#### Math 362: Mathematical Statistics II

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# Chapter 13. Randomized Block Designs

§ 13.1 Introduction

§ 13.2 The F Test for a Randomized Block Design

§ 13.A Appendix: Some Discussions and Extensions

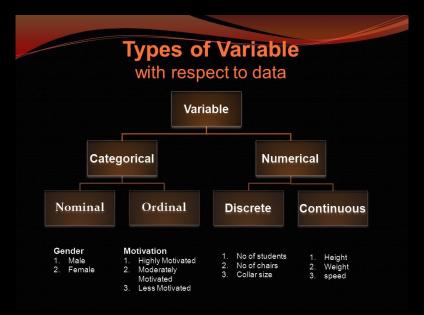
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## Chapter 13. Randomized Block Designs

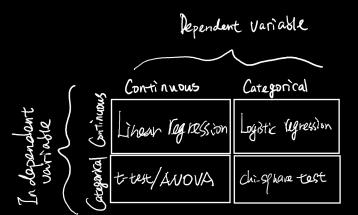
§ 13.1 Introduction

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		Numeriz XI Values	Categorisal Values			
N Semples	Sample 1	y, y, m	C11 C15			
	Sample 2	y y 2m	C21 C2P			
	::		; ;			
(	Sample 11	y y	Cn. · · · Cnp			
		m	P			



#### Indep. v.s. Dependent

#### 1. Categorical v.s. Continuous

1.1 
$$p = 1, m = 1,$$
 One-way ANOVA

 1.2  $p = 2, m = 1,$ 
 Two-way ANOVA

 1.3  $p \ge 3, m = 1,$ 
 $p$ -way ANOVA

 1.4  $p = 1, m \ge 2,$ 
 One-way MANOVA

 1.5  $p = 2, m \ge 2,$ 
 Two-way MANOVA

 1.6  $p > 3, m > 2,$ 
 $p$ -way ANOVA

#### 2. Continuous v.s. Continuous

2.1 
$$\textit{m}_{\textit{ind}} = 1, \textit{m}_{\textit{dep}} = 1,$$
 Simple linear regression 2.2  $\textit{m}_{\textit{ind}} \geq 2$  Multiple linear regression 2.3  $\textit{m}_{\textit{dep}} \geq 2$  Multivariate linear regression

<sup>&</sup>lt;sup>a</sup>MANOVA refers to the multivariate analysis of variance ANOVA refers to the univariate analysis of variance.

### E.g. One example for MANOVA<sup>1</sup>.



<sup>1</sup>http://www.sthda.com/english/wiki/
manova-test-in-r-multivariate-analysis-of-variance







**Iris Versicolor** 

Iris Setosa

Iris Virginica

> librar	> library (datasets)												
> data(	> data(iris)												
> summ	> summary(iris)												
Sepal.Length Se			al.Width Petal.Lengt		.Length	Petal. Width							
	Species												
Min.	:4.300	Min.	:2.000	Min.	:1.000	Min.	:0.100	setosa					
	:50												
1st Qu.:5.100		1st Qu.:2.800		1st Qu.:1.600		1st Qu.:0.300		versicolor					
:50													
Median:5.800		Median :3.000		Median :4.350		Median:1.300		virginica					
	:50												
Mean	:5.843	Mean	:3.057	Mean	:3.758	Mean	:1.199						
3rd Qu.:6.400		3rd Qı	3rd Qu.:3.300 3rd		Qu.:5.100 3rd Qu.:1.800								
Max.	:7.900	Max.	:4.400	Max.	:6.900	Max.	:2.500						
> my_data <- iris													
> my_d	> my data												
Sep	Sepal.Length Sepal.Width Petal.Length Petal.Width Species												
	5.1		3.5		.4	0.2	setosa						
2	4.9		3.0		.4	0.2	setosa						
3	4.7		3.2		.3	0.2	setosa						
4	4.6		3.1		.5	0.2	setosa						
5	5.0		3.6		.4	0.2	setosa						
6	5.4		3.9		.7	0.4	setosa						
7	4.6		3.4		.4	0.3	setosa						
8	5.0		3.4		.5	0.2	setosa						
9	4.4		2.9		.4	0.2	setosa						
10	4.9		3.1		.5	0.1	setosa						

```
2 > res.man <- manova(cbind(Sepal.Length, Petal.Length) ~ Species, data = iris)
3 > summary(res.man)
            Df Pillai approx F num Df den Df Pr(>F)
            2 0.9885 71.829 4 294 < 2.2e-16 ***
5 Species
6 Residuals 147
8 Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' 1
9 > # Look to see which differ
10 > summary.aov(res.man)
   Response Sepal.Length:
              Df Sum Sq Mean Sq F value Pr(>F)
13 Species 2 63.212 31.606 119.26 < 2.2e-16 ***
   Residuals 147 38.956 0.265
   Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '. ' 0.1 ' ' 1
   Response Petal.Length:
              Df Sum Sq Mean Sq F value Pr(>F)
20 Species 2 437.10 218.551 1180.2 < 2.2e-16 ***
   Residuals 147 27.22 0.185
23 Signif. codes: 0 '*** 0.001 '** 0.01 '* 0.05 '.' 0.1 ' ' 1:w
```

Concl.: Two variables are highly significantly different among species.