Exercise 5

Chutian Chen cc4515; Congcheng Yan cy2550; Mingrui Liu ml4404 3/1/2020

transplant2 <- read.table("./transplant.txt", quote="\"", comment.char="")
colnames(transplant2)<-c("time","type","s")</pre>

(1)

Because the failure rate doesn't dependent on the number of individuals.

Let the probability of individuals that die/relapse be $p_A^{(k)} = p_B^{(k)} = p$ (under H_0).

So
$$P(y^{(k)} = m, n_d^{(k)} = m_d | n_A^{(k)} = m_A, n_B^{(k)} = m_B) = {m_A \choose m} p^m (1-p)^{m_A-m} {m_B \choose m_d-m} p^{m_d-m} (1-p)^{m_B-m_d+m} = {m_A \choose m} {m_B \choose m_d-m} p^{m_d} (1-p)^{m_A+m_B-m_d}$$

$$P(n_d^{(k)} = m_d | n_A^{(k)} = m_A, n_B^{(k)} = m_B) = {\binom{m_A + m_B}{m_d}} p^{m_d} (1 - p)^{m_A + m_B - m_d}$$

So
$$P(y^{(k)} = m|n_d^{(k)} = m_d, n_A^{(k)} = m_A, n_B^{(k)} = m_B)$$

= $\frac{P(y^{(k)} = m, n_d^{(k)} = m_d|n_A^{(k)} = m_A, n_B^{(k)} = m_B)}{P(n_d^{(k)} = m_d|n_A^{(k)} = m_A, n_B^{(k)} = m_B)} = \frac{\binom{m_A}{m}\binom{m_B}{m_d-m}}{\binom{m_A+m_B}{m_d}}$

It's HyperGeometric $\left(n_A^{(k)} + n_B^{(k)}, n_A^{(k)}, n_d^{(k)}\right)$.

(2)

$$P = \frac{\binom{n_A}{y} \binom{n_B}{n_d - y}}{\binom{n}{n_d}}$$

Let $Y_i = \begin{cases} 1 & \text{if the } i \text{ th selection is under treatment A} \\ 0 & \text{otherwise} \end{cases}$

So
$$P(Y_i = 1) = \frac{n_A}{n}$$
, $P(Y_i = 1, Y_j = 1) = \frac{n_A}{n} \frac{n_A - 1}{n}$

$$E^{(k)} = E(\sum Y_i) = n_d EY = \frac{n_A^{(k)} n_d^{(k)}}{n_d^{(k)}}$$

$$Var^{(k)} = Var(\sum Y_i) = n_d Var Y + n_d(n_d - 1)Cov(Y_i, Y_j)$$

$$VarY = \frac{n_A n_B}{n^2}, \ Cov(Y_i, Y_j) = EY_i Y_j - EY_i EY_j = P(Y_i = 1, Y_j = 1) - (\frac{n_A}{n})^2 = -\frac{n_A n_B}{n^2(n-1)}$$

Then
$$Var^{(k)} = \frac{n_A n_B n_d}{n^2} - \frac{n_A n_B n_d (n_d - 1)}{n^2 (n - 1)} = \frac{n_A^{(k)} n_B^{(k)} n_d^{(k)} n_s^{(k)}}{\left(n^{(k)}\right)^2 \left(n^{(k)} - 1\right)}$$

(3)

Let condition
$$\left(n_A^{(k)}, n_B^{(k)}, n_d^{(k)}\right)$$
 be C_i .

$$\mathrm{Var}\left[y^{(k)} - E^{(k)}\right] = \mathrm{Var}[\mathbb{E}[y^{(k)} - E^{(k)}|C_i]] + \mathbb{E}[\mathrm{Var}[y^{(k)} - E^{(k)}|C_i]]$$

As we showed in (2), $\mathbb{E}[y^{(k)} - E^{(k)}|C_i] = 0$, $\text{Var}[y^{(k)} - E^{(k)}|C_i] = V^{(k)}$.

So
$$\operatorname{Var}\left[y^{(k)}-E^{(k)}\right]=E[V^{(k)}]$$

(4)

$$\operatorname{Var}\left[\sum_{k=1}^{K}\left(y^{(k)}-E^{(k)}\right)\right] = \sum_{k=1}^{K}\operatorname{Var}\left[y^{(k)}-E^{(k)}\right] + 2\sum_{i=1}^{K-1}\sum_{j=i+1}^{K}\operatorname{Cov}\left[y^{(i)}-E^{(i)},y^{(j)}-E^{(j)}\right]$$

Let the condition $\left(n_A^{(i)},n_B^{(i)},n_d^{(i)},n_A^{(j)},n_B^{(j)},n_d^{(j)}\right)$ be C_{ij}

With Law of Total Variance, we can get

$$\begin{split} &\operatorname{Cov}\left[y^{(i)} - E^{(i)}, y^{(j)} - E^{(j)}\right] \\ &= \operatorname{Cov}[\mathbb{E}[y^{(i)} - E^{(i)}|C_{ij}], \mathbb{E}[y^{(j)} - E^{(j)}|C_{ij}]] + \mathbb{E}[\operatorname{Cov}[y^{(i)} - E^{(i)}, y^{(j)} - E^{(j)}|C_{ij}]] \\ &\mathbb{E}[y^{(i)} - E^{(i)}|C_{ij}] = \mathbb{E}[y^{(i)} - E^{(i)}|C_{i}] = 0 \\ &\operatorname{So} \operatorname{Cov}[\mathbb{E}[y^{(i)} - E^{(i)}|C_{ij}], \mathbb{E}[y^{(j)} - E^{(j)}|C_{ij}]] = \operatorname{Cov}(0, 0) = 0 \\ &\operatorname{Cov}[y^{(i)} - E^{(i)}, y^{(j)} - E^{(j)}|C_{ij}] = \operatorname{Cov}[y^{(i)}, y^{(j)}|C_{ij}] \end{split}$$

Under the condition C_{ij} , $y^{(i)}$, $y^{(j)}$ are two independent hyperGeometric distributions, so the covariance of them is 0.

Then Cov
$$[y^{(i)} - E^{(i)}, y^{(j)} - E^{(j)}] = 0 \ \forall i \neq j$$

So Var $\left[\sum_{k=1}^{K} (y^{(k)} - E^{(k)})\right] = \sum_{k=1}^{K} \text{Var} \left[y^{(k)} - E^{(k)}\right] = \sum_{k=1}^{K} \mathbb{E} \left[V^{(k)}\right]$

(5)

library(dplyr)

```
n <- nrow(transplant2)</pre>
k <- nrow(transplant2 %>% filter(s==1))
na <- nrow(transplant2 %>% filter(type==1))
nb <- nrow(transplant2 %>% filter(type==2))
df <- transplant2[order(transplant2$time),]</pre>
df <- df %>% filter(s==1)
y = c()
e = c()
v = c()
for (i in 1:k) {
  e[i] \leftarrow na/(na+nb)
  v[i] <- na*nb*(na+nb-1)/(na+nb)^2/(na+nb-1)
  if (df$type[i]==1) {
    y[i] <- 1
   na <- na - 1
  } else {
    y[i] <- 0
    nb <- nb - 1
  }
z <- sum(y-e)/sqrt(sum(v))</pre>
print(pnorm(z))
```

[1] 0.2828261

We can accept H_0 at 0.05 significant level.