

# Math 362: Mathematical Statistics II

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# Chapter 12. The Analysis of Variance

# Plan

§ 12.1 Introduction

§ 12.2 The  $F$  Test

§ 12.3 Multiple Comparisons: Turkey's Method

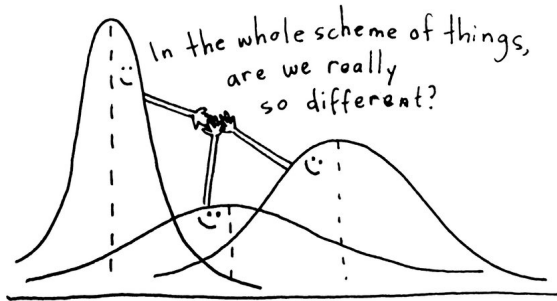
# Chapter 12. The Analysis of Variance

§ 12.1 Introduction

§ 12.2 The  $F$  Test

§ 12.3 Multiple Comparisons: Turkey's Method

## § 12.1 Introduction



E.g. 1 Study the relation between smoking and heart rates.

Generations of athletes have been cautioned that cigarette smoking impedes performance. One measure of the truth of that warning is the effect of smoking on heart rate. In one study, six nonsmokers, six light smokers, six moderate smokers, and six heavy smokers each engaged in sustained physical exercise. Table 8.1.1 lists their heart rates after they had rested for three minutes.

Show whether smoking affects heart rates at  $\alpha = 0.05$ .

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	Nonsmokers	Light Smokers	Moderate Smokers	Heavy Smokers
	69	55	66	91
	52	60	81	72
	71	78	70	81
	58	58	77	67
	59	62	57	95
	65	66	79	84
<i>Averages:</i>	62.3	63.2	71.7	81.7

Show whether smoking affects heart rates at  $\alpha = 0.05$ .

E.g. 2 A certain fraction of antibiotics injected into the bloodstream are “bound” to serum proteins. This phenomenon bears directly on the effectiveness of the medication, because the binding decreases the systemic uptake of the drug. Table below lists the binding percentages in bovine serum measured for five widely prescribed antibiotics. Which antibiotics have similar binding properties, and which are different?



E.g. 2 A certain fraction of antibiotics injected into the bloodstream are “bound” to serum proteins. This phenomenon bears directly on the effectiveness of the medication, because the binding decreases the systemic uptake of the drug. Table below lists the binding percentages in bovine serum measured for five widely prescribed antibiotics. Which antibiotics have similar binding properties, and which are different?

	Penicillin G	Tetra- cycline	Strepto- mycin	Erythro- mycin	Chloram- phenicol
	29.6	27.3	5.8	21.6	29.2
	24.3	32.6	6.2	17.4	32.8
	28.5	30.8	11.0	18.3	25.0
	32.0	34.8	8.3	19.0	24.2
$T_j$	114.4	125.5	31.3	76.3	111.2
$\bar{Y}_j$	28.6	31.4	7.8	19.1	27.8

	<i>Treatment Level</i>			
	1	2	...	$k$
	$Y_{11}$	$Y_{12}$		$Y_{1k}$
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	$\vdots$	$\vdots$	...	$\vdots$
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Sample sizes:	$n_1$	$n_2$	...	$n_k$
Sample totals:	$T_{.1}$	$T_{.2}$		$T_{.k}$
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Problem Testing

$$H_0 : \mu_1 = \mu_2 = \dots = \mu_k$$

versus

$H_1$  : not all the  $\mu_j$ 's are equal

Or testing *subhypotheses* such as

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ANOVA was developed by statistician and evolutionary biologist —



## Ronald Fisher



Statistician

Sir Ronald Aylmer Fisher FRS was a British statistician and geneticist. For his work in statistics, he has been described as "a genius who almost single-handedly created the foundations for modern statistical science" and "the single most important figure in 20th century statistics". [Wikipedia](#)

**Born:** February 17, 1890, [East Finchley, London, United Kingdom](#)

**Died:** July 29, 1962, [Adelaide, Australia](#)

**Known for:** [Fisher's principle](#), [Fisher information](#)

**Residence:** [United Kingdom, Australia](#)

**Education:** [Gonville & Caius College, University of Cambridge](#),  
[Harrow School](#)

<https://www.youtube.com/watch?v=0XsovsSnRuw>

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# Chapter 12. The Analysis of Variance

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§ 12.2 The  $F$  Test

§ 12.3 Multiple Comparisons: Turkey's Method

### Model assumptions

1. Independence of observations
2. Normality
3. Homogeneity of variances



#### Assume:

$\forall j = 1, \dots, k, \forall i = 1, \dots, n_j,$

1.  $Y_{ij}$  are independent.
2.  $Y_{ij} \sim N(\mu_j, \sigma^2)$



#### Assume:

$\forall j = 1, \dots, k, \forall i = 1, \dots, n_j,$

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# Likelihood ratio test

1. The parameter spaces are

$$\Omega = \left\{ (\mu_1, \dots, \mu_k, \sigma^2) : -\infty < \mu_1, \dots, \mu_k < \infty, \sigma^2 > 0 \right\}$$

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$$\frac{\partial \ln L(\omega)}{\partial \mu} = \frac{1}{\sigma^2} \sum_{j=1}^k \sum_{i=1}^{n_j} (y_{ij} - \mu)$$

$$\frac{\partial \ln L(\omega)}{\partial (\sigma^2)} = -\frac{n}{2\sigma^2} + \frac{1}{2\sigma^4} \sum_{j=1}^k \sum_{i=1}^{n_j} (y_{ij} - \mu)^2$$

Setting the above derivatives to zero, the solutions for  $\mu$  and  $\sigma^2$  are,

$$\frac{1}{n} \sum_{j=1}^k \sum_{i=1}^{n_j} y_{ij} = \bar{y}_{..}$$

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$$L(\hat{\omega}) = \left( \frac{n}{2\pi \sum_{j=1}^k \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_{..})^2} \right)^{n/2} \exp \left\{ -\frac{n \sum_{j=1}^k \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_{..})^2}{2 \sum_{j=1}^k \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_{..})^2} \right\}$$

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$$\lambda = \frac{L(\hat{\omega})}{L(\hat{\Omega})} = \left( \frac{\sum_{j=1}^k \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_{\cdot j})^2}{\sum_{j=1}^k \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_{\cdot \cdot})^2} \right)^{n/2}$$

⇒ Test statistic:

$$\Lambda = \frac{L(\hat{\omega})}{L(\hat{\Omega})} = \left( \frac{\sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{\cdot j})^2}{\sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{\cdot \cdot})^2} \right)^{n/2}$$

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$$\begin{aligned}
SSTOT &:= \sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{..})^2 \\
&= \sum_{j=1}^k \sum_{i=1}^{n_j} [(Y_{ij} - \bar{Y}_{.j}) + (\bar{Y}_{.j} - \bar{Y}_{..})]^2 \\
&= \sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{.j})^2 + \text{zero cross term} + \sum_{j=1}^k \sum_{i=1}^{n_j} (\bar{Y}_{.j} - \bar{Y}_{..})^2 \\
&= \underbrace{\sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{.j})^2}_{SSE} + \underbrace{\sum_{j=1}^k n_j (\bar{Y}_{.j} - \bar{Y}_{..})^2}_{SSTR}
\end{aligned}$$

↓

$$\Lambda = \left( \frac{SSE}{SSTOT} \right)^{n/2} = \left( \frac{SSE}{SSE + SSTR} \right)^{n/2} = \left( \frac{1}{1 + SSTR/SSE} \right)^{n/2}$$

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6. Critical regions: for some  $\lambda_* \in (0, 1)$  close to 0,

$$\begin{aligned}\alpha &= \mathbb{P}(\Lambda \leq \lambda_*) \\ &= \mathbb{P}\left(\frac{1}{1 + SSTR/SSE} \leq \lambda_*^{2/n}\right) \\ &= \mathbb{P}\left(\frac{SSTR}{SSE} \leq \lambda_*^{-2/n} - 1\right) \\ &= \mathbb{P}\left(\frac{SSTR/(k-1)}{SSE/(n-k)} \leq \left(\lambda_*^{-2/n} - 1\right) \frac{n-k}{k-1}\right)\end{aligned}$$

7. We will prove that under  $H_0$ ,  $\frac{SSTR/(k-1)}{SSE/(n-k)} \sim F$ -distr.  $df_1 = k-1$ ,  $df_2 = n-k$

$$\Rightarrow \left(\lambda_*^{-2/n} - 1\right) \frac{n-k}{k-1} = F_{1-\alpha, k-1, n-k}.$$

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□

## Treatment sum of squares: SSTR

Sample size: (Weights)	$n_1$	$n_2$	$\dots$	$n_k$	$n = \sum_{j=1}^k n_j$  <i>Weighted average</i>
Sample means:	$\bar{Y}_{.1}$	$\bar{Y}_{.2}$	$\dots$	$\bar{Y}_{.k}$	$\bar{Y}_{..} = \frac{1}{n} \sum_{j=1}^k n_j \bar{Y}_{.j}$
True means:	$\mu_1$	$\mu_2$	$\dots$	$\mu_k$	$\mu = \frac{1}{n} \sum_{j=1}^k n_j \mu_j$
Squares:	$(\bar{Y}_{.1} - \bar{Y}_{..})^2$	$(\bar{Y}_{.2} - \bar{Y}_{..})^2$	$\dots$	$(\bar{Y}_{.k} - \bar{Y}_{..})^2$	<b>SSTR</b>

$$SSTR := \frac{1}{n} \sum_{j=1}^k n_j (\bar{Y}_{.j} - \bar{Y}_{..})^2$$

1. When  $k = 1$ ,  $SSTR \equiv 0$ .
2. When  $k = 2$ , say  $X_1, \dots, X_n$  and  $Y_1, \dots, Y_m$ :

$$\bar{Y}_{..} = \frac{1}{m+n} (n\bar{X} + m\bar{Y})$$

$$\Rightarrow SSTR = \frac{(\bar{X} - \bar{Y})^2}{\frac{1}{m} + \frac{1}{n}}$$

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$$\begin{aligned} SSTR &= n \left[ \bar{X} - \frac{1}{n+m} (n\bar{X} + m\bar{Y}) \right]^2 + m \left[ \bar{Y} - \frac{1}{n+m} (n\bar{X} + m\bar{Y}) \right]^2 \\ &= n \left[ \frac{m(\bar{X} - \bar{Y})}{n+m} \right]^2 + m \left[ \frac{n(\bar{X} - \bar{Y})}{n+m} \right]^2 \\ &= \left[ \frac{nm^2}{(n+m)^2} + \frac{mn^2}{(n+m)^2} \right] (\bar{X} - \bar{Y})^2 \\ &= \frac{nm}{n+m} (\bar{X} - \bar{Y})^2 \end{aligned}$$

$$\Rightarrow SSTR = \frac{(\bar{X} - \bar{Y})^2}{\frac{1}{m} + \frac{1}{n}}$$

$$\begin{aligned}
SSTR &= \sum_{j=1}^k n_j (\bar{Y}_{.j} - \bar{Y}_{..})^2 = \sum_{j=1}^k n_j [(\bar{Y}_{.j} - \mu) - (\bar{Y}_{..} - \mu)]^2 \\
&= \sum_{j=1}^k n_j [(\bar{Y}_{.j} - \mu)^2 + (\bar{Y}_{..} - \mu)^2 - 2(\bar{Y}_{.j} - \mu)(\bar{Y}_{..} - \mu)] \\
&= \sum_{j=1}^k n_j (\bar{Y}_{.j} - \mu)^2 + \sum_{j=1}^k n_j (\bar{Y}_{..} - \mu)^2 - 2(\bar{Y}_{..} - \mu) \sum_{j=1}^k n_j (\bar{Y}_{.j} - \mu) \\
&= \sum_{j=1}^k n_j (\bar{Y}_{.j} - \mu)^2 + n(\bar{Y}_{..} - \mu)^2 - 2(\bar{Y}_{..} - \mu)n(\bar{Y}_{..} - \mu) \\
&= \sum_{j=1}^k n_j (\bar{Y}_{.j} - \mu)^2 - n(\bar{Y}_{..} - \mu)^2 \tag{12.2.1}
\end{aligned}$$

↓

$$SSTR = \sum_{j=1}^k n_j \left[ (\bar{Y}_{.j} - \mu_j)^2 - 2(\bar{Y}_{.j} - \mu_j)(\mu - \mu_j) + (\mu - \mu_j)^2 \right] - n(\bar{Y}_{..} - \mu)^2$$

Notice that

$$\bar{Y}_{\cdot j} \sim N(\mu_j, \sigma^2/n_j) \quad \text{and} \quad \bar{Y}_{..} \sim N(\mu_j, \sigma^2/n)$$

⇒

$$\begin{aligned} \mathbb{E}[SSTR] &= \sum_{j=1}^k n_j \left[ \frac{\sigma^2}{n_j} - 2 \times 0 + (\mu - \mu_j)^2 \right] - n \frac{\sigma^2}{n} \\ &= (k-1)\sigma^2 + \sum_{j=1}^k n_j (\mu - \mu_j)^2 \end{aligned}$$

**Remark** When  $\mu_1 = \dots = \mu_j$  then

0.1  $\mathbb{E}[SSTR] = (k-1)\sigma^2$

0.2  $MSTR := \frac{SSTR}{k-1}$  is an unbiased estimator for  $\sigma^2$ .

0.3  $SSTR/\sigma^2 \sim \text{Chi square } (df = k-1)$ .

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When  $\mu_1 = \dots = \mu_j$  then

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Case I. when  $\sigma^2$  is known.

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$$\begin{aligned}SSE &:= \sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{\cdot j})^2 \\&= \sum_{j=1}^k (n_j - 1) \left[ \frac{1}{n_j - 1} \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{\cdot j})^2 \right] \\&= \sum_{j=1}^k (n_j - 1) S_j^2\end{aligned}$$

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$$S_p^2 = \frac{SSE}{\sum_{j=1}^k (n_j - 1)} = \frac{SSE}{n - k}$$

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Proof. We have shown part (a). Part (b) is trickier. Indeed, both parts are a consequence of **Cochran's theorem**<sup>1</sup> ... □

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  - $(n - 2)S_p^2/\sigma^2 \sim \chi^2(n - 2)$
  - $\bar{X} - \bar{Y} \perp S_p^2 \iff SSTR \perp SSE$

Let's see two special cases of

**Thm.** No matter  $H_0 : \mu_1 = \dots = \mu_k$  is true or not

- $SSE/\sigma^2 = (n - k)S_p^2/\sigma^2 \sim \text{Chi square } (df = \sum_{j=1}^k (n_j - 1) = n - k)$
- $SSTR \perp SSE$ .

## Cases

- $k = 1$ , one sample case,  $S_p^2$  is sample variance Chapter 7
  - $(n - 1)S^2/\sigma^2 \sim \chi^2(n - 1)$
  - $SSTR \equiv 0$
  
- $k = 2$ , two sample case Chapter 9
  - $(n - 2)S_p^2/\sigma^2 \sim \chi^2(n - 2)$
  - $\bar{X} - \bar{Y} \perp S_p^2 \iff SSTR \perp SSE$

Under  $H_0 : \mu_1 = \dots = \mu_k$

1.  $SSTR/\sigma^2 \sim \chi^2(k-1)$

2.  $SSE/\sigma^2 \sim \chi^2(n-k)$

3.  $SSTR \perp SSE$

$$\Rightarrow F = \frac{SSTR/(k-1)}{SSE/(n-k)} \sim F(df_1 = k-1, df_2 = n-k)$$

Reject  $H_0$  if  $F \geq F_{1-\alpha, k-1, n-k}$

Under  $H_0 : \mu_1 = \dots = \mu_k$

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Reject  $H_0$  if  $F \geq F_{1-\alpha, k-1, n-k}$

# Total Sum of Squares: SSTOT

$$\text{SSTOT} = \text{SSE} + \text{SSTR}$$

$$\text{SSTOT} := \sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{..})^2$$

||

$$\sum_{j=1}^k \sum_{i=1}^{n_j} \left[ (Y_{ij} - \bar{Y}_{j\cdot}) + (\bar{Y}_{j\cdot} - \bar{Y}_{..}) \right]^2$$

||

$$\sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{j\cdot})^2 + 2 \sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{j\cdot}) (\bar{Y}_{j\cdot} - \bar{Y}_{..}) + \sum_{j=1}^k \sum_{i=1}^{n_j} (\bar{Y}_{j\cdot} - \bar{Y}_{..})^2$$

||

$$\sum_{j=1}^k \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{j\cdot})^2 + 2 \sum_{j=1}^k (\bar{Y}_{j\cdot} - \bar{Y}_{..}) \sum_{i=1}^{n_j} (Y_{ij} - \bar{Y}_{j\cdot}) + \sum_{j=1}^k n_j (\bar{Y}_{j\cdot} - \bar{Y}_{..})^2$$

||

$$\text{SSE} + 0 + \text{SSTR}$$

$$SSTOT = SSE + SSTR$$

↓

$$\frac{SSTOT}{\sigma^2} = \frac{SSE}{\sigma^2} + \frac{SSTR}{\sigma^2}$$

}

}

}

$$\chi^2(n-1) \quad \chi^2(n-k) \quad \perp \quad \chi^2(k-1)$$

Under  $H_0$

✓

Under  $H_0$

# One-way ANOVA Table

Source of Variance	Degree of Freedom (df)	Sum Square (SS)	Mean Square (MS)	F-ratio
Between Groups (Treatment)	k-1	$SSB = \sum_{j=1}^k \left( \frac{T_j^2}{n_j} \right) - \frac{T^2}{n}$ $SSB = \sum_{j=1}^k n_j (\bar{X}_j - \bar{X}_t)^2$	$MSB = \frac{SSB}{k-1}$	$F = \frac{MSB}{MSW}$
Within Groups (Error)	n-k	$SSW = \sum_{j=1}^k \sum_{i=1}^n X_{ij}^2 - \sum_{j=1}^k \left( \frac{T_j^2}{n_j} \right)$ $SSW = \sum_{j=1}^k \sum_{i=1}^n (X_{ij} - \bar{X}_j)^2$	$MSW = \frac{SSW}{n-k}$	
Total	n-1	$SST = \sum_{j=1}^k \sum_{i=1}^n X_{ij}^2 - \frac{T^2}{n}$ $SST = \sum_{j=1}^k \sum_{i=1}^n (X_{ij} - \bar{X}_t)^2$		

- $SST = SSB + SSW$

k: number of groups    n: number of samples  
df: degree of freedom

Source	df	SS	MS	F	P
Treatment	k - 1	SSTR	MSTR	$\frac{MSTR}{MSE}$	$P(F_{k-1, n-k} \geq \text{observed } F)$
Error	n - k	SSE	MSE		
Total	n - 1	SSTOT			

## Common notation

d.f.

k-1 Error sum of squares  
Mean square of error  
(Pooled sample variance)

$$SSE = SSW = SS_{within}$$
$$MSE = MSW = MS_{within} = S_p^2$$

n-k Treatment sum of squares  
Mean square of treatment

$$SSTR = SSB = SS_{between}$$
$$MSTR = MSB = MS_{between}$$

n-1 Total sum of squares:

$$SST = SSTOT$$

## Common notation

d.f.

**k-1** Error sum of squares

Mean square of error

(Pooled sample variance)

$$SSE = SSW = SS_{within}$$

$$MSE = MSW = MS_{within} = S_p^2$$

**n-k** Treatment sum of squares

Mean square of treatment

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## One way ANOVA v.s. Two sample $t$ -test

Let  $X_1, \dots, X_n$  and  $Y_1, \dots, Y_m$  be samples from  $N(\mu_X, \sigma^2)$  and  $N(\mu_Y, \sigma^2)$ , respectively.

Recall

$$1. SSTR/\sigma^2 = \frac{(\bar{X} - \bar{Y})^2}{\sigma^2 \left(\frac{1}{n} + \frac{1}{m}\right)} \sim \chi^2(1)$$

$$2. SSE/\sigma^2 = (n + m - 2)S_p^2/\sigma^2 \sim \chi^2(n + m - 2)$$

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||  
 $T^2$

$$\Rightarrow \alpha = \mathbb{P}(|T| \geq t_{\alpha/2, n+m-2}) = \mathbb{P}(T^2 \geq t_{\alpha/2, n+m-2}^2) = \mathbb{P}(F \geq F_{1-\alpha, 1, n+m-2})$$

Equivalent!

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Equivalent!

E.g. 1 Study the relation between smoking and heart rates.

Generations of athletes have been cautioned that cigarette smoking impedes performance. One measure of the truth of that warning is the effect of smoking on heart rate. In one study, six nonsmokers, six light smokers, six moderate smokers, and six heavy smokers each engaged in sustained physical exercise. Table 8.1.1 lists their heart rates after they had rested for three minutes.

Show whether smoking affects heart rates at  $\alpha = 0.05$ .

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	Nonsmokers	Light Smokers	Moderate Smokers	Heavy Smokers
	69	55	66	91
	52	60	81	72
	71	78	70	81
	58	58	77	67
	59	62	57	95
	65	66	79	84
<i>Averages:</i>	62.3	63.2	71.7	81.7

Show whether smoking affects heart rates at  $\alpha = 0.05$ .



Sol. Let  $\mu_1, \dots, \mu_4$  be the true heart rates.

Test  $H_0 : \mu_0 = \dots = \mu_4$  or not.

Critical region:

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Critical region:

Let  $\alpha = 0.05$ . For these data,  $k = 4$  and  $n = 24$ , so  $H_0 : \mu_1 = \mu_2 = \mu_3 = \mu_4$  should be rejected if

$$F = \frac{SSTR/(4-1)}{SSE/(24-4)} \geq F_{1-0.05, 4-1, 24-4} = F_{.95, 3, 20} = 3.10$$

(see Figure 12.2.2).

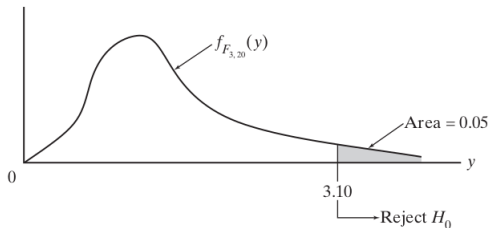


Figure 12.2.2

Computing....

## Computing....

**Table 12.2.1**

	Nonsmokers	Light Smokers	Moderate Smokers	Heavy Smokers
	69	55	66	91
	52	60	81	72
	71	78	70	81
	58	58	77	67
	59	62	57	95
	65	66	79	84
$T_{.j}$	374	379	430	490
$\bar{Y}_{.j}$	62.3	63.2	71.7	81.7

The overall sample mean,  $\bar{Y}_{..}$ , is given by

$$\begin{aligned}\bar{Y}_{..} &= \frac{1}{n} \sum_{j=1}^k T_{.j} = \frac{374 + 379 + 430 + 490}{24} \\ &= 69.7\end{aligned}$$

Therefore,

$$\begin{aligned}SSTR &= \sum_{j=1}^4 n_j (\bar{Y}_{.j} - \bar{Y}_{..})^2 = 6[(62.3 - 69.7)^2 + \dots + (81.7 - 69.7)^2] \\ &= 1464.125\end{aligned}$$

Similarly,

$$\begin{aligned}SSE &= \sum_{j=1}^4 \sum_{i=1}^6 (Y_{ij} - \bar{Y}_{.j})^2 = [(69 - 62.3)^2 + \dots + (65 - 62.3)^2] \\ &\quad + \dots + [(91 - 81.7)^2 + \dots + (84 - 81.7)^2] \\ &= 1594.833\end{aligned}$$

The observed test statistic, then, equals 6.12:

$$F = \frac{1464.125/(4 - 1)}{1594.833/(24 - 4)} = 6.12$$

Since  $6.12 > F_{.95,3,20} = 3.10$ ,  $H_0: \mu_1 = \mu_2 = \mu_3 = \mu_4$  should be rejected. These data support the contention that smoking influences a person's heart rate.

Figure 12.2.3 shows the analysis of these data summarized in the ANOVA table format. Notice that the small  $P$ -value ( $= 0.004$ ) is consistent with the conclusion that  $H_0$  should be rejected.

Source	df	SS	MS	F	P
Treatment	3	1464.125	488.04	6.12	0.004
Error	20	1594.833	79.74		
Total	23	3058.958			

**Figure 12.2.3**



```
1 > Input <-c("
2 + rates group
3 + 69 non
4 + 52 non
5 + 71 non
6 + 58 non
7 + 59 non
8 + 65 non
9 + 55 light
10 + 60 light
11 + 78 light
12 + 58 light
13 + 62 light
14 + 66 light
15 + 66 moderate
16 + 81 moderate
17 + 70 moderate
18 + 77 moderate
19 + 57 moderate
20 + 79 moderate
21 + 91 heavy
22 + 72 heavy
23 + 81 heavy
24 + 67 heavy
25 + 95 heavy
26 + 84 heavy
27 + ")
28 > Data = read.table(textConnection(Input),
29 + header=TRUE)
```

```
1 > Data
2 rates group
3 1 69 non
4 2 52 non
5 3 71 non
6 4 58 non
7 5 59 non
8 6 65 non
9 7 55 light
10 8 60 light
11 9 78 light
12 10 58 light
13 11 62 light
14 12 66 light
15 13 66 moderate
16 14 81 moderate
17 15 70 moderate
18 16 77 moderate
19 17 57 moderate
20 18 79 moderate
21 19 91 heavy
22 20 72 heavy
23 21 81 heavy
24 22 67 heavy
25 23 95 heavy
26 24 84 heavy
```



```

1 > # Check the levels
2 > levels(Data$group)
3 [1] "heavy" "light" "moderate" "non"
4 > # Order the groups
5 > Data$group <- ordered(Data$group, levels = c("non", "light", "moderate", "heavy"))
6 > levels(Data$group)
7 [1] "non" "light" "moderate" "heavy"

```

```

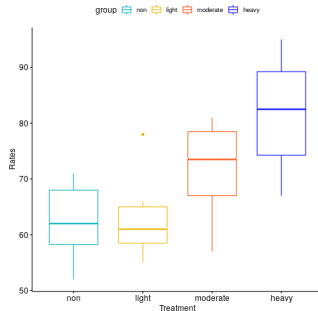
1 > # Compute summary statistics by groups
2 > # including count, mean, sd:
3 > library(dplyr) # a grammar of data manipulation
4 > group_by(Data, group) %>%
5 + summarise(
6 +   count = n(),
7 +   mean = mean(rates, na.rm = TRUE),
8 +   sd = sd(rates, na.rm = TRUE)
9 + )
10 # A tibble : 4 x 4
11   group  count mean  sd
12   <ord>  <int> <dbl> <dbl>
13 1 non      6  62.3  7.26
14 2 light    6  63.2  8.16
15 3 moderate 6  71.7  9.16
16 4 heavy    6  81.7 10.8

```

```

1 # Box plots
2 # ++++++
3 # Plot rates by group and color by group
4 library (ggpubr)
5 png("Case_12-2-1-ggboxplot.png")
6 ggboxplot(Data, x = "group", y = "rates",
7           color = "group", palette = c("#00AFBB", "#E7B800", "#FC4E07", "blue"),
8           order = c("non", "light", "moderate", "heavy"),
9           ylab = "Rates", xlab = "Treatment")
10 dev.off ()

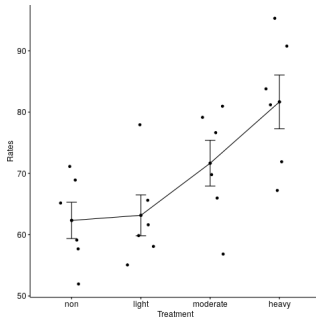
```



```

1 # Mean plots
2 # ++++++
3 # Plot rates by group
4 # Add error bars: mean_se
5 # (other values include: mean_sd, mean_ci, median_iqr, ....)
6 png("Case_12-2-1-ggline.png")
7 library(ggpubr)
8 ggline(Data, x = "group", y = "rates",
9         add = c("mean_se", "jitter "),
10        order = c("non", "light", "moderate", "heavy"),
11        ylab = "Rates", xlab = "Treatment")
12 dev.off()

```



```

1 > # Compute the analysis of variance
2 > res.aov <- aov(rates ~ group, data = Data)
3 > # Summary of the analysis
4 > summary(res.aov)
5           Df Sum Sq Mean Sq F value Pr(>F)
6 group         3  1464   488.0    6.12 0.00398 **
7 Residuals    20  1595    79.7
8 ---
9 Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

1 > # Tukey multiple multiple – comparisons
2 > TukeyHSD(res.aov)
3   Tukey multiple comparisons of means
4     95% family – wise confidence level
5
6 Fit: aov(formula = rates ~ group, data = Data)
7
8 $group
9
10      diff      lwr      upr    p adj
11 light – non  0.8333333 –13.596955 15.26362 0.9984448
12 moderate – non 9.3333333 –5.096955 23.76362 0.2978123
13 heavy – non   19.3333333 4.903045 33.76362 0.0063659
14 moderate – light 8.5000000 –5.930289 22.93029 0.3755571
15 heavy – light  18.5000000 4.069711 32.93029 0.0091463
16 heavy – moderate 10.0000000 –4.430289 24.43029 0.2438158

```

1. diff: difference between means of the two groups
2. lwr, upr: the lower and the upper end point of the C.I. at 95% (default)
3. p adj: p-value after adjustment for the multiple comparisons

Inferences

if  $p\text{-value} \leq 0.05$   $\iff$  if zero is in the C.I.

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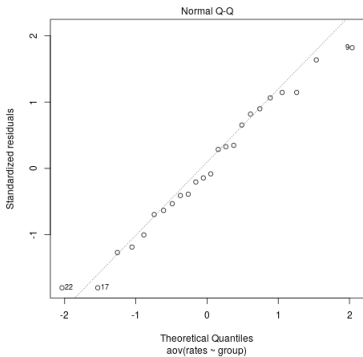
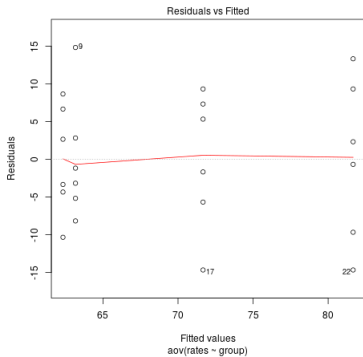


```

1 > # Or one may use multcomp package or multiple comparisons
2 > library (multcomp)
3 > summary(glht(res.aov, linfct = mcp(group = "Tukey")))
4
5 Simultaneous Tests for General Linear Hypotheses
6
7 Multiple Comparisons of Means: Tukey Contrasts
8
9
10 Fit: aov(formula = rates ~ group, data = Data)
11
12 Linear Hypotheses:
13
14 Estimate Std. Error t value Pr(>|t|)
15 light - non == 0 0.8333 5.1556 0.162 0.99844
16 moderate - non == 0 9.3333 5.1556 1.810 0.29776
17 heavy - non == 0 19.3333 5.1556 3.750 0.00629 **
18 moderate - light == 0 8.5000 5.1556 1.649 0.37544
19 heavy - light == 0 18.5000 5.1556 3.588 0.00901 **
20 heavy - moderate == 0 10.0000 5.1556 1.940 0.24382
21 ---
22 Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
  (Adjusted p values reported -- single-step method)

```

```
1 # Check ANOVA assumptions: test validity?  
2 # diagnostic plots  
3 layout(matrix(c(1,2),1,2)) # optional 1x2 graphs/page  
4 plot(res.aov,c(1,2))
```



## 1. Residuals vs Fitted: test homogeneity of variances

One can also use Levene's test for this purpose:

```
1 > # Use Levene's test to test homogeneity of variances
2 > library(car)
3 > leveneTest(rates ~ group, data = Data)
4 Levene's Test for Homogeneity of Variance (center = median)
5   Df F value Pr(>F)
6 group 3  0.3885 0.7625
7      20
```

## 2. Normal Q-Q plot: Test normality. (It should be close to diagonal line.)

One can also use Shapiro-Wilk test:

```
1 # Extract the residuals
2 > aov_residuals <- residuals(object = res.aov )
3 > # Run Shapiro-Wilk test
4 > shapiro.test(x = aov_residuals )
5
6 Shapiro-Wilk normality test
7
8 data:  aov_residuals
9 W = 0.9741, p-value = 0.7677
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9 W = 0.9741, p-value = 0.7677
```

# Non-parametric alternative to one-way ANOVA test

```
1 > # Non-parametric alternative to one-way ANOVA test
2 > # a non-parametric alternative to one-way ANOVA
3 > # is Kruskal-Wallis rank sum test, which can be
4 > # used when ANNOVA assumptions are not met.
5 > kruskal.test(rates ~ group, data = Data)
6
7 Kruskal-Wallis rank sum test
8
9 data: rates by group
10 Kruskal-Wallis chi-squared = 10.729, df = 3, p-value = 0.01329
```

See Section 4 of Chapter 14 for more details.

# Plan

§ 12.1 Introduction

§ 12.2 The  $F$  Test

§ 12.3 Multiple Comparisons: Turkey's Method



# Chapter 12. The Analysis of Variance

§ 12.1 Introduction

§ 12.2 The  $F$  Test

§ 12.3 Multiple Comparisons: Turkey's Method

## § 12.3 Multiple Comparisons: Turkey's Method



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2. The Tukey range test, the Tukey lambda distribution, the Tukey test of additivity, and the Teichmüller-Tukey lemma all bear his name.
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$N(\mu_1, \sigma^2)$	$N(\mu_2, \sigma^2)$	$\dots$	$N(\mu_k, \sigma^2)$
$Y_{11}$	$Y_{12}$	$\dots$	$Y_{1k}$
$Y_{21}$	$Y_{22}$	$\dots$	$Y_{2k}$
$\vdots$	$\vdots$	$\vdots$	$\vdots$
$Y_{r1}$	$Y_{r2}$	$\dots$	$Y_{rk}$

Goal For any  $i \neq j$ , test

$$H_0 : \mu_i = \mu_j \quad \text{v.s.} \quad H_1 : \mu_i \neq \mu_j$$

at the  $\alpha$  level of significance defined as

$$\mathbb{P} \left( \bigcup_{j=1}^{\binom{k}{2}} E_j \right) = \alpha$$

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**Goal'** Simultaneous C.I.'s for  $\binom{k}{2}$  pairs of means:

Given  $\alpha$ , find  $l_{ij}$ , the C.I. for  $\mu_i - \mu_j$  (with  $i, j = 1, \dots, k$  and  $i \neq j$ ), s.t.

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Suppose  $\mathbb{P}(E_j) = \alpha_*$ . Then

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Hence,

$$\alpha_* \approx 1 - (1 - \alpha)^{1/\binom{k}{2}}$$

E.g.,  $\alpha = 0.05$

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# Bonferroni's method

— A straightforward method

$$\mathbb{P}(\mu_i - \mu_j \in I_{ij}, \forall i \neq j)$$

$$\mathbb{P}\left(\bigcap_{i \neq j} \mu_i - \mu_j \in I_{ij}\right)$$

$$1 - \mathbb{P}\left(\bigcup_{i \neq j} \mu_i - \mu_j \notin I_{ij}\right)$$

$$1 - \sum_{i \neq j} \mathbb{P}(\mu_i - \mu_j \notin I_{ij})$$

$$1 - \binom{k}{2} \alpha_*$$

1. If we choose  $\alpha_* = \alpha / \binom{k}{2}$ ,

2. let  $I_{ij}$  be the  $(1 - \alpha_*)$  100% C.I.  $i \neq j$

↓

$$\mathbb{P}(\mu_i - \mu_j \in I_{ij}, \forall i \neq j)$$

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$$1 - \mathbb{P}\left(\bigcup_{i \neq j} \mu_i - \mu_j \notin I_{ij}\right)$$

$$\forall i$$
$$1 - \sum_{i \neq j} \mathbb{P}(\mu_i - \mu_j \notin I_{ij})$$

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**Remark** This is an approximation. The resulting C.I. are in general too wide.

The exact, and much more precise, solution is given by J.W. Turkey.

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## Tukey's HSD (honestly significant difference) test

Let's construct  $(1 - \alpha)100\%$  C.I.'s simultaneously for all pairs.

$$\begin{aligned} \mathbb{P} \left( \left| (\bar{Y}_{.i} - \mu_i) - (\bar{Y}_{.j} - \mu_j) \right| \leq \mathcal{E}, \quad \forall i \neq j \right) &= 1 - \alpha \\ &\parallel \\ \mathbb{P} \left( \max_i (\bar{Y}_{.i} - \mu_i) - \min_j (\bar{Y}_{.j} - \mu_j) \leq \mathcal{E} \right) \\ &\parallel \\ \mathbb{P} \left( \max_i \bar{Y}_{.i} - \min_j \bar{Y}_{.j} \leq \mathcal{E} \right) \end{aligned}$$

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**Def.** Let  $W_1, \dots, W_k$  be  $k$  i.i.d. r.v.'s from  $N(\mu, \sigma^2)$ . Let  $R$  denote their range:

$$R = \max_i W_i - \min_i W_i.$$

Let  $S^2$  be an unbiased estimator for  $\sigma^2$  independent of the  $W_i$ 's and based on  $\nu$  df. Define the **Studentized range**,  $Q_{k,\nu}$ , to be the ratio:

$$Q_{k,\nu} := \frac{R}{S}.$$

**Remark** 0.1 We need  $R \perp S$  to mimic Student's t-distribution.

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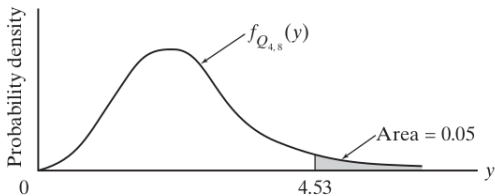
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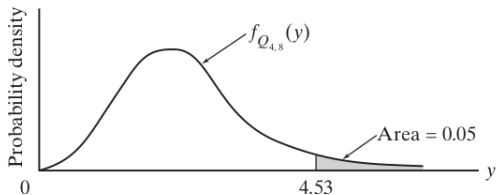
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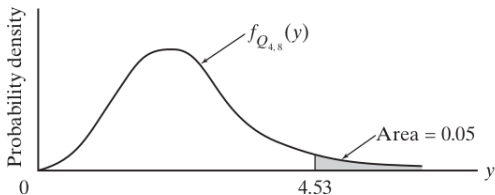
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$Q_{k,\nu} \sim$  **Studentized range distribution** with parameters  $k$  and  $\nu$ .

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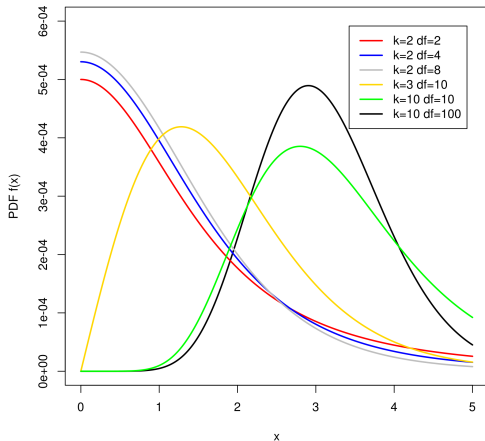
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Let's find one example that all requirements of the  $Q_{k,\nu}$  are satisfied.

1. Take  $W_j = \bar{Y}_{.j} - \mu_j, j = 1, \dots, k \implies W_j \sim N(0, \sigma^2/r)$ .

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& \mathbb{P} \left( \bar{Y}_{.i} - \bar{Y}_{.j} - \frac{Q_{\alpha, k, rk-k}}{\sqrt{r}} \sqrt{MSE} \leq \mu_i - \mu_j \leq \bar{Y}_{.i} - \bar{Y}_{.j} + \frac{Q_{\alpha, k, rk-k}}{\sqrt{r}} \sqrt{MSE}, \forall i \neq j \right)
\end{aligned}$$

Therefore, for all  $i \neq j$ , the  $100(1 - \alpha)\%$  C.I. for  $\mu_i - \mu_j$  is

$$\bar{Y}_{.i} - \bar{Y}_{.j} \pm \frac{Q_{\alpha, k, rk-k}}{\sqrt{2}} \sqrt{MSE} \sqrt{\frac{2}{r}}$$

To test  $H_0 : \mu_i = \mu_j$  for specific  $i \neq j$ , reject  $H_0$  in favor of  $H_1 : \mu_i \neq \mu_j$  if the C.I. does NOT contain 0, at the  $\alpha$  level of significance.  $\square$

**Note:** When sample sizes are not equal, use the **Tukey-Kramer method**:

$$\bar{Y}_{.i} - \bar{Y}_{.j} \pm \frac{Q_{\alpha, k, rk-k}}{\sqrt{2}} \sqrt{MSE} \sqrt{\frac{1}{r_i} + \frac{1}{r_j}}$$

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**Note:** When sample sizes are not equal, use the **Tukey-Kramer method**:

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E.g. 2 A certain fraction of antibiotics injected into the bloodstream are “bound” to serum proteins. This phenomenon bears directly on the effectiveness of the medication, because the binding decreases the systemic uptake of the drug. Table below lists the binding percentages in bovine serum measured for five widely prescribed antibiotics. Which antibiotics have similar binding properties, and which are different?

E.g. 2 A certain fraction of antibiotics injected into the bloodstream are “bound” to serum proteins. This phenomenon bears directly on the effectiveness of the medication, because the binding decreases the systemic uptake of the drug. Table below lists the binding percentages in bovine serum measured for five widely prescribed antibiotics. Which antibiotics have similar binding properties, and which are different?

	Penicillin G	Tetra- cycline	Strepto- mycin	Erythro- mycin	Chloram- phenicol
	29.6	27.3	5.8	21.6	29.2
	24.3	32.6	6.2	17.4	32.8
	28.5	30.8	11.0	18.3	25.0
	32.0	34.8	8.3	19.0	24.2
$T_j$	114.4	125.5	31.3	76.3	111.2
$\bar{Y}_j$	28.6	31.4	7.8	19.1	27.8

To answer that question requires that we make all  $\binom{5}{2} = 10$  pairwise comparisons of  $\mu_i$  versus  $\mu_j$ . First,  $MSE$  must be computed. From the entries in Table 12.3.1,

$$SSE = \sum_{j=1}^5 \sum_{i=1}^4 (Y_{ij} - \bar{Y}_{.j})^2 = 135.83$$

so  $MSE = 135.83/(20 - 5) = 9.06$ . Let  $\alpha = 0.05$ . Since  $n - k = 20 - 5 = 15$ , the appropriate cutoff from the studentized range distribution is  $Q_{.05,5,15} = 4.37$ . Therefore,  $D = 4.37/\sqrt{4} = 2.185$  and  $D\sqrt{MSE} = 6.58$ .

**Table 12.3.2**

Pairwise Difference	$\bar{Y}_{.i} - \bar{Y}_{.j}$	Tukey Interval	Conclusion
$\mu_1 - \mu_2$	-2.8	(-9.38, 3.78)	NS
$\mu_1 - \mu_3$	20.8	(14.22, 27.38)	Reject
$\mu_1 - \mu_4$	9.5	(2.92, 16.08)	Reject
$\mu_1 - \mu_5$	0.8	(-5.78, 7.38)	NS
$\mu_2 - \mu_3$	23.6	(17.02, 30.18)	Reject
$\mu_2 - \mu_4$	12.3	(5.72, 18.88)	Reject
$\mu_2 - \mu_5$	3.6	(-2.98, 10.18)	NS
$\mu_3 - \mu_4$	-11.3	(-17.88, -4.72)	Reject
$\mu_3 - \mu_5$	-20.0	(-26.58, -13.42)	Reject
$\mu_4 - \mu_5$	-8.7	(-15.28, -2.12)	Reject

```

1 > # Case Study 12.3.1
2 > # Input data first
3 > Input <- c("
4 + rates group
5 + 29.6 M1
6 + 24.3 M1
7 + 28.5 M1
8 + 32.0 M1
9 + 27.3 M2
10 + 32.6 M2
11 + 30.8 M2
12 + 34.8 M2
13 + 5.8 M3
14 + 6.2 M3
15 + 11.0 M3
16 + 8.3 M3
17 + 21.6 M4
18 + 17.4 M4
19 + 18.3 M4
20 + 19.0 M4
21 + 29.2 M5
22 + 32.8 M5
23 + 25.0 M5
24 + 24.2 M5
25 + ")
26 > Data = read.table(
27   textConnection(Input),
28   +   header=TRUE)

```

```

1 > # Compute one-way ANOVA test
2 > res.aov <- aov(rates ~ group, data = Data)
3 > # Summary of the analysis
4 > summary(res.aov)
5           Df Sum Sq Mean Sq F value Pr(>F)
6 group      4 1480.8   370.2  40.88 6.74e-08 ***
7 Residuals 15  135.8     9.1
8 ---
9 Signif. codes:  0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1

```

```

1 > # Tukey multiple pairwise-comparisons
2 > TukeyHSD(res.aov)
3 Tukey multiple comparisons of means
4 95% family-wise confidence level
5
6 Fit: aov(formula = rates ~ group, data = Data)
7
8 $group
9      diff      lwr      upr    p adj
10 M2-M1 2.775 -3.795401 9.345401 0.6928357
11 M3-M1 -20.775 -27.345401 -14.204599 0.0000006
12 M4-M1 -9.525 -16.095401 -2.954599 0.0034588
13 M5-M1 -0.800 -7.370401 5.770401 0.9952758
14 M3-M2 -23.550 -30.120401 -16.979599 0.0000001
15 M4-M2 -12.300 -18.870401 -5.729599 0.0003007
16 M5-M2 -3.575 -10.145401 2.995401 0.4737713
17 M4-M3 11.250 4.679599 17.820401 0.0007429
18 M5-M3 19.975 13.404599 26.545401 0.0000010
19 M5-M4 8.725 2.154599 15.295401 0.0071611

```

```

1 > round(TukeyHSD(res.aov)$group,2)
2           diff      lwr      upr    p adj
3 M2-M1    2.78    -3.80    9.35    0.69
4 M3-M1   -20.77   -27.35   -14.20    0.00
5 M4-M1    -9.52   -16.10    -2.95    0.00
6 M5-M1    -0.80    -7.37    5.77    1.00
7 M3-M2   -23.55   -30.12   -16.98    0.00
8 M4-M2   -12.30   -18.87    -5.73    0.00
9 M5-M2    -3.58   -10.15    3.00    0.47
10 M4-M3    11.25    4.68    17.82    0.00
11 M5-M3    19.97    13.40    26.55    0.00
12 M5-M4    8.73     2.15    15.30    0.01
13 ---
14 Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 '
    ' 1
15 (Adjusted p values reported -- single-step method)

```

**Table 12.3.2**

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$\mu_4 - \mu_5$	-8.7	(-15.28, -2.12)	Reject

```

1 > # Or one may use multcomp package or multiple comparisons
2 > library (multcomp)
3 > summary(glht(res.aov, linfct = mcp(group = "Tukey")))

```

## Simultaneous Tests for General Linear Hypotheses

### Multiple Comparisons of Means: Tukey Contrasts

```

9
10 Fit: aov(formula = rates ~ group, data = Data)

```

#### Linear Hypotheses:

	Estimate	Std. Error	t value	Pr(> t )
M2 - M1 == 0	2.775	2.128	1.304	0.69283
M3 - M1 == 0	-20.775	2.128	-9.764	< 0.001 ***
M4 - M1 == 0	-9.525	2.128	-4.477	0.00348 **
M5 - M1 == 0	-0.800	2.128	-0.376	0.99528
M3 - M2 == 0	-23.550	2.128	-11.068	< 0.001 ***
M4 - M2 == 0	-12.300	2.128	-5.781	< 0.001 ***
M5 - M2 == 0	-3.575	2.128	-1.680	0.47374
M4 - M3 == 0	11.250	2.128	5.287	< 0.001 ***
M5 - M3 == 0	19.975	2.128	9.388	< 0.001 ***
M5 - M4 == 0	8.725	2.128	4.101	0.00717 **

```

24 ---

```

```

25 Signif. codes: 0 '***' 0.001 '**' 0.01 '*' 0.05 '.' 0.1 ' ' 1
26 (Adjusted p values reported -- single-step method)

```

	Estimate	Std. Error	t value	Pr(> t )
1 M2 - M1 == 0	2.775	2.128	1.304	0.69283
2 M3 - M1 == 0	-20.775	2.128	-9.764	< 0.001 ***
3 M4 - M1 == 0	-9.525	2.128	-4.477	0.00348 **
4 M5 - M1 == 0	-0.800	2.128	-0.376	0.99527
5 M3 - M2 == 0	-23.550	2.128	-11.068	< 0.001 ***
6 M4 - M2 == 0	-12.300	2.128	-5.781	< 0.001 ***
7 M5 - M2 == 0	-3.575	2.128	-1.680	0.47371
8 M4 - M3 == 0	11.250	2.128	5.287	< 0.001 ***
9 M5 - M3 == 0	19.975	2.128	9.388	< 0.001 ***
10 M5 - M4 == 0	8.725	2.128	4.101	0.00719 **
11				

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## Two more examples of ANOVA using R

E.g. 1 <http://www.sthda.com/english/wiki/one-way-anova-test-in-r>

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