

## 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,  $\mu_{ij}$  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  $i, j$ ).

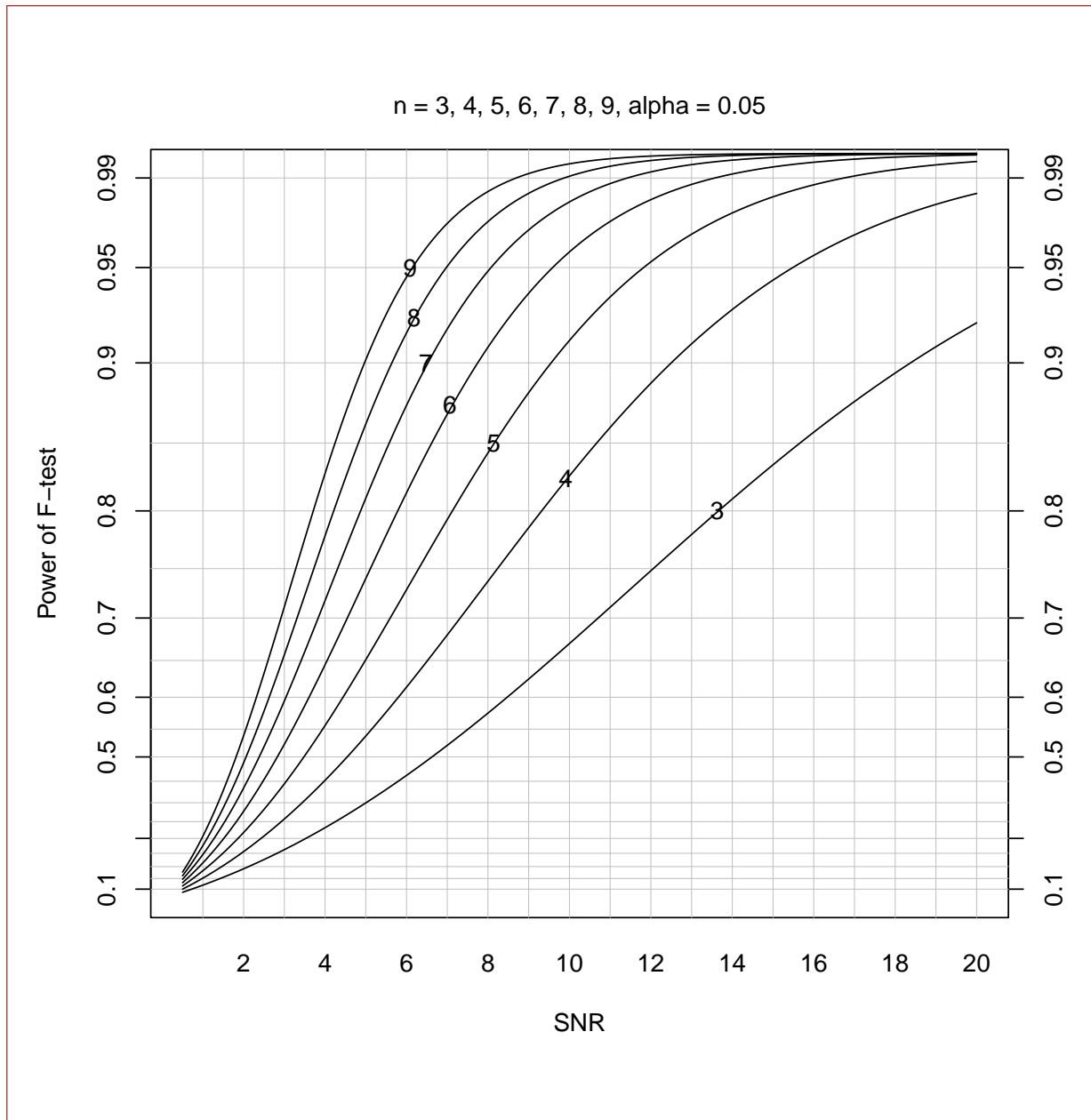
Note that in the balanced design (equal replications in each treatment group), we have  $\mathbf{X}^T \mathbf{X} = n_N$ . Moreover,  $\mathbf{K}^T \mathbf{K} = 4\mathbf{I}_2$ , so we have  $[\mathbf{K}^T (\mathbf{X}^T \mathbf{X})^{-1} \mathbf{K}]^{-1} = n/4$ . This, together with  $\mathbf{m} = \mathbf{0}$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

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

- 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	106 108	93 101 98	56	
Means ( $\bar{y}_{1j.}$ )	107.0	97.3	56	93.7
Silicious	107 110 116	63 60	40 41 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)		

- This is  $\mathbf{y}^T(\mathbf{I} - \mathbf{P}_1)\mathbf{y}$ , where  $\mathbf{P}_1$  is the orthogonal projection onto  $\text{Span}\{\mathbf{1}_n\}$ .
- 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- $\alpha$  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  $H_0: \mathbf{K}^T\mathbf{b} = \mathbf{0}$  is equal to  $n\hat{\Omega}_{jj}^{-1}(\hat{\beta}_j)^2/\hat{\sigma}^2$ . The size- $\alpha$  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 of the test in part (a) as a function of the true value of the parameter  $\beta_j$  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)){

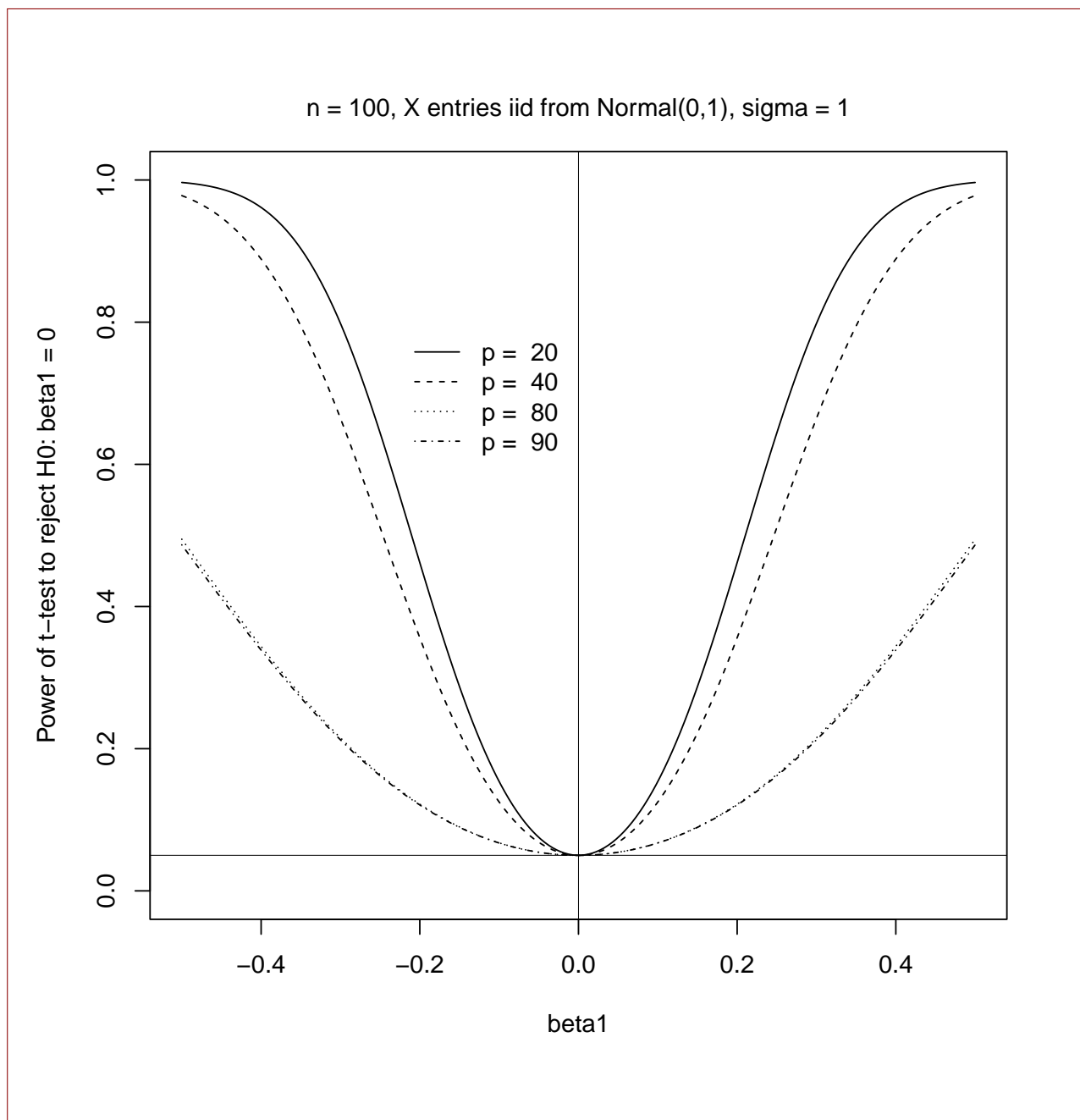
  p <- pp[j]
  X <- matrix(rnorm(n*p),n,p)
  Omega <- solve( t(X)%*%X ) * n
  Omega11 <- Omega[1,1]

  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.