ANOVA, Correlation, Linear Regression, Quantile Regression



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Analysis of Variance (ANOVA) (like what we just did but for more than two different samples)

See Chapter 7 in the Helsel and Hirsch textbook
Can also check out Brandon Foltz on Youtube:
https://www.youtube.com/watch?v=0Vj2V2qRU10
Note: We will go over One Factor ANOVA. There are many variations of ANOVA.

Note 2: This is cookbook like. Don't panic if you can't remember how many teaspoons of salt go in a given recipe. Just remember where you put the recipe and when you want to use it. Also consider when it's appropriate to use each test.

ANOVA (conceptually)

- Like a sequence of t-tests, but multiple t-tests end up compounding the type I error (end up with a higher level of error than you set up with alpha)
- Compares the overall variability among/ between means of different groups with the internal spread within each group

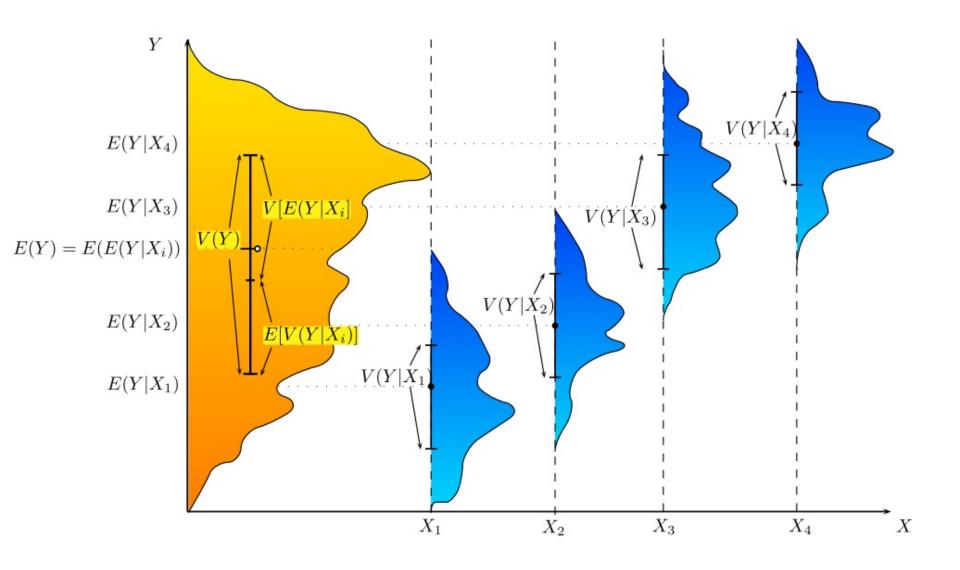


Figure 1: ANOVA: Fair fit

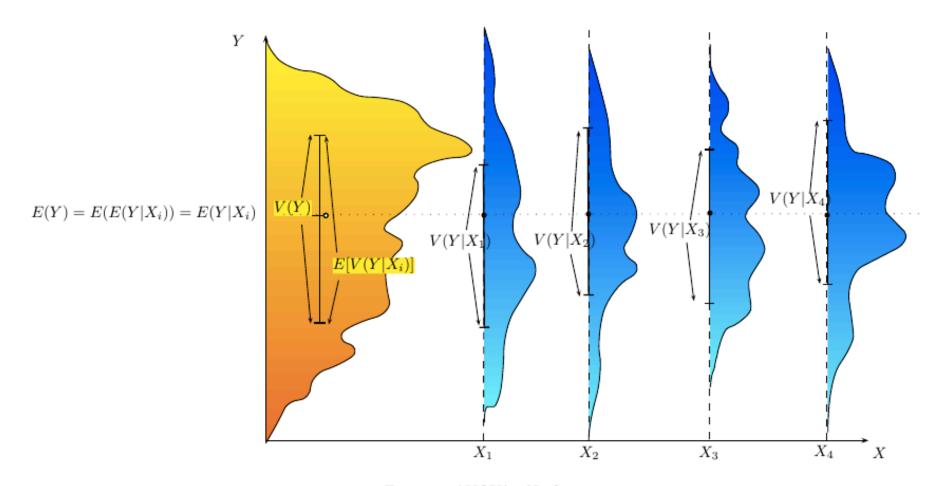


Figure 2: ANOVA : No fit

Illustration from: https://en.wikipedia.org/wiki/Analysis_of_variance

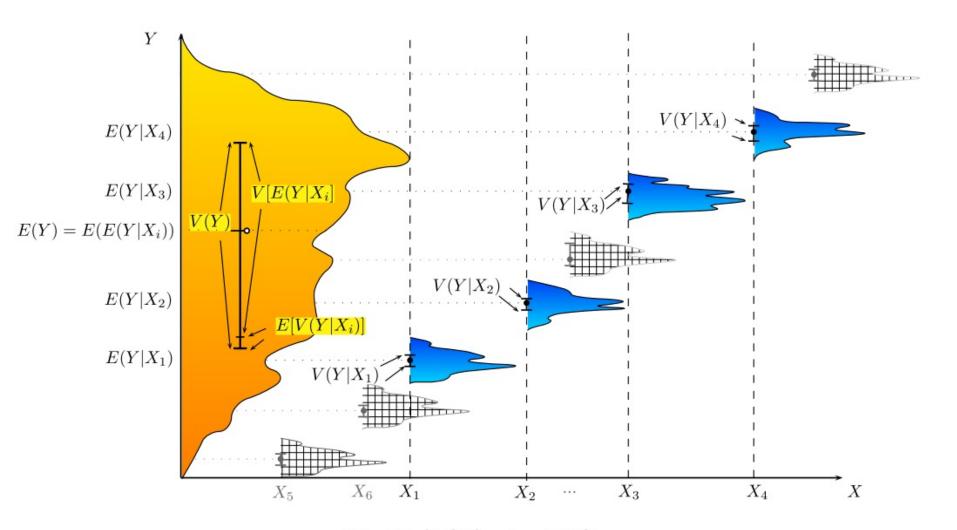
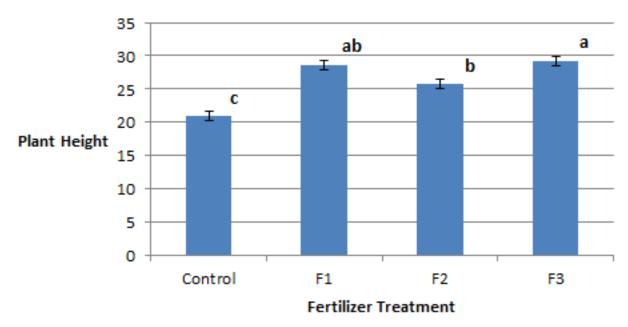


Figure 3: ANOVA: very good fit

We think plants will grow to different heights with different fertilizer treatment.

But we need a falsifiable hypothesis. What might this be?

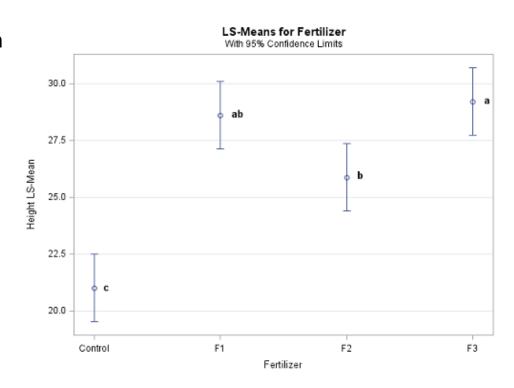
Hypothetical Greenhouse Experiment



Null Hypothesis: All groups have the same central value (mean).

The test:

- Compare the mean values of each group with overall mean for the entire data set.
- Need to analyze the variance to tell if we can really tell the mean values apart
- The F-ratio looks at the variance between groups divided by the variance within groups



How much variability is due to the different groups vs. something else (error/within group variability)

Total sum of squares = Treatment sum of squares + Error sum of squares (overall variation) = (group means – overall mean) + (variation within groups)
$$\sum_{j=1}^{k} \sum_{i=1}^{n_j} (y_{ij} - \bar{y})^2 = \sum_{j=1}^{k} n_j (\bar{y}_j - \bar{y})^2 + \sum_{j=1}^{k} \sum_{i=1}^{n_j} (y_{ij} - \bar{y}_j)^2$$

If the total sum of squares is divided by N-1, where N is the total number of observations, it equals the variance of the yij's. Thus ANOVA partitions the variance of the data into two parts, one measuring the signal and the other the noise. These parts are then compared to determine if the means are significantly different.

Where there are k different groups, with index j And each group has n_i different samples, with index i

Key Assumptions for ANOVA

If ANOVA is performed on two groups, the F statistic which results will equal the square of the two-sample t-test statistic $F=t^2$, and will have the same p-value. It is not surprising, then, that the same assumptions apply to both tests:

- All samples are random samples from their respective populations.
- All samples are independent of one another.
- 3. Departures from the group mean $(y_{ij} \overline{y}_{j})$ are normally distributed for all j groups.
- 4. All groups have equal population variance σ^2 estimated for each group by s_i^2

$$s_{j}^{2} = \frac{\sum_{i=1}^{n_{j}} (y_{ij}^{2} - \overline{y}_{j}^{2})^{2}}{n_{j}^{2} - 1}$$

See page 166 Helsel and Hirsh

Data from 3 Fertilizer Treatments

Fertilizer Treatment #

(also in lab)

		j=1	j=2	j=3	j=4
	(cm)	Control	F1	F2	F3
i=1	nt (c	21	32	22.5	28
i=2	<u>a</u> n	19.5	30.5	26	27.5
	of p	22.5	25	28	31
		21.5	27.5	27	29.5
	Height	20.5	28	26.5	30
i=6	He	21	28.6	25.2	29.2

Mean Square		<u>Formula</u>	Estimates:
Variance of yij	=	Total SS / N-1	Total variance of the data
MST	=	SST / k-1	Variance within groups +
			variance between groups.
MSE	=	SSE / N-k	Variance within groups.

N=24 (total number of observations) K=4 (number of different groups)

ANOVA Table

Source	df	SS	MS	F	p-value
Treatment	(k-1)	SST	MST	MST/MSE	р
Error	(N-k)	SSE	MSE		
Total	N-1	Total SS			

	j=1	j=2	j=3	j=4
	Control	F1	F2	F3
i=1	21	32	22.5	28
i=2	19.5	30.5	26	27.5
	22.5	25	28	31
	21.5	27.5	27	29.5
	20.5	28	26.5	30
i=6	21	28.6	25.2	29.2

<u>Mean Square</u>		<u>Formula</u>
Variance of yij	=	Total SS / N-1
MST	=	SST / k-1
MSE	=	SSE / N-k
N=24		
K=4		

Estimates:
Total variance of the data
Variance within groups +
variance between groups.
Variance within groups.

ANOVA

Source	df	ss	MS	F

	j=1	j=2	j=3	j=4
	Control	F1	F2	F3
i=1	21	32	22.5	28
i=2	19.5	30.5	26	27.5
	22.5	25	28	31
	21.5	27.5	27	29.5
	20.5	28	26.5	30
i=6	21	28.6	25.2	29.2

Mean Square		<u>Formula</u>	Estima
Variance of yij	=	Total SS / N-1	Total v
MST	=	SST / k-1	Variano
			va
MSE	=	SSE / N-k	Variano
N=24			

Estimates: Total variance of the data Variance within groups + variance between groups.

Variance within groups.

ANOVA

Source	df	ss	MS	F
Treatment	k-1=3			
Error	N-k=20			
Total	N-1=23			

	j=1	j=2	j=3	j=4
	Control	F1	F2	F3
i=1	21	32	22.5	28
i=2	19.5	30.5	26	27.5
	22.5	25	28	31
	21.5	27.5	27	29.5
	20.5	28	26.5	30
i=6	21	28.6	25.2	29.2

$$SST = \sum_{j=1}^{k} n_{j} (\overline{y}_{j} - \overline{y})^{2}$$

SSTrt =
$$6 * (21.0 - 26.1667)^2 + 6 * (28.6 - 26.1667)^2 + 6 * (25.8667 - 26.1667)^2 + 6 * (29.2 - 26.1667)^2 = 251.44$$
ANOVA

Source	df	ss	MS	F
Treatment	k-1=3	251.44		
Error	N-k=20			
Total	N-1=23			

	j=1	j=2	j=3 `	j=4
i=1 i=2	Control	F1	F2	F3
	21	32	22.5	28
	19.5	30.5	26	27.5
	22.5	25	28	31
	21.5	27.5	27	29.5
	20.5	28	26.5	30
i=6	21	28.6	25.2	29.2

Mean Square		<u>Formula</u>	<u>Estim</u>
Variance of yij	=	Total SS / N-1	Total v
MST	=	SST / k-1	Variano
			va
MSE	=	SSE / N-k	Variano

$$SSE = \sum_{i=1}^{k} \sum_{i=1}^{n_j} (y_{ij} - \overline{y}_j)^2$$

ANOVA

Source	df	ss	MS	F
Treatment	k-1=3	251.44		
Error	N-k=20	61.033		
Total	N-1=23			

	j=1	j=2	j=3	j=4
	Control	F1	F2	F3
i=1	21	32	22.5	28
i=2	19.5	30.5	26	27.5
	22.5	25	28	31
	21.5	27.5	27	29.5
	20.5	28	26.5	30
i=6	21	28.6	25.2	29.2

Total SS =
$$\sum_{j=1}^{k} \sum_{i=1}^{n_j} (y_{ij} - \overline{y})^2$$

+

Error sum of squares

(overall variation)

(group means – overall mean)

(variation within groups)

Recall: If you calculate two SS, you can get the third easily.

Source	df	ss	MS	F
Treatment	k-1=3	251.44		
Error	N-k=20	61.033		
Total	N-1=23	312.47		

	j=1	j=2	j=3	j=4
	Control	F1	F2	F3
i=1	21	32	22.5	28
i=2	19.5	30.5	26	27.5
	22.5	25	28	31
	21.5	27.5	27	29.5
	20.5	28	26.5	30
i=6	21	28.6	25.2	29.2

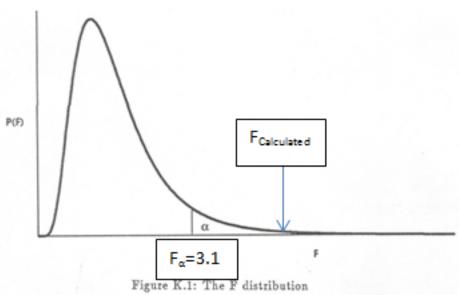
Mean Square		<u>Formula</u>	Estimates:
Variance of yij	=	Total SS / N-1	Total variance of the data
MST	=	SST $/ k-1$	Variance within groups +
			variance between groups
MSE	=	SSE / N-k	Variance within groups.

$$F = \frac{MS_{Trt}}{MS_{Error}} = \frac{83.813}{3.052} = 27.46$$

ANOVA

Source	df	ss	MS	F
Treatment	k-1=3	251.44	83.813	27.46
Error	N-k=20	61.033	3.052	
Total	N-1=23	312.47		

Look F up with a table or a software program. If it's p value is less than your rejection value, you reject the null.



We reject the null when the between treatment variance is significantly more than the within treatment variance.

One factor analysis of variance

Situation Several groups of data are to be compared, to determine if their means are significantly different. Each group is assumed to have a normal distribution around its mean. All groups have the same variance.

Computation The treatment mean square and error mean square are computed as their sum of squares divided by their degrees of freedom (df). When the treatment mean square is larger than the error mean square as measured by an F-test, the group means are significantly different.

$$MSE = \frac{\sum_{j=1}^{k} \sum_{i=1}^{n_{j}} (y_{ij} - \overline{y}_{j})^{2}}{N - k}$$
 where N-k = error degrees of freedom

Tied data No alterations necessary.

Test Statistic The test statistic F: From Helsel and Hirsh

F = MST / MSE

Decision Rule To reject H₀: the mean of every group is identical, versus H₁: at least one mean differs.

Reject H₀ if $F \ge F_{1-\alpha, k-1, N-k}$ the 1- α quantile of an F distribution with k-1 and N-k degrees of freedom; otherwise do not reject H₀.

If you reject the null, how do you know which sample (or samples) differ(s) from the rest?

- Multiple Approaches Exist
- Check what your software is using and cite it appropriately
- Don't just use multiple t-tests because you can inflate your type I error
- Most common (recommended) is Tukey's test:

Two group means
$$\bar{y}_i$$
 and \bar{y}_j can be considered different if $\left| \overline{y}_i - \overline{y}_j \right| > q_{(1-\alpha), k, N-k} \cdot \sqrt{MSE / n}$

where

is the studentized range statistic from Neter, Wasserman and Kutner (1985),

Also online and in stats books as Tukey q values is the overall significance level,

α

is the number of treatment group means compared, k

N–k are the degrees of freedom for the MSE, and

is the sample size per group. n

See page 198 of H&H or

https://onlinecourses.science.psu.edu/stat502/node/143

Also see:

https://www.mathworks.com/help/stats/analysis-of-variance-and-covariance.html

See Ch 7 H&H for Non-parametric

- ANOVA is like a t-test between three or more groups of data, and as such is restricted to the same assumptions as the t-test
- The Kruskal-Wallis test is similar to the rank-sum test, but is extended to more than two groups and compares the medians.
- There are many more advanced versions of the ANOVA (e.g., Multi-factor and ANCOVA), which you may wish to read about, but we will not go into them in this class

Lab 3.1 teaches ANOVA

- Teaches you how to do the fertilizer problem we just did in python code
- Relevant to the first problem on homework 3

Correlation Analysis

Correlation Coefficient:

For two samples X and Y with sample size n, consider the quantity:

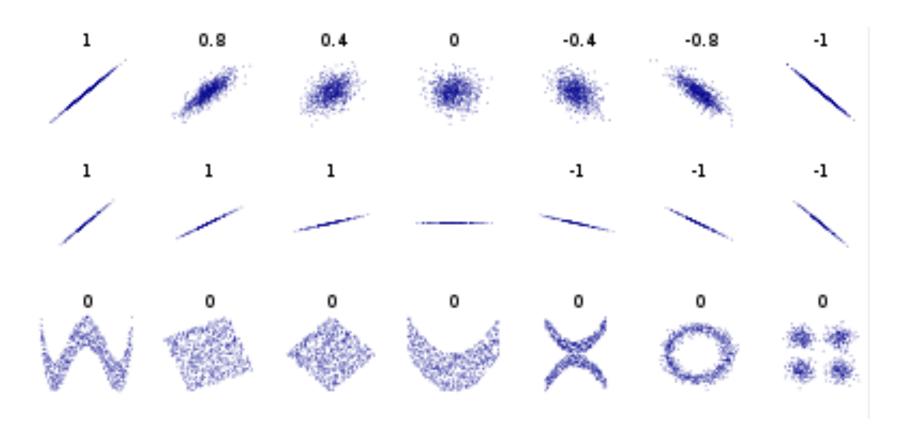
$$s_{xy} = \sum_{i=1}^{n} (x_i - \bar{x})(y_i - \bar{y})$$

If the anomalies of X and Y are mostly in the same direction at the same time, this will be a large positive value. If the anomalies are mostly of opposite sign at the same time, this will be a large negative value. If there is essentially no pattern, this value will be small.

We can scale this quantity so it varies between -1.0 and 1.0 to define the correlation coefficient r:

$$r = \frac{s_{xy}}{\left(\sqrt{\sum(x_i - \bar{x})^2}\right)(\sqrt{\sum(y_i - \bar{y})^2})}$$

Example Data Sets Plotted as (x,y) Pairs with Associated Correlation Coefficients



http://en.wikipedia.org/wiki/Pearson_product-moment_correlation_coefficient

Population vs. Sample Statistics

This is TRUE correlation, summed over all values $\rho = R = = \frac{\sum_{i=1}^{n} (X_i - \bar{X})(Y_i - \bar{Y})}{\sqrt{\sum_{i=1}^{n} (X_i - \bar{X})^2} \sqrt{\sum_{i=1}^{n} (Y_i - \bar{Y})^2}}$

So the sample statistic r from the previous slide is an estimator for the population statistic R above.

$$r = \frac{s_{xy}}{\left(\sqrt{\sum(x_i - \bar{x})^2}\right)\left(\sqrt{\sum(y_i - \bar{y})^2}\right)} \qquad s_{xy} = \sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})$$

Use, for example, the CORREL function in Excel to calculate the sample correlation coefficient.

In matlab, use corrcoef (see the lab)

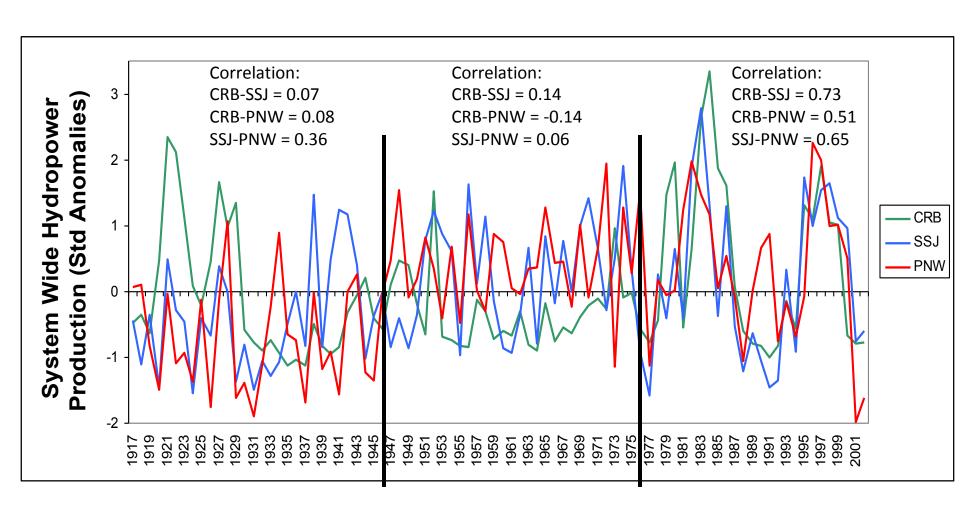
This is calculated for our dataset, sometimes also represented by R.

Some Important Characteristics of the Correlation Coefficient

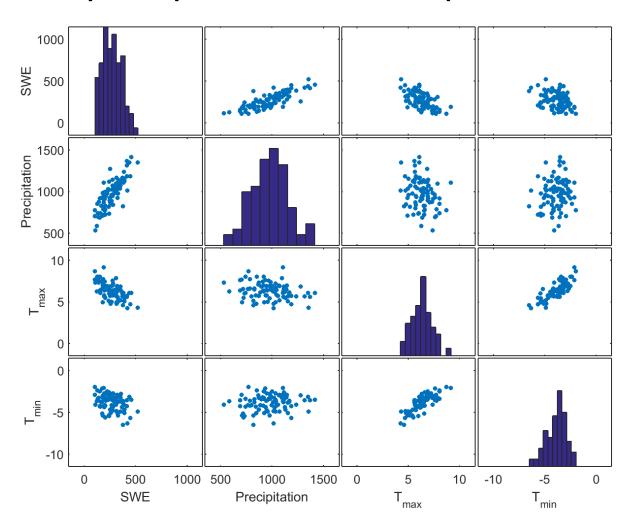
- 1. Value of **r** does not depend on labeling of "x" and "y".
- 2. The value of \mathbf{r} is independent of units (and is also invariant for certain kinds of linear transformations).
- **3. r** lies in the interval [-1,1].
- **4.** $\mathbf{r} = 1$ if and only if the x,y pairs lie on a straight line with positive slope, and $\mathbf{r} = -1$ if and only if the x,y pairs lie on a straight line with negative slope. If the slope is zero, then $\mathbf{r} = 0$. (Note this a "problem" in that if the relationship is very weak, but strongly linear, the relationship is still reported as being very strong. Solution, plot the data!)
- 5. The square of the sample correlation coefficient is R², the coefficient of determination for a simple linear regression model between the two variables.

Example:

CRB = Colorado River Basin SSJ = Sacramento-San Joaquin (California) PNW = Pacific Northwest



Example: How much of change in snow water equivalent is explained by changes in precipitation vs. temperature?



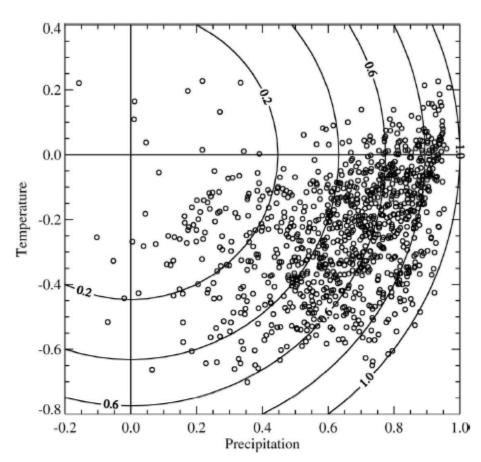


FIG. 3. Each small circle marks the correlations between 1 Apr SWE at one of the 995 snow course locations and the reference time series of Nov-Mar precipitation (x axis) and temperature (y axis). Contours indicate the quantity ($r_T^2 + r_P^2$), an approximation of the variance explained.

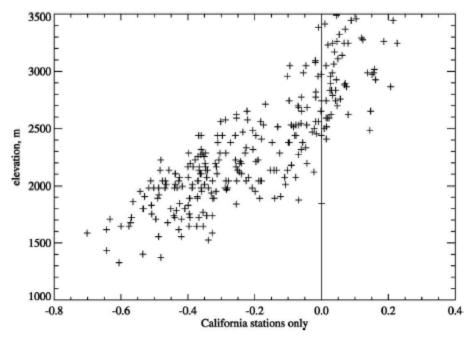
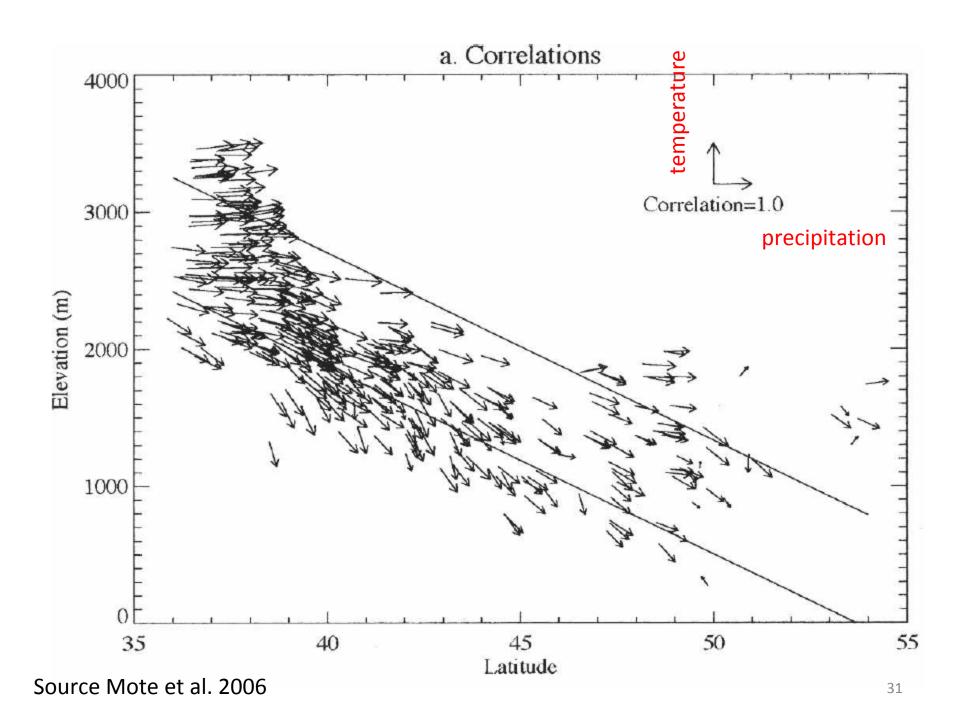


FIG. 4. Correlation between 1 Apr SWE and Nov-Mar temperature at each snow course in California, plotted as a function of snow course elevation.

 $\langle S \rangle = a_p \langle P \rangle + a_T \langle T \rangle$ are shown in Fig. 5 for the period 1960–2002. As noted previously for the periods of record 1950–97 (Mote et al. 2005) and 1916–2003 (Hamlet et al. 2005), observed trends in 1 April SWE over the period of record 1960–2002 are also predominantly pagetives in fact, the fraction of sites begins



Hypothesis Testing:

Null Hypothesis:

$$r = \frac{s_{xy}}{\left(\sqrt{\sum(x_i - \bar{x})^2}\right)(\sqrt{\sum(y_i - \bar{y})^2})}$$

$$\rho = 0$$

R and r are used interchangeably here. In Devore, this tests the certainty that your calculated correlation is the true correlation

Test Statistic:

$$t = \frac{R\sqrt{n-2}}{\sqrt{1-R^2}}$$
 n = Number of samples

Alternate Hypotheses:

$$\rho > 0$$

$$\rho < 0$$

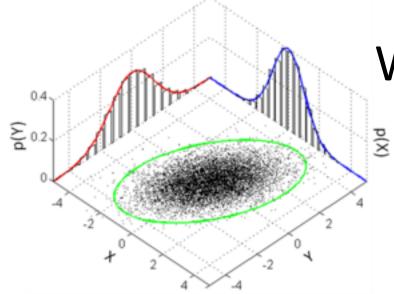
$$\rho \neq 0$$

$$t \geq t_{\alpha,n-2}$$

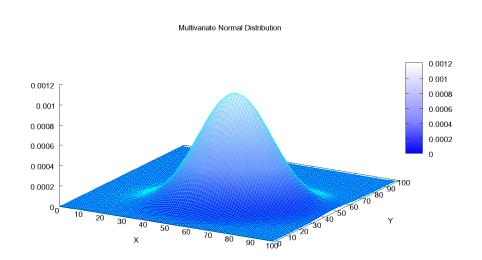
$$t \leq -t_{\alpha,n-2}$$

$$t \leq -t_{\frac{\alpha}{2},n-2} \quad OR \quad t \geq t_{\frac{\alpha}{2},n-2}$$

And these are read from the same tables as the t-test, student t distribution.



Examples of bivariate normal (from https://en.wikipedia.org/wiki/Multivariate_normal_distribution)



Why this test statistic?

We're testing for the absence of correlation, but X and Y may be normally distributed and uncorrelated but _not_ independent (recall happy face graphs – anything symmetric).

We assume that both X and Y are random, with a bivariate normal probability distribution (see Section 5.2 in Devore).

After you assume this, you can do a lot of math about drawing numbers from this distribution and chances of getting various correlations among those numbers.

https://en.wikipedia.org/wiki/
Pearson_productmoment_correlation_coefficient#Testing
_using_Student.27s_t-distribution 33

Hypothesis Testing Part II:

See page 493-494 in Devore

On canvas as

Null Hypothesis:

Devore_correlation.pdf

$$\rho = \rho_0$$

If we say that correlation is a specific number, say you want to be sure you have better than 0.5 correlation

Test Statistic:

$$Let V = \frac{1}{2} \ln \left(\frac{1+R}{1-R} \right)$$

$$ztest = \frac{V - 1/2 \cdot \ln{(\frac{1 + \rho_0}{1 - \rho_0})}}{1/\sqrt{n - 3}}$$

Alternate Hypotheses:

Rejection Region:

$$\rho > \rho_0
\rho < \rho_0
\rho \neq \rho_0$$

$$ztest > z_{\alpha}$$

$$ztest < -z_{\alpha}$$

$$ztest < -z_{\alpha/2} OR ztest > z_{\alpha/2}$$

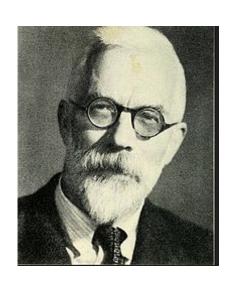
Why this formula?

Fisher figured this out in 1915.

https://en.wikipedia.org/wiki/Fisher transformation

Basically the key idea is to transform the sample correlation coefficient in such a way that the transformed variable would be normally distributed.

You can read these articles if you want to see the math.



$$Let V = \frac{1}{2} \ln \left(\frac{1+R}{1-R} \right)$$

$$ztest = \frac{V - 1/2 \cdot \ln{(\frac{1 + \rho_0}{1 - \rho_0})}}{1/\sqrt{n - 3}}$$

Regression Models

Homework 3 asks you to develop regression models related to streamflow timeseries.

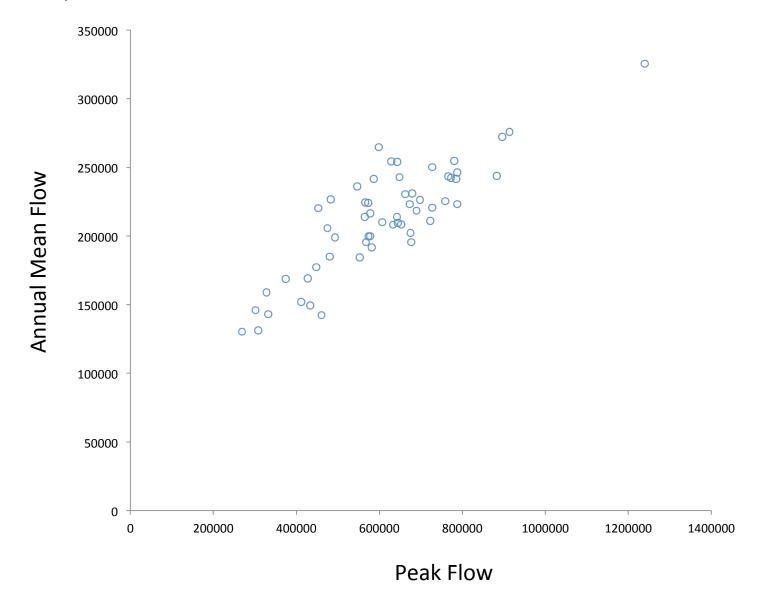
Least Squares Linear Regression:

In this approach we posit a linear relationship between an "independent" or "explanatory" variable $\,x\,$ and some "dependent" variable $\,y\,$:

$$y = B_0 + B_1 x$$

The first step in this process is to check whether a linear model approximation is reasonable. A good way to do this is to make a scatter plot of the available data:

Example from Homework 3, Problem 2: Columbia River flow



Fitting of Parameters:

The parameters: B_0 and B_1

Are selected so that the sum of the squared errors of the model are minimized for the available data. I.e. minimize:

$$\sum_{i=1}^{n} (y_i - (B_0 + B_1 x_i))^2$$

Taking partial derivatives with respect to $m B_0$ and $m B_1$ and setting equal to zero yields :

$$nB_0 + \left(\sum_{i=1}^n x_i\right) B_1 = \left(\sum_{i=1}^n y_i\right)$$
$$\left(\sum_{i=1}^n x_i\right) B_0 + \left(\sum_{i=1}^n x_i^2\right) B_1 = \left(\sum_{i=1}^n x_i y_i\right)$$

Solving for B_0 and B_1 yields:

$$B_1 = \frac{n \left(\sum_{i=1}^n x_i y_i\right) - \left(\sum_{i=1}^n x_i\right) \left(\sum_{i=1}^n y_i\right)}{n \left(\sum_{i=1}^n x_i^2\right) - \left(\sum_{i=1}^n x_i\right)^2}$$

$$B_0 = \frac{(\sum_{i=1}^n y_i) - B_1(\sum_{i=1}^n x_i)}{n} = \bar{y} - B_1 \bar{x}$$

$$Let \ \hat{y}_i = B_0 + B_1 x_i$$

Then the quantity $(y_i - \widehat{y}_i)$ is called the "ith residual".

Let:

$SSE = Sum \ of \ Squared \ Errors$

$$SSE = \sum_{i=1}^{n} (y_i - \hat{y}_i)^2$$

$$\sigma^2 = s^2 = \frac{SSE}{(n-2)}$$

$$\sigma = \sqrt{\frac{SSE}{(n-2)}}$$

s is also called the "standard error" of the regression model.

SST = Total Sum of Squares

$$Let SST = \sum_{i=1}^{n} (y_i - \bar{y})^2$$

How much variance is there about the mean.

$$R^2 = 1 - \frac{SSE}{SST}$$

 R^2 is often described as the fraction of the variance explained by the model. If the model is no better than predicting the mean, then the variance explained would be zero. A perfect model (i.e. SSE=0) is said to explain 100% of the variance.

Note similarity here to the ANOVA formulation. You can often see people using ANOVA analysis to discuss the variance explained by a specific grouping or classification. ANOVA is sometimes considered a special case of linear regression.

Confidence Bounds on Regression Parameters:

The variance of the regression parameter \hat{B}_1 is a function of the standard error and the "spread" of the x values.

$$s_{B_1}^2 = \frac{s^2}{\sum_{i=1}^n (x_i - \bar{x})^2}$$

And $\frac{(\hat{B_1} - B_1)}{s_{B_1}}$ is T distributed with n-2 degrees of freedom.

So a confidence interval for $\, B_{1} \,$ is:

$$\hat{B}_1 \pm t_{\frac{\alpha}{2},n-2} \cdot s_{B_1}$$

Hypothesis test for the estimator \widehat{B}_1

Asking, "Is there really a slope to my regression line?"

Null Hypothesis:
$$\widehat{B}_1 = B_1$$

 α = Probability of a Type I error, number of degrees of freedom = (n-2)

Test statistic:
$$t = \frac{(\hat{B}_1 - B_1)}{s_{B_1}}$$

Alternate Hypothesis: Rejection Region:

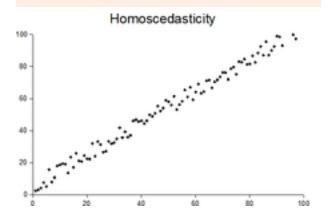
$$\hat{B}_1 > B_1$$
 $t \ge t_{\alpha,n-2}$ $\hat{B}_1 < B_1$ $t \le -t_{\alpha,n-2}$ $\hat{B}_1 \ne B_1$ $t \le -t_{\alpha,n-2}$ $t \le -t_{\alpha,n-2}$ $t \le -t_{\alpha,n-2}$ $t \le -t_{\alpha,n-2}$ $t \le -t_{\alpha,n-2}$

Estimating the Trend Using a Least Squares Linear Model:

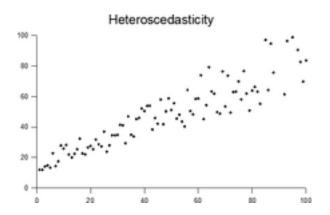
Some conditions that should be met for good results (see Helsel and Hirsch for more)

- Data should not be strongly auto correlated (more on this next week),
- •There shouldn't be any dramatic expansion in the variance over time.
- •A linear model should fit reasonably well (use a scatter plot to confirm)
- •The residuals for the linear model should be approximately normally distributed and shouldn't have large trends in them (plot these to get a sense of whether there are problems).

Homoscedasticity: random variables in a sequence have the same finite variance.



Heteroscedasticity: subpopulations have different variance from others.



Estimating the Trend Using a Least Squares Linear Model:

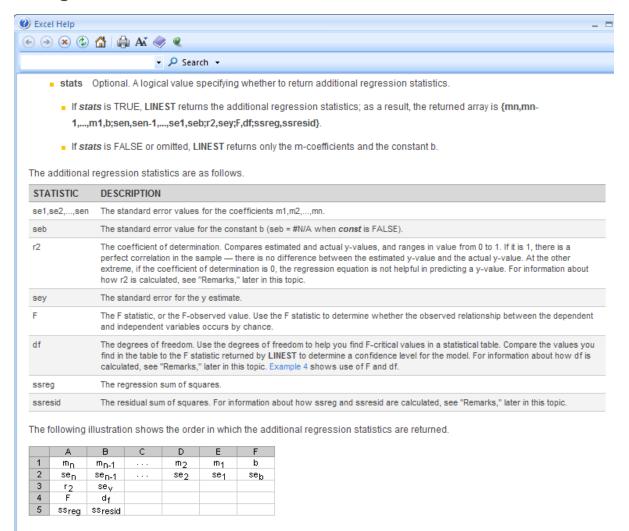
Some conditions that should be met for good results (see Helsel and Hirsch for more)

- Data should not be strongly auto correlated,
- •There shouldn't be any dramatic expansion in the variance over time.
- •A linear model should fit reasonably well (use a scatter plot to confirm)
- •The residuals for the linear model should be approximately normally distributed and shouldn't have large trends in them (plot these to get a sense of whether there are problems).

Procedures:

- •Calculate $\mathbf{B_1}$ (the trend) in the normal manner. (What are the units?)
- •Use hypothesis tests on $\, {\bf B_1} \,$ to see whether the trend is significantly different from 0 (i.e. no trend).
- •Use the confidence interval around the estimate of $\,B_{1}\,$ to express the uncertainty in the trend.

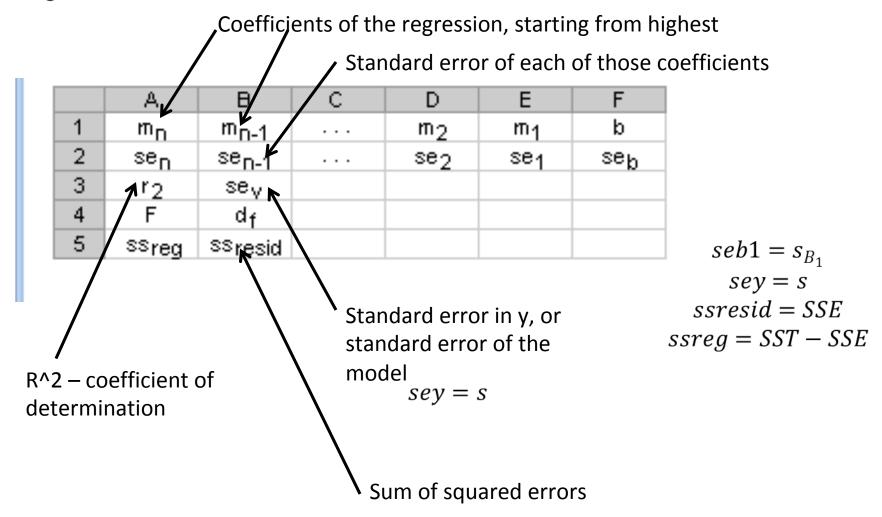
Using the LINEST Function in Excel:



 $seb1 = s_{B_1}$ sey = s ssresid = SSEssreg = SST - SSE

Note that LINEST function produces a table of output, so select a group of cells before typing in the formula and complete your formula entry as an array (ctrl-shift-enter).

Using the LINEST Function in Excel:



Note that LINEST function produces a table of output, so select a group of cells before typing in the formula and complete your formula entry as an array (ctrl-shift-enter).

LINEST Examples:

See example spreadsheet LINEST_examples.xlsx in tools folder on website

Constructing a Confidence Interval for the Predicted Values of Y: (see Regression_conf_intervals_Devore_p483.pdf or https://www.ma.utexas.edu/users/mks/statmistakes/ClvsPI.html)

Let x^* be a particular value of x

$$Let y^* = B_0 + B_1 x^*$$

$$Var[Y - (B_0 + B_1 x^*)]$$

= $Var(Y) + Var(B_0 + B_1 x^*)$

Var(Y) is the variance of Y relative to the model

How well did your model actually fit the data

(That is, the standard error of the model σ squared.)

and
$$Var(B_0 + B_1x^*)$$

is the variance of the model prediction.

How far away from your data mean are you

$SSE = Sum \ of \ Squared \ Errors$

$$SSE = \sum_{i=1}^{n} (y_i - \hat{y}_i)^2$$

$$\sigma^2 = s^2 = \frac{SSE}{(n-2)}$$

$$\sigma = \sqrt{\frac{SSE}{(n-2)}}$$

s is also called the "standard error" of the regression model.

The combined variance of the error of prediction at x^* can be shown to be:

$$\sigma_{E_p}^2(x^*) = var\left(y-y^*\right) =$$

$$= s^{2} \left[1 + \frac{1}{n} + \frac{n(x^{*} - \bar{x})^{2}}{n \sum_{i=1}^{n} x_{i}^{2} + (\sum_{i=1}^{n} x_{i})^{2}} \right]$$

Note: Xbar and Xi refer to the ORIGINAL data used to make the model. S is the original standard error.

And the statistic:

$$T = \frac{(y - y^*)}{\sigma_{E_p}(x^*)}$$

has a t distribution with n-2 degrees of freedom. Note: TINV in excel looks up a t-table. Also, check out tinv.m in matlab (built in function that looks up the t-table).

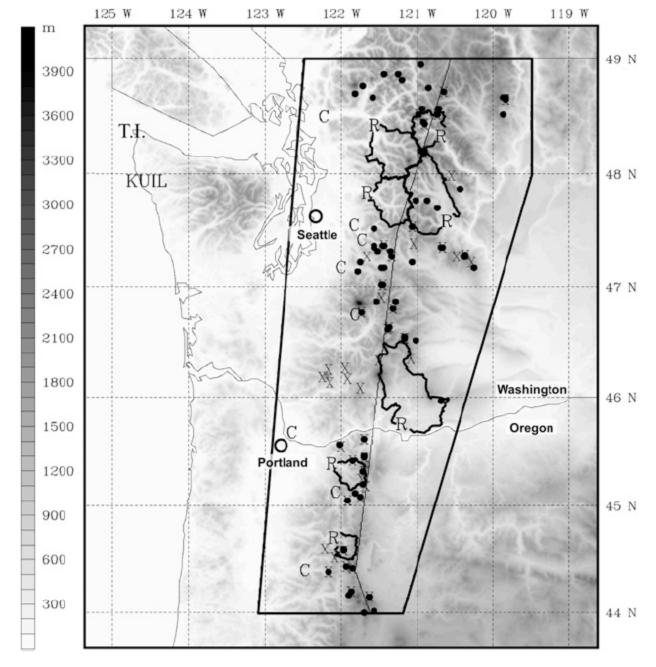
Thus a $(1-\alpha)$ confidence interval for

y at an arbitrary value of x^* is:

$$y^* \pm t_{\frac{\alpha}{2},n-2} \cdot \sigma_{E_p}(x^*)$$

Note that the uncertainty is a function of x^* and the farther away from \bar{x} we find ourselves the larger the uncertainty in the prediction of y!

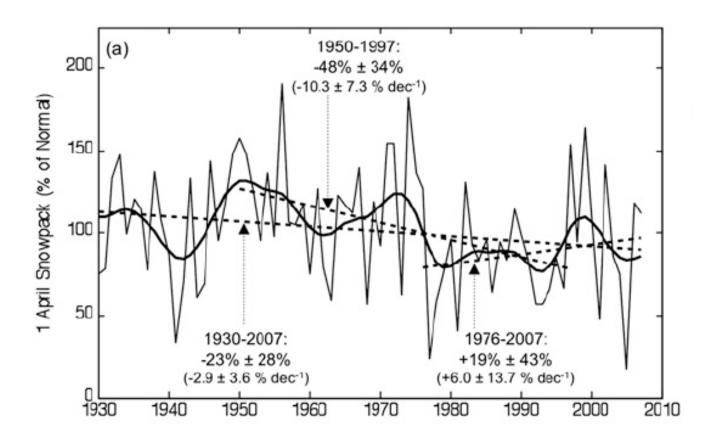
(Key thing to remember, these are not constant and vary with the location you want to predict.)



Stoelinga et al. 2010: Cascades Snowpack trends

FIG. 1. Map of study area. Heavy solid polygon defines "Cascade Mountains" for the purposes of this study. The thin solid line di-

What is shown here: Confidence values in predicted snow values based on a linear fit? Or uncertainties in the fitted slope of the line?



Snowpack trends for different periods: Note importance of the confidence intervals in those trends (from Stoelinga et al. 2010)

Non-Parametric Quantile Regression

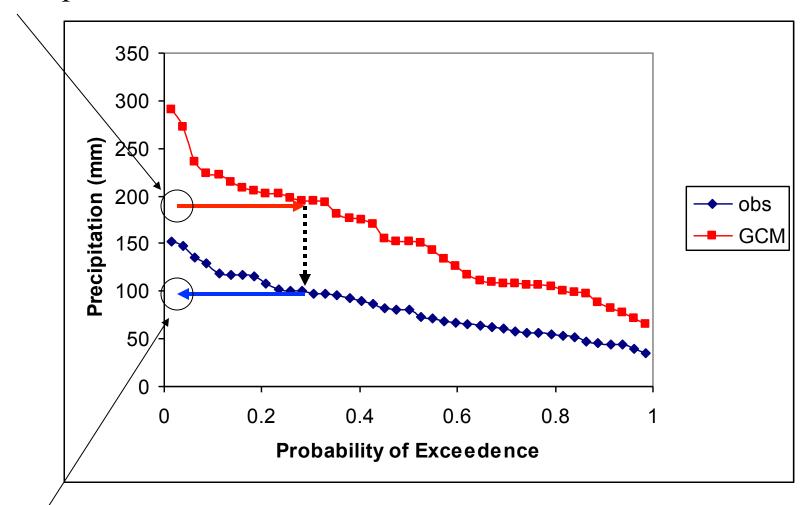
Non-Parametric Regression:

Non-parametric regression approaches have many advantages:

- •Do not require that the underlying probability distributions are known or have any particular form.
- •A linear relationship between the two variables is not required.
- •The time series of the data need not be the same (or even from the same time period) in the explanatory and dependent variables. That is, paired data is not required (although in many cases it is desirable).

We presume that relative ranking and frequencies of events are correct, even if actual values don't match up in a linear way.

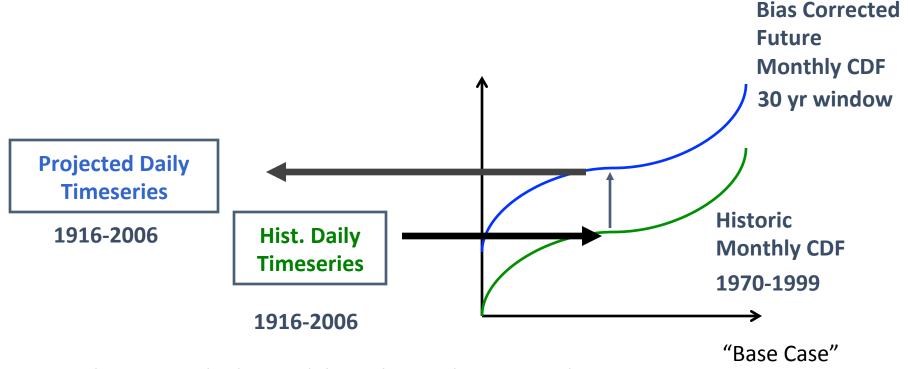
GCM Input = 190



Bias Corrected Output = 100

Hybrid Downscaling Method

Performed for each VIC grid cell:



- Used to correct hydro models in places where groundwater important
- Can be biased in absolute values but still have good change signals

Excel Example of Quantile Mapping Process:

See example spreadsheet quantile_regression. Xlsx

In tools folder

We will also work with this in today's lab.