: T by mz, (t, i) entry = P(x_{1:t}, Z_{t} = i)
 T = length(x)
 mz = para$mz
 A = para A
 B = para\$B
 w = para$w
 beta = matrix(1, T, mz)
  # The last row of beta is all 1.
  # Recursively compute the previous rows of beta
  for(t in (T-1):1){
   tmp = as.matrix(beta[t+1, ] * B[, x[t+1]]) # make tmp a column vector
   beta[t, ] = t(A %*% tmp)
 }
 return(beta)
```

Define BW.onestep function

```
BW.onestep = function(x, para){
    # Input:
```

```
# x: T-by-1 observation sequence
  # para: mx, mz, and current para values for
      A: initial estimate for mz-by-mz transition matrix
      B: initial estimate for mz-by-mx emission matrix
       w: initial estimate for mz-by-1 initial distribution over Z_1
  # Output the updated parameters after one iteration
  \# We DO NOT update the initial distribution w
  T = length(x)
  mz = para$mz
  mx = para$mx
  A = para A
  B = para$B
  w = para$w
  alp = forward.prob(x, para)
  beta = backward.prob(x, para)
  myGamma = array(0, dim=c(mz, mz, T-1))
  ## MY CODE:
  ## Compute gamma_t(i,j) P(Z[t] = i, Z[t+1]=j),
  ## for t=1:T-1, i=1:mz, j=1:mz,
  ## which are stored an array, myGamma
  for(t in 1:(T-1)){
    for(i in 1:mz){
      for(j in 1:mz){
        myGamma[i,j,t] = alp[t,i]*A[i,j]*B[j,x[(t+1)]]*beta[(t+1),j]
    }
  }
  # M-step for parameter A
  A = rowSums(myGamma, dims = 2)
  A = A/rowSums(A)
  # M-step for parameter B
  tmp = apply(myGamma, c(1, 3), sum) # mz-by-(T-1)
  tmp = cbind(tmp, colSums(myGamma[, , T-1]))
  for(1 in 1:mx){
   B[, 1] = rowSums(tmp[, which(x==1)])
  B = B/rowSums(B)
  para$A = A
  para$B = B
  return(para)
```

Define myBM function

```
myBW = function(x, para, n.iter = 100){
    # Input:
    # x: T-by-1 observation sequence
```

```
# para: initial parameter value
# Output updated para value (A and B; we do not update w)

for(i in 1:n.iter){
   para = BW.onestep(x, para)
}
return(para)
}
```

The Viterbi Algorihtm

```
myViterbi = function(x, para){
  # Output: most likely sequence of Z (T-by-1)
  T = length(x)
  mz = para$mz
  A = para A
  B = para$B
  w = para$w
  log.A = log(A)
  log.w = log(w)
  log.B = log(B)
  # Compute delta (in log-scale)
  delta = matrix(0, T, mz)
  # fill in the first row of delta
  delta[1, ] = log.w + log.B[, x[1]]
  ## MY CODE:
  ## Recursively compute the remaining rows of delta
  for(t in 1:(T-1)){
   tmp = NULL
   for(j in 1:mz){
      tmp = rbind(tmp, delta[t,j] + log.A[j,])
   log.max = apply(tmp, 2, function(x) max(x))
    delta[(t+1), ] = log.max + log.B[,x[(t+1)]]
  }
  # Compute most prob sequence Z
  Z = rep(0, T)
  # start with the last entry of Z
  Z[T] = which.max(delta[T, ])
  ## MY CODE:
  ## Recursively compute the remaining entries of Z
  for(t in T:2){
    Z[t-1] = which.max(delta[(t-1), ] + log.A[, Z[t]])
 return(Z)
```

}

Test My Algorithm

```
data = genHMM(para0, T)
mz = 2
mx = 3
ini.w = rep(1, mz); ini.w = ini.w / sum(ini.w)
ini.A = matrix(1, 2, 2); ini.A = ini.A / rowSums(ini.A)
ini.B = matrix(1:6, 2, 3); ini.B = ini.B / rowSums(ini.B)
ini.para = list(mz = 2, mx = 3, w = ini.w,
                A = ini.A, B = ini.B)
myout = myBW(data, ini.para, n.iter = 100)
myout.Z = myViterbi(data, myout)
myout.Z[myout.Z==1] = 'A'
myout.Z[myout.Z==2] = 'B'
library(HMM)
hmm0 = initHMM(c("A", "B"), c(1, 2, 3),
              startProbs = ini.w,
              transProbs = ini.A,
              emissionProbs = ini.B)
Rout = baumWelch(hmm0, data, maxIterations=100, delta=1E-9, pseudoCount=0)
Rout.Z = viterbi(Rout$hmm, data)
myout$A
             [,1]
                       [,2]
## [1,] 0.4526640 0.5473360
## [2,] 0.4564762 0.5435238
Rout$hmm$transProbs
##
## from
##
      A 0.4526640 0.5473360
      B 0.4564762 0.5435238
They're the same.
myout$B
             [,1]
                       [,2]
## [1,] 0.2454993 0.2414352 0.5130655
## [2,] 0.3639114 0.2388004 0.3972881
Rout$hmm$emissionProbs
         symbols
## states
                  1
        A 0.2454993 0.2414352 0.5130655
       B 0.3639114 0.2388004 0.3972881
##
They're the same.
```

```
cbind(Rout.Z, myout.Z)[c(1:10, 180:200), ]
##
         Rout.Z myout.Z
                 "A"
##
    [1,] "A"
                 "B"
##
    [2,] "B"
    [3,] "A"
                 "A"
##
##
    [4,] "B"
                 "B"
    [5,] "A"
                 "A"
##
    [6,] "A"
                 "A"
##
                 "B"
    [7,] "B"
##
    [8,] "B"
                 "B"
##
##
    [9,] "B"
                 "B"
## [10,] "A"
                 "A"
## [11,]
         "A"
                 "A"
  [12,] "B"
                 "B"
##
                 "B"
## [13,] "B"
                 "B"
## [14,] "B"
## [15,] "A"
                 "A"
## [16,] "A"
                 "A"
## [17,] "B"
                 "B"
## [18,] "A"
                 "A"
## [19,] "B"
                 "B"
## [20,] "B"
                 "B"
## [21,] "A"
                 "A"
## [22,] "B"
                 "B"
## [23,] "B"
                 "B"
                 "B"
## [24,] "B"
## [25,] "A"
                 "A"
## [26,] "B"
                 "B"
## [27,] "B"
                 "B"
                 "A"
## [28,] "A"
## [29,] "B"
                 "B"
                 "B"
## [30,] "B"
## [31,] "A"
                 "A"
sum(Rout.Z != myout.Z)
```

[1] 0

The output from my Viterbi algorithm and the one from HMM are exactly the same.