

1 a)

$$\text{HTN: } t = \frac{\hat{\beta}_1}{SE(\hat{\beta}_1)} = \frac{0.111}{0.031} \approx \boxed{3.58} > 1.96$$

So, reject the null. Thus, there is evidence that a history of hypertension is associated with an increased probability of death.

$$\text{Smoke: } t = \frac{\hat{\beta}_2}{SE(\hat{\beta}_2)} = \frac{0.234}{0.072} = \boxed{3.25} > 1.96$$

So, reject the null. Thus, there is evidence that a history of smoking is associated with an increased probability of death.

$$\text{Age: } t = \frac{\hat{\beta}_3}{SE(\hat{\beta}_3)} = \frac{0.100}{0.100} = \boxed{1.00} < 1.96$$

So, fail to reject the null. Thus, there is no evidence that advanced age is associated with an increased probability of death.

$$b) \text{ OR} = \exp(d\hat{\beta}_3) = \exp(0.1 \cdot 2) \approx \boxed{1.22}$$

Remember, it increments by 5 yr.

Thus, the odds of sudden death is 1.22 higher in women who are 10 yr. older.

$$95\% \text{ CI} = \exp\left(\underbrace{d \cdot \hat{\beta}_3}_{0.2} \pm 1.96 \cdot \underbrace{d \cdot SE(\hat{\beta}_3)}_{2 \cdot 0.1}\right) = \boxed{(0.825, 1.808)}$$

c) According to Dr. Zou during review.

In the situation where you have a case-control study, you cannot use the fitted logistic regression for prediction. Since you have selected on a specific outcome, the number of controls $\frac{1}{2}$ cases are not as they are present in the population. In this case, the intercept is a result of our design $\frac{1}{2}$ is not applicable to the population as it will tend to overestimate the probability of death.

1 d) According to Dr. Zou, there are two correct ways of phrasing this answer.

Method 1: We want to test $H_0: \beta_1 = \beta_2 \Leftrightarrow H_0: \beta_1 - \beta_2 = 0$.

$$\text{We could use a score test, } \frac{(\hat{\beta}_1 - \hat{\beta}_2)}{SE(\hat{\beta}_1 - \hat{\beta}_2)} = \frac{(\hat{\beta}_1 - \hat{\beta}_2)}{\sqrt{\text{Var}(\hat{\beta}_1 - \hat{\beta}_2)}} = \frac{(\hat{\beta}_1 - \hat{\beta}_2)}{\sqrt{\text{Var}(\hat{\beta}_1) + \text{Var}(\hat{\beta}_2) - 2\text{Cov}(\hat{\beta}_1, \hat{\beta}_2)}}$$

We don't have $\text{Cov}(\hat{\beta}_1, \hat{\beta}_2)$, we were only given $\text{Var}(\hat{\beta}_1) = SE(\hat{\beta}_1)^2$ and $\text{Var}(\hat{\beta}_2) = SE(\hat{\beta}_2)^2$.

So, we are shit out of luck. ☹

Method 2: Again, want to test $H_0: \beta_1 = \beta_2 \Leftrightarrow H_0: \beta_1 - \beta_2 = 0$.

We could use a likelihood ratio (LR) test. This involves computing a log-likelihood ratio for the full model ($\text{logit}(\hat{p}) = \hat{\beta}_0 + \hat{\beta}_1 \cdot \text{HTN} + \hat{\beta}_2 \cdot \text{Smoke} + \hat{\beta}_3 \cdot \text{Age}$) and a log-likelihood ratio for the reduced model ($\text{logit}(\hat{p}) = \hat{\beta}_0^* + \hat{\beta}_1^* (\text{HTN} + \text{Smoke}) + \hat{\beta}_2^* \cdot \text{Age}$).

Would calculate $-2LR(\text{reduced}) - (-2LR(\text{full})) \sim \chi^2_{df(\text{full}) - df(\text{reduced})} \equiv \chi^2_1$

However, again we were not given the likelihood ratios for the full & reduced models, so no bueno.

1 e) Without info regarding the distribution of age in the controls, the investigator cannot make any conclusion regarding risk of death.

2 a)

$$\text{First, know } \text{Var}(\hat{p}) = \text{Var}\left(\frac{X}{n}\right) = \frac{1}{n^2} \text{Var}(X) = \frac{1}{n^2} n \hat{p}(1-\hat{p}) = \frac{\hat{p}(1-\hat{p})}{n}$$

Will be able to use this eqn. for variance to derive 95% CI's.

Point Estimates: $\hat{p}_0 = P(\text{outcome} = 1 | \text{intervention} = 0) = \frac{14+5}{75} \approx \boxed{0.253}$

$$\hat{p}_1 = P(\text{outcome} = 1 | \text{intervention} = 1) = \frac{20+10}{75} = \boxed{0.40}$$

$$95\% \text{ CI}(\hat{p}_0) = 0.253 \pm 1.96 \sqrt{\frac{0.253(1-0.253)}{75}} = \boxed{(0.155, 0.351)}$$

$$95\% \text{ CI}(\hat{p}_1) = 0.40 \pm 1.96 \sqrt{\frac{0.40(1-0.40)}{75}} = \boxed{(0.289, 0.511)}$$

2 b) Can compute an OR, RR, or RD (risk difference, same as difference in proportions).
Here, will compute an RD.

$$\hat{RD} = \hat{p}_1 - \hat{p}_0 \text{ where } \hat{p}_0 = P(\text{outcome} = 1 | \text{intervention} = 0) = 0.253 \text{ (last part)}$$

$$\hat{p}_1 = P(\text{outcome} = 1 | \text{intervention} = 1) = 0.40 \text{ (last part)}$$

$$\Rightarrow \hat{RD} = 0.4 - 0.253 = 0.247$$

$$95\% \text{ CI}(\hat{RD}) = (\hat{p}_1 - \hat{p}_0) \pm 1.96 \sqrt{\frac{\hat{p}_1(1-\hat{p}_1)}{n_1} + \frac{\hat{p}_0(1-\hat{p}_0)}{n_0}}$$

$$= 0.247 \pm 1.96 \sqrt{\frac{0.4(1-0.4)}{75} + \frac{0.253(1-0.253)}{75}}$$

$$= \boxed{(0.099, 0.395)}$$

The patients who received ^{the} intervention had almost 25 additional cases (out of 100) of ^{persons} HbA1c below 7.5% when compared to patients who received only usual care.

2 c) Taking difference in proportions from 2b) have,

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$$H_0: p_1 - p_0 = 0$$

$$Z = \frac{\hat{p}_1 - \hat{p}_0}{\sqrt{\frac{\hat{p}_1(1-\hat{p}_1)}{n_1} + \frac{\hat{p}_0(1-\hat{p}_0)}{n_2}}} = \frac{0.4 - 0.253}{\sqrt{\frac{0.4(1-0.4)}{75} + \frac{0.253(1-0.253)}{75}}} \approx 1.94 < 1.96$$

Using `2 * pnorm(1.943, lower.tail=F)` in R returns $p = 0.052$

Assuming an $\alpha = 0.05$, we fail to reject the null hypothesis & conclude that there is no evidence that the intervention is effective in lowering BS.

2 d) Have $\text{logit}(p) = \beta_0 + \beta_1 \cdot \text{intervention} + \beta_2 \text{sex} + \beta_3 \cdot \text{intervention} \cdot \text{sex} + \varepsilon$

$$\text{Have intervention} = \begin{cases} 1, & \text{intervention} \\ 0, & \text{usual care} \end{cases}$$

$$\text{sex} = \begin{cases} 0, & \text{female} \\ 1, & \text{male} \end{cases}$$

Testing: $H_0: \beta_3 = 0$

$H_1: \beta_3 > 0$ (Know β_3 must be positive since β_3 exists if $\text{sex} = 1 \wedge \text{intervention} = 1$ in order for an increased logit(p) - our hypothesis.

Fitting the above model in R gives: $\hat{\beta}_3 = 0.2029$
 $p = 0.7903$ (for a two-sided test)

used,

`glm(Outcome ~ Intervention * sex,
data=df, family='binomial')`

However, we need the p-value for a one-sided test. For symmetric distributions, if two-tailed p-value > 0.5 , then one-tailed p-value = $1 - \frac{(\text{two-tailed } p)}{2}$.

If two-tailed p-value < 0.5 , then one-tailed p-value = $\frac{(\text{two-tailed } p)}{2}$.

Here $p_2 = 0.7903 > 0.5$, so $p_1 = 1 - \frac{0.7903}{2} \approx 0.605$

Since $p = 0.605 > \alpha = 0.05$, we fail to reject the null. There is no evidence that males respond better to the intervention.

2 e) Simply fit the model $\text{logit}(p) = \beta_0 + \beta_1 \cdot \text{intervention} + \beta_2 \cdot \text{sex} + \varepsilon$
 using `glm(Outcome ~ intervention + sex, data = df, family = "binomial")`
 in R to get a sex adjusted estimate of $\hat{\beta}_1 = 0.7165$ with associated
 $SE(\hat{\beta}_1) = 0.3634$.

$$\text{Then, } 95\% \text{ CI}(\beta_1) = \hat{\beta}_1 \pm 1.96 \cdot SE(\hat{\beta}_1) = 0.7165 \pm \overbrace{1.96 \cdot 0.3634}^{0.712264}$$

$$= \boxed{(0.0042, 1.4288)}$$

The confidence interval borderline contains the 0 estimate. However, at a significance level of $\alpha = 0.05$, these results would still be considered "statistically significant," and we would conclude that the intervention, when adjusted for differences in sex, has a positive effect on the outcome (increasing the probability of low blood sugar).

3 a) $H_0: \mu_1 = \mu_2 = \mu_3 = 0$ vs. $H_1: \mu_i \neq 0$ for at least one $i=1,2,3$

ANOVA	SS	df	F	General formulas for I groups
Between	$\sum_{i=1}^I \sum_{j=1}^{n_i} [\bar{y}_i - \bar{y}]^2$	$i-1$	$(SS_{\text{between}}/i-1)$	
Within	$\sum_{i=1}^I \sum_{j=1}^{n_i} [y_{ij} - \bar{y}_i]^2$	$N-i$	$(SS_{\text{within}}/n-i)$	
Total	$SS_{\text{between}} + SS_{\text{within}}$	$N-1$		

ANOVA	SS	df	F
Between	70.94	2	52.24
Within	201.57	297	
Total	272.51	299	

where $\bar{y} = \text{grand mean} = \frac{3.78(100) + 3.23(100) + 2.59(100)}{300} = \frac{3.78 + 3.23 + 2.59}{3} = 3.2$

Then, $SS_{\text{between}} = \sum_{i=1}^3 \sum_{j=1}^{100} [\bar{y}_i - \bar{y}]^2 = \underbrace{100}_{n_1} \underbrace{(3.78 - 3.2)^2}_{\bar{y}_1 - \bar{y}} + \underbrace{100}_{n_2} \underbrace{(3.23 - 3.2)^2}_{\bar{y}_2 - \bar{y}} + \underbrace{100}_{n_3} \underbrace{(2.59 - 3.2)^2}_{\bar{y}_3 - \bar{y}}$
 $= 70.94$

$SS_{\text{within}} = \sum_{i=1}^3 \sum_{j=1}^{100} [y_{ij} - \bar{y}_i]^2 = (n_1 - 1) \cdot SD_1^2 + (n_2 - 1) \cdot SD_2^2 + (n_3 - 1) \cdot SD_3^2$
 $= 99(0.79)^2 + 99(0.86)^2 + 99(0.82)^2$

$$\approx 201.57$$

$$SS_{\text{Total}} = SS_{\text{between}} + SS_{\text{within}} = 70.94 + 201.57 = 272.51$$

$$df_{\text{between}} = 3 - 1 = 2, \quad df_{\text{within}} = 300 - 3 = 297, \quad df_{\text{total}} = 300 - 1 = 299$$

$$F = \frac{SS_{\text{between}} / (i-1)}{SS_{\text{within}} / \left[\left(\sum_{i=1}^I n_i \right) - i \right]} = \frac{(70.94/2)}{(201.57/297)} \approx 52.24 \sim F_{2,297}$$

$$p\text{-value} = pf(52.24, df1=2, df2=297, \text{lower.tail}=F) = 1.61 \times 10^{-19} \text{ in R.}$$

Since $p = 1.61 \times 10^{-19} < \alpha = 0.05$, we reject the null hypothesis and conclude that the group means are not all identical (at least one is different).

3 b)

From given construct, have:

$$Y = X\beta + \epsilon \text{ where,}$$

$$Y: 300 \times 1$$

$$X: 300 \times 2$$

$$\beta: 2 \times 1$$

$$\epsilon: 300 \times 1$$

$$X = \begin{bmatrix} \vdots & 1 \\ \vdots & 2 \\ \vdots & 2 \\ \vdots & 3 \\ \vdots & 3 \end{bmatrix} \begin{matrix} \left. \begin{matrix} \vdots \\ \vdots \\ \vdots \end{matrix} \right\} 100 \times \\ \left. \begin{matrix} 1 \\ 2 \\ 2 \end{matrix} \right\} 100 \times \\ \left. \begin{matrix} 3 \\ 3 \end{matrix} \right\} 100 \times \end{matrix}$$

Design matrix

$$\beta = \begin{pmatrix} \alpha_1 \\ \alpha_2 \end{pmatrix}$$

Then, three steps:

① Find parameter estimates.

Fit following model in R, $\text{lm}(y \sim x)$ where $y = c(3.78, 3.23, 2.59)$
 \uparrow
 vector

and $x = 1:3$

\uparrow
 vector containing 1, 2, and 3

Will return parameter estimates of, $\hat{\alpha}_1 = 4.39$ and $\hat{\alpha}_2 = -0.595$

② Find MSE, $\hat{\sigma}^2$

According to the GG3 textbook, on pg. 325,

$$\hat{\sigma}^2 = \text{SS}_{\text{within}} / N - i = \underbrace{201.57}_{\text{on previous pg, we wrote this for df}} / 297 \approx \boxed{0.679}$$

\nwarrow MSE

③ Find SEE (standard error of the estimates).

Need $SE(\hat{\alpha}_1)$ and $SE(\hat{\alpha}_2)$.

Use formula $SEE = \sqrt{\hat{\sigma}^2 \text{diag}(X'X)^{-1}}$. Code X in R using $X = \text{cbind}(\text{rep}(1, 300), \text{c}(\text{rep}(1, 100), \text{rep}(2, 100), \text{rep}(3, 100)))$

Then, use $(\underbrace{t(X)}_{\text{transpose of X}} \%*\% X)^{\underbrace{(-1)}_{\text{inverse}}}$ to get inverse SSCP, $(X'X)^{-1}$

means diagonal entries of matrix $(X'X)^{-1}$

Then, use $\text{sqr} + (0.679 * (X'X)^{-1})$ and grab diagonal entries. These will be $SE(\hat{\alpha}_1)$ and $SE(\hat{\alpha}_2)$. You will find $SE(\hat{\alpha}_1) = 0.0476$ and $SE(\hat{\alpha}_2) = 0.022$.

3 c) The given hypothesis of $H_0: \beta_2 = \beta_3 = 0$ is equivalent to testing $\mu_1 = \mu_2 = \mu_3$ because if $\beta_2 = \beta_3 = 0 \Rightarrow \mu_1 = \beta_1, \mu_2 = \beta_1$, and $\mu_3 = \beta_1$.

This is equivalent to the F test in the one-way ANOVA table in part a).

Thus, $F \approx 52.24 \sim F_{2,297}$

p-value = $\text{pf}(52.24, \text{df1}=2, \text{df2}=297, \text{lower.tail}=F) = 1.61 \times 10^{-19}$ } in R

Since $p = 1.61 \times 10^{-19} < \alpha = 0.05$, we reject the null hypothesis and conclude that at least one β_i (for $i=2,3$) is non-zero.

3 e) $H_0: \mu_1 = \mu_2 = \mu_3 = 0$ vs. $H_1: \mu_i \neq 0$ for at least one $i = 1, 2, 3$

ANOVA	SS	df	F
Between	$\sum_{i=1}^I \sum_{j=1}^{n_i} [\bar{y}_i - \bar{y}]^2$	$i-1$	$(SS_{\text{between}}/i-1)$
Within	$\sum_{i=1}^I \sum_{j=1}^{n_i} [y_{ij} - \bar{y}_i]^2$	$N-i$	$(SS_{\text{within}}/n-i)$
Total	$SS_{\text{between}} + SS_{\text{within}}$	$N-1$	

General formulas for I groups

ANOVA	SS	df	F
Between	70.94	2	52.24
Within	201.57	297	
Total	272.51	299	

where \bar{y} = grand mean = $\frac{3.78(100) + 3.23(100) + 2.59(100)}{300} = \frac{3.78 + 3.23 + 2.59}{3} = 3.2$

Then, $SS_{\text{between}} = \sum_{i=1}^3 \sum_{j=1}^{100} [\bar{y}_i - \bar{y}]^2 = \underbrace{100}_{n_1} \underbrace{(3.78 - 3.2)^2}_{\bar{y}_1 - \bar{y}} + \underbrace{100}_{n_2} \underbrace{(3.23 - 3.2)^2}_{\bar{y}_2 - \bar{y}} + \underbrace{100}_{n_3} \underbrace{(2.59 - 3.2)^2}_{\bar{y}_3 - \bar{y}}$
 $= 70.94$

$SS_{\text{within}} = \sum_{i=1}^3 \sum_{j=1}^{100} [y_{ij} - \bar{y}_i]^2 = (n_1 - 1) \cdot SD_1^2 + (n_2 - 1) \cdot SD_2^2 + (n_3 - 1) \cdot SD_3^2$
 $= 99(0.79)^2 + 99(0.86)^2 + 99(0.82)^2$
 ≈ 201.57

$SS_{\text{Total}} = SS_{\text{between}} + SS_{\text{within}} = 70.94 + 201.57 = 272.51$

$df_{\text{between}} = 3 - 1 = 2$, $df_{\text{within}} = 300 - 3 = 297$, $df_{\text{total}} = 300 - 1 = 299$

$F = \frac{SS_{\text{between}} / (i-1)}{SS_{\text{within}} / [(\sum_{i=1}^I n_i) - i]} = \frac{(70.94/2)}{(201.57/297)} \approx 52.24 \sim F_{2,297}$

$p\text{-value} = pf(52.24, df_1 = 2, df_2 = 297, \text{lower.tail} = F) = \frac{3.62 \times 10^{-20}}{1.61 \times 10^{-19}}$ in R.

Since $p = \frac{3.62 \times 10^{-20}}{1.61 \times 10^{-19}} < \alpha = 0.05$, we reject the null hypothesis and conclude that the group means are not all identical (at least one is different).

3 b)

From given construct, have:

$$Y = X\beta + \epsilon \text{ where,}$$

$$Y: 300 \times 1$$

$$X: 300 \times 2$$

$$\beta: 2 \times 1$$

$$\epsilon: 300 \times 1$$

$$X = \begin{bmatrix} \vdots & \vdots \\ 1 & 2 \\ \vdots & \vdots \\ 2 & 2 \\ \vdots & \vdots \\ 3 & 3 \\ \vdots & \vdots \\ 3 & 3 \end{bmatrix} \begin{matrix} 100 \times \\ 100 \times \\ 100 \times \end{matrix}$$

Design matrix

$$\beta = \begin{pmatrix} \alpha_1 \\ \alpha_2 \end{pmatrix}$$

$$(X'X)^{-1} = \begin{pmatrix} -0.023333 & -0.010 \\ -0.010 & 0.005 \end{pmatrix}$$

Then, three steps:

① Find parameter estimates.

$$\text{lm}(\tilde{y} \sim \alpha_1 + \alpha_2 \tilde{x}) \dots (*)$$

Fit following model in R, $\text{lm}(\tilde{y} \sim \alpha_1 + \alpha_2 \tilde{x})$ where $\tilde{y} = c(3.78, 3.23, 2.59)$
and $\tilde{x} = c(1, 2, 3)$

Will return parameter estimates of, $\hat{\alpha}_1 = 4.39$ and $\hat{\alpha}_2 = -0.595$

\downarrow SSE of (*)
 $RSS = 0.00135$

② Find MSE, $\hat{\sigma}^2$

According to the G63 textbook, on pg. 325,

$$SSE = SS_{\text{within}} + 100 * RSS = 201.7088$$

$$\hat{\sigma}^2 = \frac{SSE}{N-p} = \frac{201.7088}{298} \approx 0.67688$$

on previous pg.
do not write this
+ 100 * RSS

\swarrow SSE of the actual model

③ Find SEE (standard error of the estimates).

Need $SE(\hat{\alpha}_1)$ and $SE(\hat{\alpha}_2)$.

Use formula $SEE = \sqrt{\hat{\sigma}^2 \text{diag}(X'X)^{-1}}$. Code X in R using $X = \text{cbind}(\text{rep}(1, 300), \text{c}(\text{rep}(1, 100), \text{rep}(2, 100), \text{rep}(3, 100)))$

means diagonal entries of matrix $(X'X)^{-1}$

Then, use $(t(X) \% * \% X)^{-1}$ to get inverse SSCP, $(X'X)^{-1}$
transpose of X

Then, use $\text{sqrt}(0.677 \cdot (X'X)^{-1})$ and grab diagonal entries. These will be $SE(\hat{\alpha}_1)$ and $SE(\hat{\alpha}_2)$. You will find $SE(\hat{\alpha}_1) = 0.476$ and $SE(\hat{\alpha}_2) = 0.22$.

0.4757

0.05878