



# STA302 METHODS OF DATA ANALYSIS I

MODULE 3: ASSUMPTIONS OF LINEAR REGRESSION MODELS

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# MODULE 3 OUTLINE

1. Introduction to Linear Regression Assumptions
2. Verifying Assumptions using Residual Plots
3. Additional Conditions for Multiple Linear Models

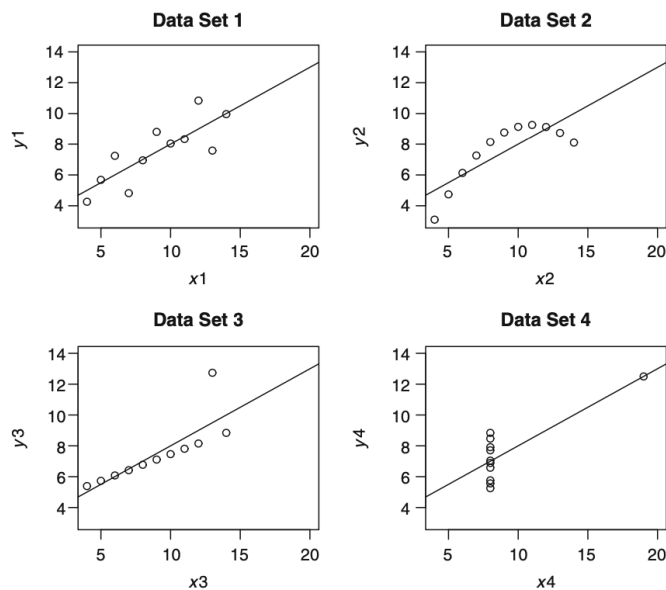


Figure 3.1 Plots of Anscombe's four data sets

From Sheather 's A Modern Approach to Regression with R, pg 46

```
> lm(y1 ~ x1, data=anscombe)
```

Call:  
lm(formula = y1 ~ x1, data = anscombe)

Coefficients:  
(Intercept)            x1  
          3.0001        0.5001

```
> lm(y2 ~ x2, data=anscombe)
```

Call:  
lm(formula = y2 ~ x2, data = anscombe)

Coefficients:  
(Intercept)            x2  
          3.001        0.500

```
> lm(y3 ~ x3, data=anscombe)
```

Call:  
lm(formula = y3 ~ x3, data = anscombe)

Coefficients:  
(Intercept)            x3  
          3.0025        0.4997

```
> lm(y4 ~ x4, data=anscombe)
```

Call:  
lm(formula = y4 ~ x4, data = anscombe)

Coefficients:  
(Intercept)            x4  
          3.0017        0.4999

```
> anscombe <- read.table("anscombe.txt", header=T)
> anscombe
```

	case	x1	x2	x3	x4	y1	y2	y3	y4
1	1	10	10	10	8	8.04	9.14	7.46	6.58
2	2	8	8	8	8	6.95	8.14	6.77	5.76
3	3	13	13	13	8	7.58	8.74	12.74	7.71
4	4	9	9	9	8	8.81	8.77	7.11	8.84
5	5	11	11	11	8	8.33	9.26	7.81	8.47
6	6	14	14	14	8	9.96	8.10	8.84	7.04
7	7	6	6	6	8	7.24	6.13	6.08	5.25
8	8	4	4	4	19	4.26	3.10	5.39	12.50
9	9	12	12	12	8	10.84	9.13	8.15	5.56
10	10	7	7	7	8	4.82	7.26	6.42	7.91
11	11	5	5	5	8	5.68	4.74	5.73	6.89

# ANSCOMBE'S DATASETS

# LINEAR REGRESSION ASSUMPTIONS

1. **Linearity** of the Relationship (also known as Mean Zero Errors) assumption

$$E(\varepsilon | X) = 0 \text{ or } E(Y|X) = X\beta \text{ or } Y = X\beta + \varepsilon$$

2. **Uncorrelated Errors** (sometimes referred to as Independence) assumption

$$\text{Cov}(\varepsilon_i, \varepsilon_j) = 0 \text{ or } \text{Cov}(y_i, y_j) = 0$$

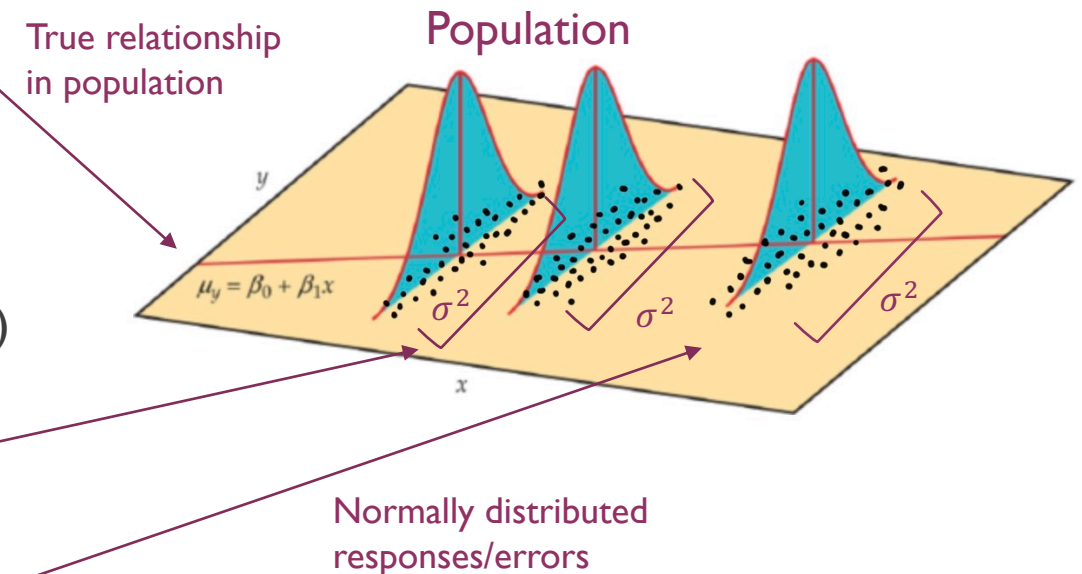
3. **Constant Error Variance** (also known as Homoskedasticity) assumption

$$\text{Var}(\varepsilon|X) = \sigma^2 I \text{ or } \text{Var}(\varepsilon_i|X) = \text{Var}(y_i|X) = \sigma^2$$

4. **Normal Errors** assumption

$$\varepsilon|X \sim N_n(0, \sigma^2 I) \text{ or } Y|X \sim N_n(X\beta, \sigma^2 I) \text{ or } \varepsilon_i \sim N(0, \sigma^2)$$

CC BY-NC-SA 3.0 image by Diane Kiernan in Natural Resources Biometrics



# MORE ON LINEAR REGRESSION ASSUMPTIONS

## *Assumptions relate to properties of the population, not the sample*

- When fitting a model, we implicitly make these assumptions EVERY time

### Uncorrelated Errors Assumption

- Each data point in population must not be related or connected to any other data point
  - i.e. knowing information about one does not give any information about another
- *Examples of violation:* stock price data, weather data, measurements on the same person, etc.

### Linearity/Mean Zero Error Assumption

- Implies two things about the **population relationship**:
  - The true relationship is linear in the coefficients
  - The true relationship is **exactly**  $Y = X\beta + \varepsilon$  with
    - no predictors omitted from  $X$  that should be present,
    - no predictors included in  $X$  that should not be present, and
    - no predictors in  $X$  that are in the wrong functional form
- *Examples of violation:* omitting a predictor known to influence response; fitting a linear model when the truth is  $y_i = \log(\beta_0 + \beta_1 x_i + \varepsilon_i)$ ; including  $x$  when it should be  $x^2$ ; etc.

# MORE ON LINEAR REGRESSION ASSUMPTIONS

## Constant Variance Assumption

- Each conditional distribution must have the same spread
  - So only difference between each one is that the mean changed by a specific amount

## Normality Assumption

- Each conditional distribution must have the same shape
- Harder to verify in small samples
- Not needed to estimate coefficients by least squares

## Importance of Assumptions

- Linearity ensures we estimate coefficients unbiasedly
- Uncorrelated errors ensure correct precision of estimates
- Constant variance ensures we obtain reasonable estimates of variability for all conditional means
- Normality allows us to utilize properties of Normal random variables for inferential purposes (e.g. computing confidence intervals)
- **Nothing stops us from fitting an invalid model with violated assumptions**

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2. Verifying Assumptions using Residual Plots
3. Additional Conditions for Multiple Linear Models

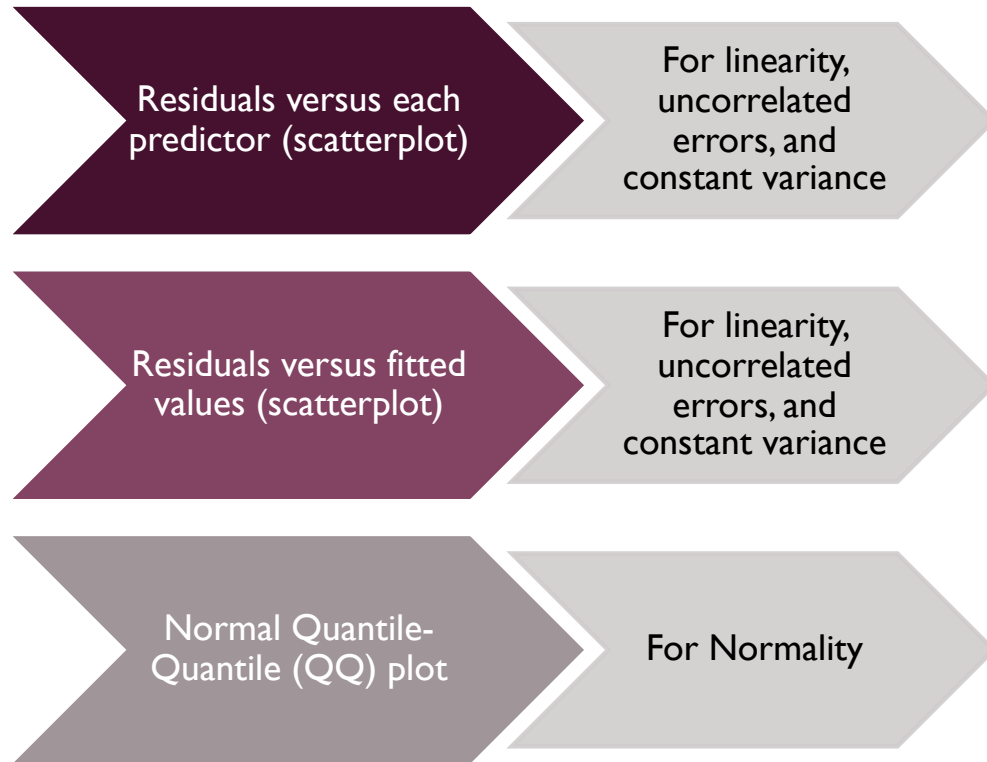
# CHECK ASSUMPTIONS USING RESIDUALS

Population Error Assumptions:  $\varepsilon|X \sim N_n(\mathbf{0}, \sigma^2 I)$

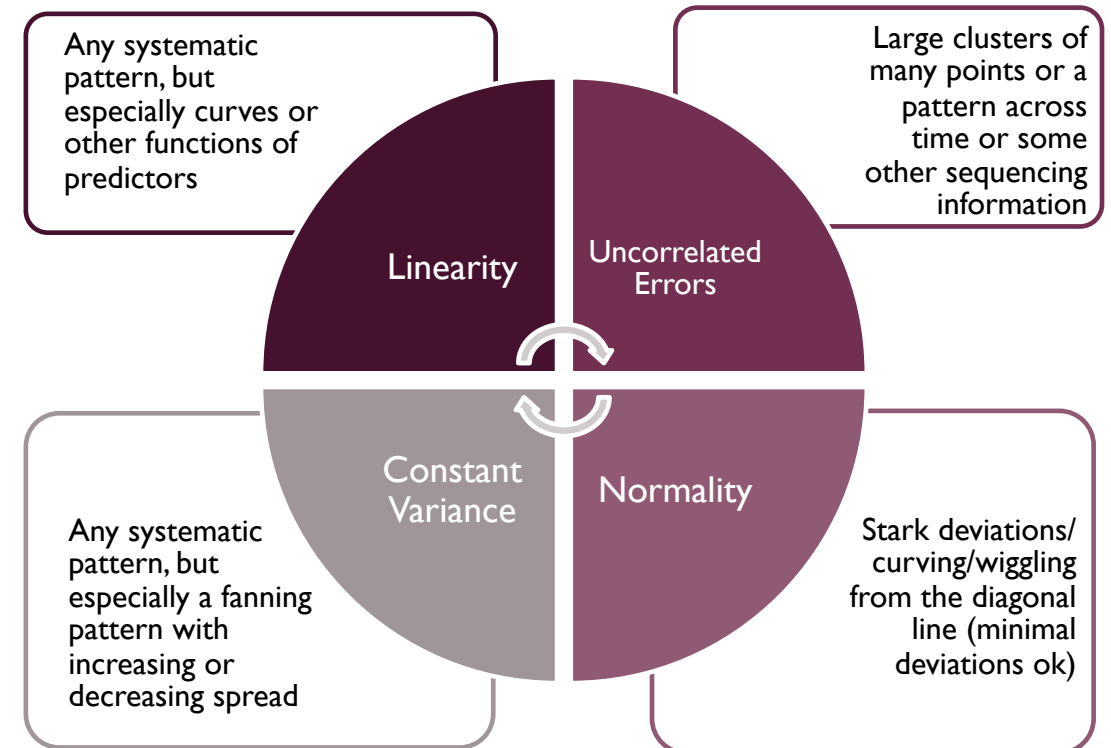
- Residuals are sample analogues of errors
  - $\varepsilon_i = y_i - E(y_i|x_i)$  whereas  $\hat{\varepsilon}_i = y_i - \hat{E}(y_i|x_i)$
  - Assuming sample was collected appropriately from population
- Residuals capture noise leftover after estimating the trend between  $Y$  and  $X$ .
  - If estimated coefficient close to the truth (unbiased) then  $\hat{\varepsilon} = Y - \hat{Y} = X(\beta - \hat{\beta}) + \varepsilon \approx \varepsilon$
- If a violation of the error assumptions occurs, we should be able to see it in the residuals
- E.g. the true relationship we want to estimate is
$$y_i = \beta_0 + \beta_1 x_{i1} + \beta_2 x_{i1}^2 + \varepsilon_i$$
- But model fit in sample data was different
$$\hat{y}_i = \hat{\beta}_0 + \hat{\beta}_1 x_{i1}$$
- The residuals would pick up the linearity issue since
$$\hat{\varepsilon}_i = y_i - \hat{y}_i = (\beta_0 - \hat{\beta}_0) + (\beta_1 - \hat{\beta}_1)x_{i1} + \beta_2 x_{i1}^2 + \varepsilon_i \approx \beta_2 x_{i1}^2 + \varepsilon_i$$



# LOOK FOR PATTERNS IN RESIDUAL PLOTS



Random bands of residuals indicates no violations



# HOW TO MAKE RESIDUAL PLOTS

## Extracting Components from Model

1. Fit the model to your data:

```
> model1 <- lm(y1 ~ x1, data = anscombe)
```

2. Extract the fitted/predicted values ( $\hat{y}_i$ ):

```
> y_hat <- fitted(model1)
> y_hat
```

	1	2	3	4	5
	8.001000	7.000818	9.501273	7.500909	8.501091
	6	7	8	9	10
	10.001364	6.000636	5.000455	9.001182	6.500727
	11				
	5.500545				

3. Extract the residuals from the model ( $\hat{e}_i$ ):

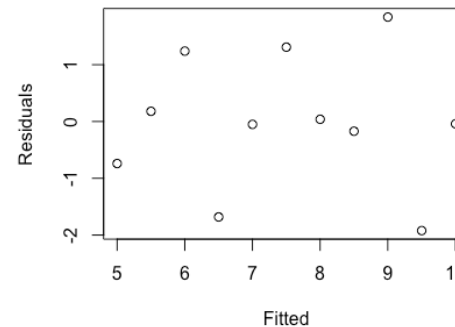
```
> e_hat <- resid(model1)
> e_hat
```

	1	2	3	4
	0.03900000	-0.05081818	-1.92127273	1.30909091
	5	6	7	8
	-0.17109091	-0.04136364	1.23936364	-0.74045455
	9	10	11	
	1.83881818	-1.68072727	0.17945455	

## Creating Residual Plots

```
> plot(x = y_hat, y = e_hat, main = "Residuals vs Fitted",
       ylab = "Residuals", xlab = "Fitted")
```

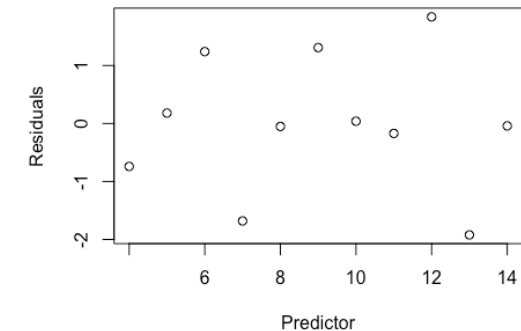
Residuals vs Fitted



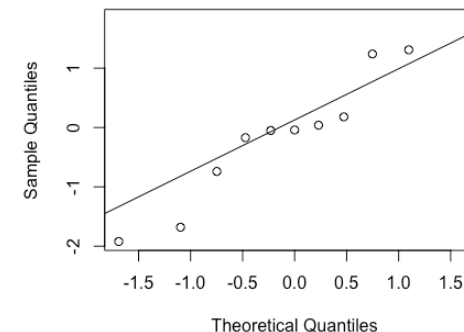
```
> qqnorm(e_hat)
> qqline(e_hat)
```

```
> plot(x = anscombe$x1, y = e_hat, main = "Residuals vs Predictor",
       ylab = "Residuals", xlab = "Predictor")
```

Residuals vs Predictor



Normal Q-Q Plot



# EXAMPLE OF NO DISTINCT PATTERNS

```
> nyc <- read.csv("nyc.csv", header=T)
> head(nyc)
```

Case	Restaurant	Price	Food	Decor	Service	East
1	1 Daniella Ristorante	43	22	18	20	0
2	2 Tello's Ristorante	32	20	19	19	0
3	3 Biricchino	34	21	13	18	0
4	4 Bottino	41	20	20	17	0
5	5 Da Umberto	54	24	19	21	0
6	6 Le Madri	52	22	22	21	0

Identifiers

Numerical predictors

Indicator variable  
1 = East location  
0 = West location

```
> model <- lm(Price ~ Food + Decor + Service + East, data=nyc)
> model
```

Call:

```
lm(formula = Price ~ Food + Decor + Service + East, data = nyc)
```

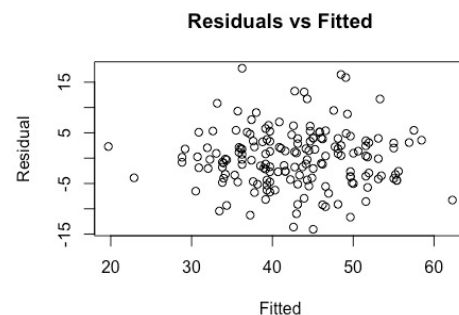
Coefficients:

```
(Intercept) -24.023800 Food 1.538120 Decor 1.910087 Service -0.002727 East 2.068050
```

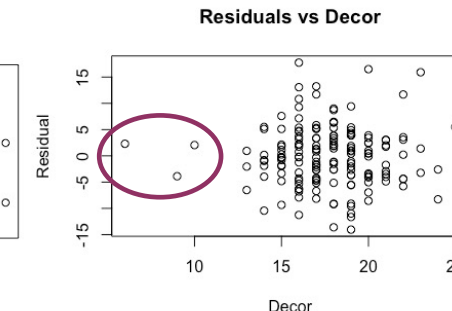
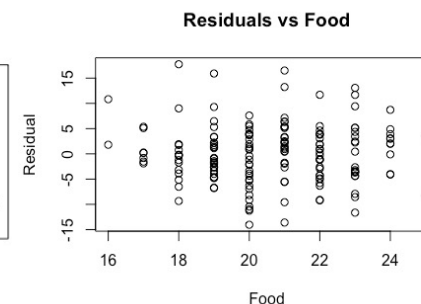
When East = 1, add this to intercept

"For a restaurant with a fixed Décor and Service rating and location, the mean Price increases by \$1.54 for a one-rating increase in Food"

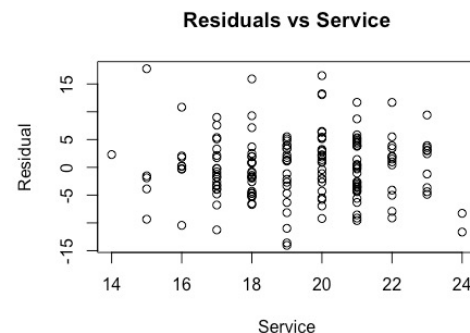
```
> e_hat <- resid(model)
> y_hat <- fitted(model)
> plot(e_hat ~ y_hat, main="Residuals vs Fitted", xlab="Fitted", ylab="Residual")
```



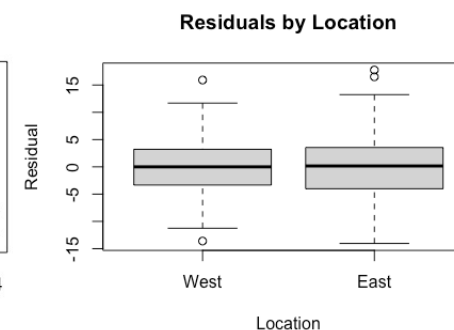
```
> plot(e_hat ~ nyc$Food, main="Residuals vs Food", xlab="Food", ylab="Residual")
> plot(e_hat ~ nyc$Decor, main="Residuals vs Decor", xlab="Decor", ylab="Residual")
```



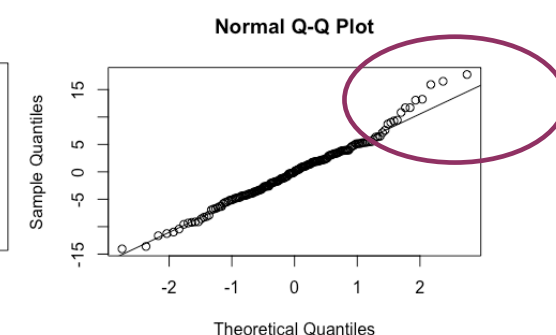
```
> plot(e_hat ~ nyc$Service, main="Residuals vs Service", xlab="Service", ylab="Residual")
```



```
> boxplot(e_hat ~ nyc$East, main="Residuals by Location", xlab="Location", ylab="Residual", names=c("West", "East"))
```



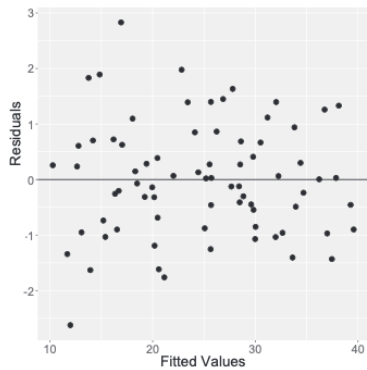
```
> qqnorm(e_hat)
> qqline(e_hat)
```



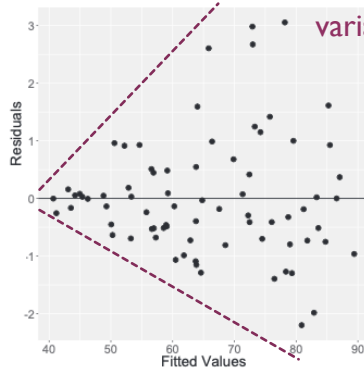
# EXAMPLES OF DISTINCT PATTERNS

From Young's Handbook of Regression Methods, pg 55

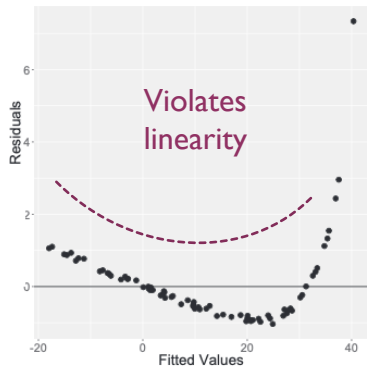
Violates constant variance



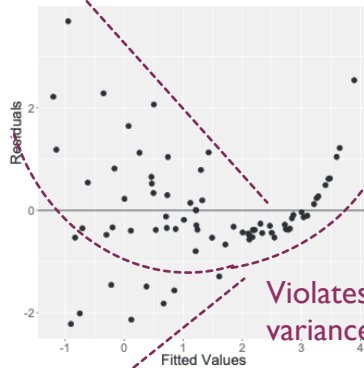
(a)



(b)

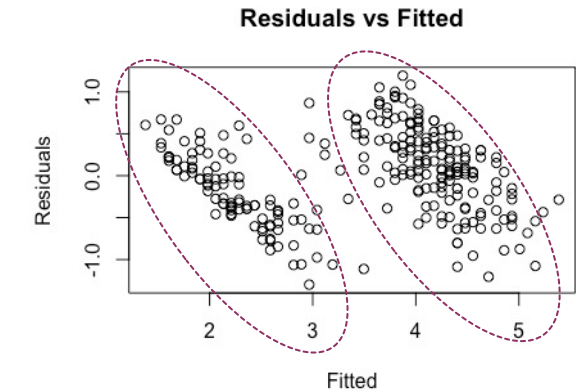
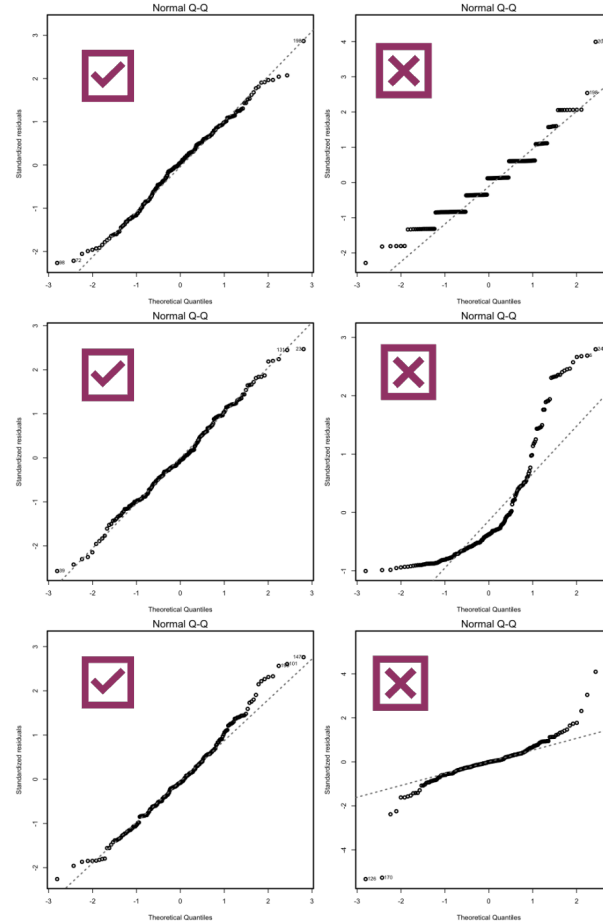


(c)

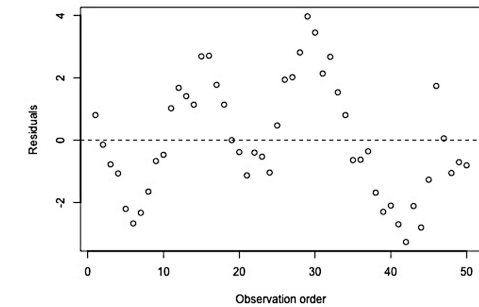


(d)

Violates constant variance and linearity



Violates uncorrelated errors



<https://online.stat.psu.edu/stat501/book/export/html/915>

# EXPLORE AND UNDERSTAND THE DATA

- Assumptions are formally checked using residual plots but knowing the data can also help
- Always conduct an **exploratory data analysis** before fitting a model
  - Skew in response variable → probably will have an issue with Normality or Linearity
  - Skews in predictor variables → may see an issue with Linearity
  - Think about underlying characteristic → may help deciding how to include predictor
- Existing literature informs your knowledge about true relationship
- Thinking about the data collection and population important too
  - *“Data was collected by voluntary response survey...”*
    - Means likely an issue with linearity assumption
  - *“Each variable was measured every day for a month...”*
    - Means likely an issue with uncorrelated errors
  - *“Neighbourhoods were randomly sampled and all subjects from selected neighbourhoods were included in the study on income inequality...”*
    - Means likely an issue with uncorrelated errors

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1. Introduction to Linear Regression Assumptions
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# ADDITIONAL CONDITIONS FOR MLR

- Recall: the interpretation of coefficients involved **holding other predictors fixed**
  - MLR estimates relationship using predictors jointly
- Certain relationships 1) between  $Y$  and  $X$ , and 2) between predictors must be identified
- In either case, presence causes residual plots to become unreliable
  - Plots can be used to say that the model is not valid
  - **Patterns in plots cannot be used to identify a specific violation and can give misleading conclusions**
- Two conditions must be checked in MLR before using residual plots

1. **Conditional mean response condition:** the mean responses are a single function of a linear combination involving  $\beta$

$$E(Y_i | \mathbf{X} = \mathbf{x}_i) = g(\beta_0 + \beta_1 x_{i1} + \cdots + \beta_p x_{ip})$$

- $E(Y|X) = \log(\beta_0 + \beta_1 x_i)$  satisfies this condition
- $E(Y|X) = \beta_1 x_{i1} / \beta_2 x_{i2} = g_1(x_1) / g_2(x_2)$  violates it

2. **Conditional mean predictor condition:** the mean of each predictor is related to each other predictor in no more complicated way than linearly

$$E(X_i | X_j) = \alpha_0 + \alpha_1 X_j$$

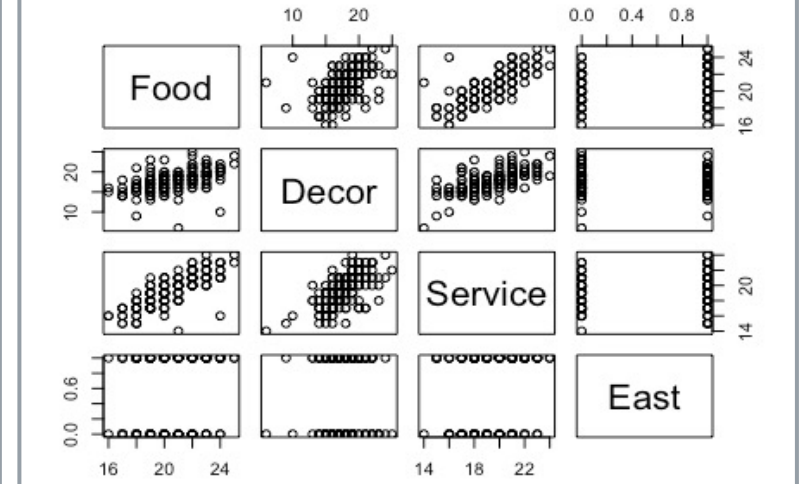
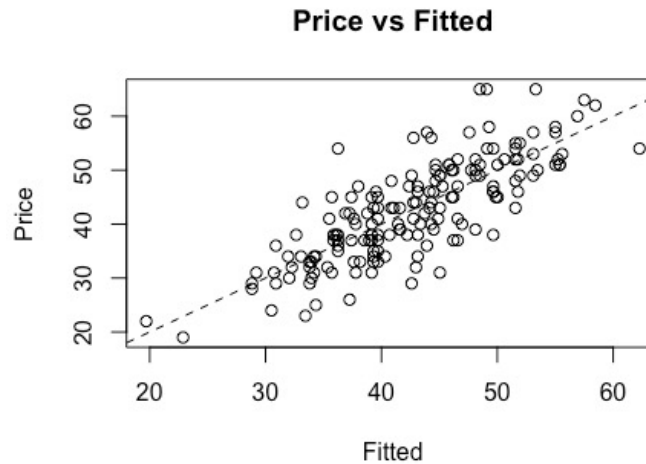
- Linear or no relationship satisfy condition; anything else violates

## Response versus Fitted Values

```
> plot(x = y_hat, y = nyc$Price, main="Price vs Fitted",  
      xlab="Fitted", ylab="Price")  
> abline(a = 0, b = 1, lty=2)
```

## Pairwise Scatterplots

```
> pairs(nyc[,4:7])
```



# HOW ARE CONDITIONS CHECKED

## 1. Conditional mean response

Scatterplot of Response  
versus Fitted values

*Look for random diagonal  
scatter or an easily  
identifiable non-linear trend*

## 2. Conditional mean predictors

All pairwise scatterplots  
of predictors

*Look for lack of curves or  
other non-linear patterns*



# CONDITION 1 DOES NOT HOLD

- Population (i.e., true relationships):

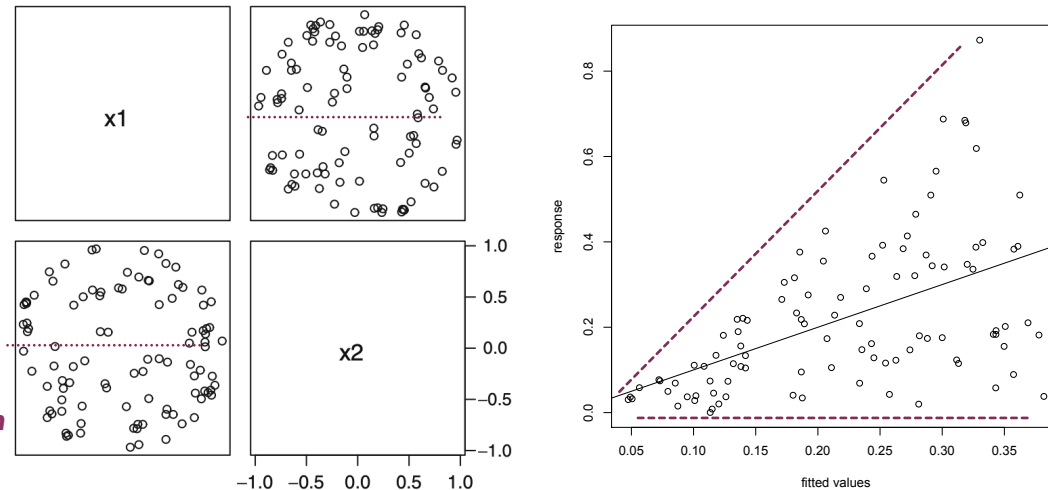
- Condition 2 holds:

$$E(X_i|X_j) = \alpha_0 + \alpha_1 X_j \quad \xrightarrow{\text{No non-linear pattern}}$$

- Condition 1 fails:

$$E(Y|X) = \frac{|x_1|}{2 + (1.5 + x_2)^2} = \frac{g(x_1)}{g(x_2)}$$

Linear model  
is not  
appropriate  
for this situation



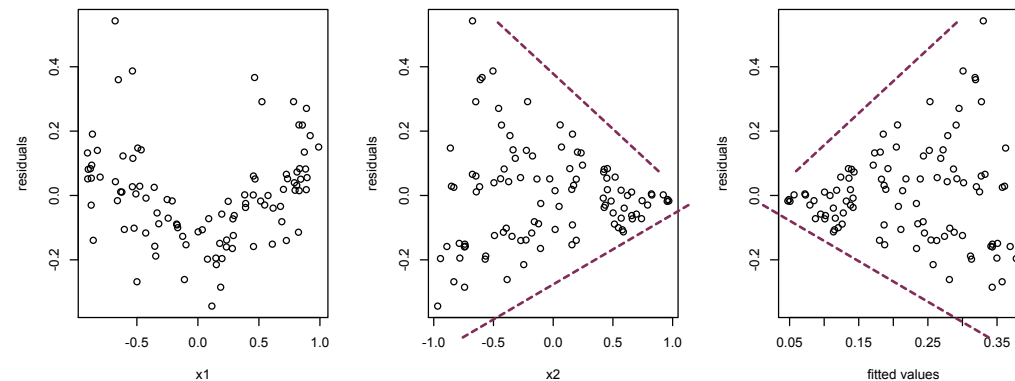
Not a single identifiable  
pattern/function and  
not random scatter

- Data are simulated from above so that  
constant variance is not violated

- Model fit in the sample:

$$Y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + e$$

- Can look at plots for conditions and residual  
plots for assumptions



Pattern implies non-constant  
variance, yet data generated  
to ensure assumptions hold

# CONDITION 2 DOES NOT HOLD

- Population (i.e., true relationship):

- Condition 1 hold:

$$Y = x_1 + 3x_2^2 + \varepsilon$$

- Condition 2 fails:

$$E(x_2|x_1) = \sin(x_1)$$

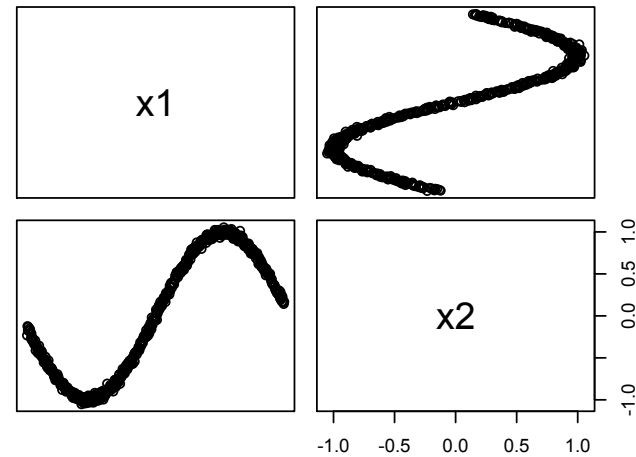
*Periodic pattern visible,  
a non-linear relationship*

- Data are simulated from above

- Model fit in the sample:

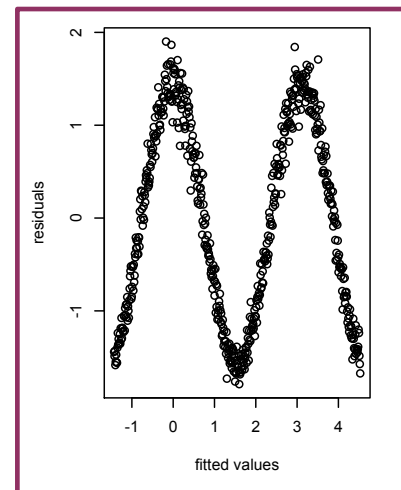
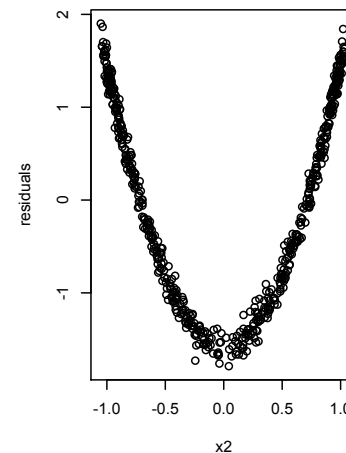
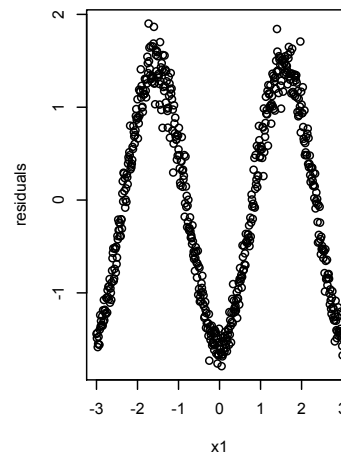
$$Y = \beta_0 + \beta_1 x_1 + \beta_2 x_2 + e$$

- We should see that **linearity is violated**  
because we are missing a squared term



*Quadratic curve in residual vs  $x_2$  plot  
highlights the missing square term from  
true relationship*

*Without checking conditions,  
would imply also missing a  
sinusoidal/periodic term as well  
→ Due to failure of condition 2*



# MODULE TAKE-AWAYS

1. What are the assumptions of linear regression and what do they mean?
2. What tools are used to assess whether each assumption holds?
3. What do we look for to know whether each assumption holds?
4. Why is it important to check the additional conditions in multiple linear models?
5. What other aspects of data, data collection or population characteristics tell us assumption may not hold?