

Linear Regression II

Presented by David John Baker
November 2019



Why Linear Regression II ?

- Today we take second pass at Linear Regression
- Goals: Have better understanding of core components of model
- Proceed through entire checklist of running linear regression model
- Clearly define more specific terms and explain how they relate to NHST



Linear Regression

- Can you predict X given Y if we assume a linear relationship between the variables in question?
- Dependant variable is continuous
- Independent variables can be either continuous or categorical



Format

- Assumptions
- Fitting
- Multiple Models
- Interpretation



Common statistical tests are linear models

Last updated: 29 June, 2019. Also check out the [R version!](#)

See worked examples and more details at the accompanying notebook: <https://github.com/eigenfoo/tests-as-linear>

	Common name	Function in scipy.stats	Equivalent linear model in smf.ols	Exact?	The linear model in words	Icon
Simple Regression: ($y \sim 1 + x$)	y is independent of x P: One-sample t-test N: Wilcoxon signed-rank	scipy.stats.ttest_1samp(y) scipy.stats.wilcoxon(y)	smf.ols("y ~ 1", data) smf.ols("y ~ 1", signed_rank(data))	✓ for $N \geq 14$	One number (intercept, i.e., the mean) predicts y. - (Same, but it predicts the <i>signed rank</i> of y.)	
	P: Paired-sample t-test N: Wilcoxon matched pairs	scipy.stats.ttest_rel(y1, y2) scipy.stats.wilcoxon(y1, y2)	smf.ols("y2_sub_y1 ~ 1", data) smf.ols("y2_sub_y1 ~ 1", signed_rank(data))	✓ for $N \geq 14$	One intercept predicts the pairwise y2-y1 differences. - (Same, but it predicts the <i>signed rank</i> of y2-y1.)	
	y ~ continuous x P: Pearson correlation N: Spearman correlation	scipy.stats.pearsonr(x, y) scipy.stats.spearmanr(x, y)	smf.ols("y ~ 1 + x", data) smf.ols("y ~ 1 + x", rank(data))	✓ for $N \geq 10$	One intercept plus x multiplied by a number (slope) predicts y. - (Same, but with <i>ranked x and y</i>)	
	y ~ discrete x P: Two-sample t-test P: Welch's t-test N: Mann-Whitney U	scipy.stats.ttest_ind(y1, y2) N/A in Python, but see R version scipy.stats.mannwhitneyu(y1, y1)	smf.ols("y ~ 1 + group", data) ^A N/A in Python, but see R version smf.ols("y ~ 1 + group", signed_rank(data)) ^A	✓ ✓ for $N \geq 11$	An intercept for group 1 (plus a difference if group 2) predicts y. - (Same, but with one variance <i>per group</i> instead of one common.) - (Same, but it predicts the <i>signed rank</i> of y.)	
Multiple regression: ($y \sim 1 + x_1 + x_2 + \dots$)	P: One-way ANOVA N: Kruskal-Wallis	scipy.stats.f_oneway(a, b, c) scipy.stats.kruskal(a, b, c)	smf.ols("y ~ 1 + G2 + G3 + ... + GN") ^A smf.ols(rank(y) ~ 1 + G2 + G3 + ... + GN) ^A	✓ for $N \geq 11$	An intercept for group 1 (plus a difference if group $\neq 1$) predicts y. - (Same, but it predicts the <i>rank</i> of y.)	
	P: One-way ANCOVA	N/A in Python, but see R version	smf.ols("y ~ 1 + G2 + G3 + ... + GN + x", data) ^A	✓	- (Same, but plus a slope on x.) <i>Note: this is discrete AND continuous. ANCOVAs are ANOVAs with a continuous x.</i>	
	P: Two-way ANOVA	N/A in Python, but see R version	smf.ols("y ~ 1 + G2 + G3 + ... + GN + S2 + S3 + ... + SK + G2*S2+G3*S3+...+GN*SK", data)	✓	Interaction term: changing sex changes the y ~ group parameters. <i>Note: G2 to GN is an indicator (0 or 1) for each non-intercept levels of the group variable. Similarly for S2 to SK for sex. The first line (with G) is main effect of group, the second (with S) for sex and the third is the group * sex interaction. For two levels (e.g. male/female), line 2 would just be "S2" and line 3 would be "S2 multiplied with each G.</i>	[Coming]
	Counts ~ discrete x N: Chi-square test	scipy.stats.chisquare(data)	Equivalent log-linear model sm.GLM(y ~ 1 + G2 + G3 + ... + GN + S2 + S3 + ... + SK + G2*S2+G3*S3+...+GN*SK, family=...) ^A	✓	Interaction term: (Same as Two-way ANOVA.) <i>Note: Run glm using the following arguments: glm(model, family=poisson()) As linear-model, the Chi-square test is log(y) = log(β0) + log(β1) + log(αβ) where α and βi are proportions. See more info in the accompanying notebook</i>	Same as Two-way ANOVA
	N: Goodness of fit	scipy.stats.chi2_contingency(data)	sm.GLM(y ~ 1 + G2 + G3 + ... + GN, family=...) ^A	✓	(Same as One-way ANOVA and see Chi-Square note.)	1W-ANOVA

List of common parametric (P) non-parametric (N) tests and equivalent linear models. The notation $y \sim 1 + x$ is R shorthand for $y = 1 + b + a \cdot x$ which most of us learned in school. Models in similar colors are highly similar, but really, notice how similar they *all* are across colors! For non-parametric models, the linear models are reasonable approximations for non-small sample sizes (see "Exact" column and click links to see simulations). Other less accurate approximations exist, e.g., Wilcoxon for the sign test and Goodness-of-fit for the binomial test. The signed rank function is `signed_rank(df) = np.sign(df) * df.rank()`. The variables G_i and S_i are "dummy coded" indicator variables (either 0 or 1) exploiting the fact that when $\Delta x = 1$ between categories the difference equals the slope. Subscripts (e.g., G₂ or y₁) indicate different columns in data. lm requires long-format data for all non-continuous models. All of this is exposed in greater detail and worked examples at <https://eigenfoo.xyz/tests-as-linear>.

^A See the note to the two-way ANOVA for explanation of the notation.



Jonas Kristoffer Lindeløv, George Ho
<https://lindeloev.net> <https://eigenfoo.xyz>

<https://eigenfoo.xyz/tests-as-linear/>

Basic Linear Regression Assumptions

1. Independence of Data Points
2. Normality of Residuals
3. Homoscedasticity

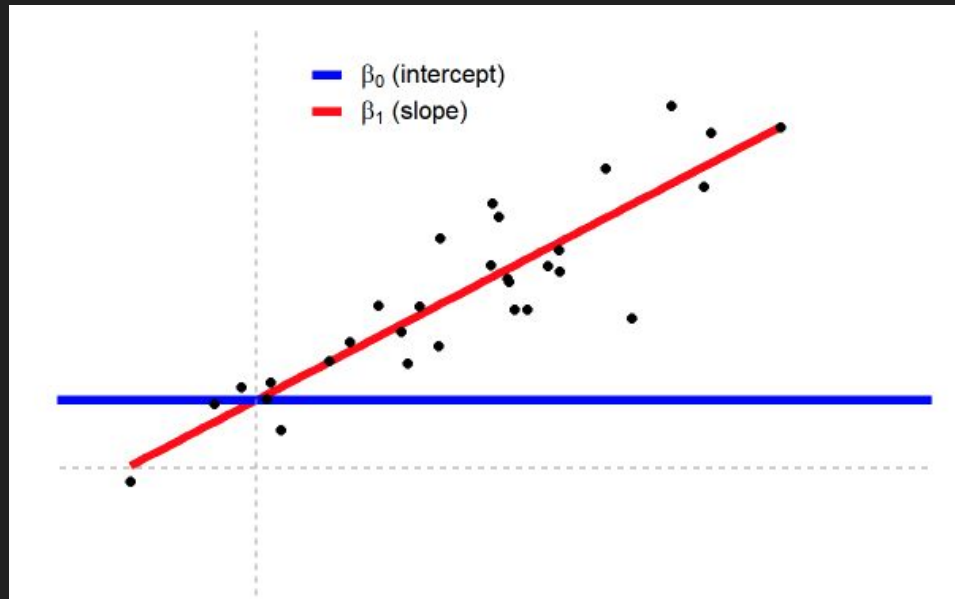
Linear Regression Assumptions

(0. Linearity)

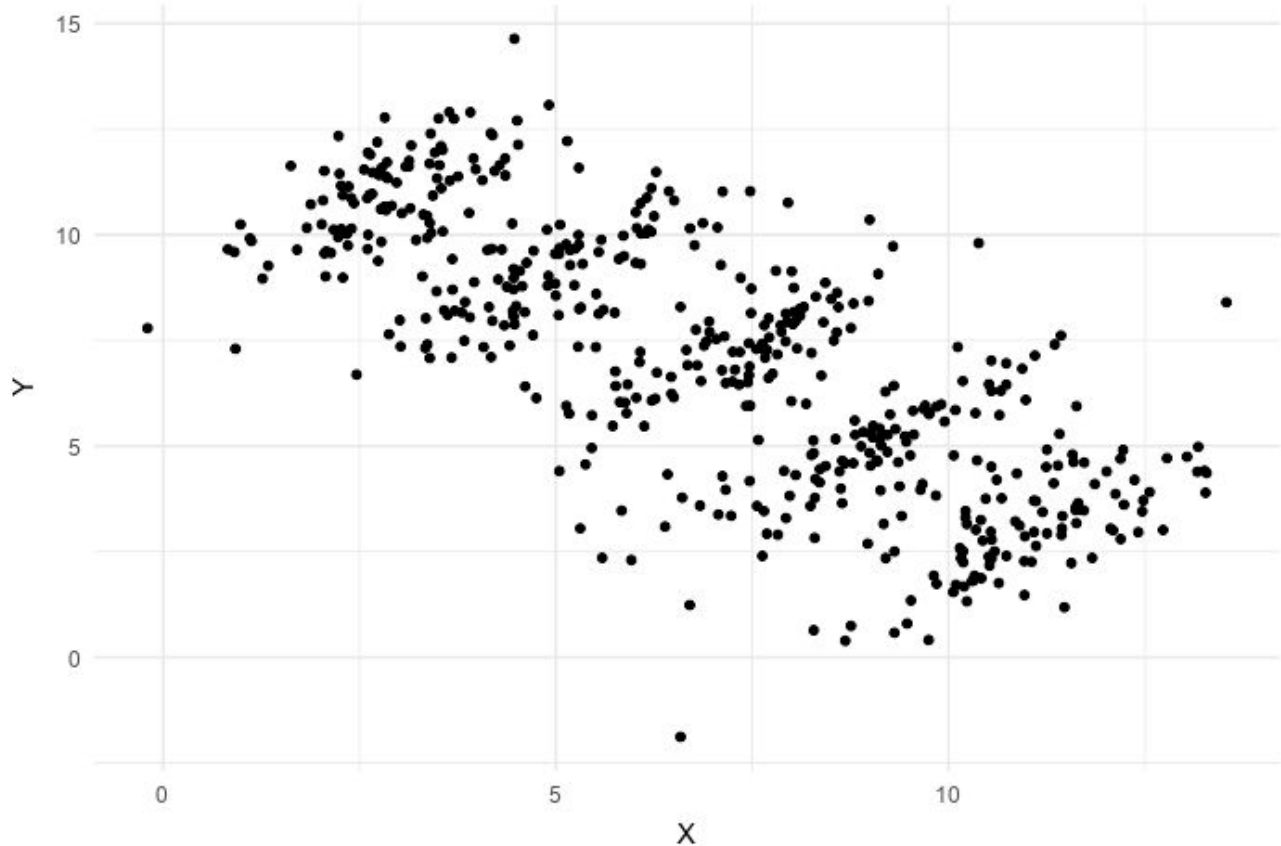
1. Independence of Data Points
2. Normality of Residuals
3. Homoscedasticity
4. Multicollinearity
5. You DO NOT need Normally distributed variables, you DO NEED to Run Diagnostic Plots

Independence of Data Points

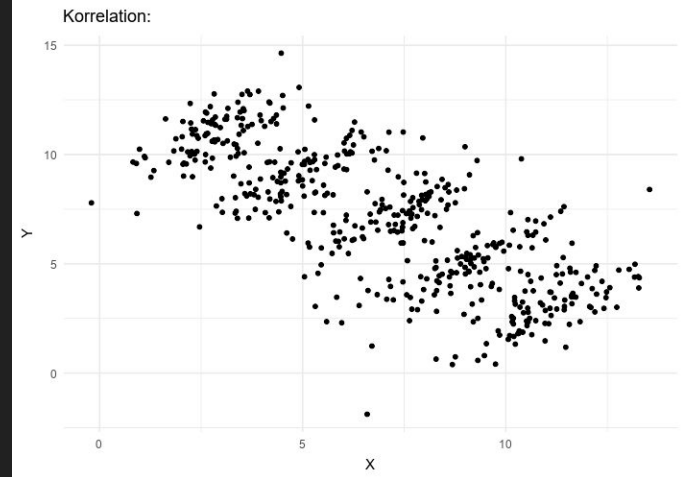
- Each data point needs to be independently sampled from the parent population
- AKA Data points shouldn't be able to influence/talk to one another!!!



Korrelation:



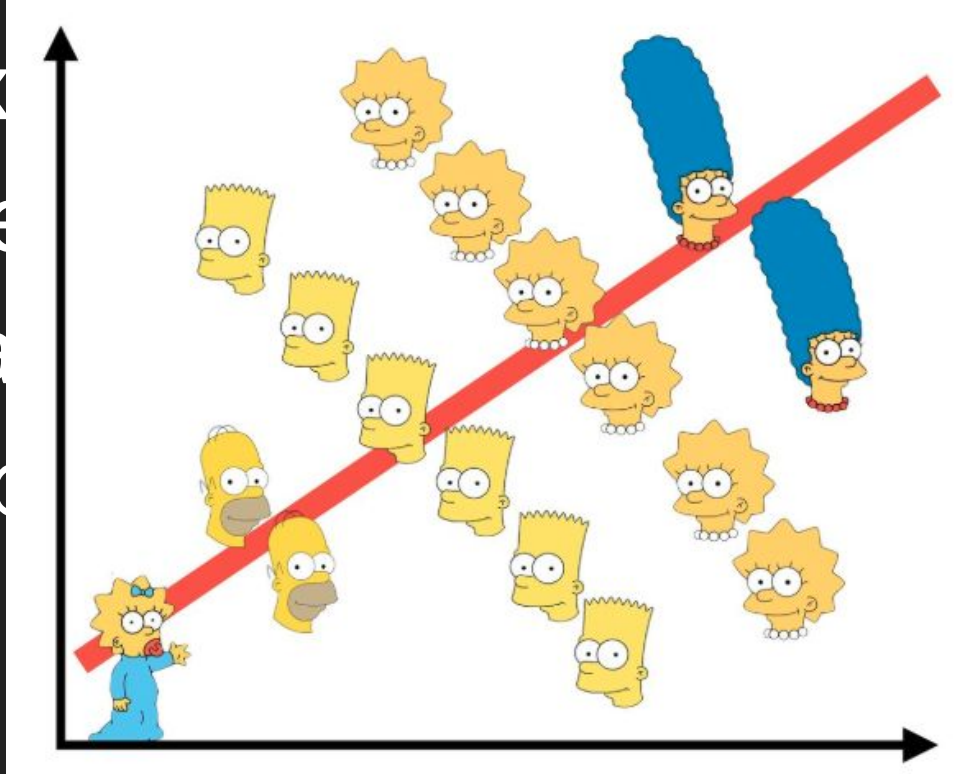
Think of situation in your area of expertise where you might come across a violation of the the assumption of independence.



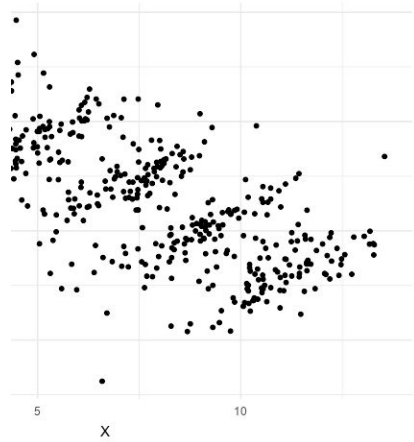
Think of situation in your area

of ex
come
the a
indep

at
ne



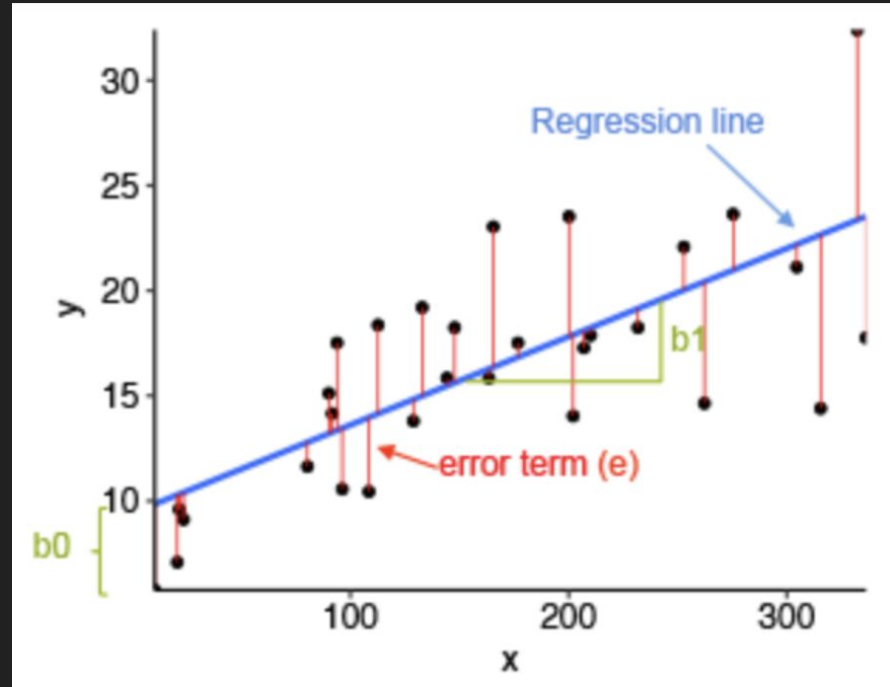
0



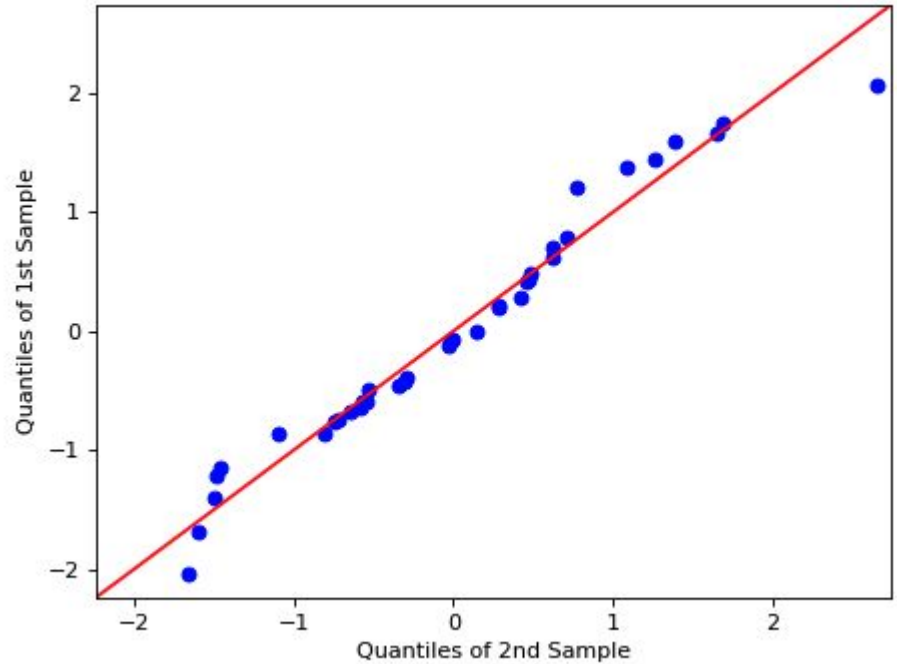
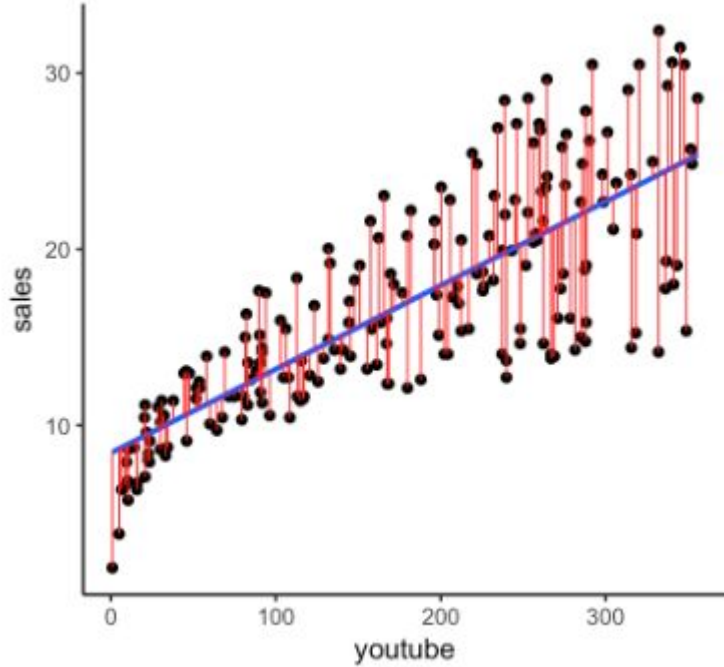
Normally Distributed Residuals

Residuals, or the distances between your regression line and observed data, when plotted should create a normal distribution

Inspect this with a QQ plot



Normally Distributed Residuals



Homoscedasticity

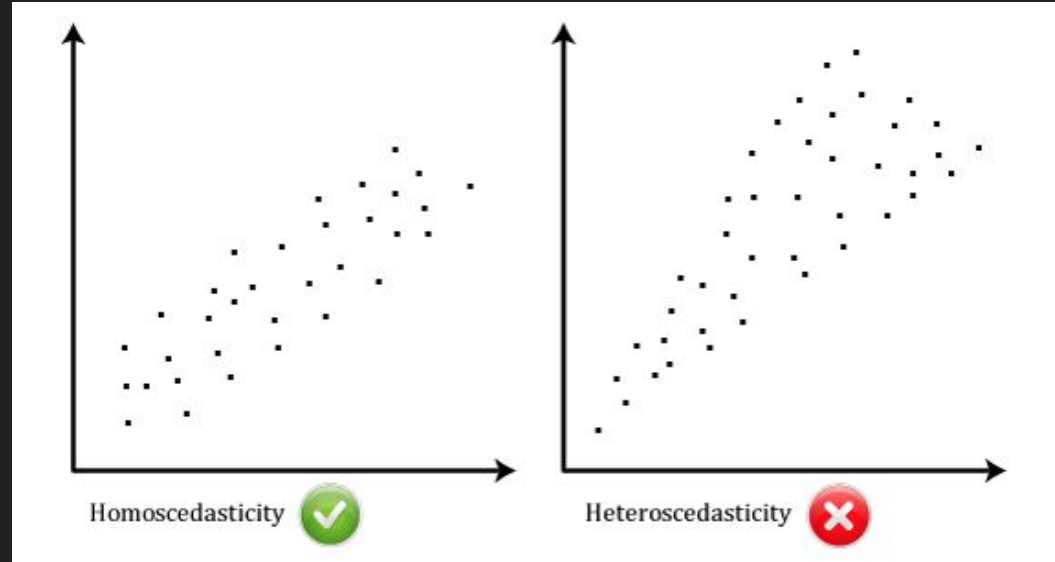
Homoscedasticity (ho-mo-ske-das-ti-ci-tee)

Heteroscedasticity (het-er-o-ske-das-tis-i-tee)

Homoscedasticity (ho-mo-ske-das-ti-ci-tee)

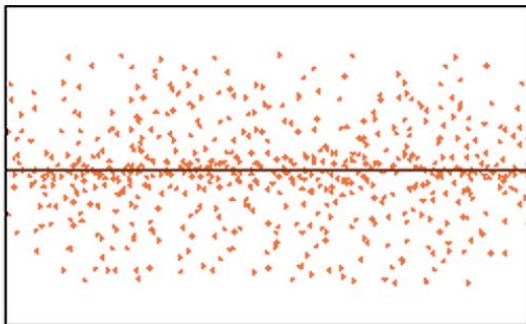
Homogeneity of Variance

- Variance around errors is uniform



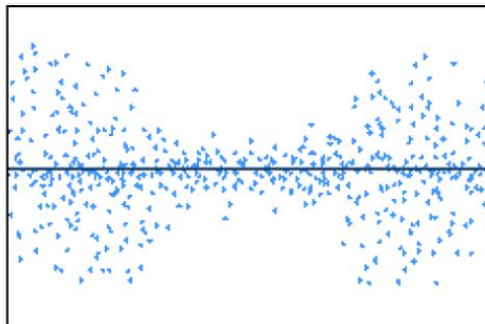
Residual Plots

Homoscedasticity



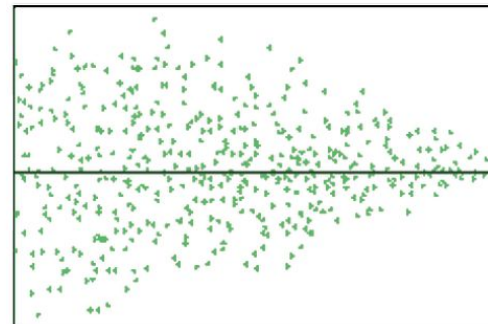
Random Cloud (No Discernible Pattern)

Heteroscedasticity



Bow Tie Shape (Pattern)

Heteroscedasticity



Fan Shape (Pattern)

Linear Regression Checklist

- **Plot all variables**
- **Check for Multicollinearity**
- **Check for Outliers (Univariate and Multivariate)**



Regression Output

Out[12]:

OLS Regression Results

Dep. Variable:	Sepal.Length	R-squared:	0.867
Model:	OLS	Adj. R-squared:	0.862
Method:	Least Squares	F-statistic:	155.8
Date:	Mon, 18 Nov 2019	Prob (F-statistic):	3.86e-60
Time:	11:57:26	Log-Likelihood:	-32.558
No. Observations:	150	AIC:	79.12
Df Residuals:	143	BIC:	100.2
Df Model:	6		
Covariance Type:	nonrobust		



Regression Output

R-Squared:

Coefficient of
Determination

0 -- 1

Variance explained

Literally, r , squared

//

Out[12]:

OLS Regression Results

Dep. Variable:	Sepal.Length	R-squared:	0.867
Model:	OLS	Adj. R-squared:	0.862
Method:	Least Squares	F-statistic:	155.8
Date:	Mon, 18 Nov 2019	Prob (F-statistic):	3.86e-60
Time:	11:57:26	Log-Likelihood:	-32.558
No. Observations:	150	AIC:	79.12
Df Residuals:	143	BIC:	100.2
Df Model:	6		
Covariance Type:	nonrobust		

Regression Output

F statistic:

Omnibus test,

Just like ANOVA

Variance / Error

//

Out[12]:

OLS Regression Results

Dep. Variable:	Sepal.Length	R-squared:	0.867
Model:	OLS	Adjusted R-squared:	0.860
Method:	Least Squares	F-statistic:	155.8
Date:	Mon, 18 Nov 2019	Prob (F-statistic):	3.86e-60
Time:	11:57:26	Log-Likelihood:	-32.558
No. Observations:	150	AIC:	79.12
Df Residuals:	143	BIC:	100.2
Df Model:	6		
Covariance Type:	nonrobust		

Regression Output

Model Fit Comparison Metrics

In [statistics](#), the **likelihood function** (often simply called the **likelihood**) expresses how likely particular values of [statistical model parameters](#) are for a given [sample of data](#).^[a]

Out[121]:

Results

Sepal.Length	R-squared:	0.867
OLS	Adj. R-squared:	0.862
Least Squares	F-statistic:	155.8
Date:	Mon, 18 Nov 2019	Prob (F-statistic): 3.86e-60
Time:	11:57:26	Log-Likelihood: -32.558
No. Observations:	150	AIC: 79.12
Df Residuals:	143	BIC: 100.2
Df Model:	6	
Covariance Type:	nonrobust	

Regression Output

Model Fit Comparison Metrics

In [statistics](#), the **likelihood function** (often simply called the **likelihood**) expresses how likely particular values of [statistical model parameters](#) are for a given [sample of data](#).^[a]

The **Akaike information criterion (AIC)** is an [estimator](#) of [out-of-sample](#) prediction error and thereby relative quality of [statistical models](#) for a given set of data.^{[1][2]} Given a

In [statistics](#), the **Bayesian information criterion (BIC)** or **Schwarz information criterion** (also **SIC**, **SBC**, **SBIC**) is a criterion for [model selection](#) among a finite set of models; the model with the lowest BIC is preferred. It is based, in part, on the [likelihood function](#) and it is closely related to the [Akaike information criterion \(AIC\)](#).

Out[121]:

Results

Sepal.Length	R-squared:	0.867
OLS	Adj. R-squared:	0.862
Least Squares	F-statistic:	155.8
Mon, 18 Nov 2019	Prob (F-statistic):	3.86e-60
11:57:26	Log-Likelihood:	-32.558
150	AIC:	79.12
143	BIC:	100.2
6		
nonrobust		

Regression Output -- Unstandardized Beta

For every change in the indicator variable, you have y changes in the dependant variable, all others equal

	coef	std err	t	P> t	[0.025	0.975]
Unnamed: 0	-3.606e-05	0.002	-0.020	0.984	-0.004	0.003
Sepal.Width	0.4960	0.086	5.737	0.000	0.325	0.667
Petal.Length	0.8290	0.070	11.888	0.000	0.691	0.967
Petal.Width	-0.3150	0.152	-2.075	0.040	-0.615	-0.015
Species_setosa	2.1722	0.285	7.628	0.000	1.609	2.735
Species_versicolour	1.4511	0.327	4.443	0.000	0.805	2.097
Species_virginica	1.1531	0.441	2.614	0.010	0.281	2.025

Regression Output -- Standardized Beta

Everything on z score

Need to standardize
Variables ahead
Of time!!!



	coef	std err	t	P> t	[0.025	0.975]
Unnamed: 0	-1.606e-05	0.002	-0.020	0.984	-0.004	0.003
Sepal.Width	0.4960	0.086	5.737	0.000	0.325	0.667
Petal.Length	0.8290	0.070	11.888	0.000	0.691	0.967
Petal.Width	-0.3151	0.152	-2.075	0.040	-0.615	-0.015
Species_setosa	2.1722	0.285	7.628	0.000	1.609	2.735
Species_versicolour	1.4511	0.327	4.443	0.000	0.805	2.097
Species_virginica	1.1531	0.441	2.614	0.010	0.281	2.025

Regression Output

	coef	std err	t	P> t	[0.025	0.975]
Unnamed: 0	-3.606e-05	0.002	-0.020	0.984	-0.004	0.003
Sepal.Width	0.4960	0.086	5.737	0.000	0.325	0.667
Petal.Length	0.8290	0.070	11.888	0.000	0.691	0.967
Petal.Width	-0.3150	0.152	-2.075	0.040	-0.615	-0.015
Species_setosa	2.1722	0.285	7.628	0.000	1.609	2.735
Species_versicolor	1.4511	0.327	4.443	0.000	0.805	2.097
Species_virginica	1.1531	0.441	2.614	0.010	0.281	2.025

Regression Output

	coef	std err	t	P> t	[0.025	0.975]
Unnamed: 0	-3.606e-05	0.002	-0.020	0.984	-0.004	0.003
Sepal.Width	0.4960	0.086	5.737	0.000	0.325	0.667
Petal.Length	0.8290	0.070	11.888	0.000	0.691	0.967
Petal.Width	-0.3150	0.152	-2.075	0.040	-0.615	-0.015
Species_setosa	2.1722	0.285	7.628	0.000	1.609	2.735
Species_versicolor	1.4511	0.327	4.443	0.000	0.805	2.097
Species_virginica	1.1531	0.441	2.614	0.010	0.281	2.025

Regression Output

	coef	std err	t	P> t	[0.025	0.975]
Unnamed: 0	-3.606e-05	0.002	-0.020	0.984	-0.004	0.003
Sepal.Width	0.4960	0.086	5.737	0.000	0.325	0.667
Petal.Length	0.8290	0.070	11.888	0.000	0.691	0.967
Petal.Width	-0.3150	0.152	-2.075	0.040	-0.615	-0.015
Species_setosa	2.1722	0.285	7.628	0.000	1.609	2.735
Species_versicolor	1.4511	0.327	4.443	0.000	0.805	2.097
Species_virginica	1.1531	0.441	2.614	0.010	0.281	2.025



Regression Output

	coef	std err	t	P> t	[0.025	0.975]
Unnamed: 0	-3.606e-05	0.002	-0.020	0.984	-0.004	0.003
Sepal.Width	0.4960	0.086	5.737	0.000	0.325	0.667
Petal.Length	0.8290	0.070	11.888	0.000	0.691	0.967
Petal.Width	-0.3150	0.152	-2.075	0.040	-0.615	-0.015
Species_setosa	2.1722	0.285	7.628	0.000	1.609	2.735
Species_versicolor	1.4511	0.327	4.443	0.000	0.805	2.097
Species_virginica	1.1531	0.441	2.614	0.010	0.281	2.025



Regression Output

Omnibus:	0.414	Durbin-Watson:	1.966
Prob(Omnibus):	0.813	Jarque-Bera (JB):	0.567
Skew:	-0.060	Prob(JB):	0.753
Kurtosis:	2.723	Cond. No.	1.98e+03

In [statistics](#), the **Durbin–Watson statistic** is a [test statistic](#) used to detect the presence of [autocorrelation](#) at lag 1 in the [residuals](#) (prediction errors) from a [regression analysis](#). It is named after [James Durbin](#) and [Geoffrey Watson](#). The [small](#)

Regression Output

Omnibus:	0.414	Durbin-Watson:	1.966
Prob(Omnibus):	0.813	Jarque-Bera (JB):	0.567
Skew:	-0.060	Prob(JB):	0.753
Kurtosis:	2.723	Cond. No.	1.98e+03

In [statistics](#), the **Jarque–Bera test** is a [goodness-of-fit](#) test of whether sample data have the [skewness](#) and [kurtosis](#) matching a [normal distribution](#). The test is named after [Carlos Jarque](#) and [Anil K. Bera](#). The test statistic is always nonnegative. If it is far from zero, it signals the data do not have a normal distribution.



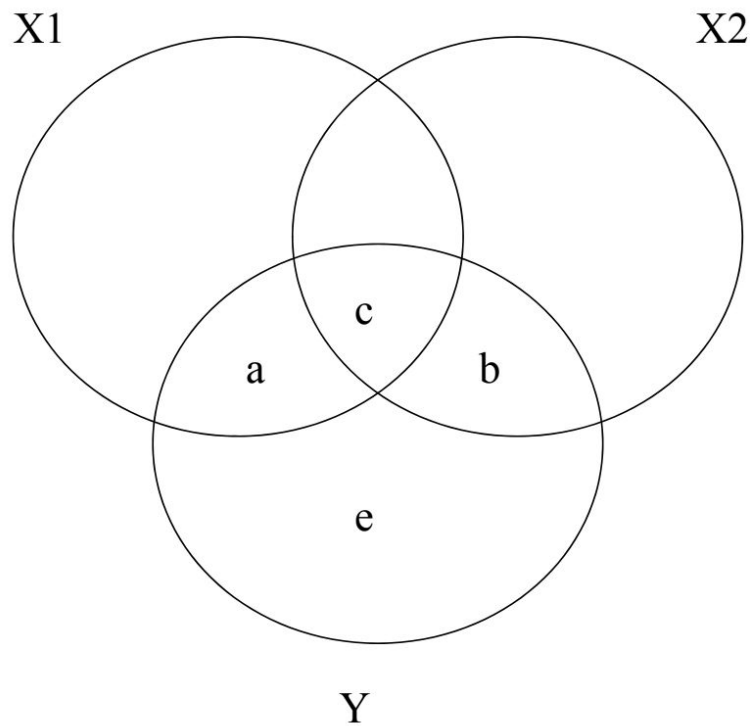
Regression Output

Omnibus:	0.414	Durbin-Watson:	1.966
Prob(Omnibus):	0.813	Jarque-Bera (JB):	0.567
Skew:	-0.060	Prob(JB):	0.753
Kurtosis:	2.723	Cond. No.	1.98e+03

In the field of [numerical analysis](#), the **condition number** of a function measures how much the output value of the function can change for a small change in the input argument. This is used to measure how [sensitive](#) a function is to changes or errors in the input, and how much error in the output

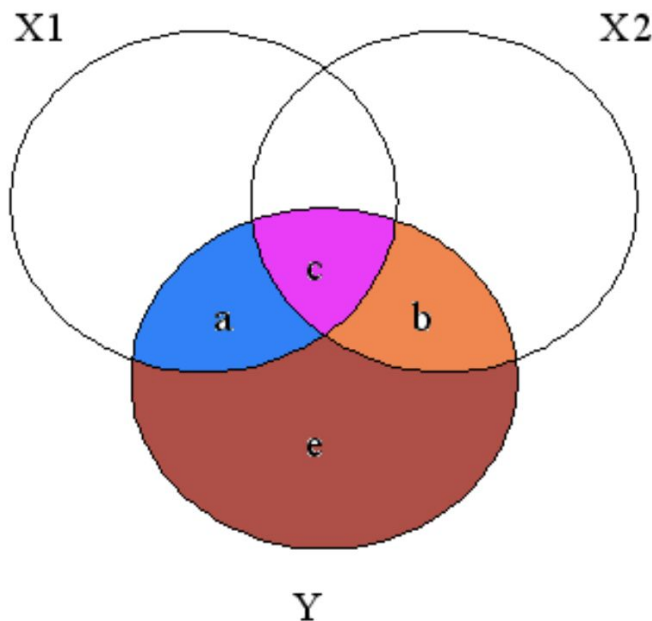


Regression Output



Regression Output

Concept of multiple R^2 for the full regression model



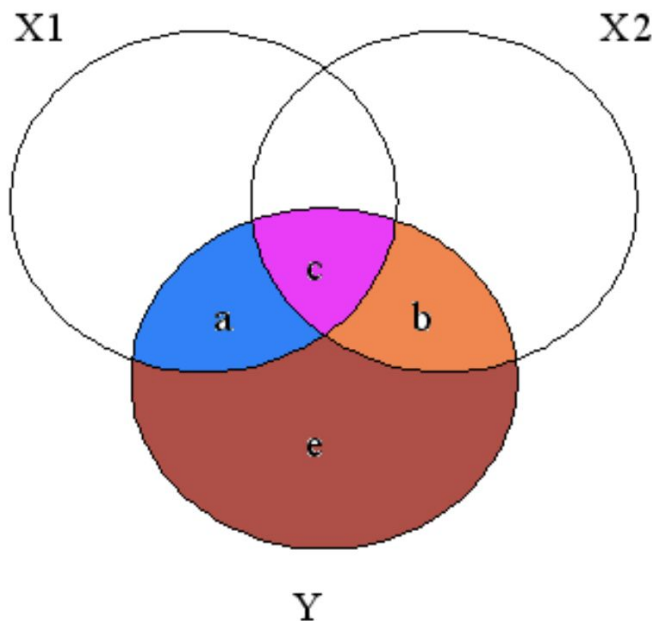
$$r_{Y \cdot 12}^2 = \frac{a + b + c}{a + b + c + e} = a + b + c$$

What of Y can be accounted for by X1, X2, and any redundancy among X1 and X2?

Area c is not unique to either X1 or X2 alone, but contributes to the full R^2

Regression Output

Concept of multiple R^2 for the full regression model



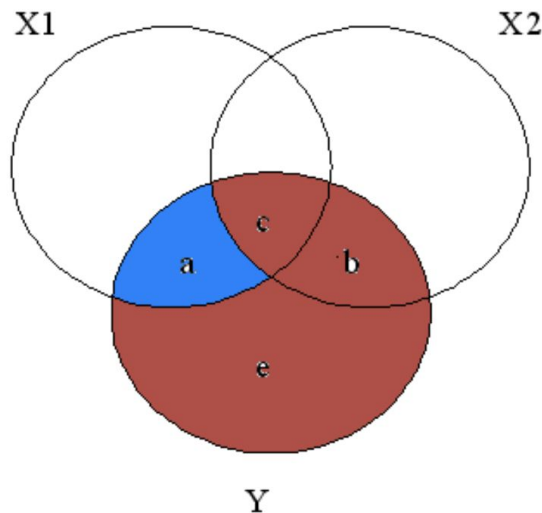
$$r_{Y \cdot 12}^2 = \frac{a + b + c}{a + b + c + e} = a + b + c$$

What of Y can be accounted for by X1, X2, and any redundancy among X1 and X2?

Area c is not unique to either X1 or X2 alone, but contributes to the full R^2

Regression Output

Influence of predictor X1 when X2 is already in the model: variance shared between X1 & Y **beyond** that accounted for by X2



“part” or
“semipartial”
correlation of
X1 with Y

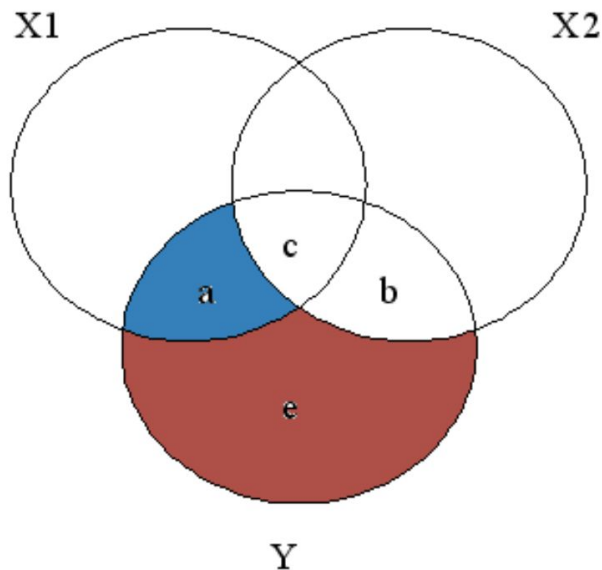
$$r_{1(Y \cdot 2)}^2 = \frac{a}{a + b + c + e} = a$$

Part correlation: the correlation
of Y with that part of X1 that
is independent of X2

The squared semipartial correlation is ONE way to determine how much influence (i.e., importance) each predictor has to the full equation

Regression Output

Partial correlation: X1 with Y when X2's overlap with Y AND X1 is controlled



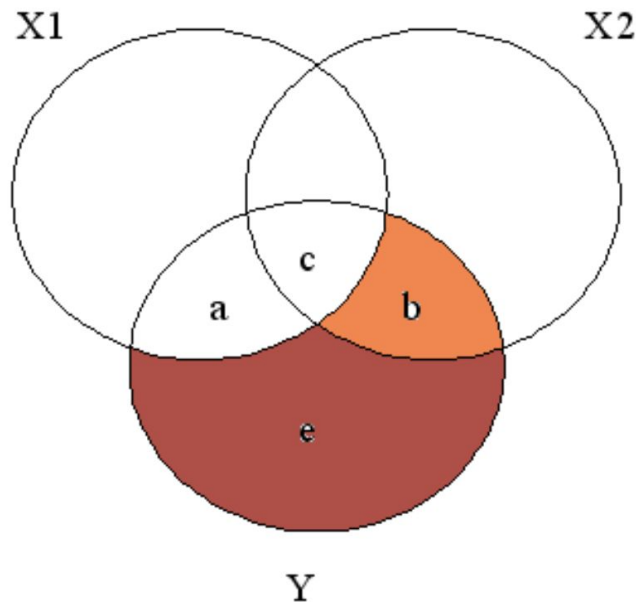
$$r_{1Y \cdot 2}^2 = \frac{a}{a + e}$$

Influence of b and c
are removed from
consideration altogether

Partial correlation: the
correlation between the
residual of one relationship
(e_{2Y}) and the residual of
another (e_{12}) **in the
variance of Y.**

Regression Output

Partial correlation: X2 with Y when X1's overlap with Y AND X2 is controlled



$$r_{2Y \cdot 1}^2 = \frac{b}{b + e}$$

Influence of a and c
are removed from
consideration altogether

Partial correlation: the
correlation between the residual
of one relationship (e_{1Y})
and the residual of another (e_{12})
in the variance of Y.