# More on linear regression analysis

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## What we are going to learn

• Interaction effects

- File: birthsmokers.txt
- **N** = 32 births
- Baby's birth weight is related mother's smoking habits during pregnancy
- Response (y): birth weight in grams of baby
- Potential predictor (x<sub>1</sub>):
  - **smoking** status of mother (yes or no)
  - Potential predictor  $(x_2)$ : length of **gestation** in weeks

• The model:

$$y_i = b_0 + b_1 x_{i1} + b_2 x_{i2} + e_i$$

- $y_i$  is the birth weight of baby *i* in grams
- $x_{i1}$  is the length of gestation of baby *i* in weeks
- $x_{l2} = 1$ , if baby *i*'s mother smoked and  $x_{l2} = 0$ , if not
- $\varepsilon_i$  error terms follow a normal distribution with mean 0 and equal variance  $\sigma^2$ .

• The model:

$$y_i = b_0 + b_1 x_{i1} + b_2 x_{i2} + e_i$$

The rehression equation based on actual data

#### Weight = -2390 + 143\*Gest – 245\*Smoking

The rehression equation based on actual data
 Weight = -2390 + 143\*Gest – 245\*Smoking



#### No interaction effects!

#### **Interaction effects**

- Depression.txt, 36 patients with depression
- Comparing the effectiveness of 3 treatments for severe depression (A, B, and C)
- y<sub>i</sub> = measure of the effectiveness of the treatment for individual *i*
- $x_{i1}$  = age (in years) of individual *i*
- x<sub>12</sub> = 0 if individual *i* received treatment A, 1 if treatment B, 2 if treatment C

#### **Interaction effects**



#### **Interaction model**

• The "main effect" regression model

$$y_i = b_0 + b_1 x_{i1} + b_2 x_{i2} + e_i$$

• The "interaction effect" regression model

$$y_i = b_0 + b_1 x_{i1} + b_2 x_{i2} + b_3 x_{i1} x_{i2} + e_i$$

#### **R** codes

```
dep =
read.table("/Users/tuannguyen/Documents/_Viet
nam2012/Can Tho /Datasets/depression.txt",
header=T)
```

```
attach(dep)
```

```
res = lm(y \sim age+TRT)
```

```
summary(res)
```

#### **Results of main effect model**

Coefficients:

	Est	tim	ate	Std.	Erı	ror t	C V	value	e Pr	(> t	)		
(Intercept	) 32	. 54	335	3	8.581	105	S	9.088	2.	23e-1	LO	***	5
age	0	. 66	446	C	0.069	978	S	9.522	7.	42e-1	L1	***	5
TRTB	-9	. 80	758	2	2.464	471	-3	3.979	0.	00037	71	***	5
TRTC	-10	. 25	276	2	2.465	542	-4	1.159	0.	00022	24	***	5
Signif. co `′1	des:	0	`***	<b>''</b> 0.	001	`**'	C	0.01	\*/	0.05	5 `	• • *	0.1

Residual standard error: 6.035 on 32 degrees of freedom Multiple R-squared: 0.784, Adjusted R-squared: 0.7637 F-statistic: 38.71 on 3 and 32 DF, p-value: 9.287e-11

#### Interpretation of the main model

(Intercept)	32.54335	3.58105	9.088	2.23e-10	***
age	0.66446	0.06978	9.522	7.42e-11	***
TRTB	-9.80758	2.46471	-3.979	0.000371	***
TRTC	-10.25276	2.46542	-4.159	0.000224	***

- Treatment A: y = 32.54 + 0.66\*age
- Treatment B: y = 32.54 + 0.66\*age 9.81= 22.73 + 0.66\*age

Treatment C: y = 32.54 + 0.66\*age - 10.25= 22.29 + 0.66\*age

# The model in graphical format



#### **Residual analysis of "main effect" model**





#### **R** codes for interaction model

```
dep =
read.table("/Users/tuannguyen/Documents/_Viet
nam2012/Can Tho /Datasets/depression.txt",
header=T)
```

```
attach(dep)
int = lm(y ~ age+TRT+age:TRT)
summary(int)
```

#### **Results of interaction effect model**

Coefficients:

	Estimate	Std. Error	t value	Pr(> t )	
(Intercept)	47.51559	3.82523	12.422	2.34e-13	***
age	0.33051	0.08149	4.056	0.000328	***
TRTB	-18.59739	5.41573	-3.434	0.001759	**
TRTC	-41.30421	5.08453	-8.124	4.56e-09	***
age:TRTB	0.19318	0.11660	1.657	0.108001	
age:TRTC	0.70288	0.10896	6.451	3.98e-07	***
Signif. code ` ' 1	es: 0 `***	′′ 0.001 ` <b>*</b> *	*′ 0.01	*′ 0.05	`.′ 0.1

Residual standard error: 3.925 on 30 degrees of freedom Multiple R-squared: 0.9143, Adjusted R-squared: 0.9001 F-statistic: 64.04 on 5 and 30 DF, p-value: 4.264e-15

## "Interpretation" of model

Estimate Std. Error t value Pr(> t )							
(Intercept)	47.51559	3.82523	12.422	2.34e-13	***		
age	0.33051	0.08149	4.056	0.000328	***		
TRTB	-18.59739	5.41573	-3.434	0.001759	**		
TRTC	-41.30421	5.08453	-8.124	4.56e-09	***		
age:TRTB	0.19318	0.11660	1.657	0.108001			
age:TRTC	0.70288	0.10896	6.451	3.98e-07	***		

The model is:

If treatment=A, then  $y = 47.51 + 0.33^{*}age$ 

If treatment=B, then  $y = 47.51 + 0.33^{\circ}age - 18.60 + 0.19^{\circ}age$ 

= 28.91 + 0.52\*age

If treatment=C, then  $y = 47.51 + 0.33^{\circ}age - 41.30 + 0.70^{\circ}age$ 

= 6.21 + 1.03\*age

#### "Interpretation" of model

y<sub>A</sub> = 47.51 + 0.33\*age

- y<sub>B</sub> = 28.91 + 0.52\*age
- y<sub>c</sub> = 6.21 + 1.03\*age



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#### **Residual analysis**





# Normal plot

