STAT7400 HW6, 2017

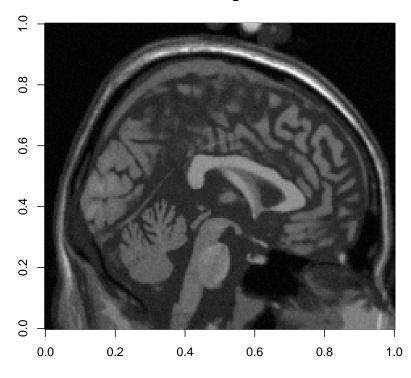
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 $Mar\ 5,\ 2017$

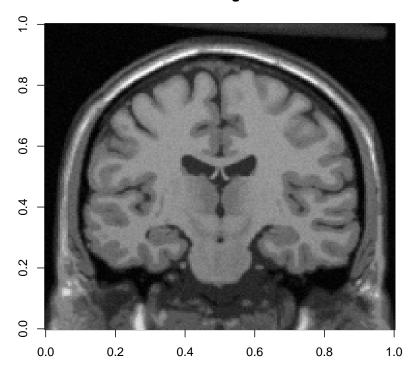
Problem 2

(a) Three middle slice graphs along different axes are given below.

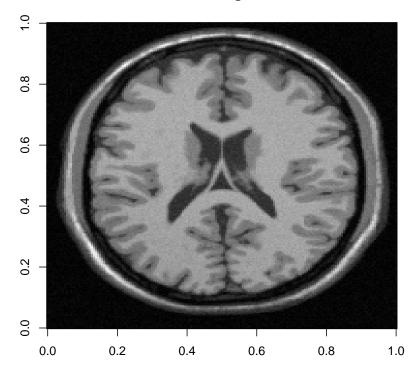
Middle slice along with X-axis



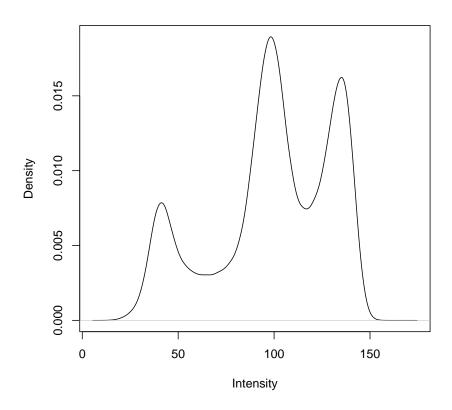
Middle slice along with Y-axis



Middle slice along with Z-axis



(b) The plot of the estimate of the density of the image intensities for voxels in the brain is given below.

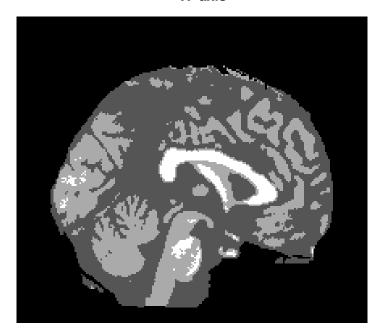


(c) The initial valuee are set by function initOtsuin package mritc.

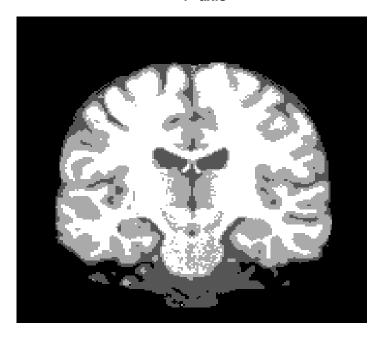
```
p.new <- colSums.Ez / sum(colSums.Ez)</pre>
      ## pack up result
      list (mu = mu.new, sigma = sigma.new, p = p.new)
+ }
> diff <- function(x, y) {</pre>
      max(abs(x - y)) / (1 + max(abs(x), abs(y)))
+ }
> converge.check <- function(theta.new, theta,
                           err, iter, iter.max) {
      diff.mu <- diff(theta.new$mu, theta$mu)</pre>
      diff.sigma <- diff(theta.new$sigma, theta$sigma)
      diff.p <- diff(theta.new$p, theta$p)</pre>
      flag <- 0
      if (diff.mu < err[1] && diff.sigma < err[2]</pre>
          && diff.p < err[3] || iter > iter.max)
+
          flag <- 1
+
      flag
+ }
> EMmix <- function(x, theta, err, iter.max) {
      iter <- 0
      repeat {
          iter <- iter + 1
+
          theta.new <- EMmix1(x, theta)</pre>
          flag <- converge.check(theta.new, theta, err, iter, iter.max)
          if (flag) break
          theta <- theta.new
      list(mu = theta.new\$mu, sigma = theta.new\$sigma, p = theta.new\$p)
+ }
> library(mritc)
> init <- initOtsu(T1.mask, 2)</pre>
> theta <- list(mu=init$mu, sigma = init$sigma, p = init$prop)</pre>
> err <- rep(1e-08, 3)
> iter.max <- 200
> theta.est1 <- EMmix(T1.mask, theta, err, iter.max)</pre>
> data.frame(mu = theta.est1$mu, sigma = theta.est1$sigma, p = theta.est1$p)
         mu
                 sigma
1 45.76594 10.223615 0.1766328
2 98.98909 13.719448 0.5502270
3 133.95977 6.460461 0.2731402
```

(d) Based on the classification results, three graphs are shown below.

Middle slice along with X-axis



Middle slice along with Y-axis



Middle slice along with Z-axis



- (e) By summaryRprof, we can see that most time is spent on EMmix1 and outer.
 - > Rprof(tmp <- tempfile())</pre>
 - > theta.est1 <- EMmix(T1.mask, theta, err, iter.max)</pre>
 - > Rprof()
 - > summaryRprof(tmp)

\$by.self

	self.time	self.pct	total.time	total.pct
".External"	57.84	34.88	57.84	34.88
"array"	24.32	14.67	24.32	14.67
"dnorm"	19.70	11.88	77.54	46.76
"aperm.default"	11.84	7.14	11.84	7.14
"_"	11.08	6.68	11.08	6.68
"outer"	9.30	5.61	108.80	65.61
"sweep"	8.20	4.95	50.72	30.59
"FUN"	7.62	4.60	99.50	60.00
"rowSums"	5.80	3.50	5.80	3.50
" * "	4.58	2.76	4.58	2.76
"colSums"	3.30	1.99	6.30	3.80

п∧п	1.68	1.01	1.68	1.01
"aperm"	0.52	0.31	36.68	22.12
" % % "	0.02	0.01	0.02	0.01
"parent.frame"	0.02	0.01	0.02	0.01
\$by.total				
	total.time	total.pct	self.time	self.pct
" <anonymous>"</anonymous>	165.82	100.00	0.00	0.00
"doTryCatch"	165.82	100.00	0.00	0.00
"EMmix"	165.82	100.00	0.00	0.00
"EMmix1"	165.82	100.00	0.00	0.00
"eval"	165.82	100.00	0.00	0.00
"evalFunc"	165.82	100.00	0.00	0.00
"try"	165.82	100.00	0.00	0.00
"tryCatch"	165.82	100.00	0.00	0.00
"tryCatchList"	165.82	100.00	0.00	0.00
"tryCatchOne"	165.82	100.00	0.00	0.00
"withVisible"	165.82	100.00	0.00	0.00
"outer"	108.80	65.61	9.30	5.61
"FUN"	99.50	60.00	7.62	4.60
"dnorm"	77.54	46.76	19.70	11.88
".External"	57.84	34.88	57.84	34.88
"sweep"	50.72	30.59	8.20	4.95
"aperm"	36.68	22.12	0.52	0.31
"array"	24.32	14.67	24.32	14.67
"aperm.default"	11.84	7.14	11.84	7.14
"_"	11.08	6.68	11.08	6.68
"colSums"	6.30	3.80	3.30	1.99
"rowSums"	5.80	3.50	5.80	3.50
" * "	4.58	2.76	4.58	2.76
"is.data.frame"	3.00	1.81	0.00	0.00
11 ∧ 11	1.68	1.01	1.68	1.01
" 응 응 "	0.02	0.01	0.02	0.01
"parent.frame"	0.02	0.01	0.02	0.01
"match.fun"	0.02	0.01	0.00	0.00
	_			

\$sample.interval

[1] 0.02

\$sampling.time
[1] 165.82

(f) Since the intensities can take on only a small number of distinct values, we should only compute the values for unique intensity values and then multiply these values by the counts of each unique intensity value. The improved EM function is as

follows.

```
> EMmix1c <- function(x.c, c, theta) {
      mu <- theta$mu
      sigma <- theta$sigma
      p <- theta$p
      M \leftarrow length(mu)
      ## E step
      Ez \leftarrow outer(x.c, 1 : M, function(x, i) p[i] *
                        dnorm(x, mu[i], sigma[i]))
      Ez \leftarrow sweep(Ez, 1, rowSums(Ez), "/")
      Ez.c <- sweep(Ez, 1, c, "*")</pre>
      colSums.Ez.c <- colSums(Ez.c)</pre>
+
      ## M step
+
      xp.c <- sweep(Ez.c, 1, x.c, "*")
      mu.new <- colSums(xp.c) / colSums.Ez.c</pre>
      sqRes.c \leftarrow outer(x.c, mu.new, function(x, m) (x - m) ^ 2)
      sigma.new <- sqrt(colSums(Ez.c * sqRes.c) / colSums.Ez.c)</pre>
      p.new <- colSums.Ez.c / sum(colSums.Ez.c)</pre>
      ## pack up result
+
      list (mu = mu.new, sigma = sigma.new, p = p.new)
+ }
> EMmixc <- function(x, theta, err, iter.max) {
      iter <- 0
      c \leftarrow table(x)
      x.c \leftarrow sort(unique(x))
      repeat {
           iter <- iter + 1
           theta.new <- EMmix1c(x.c, c, theta)
           flag <- converge.check(theta.new, theta, err, iter, iter.max)
           if (flag) break
           theta <- theta.new
      theta.new
+ }
```

Then we fit the normal mixture model again with the improve EM algorithm, the results are exactly the same as before.

```
mu.c sigma.c p.c
1 45.76594 10.223615 0.1766328
2 98.98909 13.719448 0.5502270
3 133.95977 6.460461 0.2731402
```

By summaryRprof, we can see that the improved algorithm is quite faster than before.

- > Rprof(tmp <- tempfile())</pre>
- > theta.est2 <- EMmixc(T1.mask, theta, err, iter.max)</pre>
- > Rprof()
- > summaryRprof(tmp)

\$by.self

	self.time	self.pct	total.time	total.pct
"as.character"	0.30	57.69	0.30	57.69
"table"	0.08	15.38	0.44	84.62
"match"	0.04	7.69	0.04	7.69
"unique.default"	0.04	7.69	0.04	7.69
"sweep"	0.02	3.85	0.06	11.54
"any"	0.02	3.85	0.02	3.85
"match.fun"	0.02	3.85	0.02	3.85

\$by.total

	total.time	total.pct	self.time	self.pct
" <anonymous>"</anonymous>	0.52	100.00	0.00	0.00
"doTryCatch"	0.52	100.00	0.00	0.00
"EMmixc"	0.52	100.00	0.00	0.00
"eval"	0.52	100.00	0.00	0.00
"evalFunc"	0.52	100.00	0.00	0.00
"try"	0.52	100.00	0.00	0.00
"tryCatch"	0.52	100.00	0.00	0.00
"tryCatchList"	0.52	100.00	0.00	0.00
"tryCatchOne"	0.52	100.00	0.00	0.00
"withVisible"	0.52	100.00	0.00	0.00
"table"	0.44	84.62	0.08	15.38
"factor"	0.36	69.23	0.00	0.00
"as.character"	0.30	57.69	0.30	57.69
"sweep"	0.06	11.54	0.02	3.85
"EMmix1c"	0.06	11.54	0.00	0.00
"match"	0.04	7.69	0.04	7.69
"unique.default"	0.04	7.69	0.04	7.69
"aperm.default"	0.04	7.69	0.00	0.00
"aperm"	0.04	7.69	0.00	0.00
"order"	0.04	7.69	0.00	0.00
"unique"	0.04	7.69	0.00	0.00

"any"	0.02	3.85	0.02	3.85
"match.fun"	0.02	3.85	0.02	3.85
"lapply"	0.02	3.85	0.00	0.00
"sort"	0.02	3.85	0.00	0.00
"unlist"	0.02	3.85	0.00	0.00

\$sample.interval
[1] 0.02

\$sampling.time
[1] 0.52