Solutions to Homework 1

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Problem 1

a. Let S be the event that an email is spam and M be the event that an email is marked as spam. Then

$$P(M) = P(M|S)P(S) + P(M|S^c)P(S^c) = (0.92)(0.08) + (0.03)(0.92) = 0.1012$$

```
(0.92)*(0.08) + (0.03)*(0.92)
```

[1] 0.1012

b. We can calculate P(S|M) via Bayes' theorem:

$$P(S|M) = \frac{P(M|S)P(S)}{P(M|S)P(S) + P(M|S^c)P(S^c)} = 0.73$$

```
(0.92 * 0.08)/((0.92)*(0.08) + (0.03)*(0.92))
```

[1] 0.7272727

Problem 2

```
# Making the data matrix
vals \leftarrow c(0.018, 0.035, 0.031, 0.008, 0.018, 0.002, 0.112, 0.064, 0.032, 0.069, 0.001, 0.066, 0.094,
joint <- matrix(vals, nrow = 5, byrow=TRUE, dimnames = list(c("farm", "operatives", "craf</pre>
joint
                farm operatives craftsmen sales professional
               0.018 0.035
                                0.031 0.008
## farm
                                                      0.018
## operatives 0.002
                         0.112
                                   0.064 0.032
                                                      0.069
               0.001
                         0.066
                                   0.094 0.032
                                                      0.084
## craftsmen
## sales
               0.001
                        0.018
                                   0.019 0.010
                                                      0.051
## professional 0.001
                       0.029
                                   0.032 0.043
                                                      0.130
```

a) Let F be the father's occupation and S the son's. Then the marginal probability distribution of the father's occupation is given by $P(F = f_i) = \sum_{s_i \in S} P(F = f_i, S = s_i)$. The marginal probability distribution of the father's occupation:

```
mfather <- numeric(length = 5)
rvarb <- c("farm", "operatives", "craftsmen", "sales", "professional")
for(i in 1:5)
{
    mfather[i] = sum(joint[i,])
}
names(mfather) <- rvarb
mfather</pre>
```

```
## farm operatives craftsmen sales professional
## 0.110 0.279 0.277 0.099 0.235
```

b) The marginal probability distribution of the son's occupation:

```
mson <- numeric(length = 5)
rvarb <- c("farm", "operatives", "craftsmen", "sales", "professional")
for(i in 1:5)
{
    mson[i] = sum(joint[,i])
}
names(mson) <- rvarb
mson</pre>
```

```
## farm operatives craftsmen sales professional
## 0.023 0.260 0.240 0.125 0.352
```

c) The conditional distribution $P(S = s_i | F = f_i)$ is given as

$$P(S = s_i | F = f_i) = \frac{P(S = s_i, F = f_i)}{P(F = f_i)}$$

Hence the conditional distribution of the son's occupation given that the father is a farmer is as follows:

```
cson <- joint["farm",]/mfather["farm"]
cson</pre>
```

```
## farm operatives craftsmen sales professional
## 0.16363636 0.31818182 0.28181818 0.07272727 0.16363636
```

d) Similarly, the father's conditional distribution given that the son is a farmer is as follows:

```
cfather <- joint[,"farm"]/mson["farm"]
cfather</pre>
```

```
## farm operatives craftsmen sales professional
## 0.78260870 0.08695652 0.04347826 0.04347826 0.04347826
```

Problem 3

Set Y $N(\mu, \sigma)$ and $Z = \frac{X-\mu}{\sigma}$. Then we consider the cumulative distribution function $F_Z(z)$, which is given as follows:

$$F_Z(z) = P(Z \le z) = P(\frac{X - \mu}{\sigma} \le z) = P(X \le \sigma z + \mu) = \int_{-\infty}^{\sigma z + \mu} \frac{1}{\sigma \sqrt{2\pi}} e^{-\frac{1}{2} \left(\frac{x - \mu}{\sigma}\right)^2} dx$$

Then we know that the density function $f_Z(z) = F'_z(z)$ so we have that

$$f_Z(Z) = \frac{d}{dz} \int_{-\infty}^{\sigma z + \mu} \frac{1}{\sigma \sqrt{2\pi}} e^{-\frac{1}{2} \left(\frac{x - \mu}{\sigma}\right)^2} dx = \frac{\sigma}{\sigma \sqrt{2\pi}} e^{-\frac{1}{2} \left(\frac{\sigma z + \mu - \mu}{\sigma}\right)^2} dx = \frac{1}{\sigma \sqrt{2\pi}} e^{-\frac{1}{2}z^2} dx$$

Thus Z has a Gaussian density function with $\mu = 0$ and $\sigma = 1$ and so Z N(0,1)

Problem 4

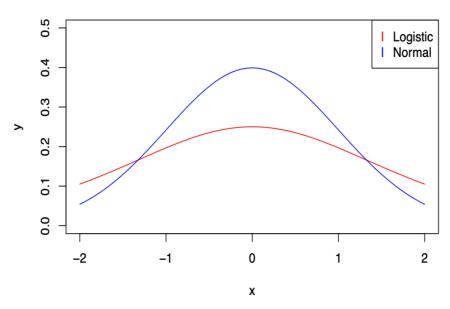
From the conditional independence formula we have that

$$\begin{split} P(X_1 = x_1, \dots, Y_m = y_m | \mu_X, \sigma_X, \mu_Y, \sigma_Y) \\ &= \prod_{i=1}^n P(X_i = x_i | \mu_X, \sigma_X, \mu_Y, \sigma_Y) \prod_{i=1}^m P(Y_i = y_i | \mu_X, \sigma_X, \mu_Y, \sigma_Y) \\ &= \prod_{i=1}^n P(X_i = x_i | \mu_X, \sigma_X) \prod_{i=1}^m P(Y_i = y_i | \mu_Y, \sigma_Y) \\ &= \frac{1}{\sigma_X^n (2\pi)^{n/2}} e^{-\frac{1}{2} \frac{\sum_{i=1}^n (x_i - \mu_X)^2}{\sigma_X^2}} \frac{1}{\sigma_Y^n (2\pi)^{m/2}} e^{-\frac{1}{2} \frac{\sum_{i=1}^m (y_i - \mu_Y)^2}{\sigma_Y^2}} \\ &= \frac{1}{\sigma_X^n \sigma_Y^n (2\pi)^{(n+m)/2}} e^{-\frac{1}{2} \left(\frac{\sum_{i=1}^n (x_i - \mu_X)^2}{\sigma_X^2} + \frac{\sum_{i=1}^m (y_i - \mu_Y)^2}{\sigma_Y^2} \right)} \end{split}$$

Problem 5

Let
$$F(x) = \frac{e^x}{1+e^x} = \frac{1}{1+e^{-x}}$$
. Then $f(x) = F'(x) = \frac{e^{-x}}{(1+e^{-x})^2}$

Comparing pdfs of logistic and normal distributions



The logistic distribution is more likely to sample extreme values as the probabilities at the extremes are higher than those given by the normal distribution.

Problem 6

Here we calculate the posterior distribution given n = 20 and y = 8:

```
require(knitr)

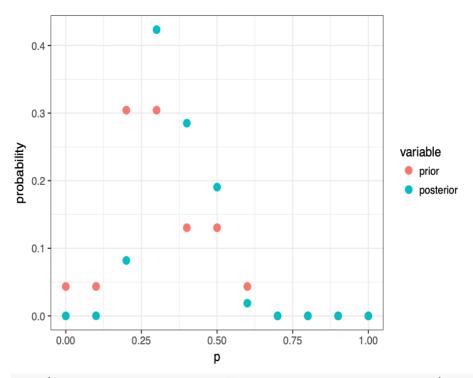
## Loading required package: knitr

#Entering data and the prior probabilities
priorvalues <- c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1)
priorprob <- c(1/23, 1/23, 7/23, 7/23, 3/23, 3/23, 1/23, 0/23, 0/23, 0/23)

n <- 20
y <- 8

#vector for storing results
jointprob <- numeric(length = length(priorvalues))</pre>
```

```
for(i in 1:length(priorvalues))
  #compute Binomial probability given value of p - likelihood
  binomprob <- dbinom(y, n, p = priorvalues[i])</pre>
  #compute joint probability - posterior
  jointprob[i] <- binomprob * priorprob[i]</pre>
#compute marginal probability of y
pofy <- sum(jointprob)</pre>
#compute posterior probabilities
posteriorprob <- jointprob/pofy</pre>
#visualizing the posterior
#put posterior probabilities in one matrix object for easy viewing
allnighterposterior <- as.data.frame(cbind(priorvalues, priorprob, posteriorprob))</pre>
names(allnighterposterior) <- c("p", "prior", "posterior")</pre>
#list the final posterior distribution, based on our prior derived in class
#allnighterposterior
#plot the prior and posterior probabilities
require(ggplot2)
## Loading required package: ggplot2
require(reshape2)
## Loading required package: reshape2
allnighterposterior_all <- melt(allnighterposterior, id = "p")</pre>
ggplot(allnighterposterior_all, aes(x = p, y = value, colour = variable)) +
  geom_point(size = 3) +
  xlab("p") + ylab("probability") +
 theme_bw(base_size = 12, base_family = "")
```



kable(allnighterposterior, caption="Posterior using one-step updating")

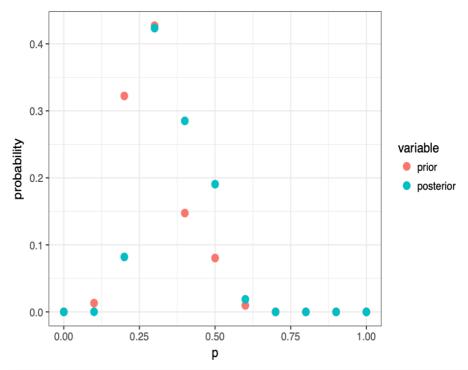
Table 1: Posterior using one-step updating

p	prior	posterior
0.0	0.0434783	0.0000000
0.1	0.0434783	0.0001881
0.2	0.3043478	0.0820219
0.3	0.3043478	0.4234056
0.4	0.1304348	0.2850547
0.5	0.1304348	0.1905607
0.6	0.0434783	0.0187690
0.7	0.0000000	0.0000000
0.8	0.0000000	0.0000000
0.9	0.0000000	0.0000000
1.0	0.0000000	0.0000000

In the following code, we assign the posterior calculated in class as prior and compute the posterior for n=10 and y=5.

```
#Calculating posterior (n=10, y=3) from the prior distribution given in class
require(knitr)
priorvalues \leftarrow c(0, .1, .2, .3, .4, .5, .6, .7, .8, .9, 1)
priorprob <- c(1/23, 1/23, 7/23, 7/23, 3/23, 3/23, 1/23, 0/23, 0/23, 0/23, 0/23)
n <- 10
y <- 3
#vector for storing results
jointprob <- numeric(length = length(priorvalues))</pre>
for(i in 1:length(priorvalues))
  #compute Binomial probability given value of p - likelihood
  binomprob <- dbinom(y, n, p = priorvalues[i])</pre>
  #compute joint probability - posterior
  jointprob[i] <- binomprob * priorprob[i]</pre>
#compute marginal probability of y
pofy <- sum(jointprob)</pre>
#compute posterior probabilities
posteriorprob <- jointprob/pofy</pre>
#Now we will do a sequential update by settling the prior probabilities to the compute
priorprob <- posteriorprob</pre>
n <- 10
y <- 5
jointprob <- numeric(length = length(priorvalues))</pre>
for(i in 1:length(priorvalues))
{
  #compute Binomial probability given value of p - likelihood
  binomprob <- dbinom(y, n, p = priorvalues[i])</pre>
```

```
#compute joint probability - posterior
  jointprob[i] <- binomprob * priorprob[i]</pre>
\#compute\ marginal\ probability\ of\ y
pofy <- sum(jointprob)</pre>
#compute posterior probabilities
posteriorprob <- jointprob/pofy</pre>
allnighterposterior <- as.data.frame(cbind(priorvalues, priorprob, posteriorprob))</pre>
names(allnighterposterior) <- c("p", "prior", "posterior")</pre>
#list the final posterior distribution, based on our prior derived in class
\#allnighterposterior
#plot the prior and posterior probabilities
require(ggplot2)
require(reshape2)
allnighterposterior_all <- melt(allnighterposterior, id = "p")</pre>
ggplot(allnighterposterior_all, aes(x = p, y = value, colour = variable)) +
 geom_point(size = 3) +
  xlab("p") + ylab("probability") +
 theme_bw(base_size = 12, base_family = "")
```



kable(allnighterposterior, caption="Posterior using sequential updating")

Table 2: Posterior using sequential updating

p	prior	posterior
0.0	0.0000000	0.0000000
0.1	0.0131236	0.0001881
0.2	0.3222345	0.0820219
0.3	0.4270731	0.4234056
0.4	0.1474735	0.2850547
0.5	0.0803851	0.1905607
0.6	0.0097102	0.0187690
0.7	0.0000000	0.0000000
0.8	0.0000000	0.0000000
0.9	0.0000000	0.0000000
1.0	0.0000000	0.0000000

Thus we see that the posterior with the bigger data set calculated in one update is the same as the posterior calculated by sequential updates of 10 at a time. This is because we assume that the trials are independent so the fact that out of 10 people 3 stayed up last year does not affect that out of 10 other people, 5 stayed up last year.