

STAT 714 hw 7

Likelihood ratio test (F test) for general linear hypothesis

1. Let $Y_{ijk} = \mu_{ij} + \varepsilon_{ijk}$, $\varepsilon_{ijk} \stackrel{\text{ind}}{\sim} \text{Normal}(0, \sigma^2)$ for $i = 1, \dots, a$ and $j = 1, \dots, b$, $k = 1, \dots, n_{ij}$. In the model, μ_{ij} represents the mean response of experimental units under treatment level i of factor A and treatment level j of factor B , for $i = 1, \dots, a$ and $j = 1, \dots, b$. This is called a two-way factorial design.

(a) Write the model in matrix form $\mathbf{y} = \mathbf{X}\mathbf{b} + \mathbf{e}$.

(b) Assume $a = b = 2$, so that each factor has only two treatment levels. Consider testing the hypothesis $H_0: \mu_{ik} - \mu_{jk} = \mu_{im} - \mu_{jm}$ for all i, j, k, m .

i. Give an interpretation of the null hypothesis.

The null hypothesis states that there is “no interaction” between the two factors; that is, the affect on the response mean of one factor does not depend on the level of the other factor.

ii. Give H_0 in the form $H_0: \mathbf{K}^T\mathbf{b} = \mathbf{m}$.

Let $\mathbf{m} = \mathbf{0}$ and set $\mathbf{K}^T = [1 \ -1 \ -1 \ 1]$ or $\mathbf{K}^T = [-1 \ 1 \ 1 \ -1]$, or any scalar multiple of this.

iii. Let $n_{11} = 5$, $n_{12} = 3$, $n_{21} = 5$, and $n_{22} = 4$ and suppose $\sigma = 1/3$. Give the power of the likelihood ratio test of H_0 when $\mu_{11} = 1$, $\mu_{12} = 2$, $\mu_{21} = 1$, and $\mu_{22} = 3$. Use significance level $\alpha = 0.05$.

```

nn <- c(5,3,5,4)
mu <- c(1,2,1,3)
sigma <- 1/3

a <- 2
b <- 2
N <- sum(nn)

# build X
X <- matrix(0,N,a*b)
m <- 1
for(i in 1:a)
  for(j in 1:b){

    k <- b*(i-1) + j
    X[m:(m + nn[k] - 1),k] <- rep(1,nn[k])
    m <- m + nn[k]

  }

# generate y
e <- rnorm(N,0,sigma)
y <- as.numeric(X %*% mu) + e

# construct K
K <- c(1,-1,-1,1)

# compute noncentrality parameter
Hinv <- solve( t(K) %*% solve(t(X) %*% X ) %*% K)
ncp <- as.numeric(t(t(K) %*% mu) %*% Hinv %*% t(K) %*% mu / sigma^2)

# compute power
alpha <- 0.05
powF <- 1 - pf(qf(1 - alpha,df1=1, df2=N-4),df1=1, df2=N-4, ncp=ncp)
powF
## [1] 0.7984592

```

- iv. Suppose one has not yet collected data, but one wants to know what number of replicates in each group will be necessary to achieve a certain statistical power. Use R to generate a plot showing the power of the likelihood ratio test of H_0 against the value of the signal-to-noise ratio $\text{SNR} = \|\mathbf{K}^T \mathbf{b}\|^2 / \sigma^2$. Include power curves under $n = 3, 4, 5, 6, 7, 8, 9$, where n is the number of replicates at each treatment level combination (so use $n_{ij} = n$ for all i, j).

Note that in the balanced design (equal replications in each treatment group), we have $\mathbf{X}^T\mathbf{X} = n_N$. Moreover, $\mathbf{K}^T\mathbf{K} = 4\mathbf{I}_2$, so we have $[\mathbf{K}^T(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{K}]^{-1} = n/4$. This, together with $\mathbf{m} = \mathbf{0}$, gives the noncentrality parameter

$$\phi = \frac{1}{\sigma^2}(\mathbf{K}^T\mathbf{b} - \mathbf{m})^T[\mathbf{K}^T(\mathbf{X}^T\mathbf{X})^{-1}\mathbf{K}]^{-1}(\mathbf{K}^T\mathbf{b} - \mathbf{m}) = \frac{n}{4\sigma^2}\|\mathbf{K}^T\mathbf{b}\|^2 = \frac{n}{4}\text{SNR}.$$

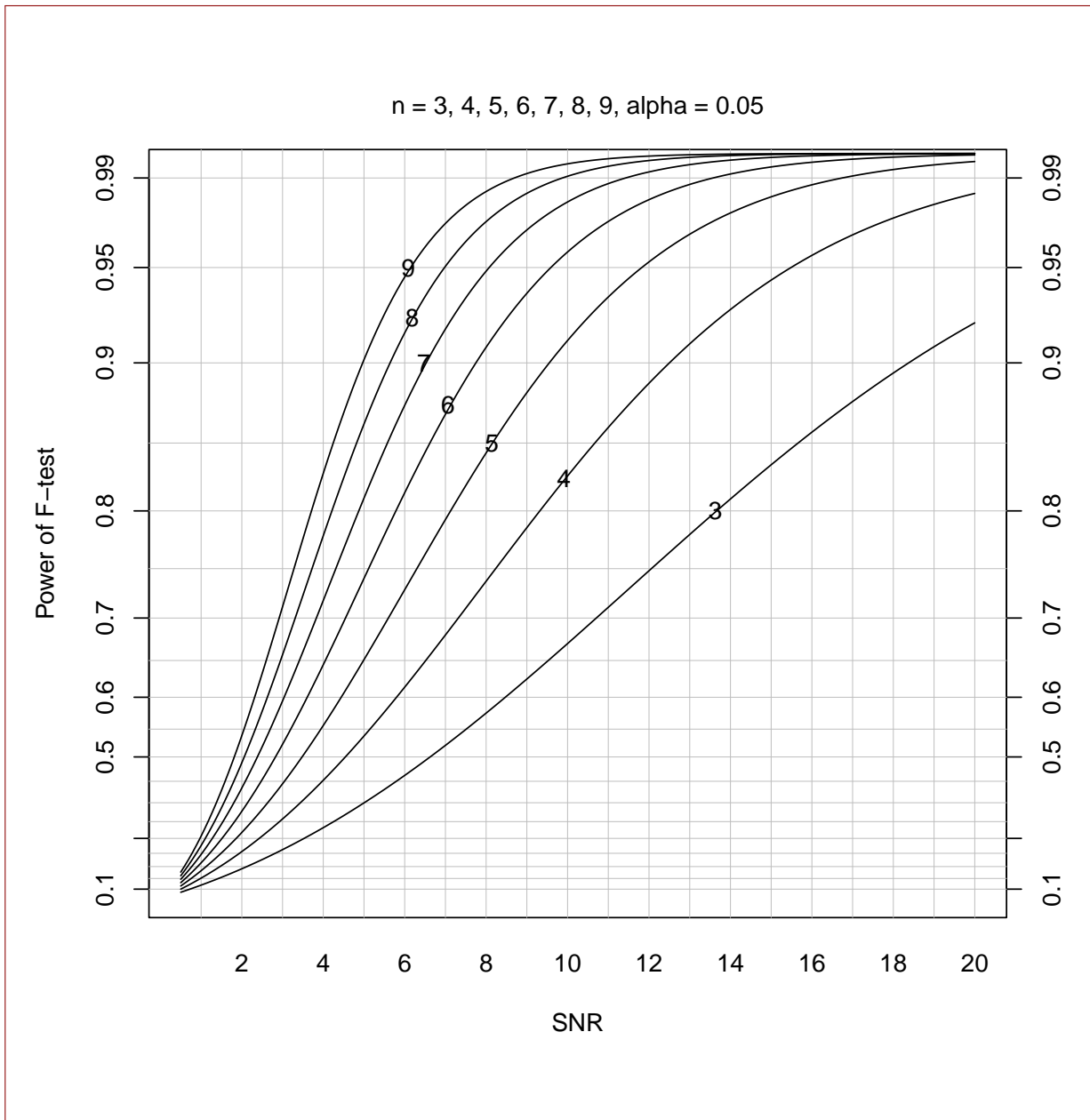
```

nn <- c(3,4,5,6,7,8,9)
snr <- seq(1/2,20,length=200)
powF <- matrix(NA,length(nn),length(snr))
for(i in 1:length(nn)){

  n <- nn[i]
  ncp <- n * snr/4
  powF[i,] <- 1-pf(qf(1-alpha,df1=1,df2=n*4-4),df1=1,df2=n*4-4,ncp=ncp)
}

plot(NA, xlim = range(snr), ylim = exp(exp(c(.1,.99))),
     yaxt = "n", xaxt = "n", ylab = "Power of F-test",xlab = "SNR")
at <- c(.1,.3,.5,.6,.7,.8,.9,.95,.99)
axis(side = 2, at = exp(exp(at)), labels = at)
axis(side = 4, at = exp(exp(at)), labels = at)
abline(h = exp(exp(c(seq(.1,.95, by = .05),.99))),lwd = .5,col = "gray")
axis(side = 1, at = seq(2,20, by = 2), tick = FALSE)
abline(v = 1:20, lwd = .5, col = "gray")
pow_at <- seq(.8,.95,length = length(nn))
for(i in 1:length(nn)){
  lines(exp(exp(powF[i,])) ~ snr)
  snr_pow <- sum(exp(exp(powF[i,])) < exp(exp(pow_at[i])))
  text(x = snr[snr_pow], y = exp(exp(pow_at[i])), label = nn[i])
}
mtext(side = 3, text = paste("n = ",paste(nn,collapse=", "),
                             ", alpha = ",alpha,sep = ""), line = 1)

```



- v. Suppose $\mu_{11} = 1$, $\mu_{12} = 2$, $\mu_{21} = 1$, and $\mu_{22} = 3$ and $\sigma = 1/3$. Use your plot to determine the necessary number of replicates per treatment group to reject H_0 with probability at least 0.90 when testing at the $\alpha = 0.05$ significance level.

```
sigma <- 1/3
mu <- c(1,2,1,3)
snr <- sum( (t(K)%*% mu)^2)/sigma^2
snr
## [1] 9
```

These settings give a signal to noise ratio of 9. According to the plot we would need

6 replicates per treatment group in order to detect an interaction at the $\alpha = 0.05$ significance level.

- (c) To test for the significance of a *main effect* of factor A , one tests $H_0: \bar{\mu}_i = \bar{\mu}_j$ for all i, j , where $\bar{\mu}_i = b^{-1} \sum_{k=1}^b \mu_{ik}$ for each $i = 1, \dots, a$. The null hypothesis for testing significance of a main effect of factor B is formulated analogously. For this part suppose $a = 3$ and $b = 2$. In answering the following, it may be helpful to draw a table like this one for yourself:

$$\begin{array}{cc} \mu_{11} & \mu_{12} \\ \mu_{21} & \mu_{22} \\ \mu_{31} & \mu_{32} \end{array}$$

For each of the following, give the matrix \mathbf{K} such that we may formulate the hypothesis of interest as $H_0: \mathbf{K}^T \mathbf{b} = \mathbf{0}$.

- i. For testing the significance of the main effect of treatment A .

We wish to test

$$H_0: (\mu_{11} + \mu_{12})/2 = (\mu_{21} + \mu_{22})/2 \text{ and } (\mu_{21} + \mu_{22})/2 = (\mu_{31} + \mu_{32})/2.$$

We can reformulate this as

$$H_0: \mu_{11} + \mu_{12} - \mu_{21} - \mu_{22} = 0 \text{ and } \mu_{21} + \mu_{22} - \mu_{31} - \mu_{32} = 0.$$

With $\mathbf{b} = [\mu_{11} \ \mu_{12} \ \mu_{21} \ \mu_{22} \ \mu_{31} \ \mu_{32}]^T$, we see that we can express the hypothesis as $H_0: \mathbf{K}^T \mathbf{b} = \mathbf{0}$, where

$$\mathbf{K}^T = \begin{bmatrix} 1 & 1 & -1 & -1 & 0 & 0 \\ 0 & 0 & 1 & 1 & -1 & -1 \end{bmatrix}.$$

- ii. For testing the significance of the main effect of treatment B .

We wish to test

$$H_0: (\mu_{11} + \mu_{21} + \mu_{31})/3 = (\mu_{12} + \mu_{22} + \mu_{32})/3.$$

We can reformulate this as

$$H_0: \mu_{11} + \mu_{21} + \mu_{31} - \mu_{12} - \mu_{22} - \mu_{32} = 0.$$

With $\mathbf{b} = [\mu_{11} \ \mu_{12} \ \mu_{21} \ \mu_{22} \ \mu_{31} \ \mu_{32}]^T$, we see that we can express the hypothesis as $H_0: \mathbf{K}^T \mathbf{b} = \mathbf{0}$, where

$$\mathbf{K}^T = [1 \ -1 \ 1 \ -1 \ 1 \ -1].$$

- iii. For testing the significance of an interaction between factors A and B . In the absence of interaction, the differences in means across the levels of one factor do not depend on the level of the other factor.

We wish to test

$$H_0: \mu_{11} - \mu_{12} = \mu_{21} - \mu_{22} \text{ and } \mu_{21} - \mu_{22} = \mu_{31} - \mu_{32}.$$

Other combinations of i, j, k, m are redundant. We can reformulate the above as

$$H_0: \mu_{11} - \mu_{12} - \mu_{21} + \mu_{22} = 0 \text{ and } \mu_{21} - \mu_{22} - \mu_{31} + \mu_{32} = 0.$$

With $\mathbf{b} = [\mu_{11} \ \mu_{12} \ \mu_{21} \ \mu_{22} \ \mu_{31} \ \mu_{32}]^T$, we see that we can express the hypothesis as $H_0: \mathbf{K}^T \mathbf{b} = \mathbf{0}$, where

$$\mathbf{K}^T = \begin{bmatrix} 1 & -1 & -1 & 1 & 0 & 0 \\ 0 & 0 & 1 & -1 & -1 & 1 \end{bmatrix}.$$

(d) Use the data in the image below scanned from [1].

Table 6.19 Tensile strength (psi) of asphaltic concrete specimens for two aggregate types with each of three kneading compaction methods

| Aggregate Type | Compaction Method | | | Aggregate Means ($\bar{y}_{i..}$) |
|-------------------------------------|-------------------|------|----------|-------------------------------------|
| | Kneading | | | |
| | Regular | Low | Very Low | |
| Basalt | 106 | 93 | 56 | |
| | 108 | 101 | | |
| | | 98 | | |
| Means ($\bar{y}_{1j.}$) | 107.0 | 97.3 | 56 | 93.7 |
| Silicious | 107 | 63 | 40 | |
| | 110 | 60 | 41 | |
| | 116 | | 44 | |
| Means ($\bar{y}_{2j.}$) | 111.0 | 61.5 | 41.7 | 72.6 |
| Compaction means ($\bar{y}_{.j}$) | 109.4 | 83.0 | 45.3 | |

Fill out the ANOVA table without using any built-in linear models functions in R.

| Source | SS | df | MS | F | p val |
|-------------|---------|-------|------|-------|--------|
| Total | (i) | (ii) | | | |
| Aggregate | (iii) | (iv) | (v) | (vi) | (vii) |
| Compaction | (viii) | (ix) | (x) | (xi) | (xii) |
| Interaction | (xiii) | (xiv) | (xv) | (xvi) | (xvii) |
| Error | (xviii) | (xix) | (xx) | | |

- i. This is $\mathbf{y}^T(\mathbf{I} - \mathbf{P}_1)\mathbf{y}$, where \mathbf{P}_1 is the orthogonal projection onto $\text{Span}\{\mathbf{1}_n\}$.
- ii. This the the rank of $\mathbf{I} - \mathbf{P}_1$.

- iii. This is the sum of squares for testing the main effect of the aggregate type, which is the value of

$$(\mathbf{K}^T \hat{\mathbf{b}} - \mathbf{m})^T [\mathbf{K}^T (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{K}]^{-1} (\mathbf{K}^T \hat{\mathbf{b}} - \mathbf{m}),$$

where \mathbf{K} is the matrix such that $H_0: \mathbf{K}^T \mathbf{b} = \mathbf{m}$.

- iv. Degrees of freedom corresponding to the main effect of the aggregate type.
- v. The is the sum of squares divided by the degrees of freedom.
- vi. The LRT test statistic for testing significance of the main effect of the aggregate type.
- vii. The p-value of the LRT test of significance of the main effect of the aggregate type.
- viii. This is the sum of squares for testing the main effect of the compaction method.
- ix. Degrees of freedom corresponding to the main effect of the compaction method.
- x. The is the sum of squares divided by the degrees of freedom.
- xi. The LRT test statistic for testing significance of the main effect of the compaction method.
- xii. The p-value of the LRT test of significance of the main effect of the compaction method.
- xiii. This is the sum of squares for testing for an interaction.
- xiv. Degrees of freedom corresponding to the interaction.
- xv. The is the sum of squares divided by the degrees of freedom.
- xvi. The LRT test statistic for testing significance of the interaction.
- xvii. The p-value of the LRT test of significance of the interaction.
- xviii. This is $\mathbf{y}^T (\mathbf{I} - \mathbf{P}_{\mathbf{X}}) \mathbf{y}$.
- xix. The rank of the matrix $\mathbf{I} - \mathbf{P}_{\mathbf{X}}$.
- xx. The sum of squares divided by the degrees of freedom.


```

y <- c(106, 108, 107, 110, 116, 93, 101, 98, 63, 60, 56, 40, 41, 44)
nn <- c(2,3,3,2,1,3);a <- 3;b <- 2;N <- sum(nn)

# build X
X <- matrix(0,N,a*b)
m <- 1
for(i in 1:a)
  for(j in 1:b){
    k <- b*(i-1) + j
    X[m:(m + nn[k] - 1),k] <- rep(1,nn[k])
    m <- m + nn[k]
  }

bhat <- solve(t(X) %*% X) %*% t(X) %*% y

# compute the sums of squares:
SST <- sum( (y - mean(y))^2)

KA <- t(rbind(c(1,1,-1,-1,0,0),c(0,0,1,1,-1,-1)))
HinvA <- solve( t(KA) %*% solve(t(X) %*% X) %*% KA )
SSA <- t( t(KA) %*% bhat) %*% HinvA %*% (t(KA) %*% bhat)

KB <- c(1,-1,1,-1,1,-1)
HinvB <- solve( t(KB) %*% solve(t(X) %*% X) %*% KB )
SSB <- t( t(KB) %*% bhat) %*% HinvB %*% (t(KB) %*% bhat)

KAB <- t(rbind(c(1,-1,-1,1,0,0),
               c(0,0,1,-1,-1,1)))
HinvAB <- solve( t(KAB) %*% solve(t(X) %*% X) %*% KAB )
SSAB <- t( t(KAB) %*% bhat) %*% HinvAB %*% (t(KAB) %*% bhat)

SSE <- sum( (y - X %*% bhat)^2 )

MSA <- SSA / 2
MSB <- SSB / 1
MSAB <- SSAB / 2
MSE <- SSE / ( N - a*b)

F_A <- MSA / MSE
F_B <- MSB / MSE
F_AB <- MSAB / MSE

qf(.999,df1=2,8)
## [1] 18.49365
qf(.999,df1=1,8)
## [1] 25.41476

```

| Source | SS | df | MS | F | p val |
|-------------|----------|----|----------|----------|---------|
| Total | 10963.21 | 13 | | | |
| Aggregate | 710.4537 | 1 | 710.4537 | 63.2686 | < 0.001 |
| Compaction | 6806.452 | 2 | 3403.226 | 303.0702 | < 0.001 |
| Interaction | 953.4492 | 2 | 476.7246 | 42.45414 | < 0.001 |
| Error | 89.83333 | 8 | 11.22917 | | |

2. Let $Y_i = \beta_1 x_{1i} + \dots + \beta_p x_{pi} + \varepsilon_i$, $\varepsilon_i \stackrel{\text{ind}}{\sim} \text{Normal}(0, \sigma^2)$ for $i = 1, \dots, n$. Assume the matrix $\mathbf{X} = (x_{ij})_{1 \leq i \leq n, 1 \leq j \leq p}$ has rank p .

(a) Show that the size- α likelihood ratio test of $H_0: \beta_j = 0$ versus $H_1: \beta_j \neq 0$ is

$$\text{Reject } H_0 \text{ if } \sqrt{n\hat{\Omega}_{jj}^{-1/2}} |\hat{\beta}_j| / \hat{\sigma} > t_{n-p, \alpha/2},$$

where $\hat{\Omega}_{jj}$ is entry j on the diagonal of $\hat{\Omega} = (n^{-1} \mathbf{X}^T \mathbf{X})^{-1}$.

Choose $\mathbf{K}^T = \mathbf{e}_j^T$, where \mathbf{e}_j is the $p \times 1$ vector with every entry equal to zero except for entry j , which is equal to 1. Then one can show that the F statistic for testing $H_0: \mathbf{K}^T \mathbf{b} = 0$ is equal to $n\hat{\Omega}_{jj}^{-1}(\hat{\beta}_j)^2 / \hat{\sigma}^2$. The size- α LRT rejects H_0 when this is greater than $F_{1, n-p, \alpha}$. Since $T \sim t_{n-p} \implies T^2 \sim F_{1, n-p}$, we have $(t_{n-p, \alpha/2})^2 = F_{1, n-p, \alpha}$, so an equivalent decision rule is $\sqrt{n\hat{\Omega}_{jj}^{-1/2}} |\hat{\beta}_j| / \hat{\sigma} > t_{n-p, \alpha/2}$

(b) Show that $\sqrt{n\hat{\Omega}_{jj}^{-1/2}} \hat{\beta}_j / \hat{\sigma} \sim t_{n-p}(\phi = \sqrt{n\hat{\Omega}_{jj}^{-1/2}} \beta_j / \sigma)$.

We have

$$\begin{aligned} \frac{\sqrt{n\hat{\Omega}_{jj}^{-1/2}} \hat{\beta}_j}{\hat{\sigma}} &= \frac{\sqrt{n\hat{\Omega}_{jj}^{-1/2}} \hat{\beta}_j / \sigma}{\sqrt{((n-p)\hat{\sigma}^2 / \sigma^2) / (n-p)}} \\ &= \frac{\sqrt{n\hat{\Omega}_{jj}^{-1/2}} (\hat{\beta}_j - \beta_j) / \sigma + \sqrt{n\hat{\Omega}_{jj}^{-1/2}} \beta_j / \sigma}{\sqrt{((n-p)\hat{\sigma}^2 / \sigma^2) / (n-p)}} \\ &= \frac{Z + \phi}{\sqrt{W / (n-p)}}, \end{aligned}$$

where $Z \sim \text{Normal}(0, 1)$, $W \sim \chi_{n-p}^2$, and $\phi = \sqrt{n\hat{\Omega}_{jj}^{-1/2}} \beta_j / \sigma$.

(c) Show that the noncentrality parameter $\phi = \sqrt{n\hat{\Omega}_{jj}^{-1/2}} \beta_j / \sigma$ can be written as

$$\phi = \frac{\beta_j}{\sigma} \|(\mathbf{I}_n - \mathbf{P}_{\mathbf{X}_{-j}}) \mathbf{x}_j\|_2,$$

where \mathbf{X}_{-j} is the matrix \mathbf{X} with column j removed and \mathbf{x}_j is column j of \mathbf{X} .

For convenience, set $j = 1$, and then partition \mathbf{X} as $\mathbf{X} = [\mathbf{x}_1 \ \mathbf{X}_{-1}]$. Then write $\mathbf{X}^T \mathbf{X}$ as a block matrix. Use the block inverse formula to obtain the $(1, 1)$ entry of $(\mathbf{X}^T \mathbf{X})^{-1}$ as $(\mathbf{x}_1^T \mathbf{x}_1 - \mathbf{x}_1^T \mathbf{X}_{-1}^T (\mathbf{X}_{-1}^T \mathbf{X}_{-1})^{-1} \mathbf{X}_{-1}^T \mathbf{x}_1)^{-1} = (\mathbf{x}_1^T (\mathbf{I}_n - \mathbf{P}_{\mathbf{X}_{-1}}) \mathbf{x}_1)^{-1}$. The answer follows.

- (d) Set $n = 100$, $\sigma = 1$ and, for $p = 20, 40, 80, 90$, generate an $n \times p$ design matrix \mathbf{X} having rows from the $\text{Normal}(\mathbf{0}, \mathbf{I}_n)$ distribution. Then plot the power curves of the test in part (a) at size 0.05 for testing $H_0: \beta_1 = 0$ versus $H_1: \beta_1 \neq 0$. Put the four power curves on the same plot.

```

#### The t-test power thingy:
rm(list=ls())

n <- 100
pp <- c(20,40,80,90)
alpha <- 0.05
sigma <- 1
beta1 <- seq(-1/2,1/2,length = 200)
pow_mat <- matrix(0,nrow = length(beta1),length(pp))
for( j in 1:length(pp)){

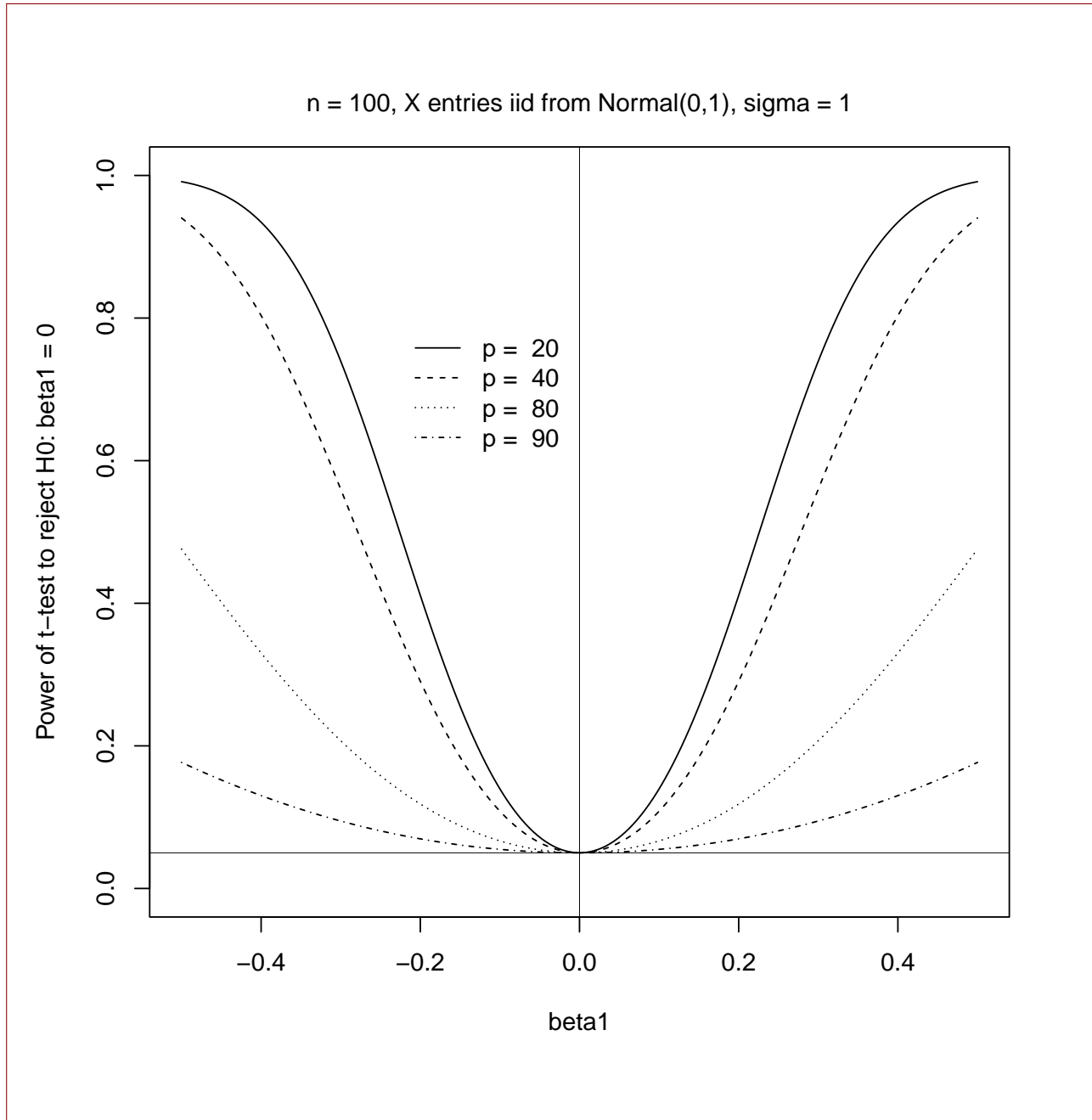
  p <- pp[j]
  X <- matrix(rnorm(n*p),n,p)
  Omega <- solve( t(X)%*%X ) * n
  Omega11 <- Omega[1,1]

  ncp <- sqrt(n) * abs(beta1) / (sigma * sqrt(Omega11) )
  t_crit <- qt(1-alpha/2,df = n - p)
  pow_mat[,j] <- 1-(pt(t_crit,df=n-p,ncp=ncp)-pt(-t_crit,df=n-p,ncp=ncp))

}

plot(NA,
  xlim = range(beta1),
  ylim = c(0,1),
  xlab = "beta1",
  ylab = "Power of t-test to reject H0: beta1 = 0")
for(j in 1:length(pp)) lines(pow_mat[,j] ~ beta1, lty = j)
abline(h = alpha, lwd = 1/2)
abline(v = 0, lwd = 1/2)
legend( x = grconvertX(from = "nfc", to = "user", .35),
  y = .8,
  legend = paste("p = ",pp),
  lty = 1:length(pp),
  bty = "n")
mtext(side = 3,
  text = paste("n = ",n,
    ", X entries iid from Normal(0,1), sigma = ",sigma,
    sep=""),
  line = 1)

```



(e) Describe the effect of having large p on the power of the test.

3. Let $\mathbf{y} = \mathbf{X}\mathbf{b} + \mathbf{e}$ and let \mathbf{K} be a $p \times s$ matrix with columns in $\text{Col } \mathbf{X}^T$ and \mathbf{m} be an $s \times 1$ vector. Let $\tilde{\mathbf{K}}$ and $\tilde{\mathbf{m}}$ be any other matrix and vector such that

$$\{\mathbf{b} : \mathbf{K}^T \mathbf{b} = \mathbf{m}\} = \{\mathbf{b} : \tilde{\mathbf{K}}^T \mathbf{b} = \tilde{\mathbf{m}}\}.$$

Show that the value of the F-statistic is the same regardless of whether one specifies the null hypothesis as $H_0: \mathbf{K}^T \mathbf{b} = \mathbf{m}$ or as $H_0: \tilde{\mathbf{K}}^T \mathbf{b} = \tilde{\mathbf{m}}$.

See book page 134.

References

- [1] R. O. Kuehl. *Design of Experiments: Statistical Principles of Research Design and Analysis*. Duxbury/Thomson Learning, 2000. Google-Books-ID: mIV2QgAACAAJ.