Simple Linear Regression goal: Quantify association between a response variable (Y) and a single explanatory variable (X)

Multiple Linear regression Goal: Quantify association between a response variable (Y) and a series of explanatory variables $(X_1, X_2, ...X_n)$

Multiple regression analysis is an extension of simple linear regression and ANOVA. It focuses on the mean response (Y) at each *combination* of the explanatory variables. The explanatory variables can be either *continuous* or *nominal* (i.e. *categorical*).

Terminology and symbols: (Meadowfoam example)

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Notation for multiple regression:

μ {*flowers* | *light*, *time* }

which reads as "the mean number of flowers, as a function of light intensity and timing."

With specific values for the explanatory variables we get:

 $\mu \{ flowers \mid light = 300, time = 24 \}$

which reads as " the mean number of flowers when light intensity is 300 units and time is 24 days prior to PFI (photoperiodic flower induction).

Multiple Linear Regression Model

Simple linear regression model assumes that a straight line describes the relationship between the mean response (Y) and X:

$$\mu \{\mathbf{Y} \mid \mathbf{X}\} = \boldsymbol{\beta}_{0} + \boldsymbol{\beta}_{1} \boldsymbol{X}$$

Multiple regression extends that approach to cases with more than one explanatory variable:

 $\mu \{Y \mid X_1, X_2\} = \beta_0 + \beta_1 X_1 + \beta_2 X_2 \quad (2 \text{ explanatory variables } [X_1 \text{ continuous; } X_2 \text{ categorical}]; \text{ parallel lines})$

 $\mu \{Y \mid X_1, X_2\} = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_1 X_2 \text{ (2 explanatory variables [X_1 and X_2 as above] with interaction term; non-parallel lines)}$

 $\mu \{Y \mid X_1, X_2\} = \beta_0 + \beta_1 \log X_1 + \beta_2 \log X_2 \text{ (2 explanatory variables [X_1 and X_2 as above] log transformed; parallel lines)}$

 $\mu \{Y \mid X_1\} = \beta_0 + \beta_1 X_1 + \beta_2 X_1^2 \quad (1 \text{ explanatory variable [continuous] with quadratic term;}$ non-monotonic association)

All those models are *linear regression models*, because the regression coefficients $(\beta_1, \beta_2, \dots, \beta_n)$ represent a linear relationship between a *function* of the explanatory variables (e.g. X_1^2) and the mean response Y.

As before, *Least-Squares* are used to estimate the regression coefficients (β_0 , β_1 , β_2 ... β_n).

The *least-squares solution* consists of the values β_0 , β_1 ,..., β_n for which the sum of squares of deviations of the observed Y-values from the corresponding values predicted by the fitted regression model *is a minimum* (i.e. the *Error SS* in the ANOVA table is minimum).

An example: Effect of two Explanatory Variables without interaction

Y is the *mean number of flowers* produced by 10 plants X_1 is *light intensity* (6 levels between 150 and 900 units) X_2 is *timing* at which light intensity is changed from a basic value of 195 units. X_2 has 2 levels: 24 days before photoperiod floral induction (PFI) or at PFI (0 day). PFI is the time at which photoperiod is increased from 8 to 16 hrs of light per day to induce flowering.

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The following regression model can be seen as describing a *plane*:

μ {flowers | light, time} = $\beta_0 + \beta_1 light + \beta_2 time$

 β_0 is the height of the plane when both light and time equal zero. β_1 is the slope of the plane as a function of *light* for any <u>fixed</u> value of time β_2 is the slope of the plane as a function of *time* for any fixed value of light

<<Fig. 9.5 Sleuth>> In 3-D

But it is more easily visualized in 2-D.

<<Fig. 9.8 Sleuth>> In 2-D

For a parallel line model (no interactions), the effect of an explanatory variable (continuous or nominal) is the *change in mean response associated with a <u>one-unit increase</u> in that explanatory variable, <u>while holding all other explanatory variables fixed</u>.*

[With a significant *interaction* between light and time, the effect of light *depends* on the level of time; see below]

Interactions

Two factors "interact" if the effect that one factor has on the mean response (Y) depends on the value of the other factor.

Example: Does the effect of light intensity on flower production depend on timing?



No interaction implies that the effect of light intensity is the same at the 2 levels of time. *The lines are parallel.* The relationship is described by a parallel lines model ($\mu \{Y \mid X_1, X_2\} =$

$$\hat{\boldsymbol{\beta}}_{0} + \boldsymbol{\beta}_{1} \boldsymbol{X}_{1} + \boldsymbol{\beta}_{2} \boldsymbol{X}_{2})$$

An interaction implies that the effect of light depends on timing of the change in light intensity (time). The lines are not parallel. The relationship is described by a model with an interaction term $(\mu \{Y | X_1, X_2\} = \beta_0 + \beta_1 X_1 + \beta_2 X_2 + \beta_3 X_1 X_2).$

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The presence of a significant interaction in a regression model affects the interpretation of main effects (here light and time). *Those main effects* <u>can no longer be interpreted independently of each other</u>.

Indicator variables (or Dummy variables) to take into account nominal factors

In the meadowfoam example, time is treated as a *categorical* variable with 2 levels (typical of ANOVA framework). How do we deal with categorical variables in Multiple <u>regression</u> models?

The methods of regression analysis can be generalized to treat categorical explanatory variables with the use of *Indicator* or *Dummy* variables. Dummy variables allow analyses of categorical variables by *comparison of several regression equations originating from a single multiple regression model*.

In the flower production example, light can be treated as a *continuous* variable (simple linear regression framework; valid if mean responses are linear).

The effect of a continuous variable is estimated by β (least squares method)

To estimate the effect of categorical variables in the multiple regression framework, a binary *indicator variable* (also called *Dummy variable*) is created to represent the presence (coded as 1) or absence (coded as 0) of each *level* of the categorical variable.

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Rule for coding indicator variables

k-1 indicator variables are used to describe the set of levels of a particular nominal variable (a *factor*). These indicator variables are included as explanatory variables in the regression model. The level corresponding to the indicator variable that is <u>not included</u> is called the *reference level*.

The coefficient of an indicator variable is the *difference* between the mean response for the indicated category (= 1) and the mean response for the reference level (=0), *at fixed values of the other explanatory variables*.

Example: A parallel line model for the flower VS light / time problem.

Light is treated as *continuous* variable Time is *nominal* (categorical with 2 levels).

Because you code time as nominal, JMP defines a dummy variable to be 0 when time = 0 day prior to PFI and 1 when time = 24 days. Let's designate that dummy variable as day24.

The single multiple regression model with the dummy variable *day24* is:

 μ {flowers | light, time} = $\beta_0 + \beta_1$ light + β_2 day24

This model, when treated by a computer program, yields the following <u>2 models</u>, which are used to estimate β_2 :

 $\begin{array}{ll} day24=0: & \mu\{flowers \mid light, time=0\} = \beta_0 + \beta_1 \, light & (reference level) \\ day24=1: & \mu\{flowers \mid light, time=1\} = (\beta_0 + \beta_2) + \beta_1 \, light & \end{array}$

Only the Y's corresponding to day24=0 are used to calculate μ {flowers | light, time=0}; only the observations matching day24=1 are used to calculate μ {flowers | light, time=1}. The difference between the 2 means provides an estimate of β_2 .

The coefficient corresponding to the effect of time is thus estimated as:

time effect = $\mu \{ flowers / light, day24 = 1 \} - \mu \{ flowers | light, day24 = 0 \}$ = $[\beta_0 + \beta_1 light + \beta_2] - [\beta_0 + \beta_1 light]$ = β_2 (where β_2 is the difference between 2 means)

Thus, the multiple regression procedure with the dummy variable day24 *incorporates the two parallel line models within a single model*: μ {flowers | light, time} = $\beta_0 + \beta_1$ light + β_2 day24

- a) This multiple regression model states that the mean number of flowers is a straight-line function of light intensity for both levels of timing, the slope of the line being β_1 .
- b) The intercept for the line when day24=0 is β_0 .
- c) The intercept for the line when day24 =1 is $\beta_0 + \beta_2$.

The coefficient of the *indicator variable*, β_2 , is the amount by which the mean number of flowers with timing =24 days exceeds that with timing = 0 day, *after accounting for (i.e. averaging) the effect of light intensity differences*.

<<Fig 9.8 Sleuth>>

Analysis of the parallel line model in JMP: flower VS light and timing

Light is continuous Time is nominal

Response: Flowers						
Summary of Fit		Lack of Fit				
RSquar e	0. 799159	Source	DF Sum	of Squares	Mean Square	FRatio
RSquare Adj	0. 780031	Lack of Fit	9	215. 31077	23. 9234	0. 4377
Root Mean Square Error	6. 441073	Pure Error	12	655. 92510	54.6604	Pr ob⊳F
Mean of Response	56. 1375	Total Error	21	871. 23588		0.8894
Observations (or Sum Worts)	2					Max RSq
						0. 8488
		Par amet er Est	timates			
		Ter m	Estimate	Std Error	t Ratio Prob>∣t	t
		Intercept	77. 385	2.998156	25.81 <.0001	
		Time[1-2]	- 6.079166	1.314779	- 4.62 0.0001	
		Intensity	- 0. 040471	0.005132	- 7.89 <.0001	

Sleuth reports a coefficient of 12.2 for the effect of Time, but JMP a coefficient of -6.1. Why?

1- The sign of the coefficient for a categorical variable depends on which *reference level* is chosen in the analysis. Sleuth used time = 0 as the reference, but JMP used time = 24.

2- JMP *does not* <u>report</u> and <u>test</u> the effect of a categorical variable by taking the difference between a given level and the *reference level*.

The parameters estimated for each level are obtained by taking the difference between the mean response at a given level and *the average of the response at all levels of the variable*.

The test on the effect of *time* labeled Time [1-2] compares the first level of *time* to the *mean of the 2 levels of time*. We obtain an estimate for *time* of - 6.1, because level 1 (change at 0 day) is contrasted to the mean of the responses obtained at the 2 light levels (so it is $\frac{1}{2}$ smaller than 12.2, the difference between the response at time=24 and time =0).

Analysis of the flower VS light and timing problem: are the lines parallel?

In multiple regression, an explanatory variable for an *interaction* is constructed as the product of the 2 explanatory variables that are thought to interact.

The model with interaction can be written as:

 $\mu\{\text{flowers} \mid \text{light}, day24\} = \beta_0 + \beta_1 \text{ light} + \beta_2 day24 + \beta_3 (\text{light x } day24)$

This model yields the following 2 models, which are used to estimate the mean response at the 2 values of day24:

day24= 0: μ {flowers | light, time=0} = $\beta_0 + \beta_1$ light (reference level) day24=1: μ {flowers | light, time=1} = ($\beta_0 + \beta_2$) + ($\beta_1 + \beta_3$) light

The single multiple regression model states that both the intercept and slope in the regression of flowers on light depend on timing (i.e. day24):

At time = 0, the intercept is β_0 and the slope β_1 .

At time =24, the intercept is $\beta_0 + \beta_2$ and the slope $\beta_1 + \beta_3$. This model allows <u>both</u> for different slopes and intercepts.

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Analysis of the interaction model flower VS light and timing in JMP

Analysis	of Va	riance		
Source	DF	Sum of Squares	Mean Square	F Ratio
Model	3	3467.2765	1155.76	26.5490
Error	20	870.6598	43.53	Prob > F
C. Total	23	4337.9362		<.0001

Effect Tests						
Source	Nparm	DF	Sum of	Squares	F Ratio	Prob > F
Time	1	1	8	86.9504	20.3742	0.0002
Intensity	1	1	25	79.7500	59.2597	<.0001
time*intensity	1	1		0.5760	0.0132	0.9096
Parameter Est	imates					
Term		E	Estimate	Std Error	t Ratio	Prob> t
Intercept			77.385	3.07118	25.20	<.0001
time[1]		-6	.079167	1.346802	-4.51	0.0002
Intensity		-0	.040471	0.005257	-7.70	<.0001
time[1]*(intensit	y-525)	-0	.000605	0.005257	-0.12	0.9096

Because there is no evidence that the coefficient for the interaction (β_3) is different from zero, we conclude that the effect of light intensity *does not* depend on the timing of change in light intensity (two-sided p-value = 0.91, from a t-test for interaction, 20 degrees of freedom).

We conclude that the parallel line model describes well the relationship between number of flowers and light intensity and timing.

Fitting your own model with Dummy variables

Letting JMP choose the reference level is not always practical; also, comparing the effect of each level of a categorical variable is not easy when there are more than 2 levels (JMP uses the difference between each level and the mean of all levels).

You can code your own Dummy variables directly in the data table, which allows you to choose your *reference level*.

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For example, a separate-means model (ANOVA) could be fitted to describe the effect of light by using 5 Dummy variables for the 6 levels of light. (Alternatively you could specify *light* as a *nominal* variable in JMP.)

Example: In the flower VS light and timing problem, you can create <u>one</u> new column (time has 2 levels) for a Dummy variable day24, that takes a value of 0 if time = 0 and 1 if time =24. Using that column to estimate the effect of timing, you get:

Estimates			
Estimate	Std Error	t Ratio	Prob> t
71.305834	3.273772	21.78	<. 0001
- 0. 040471	0.005132	- 7.89	<. 0001
12.158333	2.629557	4.62	0. 0001
	Est i mat es Est i mat e 71. 305834 - 0. 040471 12. 158333	Estimates Estimate Std Error 71.305834 3.273772 -0.040471 0.005132 12.158333 2.629557	EstimatesEstimateStd Errort Ratio71.3058343.27377221.78-0.0404710.005132-7.8912.1583332.6295574.62

Here the effect of the categorical variable time is estimated by taking the difference between the level 24 days and the *reference level* 0 day.

We conclude that increasing light intensity decreased the mean number of flowers by an estimated 4.0 flowers per plant per 100 μ mol / m²/ sec (t = - 7.89, df = 21, P < 0.0001), after accounting for the effect of timing. Beginning the light treatment 24 days prior to PFI increased mean number of flowers by 12.2 flower per plant (t = 4.62, df = 21, P = 0.0001), after accounting for variation in light intensity.

Performing the equivalent of a t-test with a dummy variable and regression

Example: Compare height of 2 populations of plants, one which received only water (n = 10) and the other a fertilizer (n=8).

Recall we compared heights between populations with a 2-sample t-test, which yielded: $t_{16} = 2.99$, P= 0.0087.

Identically, we could have created an *indicator variable* to identify the 2 treatment levels and used linear regression.

Say we created an indicator variable DUMMY coded as:

0 for the control group 1 for the fertilizer group

We then perform the simple linear regression as:

Height = $\beta_0 + \beta_1$ DUMMY

For DUMMY = 0, the predicted mean height is : height = β_0 (β_0 is the average of the control group, i.e. the response when fertilizer = 0)

For DUMMY = 1, the predicted mean height is: height = $\beta_0 + \beta_1$ (β_1 is the difference between the average of the treatment and control group)

The test for β_1 (the coefficient of DUMMY) yields t = 2.99, P = 0.0087, with β_0 = 51.91 and β_1 = 4.64.

The difference in *mean response* due to fertilizer is 4.64 (control = 51.91; fertilizer = 56.55), which is the difference between the mean response for each level of the factor (i.e. \overline{y}_1 and \overline{y}_2).

Thus, the coefficient of the *indicator variable* (β_1) is the *difference* between the mean response for the *indicated category* (=1) and the mean for the *reference category* (=0).





Regression with indicator

Polynomial Regressions

When the change in the mean of the response variable as a function of the explanatory variable is not monotonic, the straight-line model is not adequate. By including polynomial terms based on the original explanatory variables, regression can be developed for relationships that exhibit curvature.

Example: Develop a model to describe the relationship between corn yield (bu/ac) and amount of rainfall (inches) in 6 U.S. states between 1890 and 1927.

Parameter estimates:	Parameter Es				
	Ter m	Estimate	Std Error	t Ratio	Prob> t
	Intercept	23.552102	3.236462	7.28	<. 0001
	rainfall	0. 7755493	0.293864	2.64	0. 0122

Linear Fit: YIELD = 23.552 + 0.775 RAINFALL

$$R^2 = 0.16$$



Add quadratic term (i.e. rainfall²) to account for curvature (produce a new column RAINFALL X RAINFALL with CALCULATOR in JMP; or use FIT POLYNOMIAL in the Fit Y by X platform):

Polynomial Fit degree =2:

$YIELD = -5.014 + 6.004 \text{ RAINFALL} - 0.229 \text{ RAINFALL}^2 \text{ (from Fit Model Platform)}$

$R^2 = 0.26$

Parameter Estimates

Term	Estimate	Std Error	t Ratio	Prob> t
Intercept	-5.014664	11.44158	-0.44	0.6639
RAINFALL	6.004283	2.03895	2.94	0.0057
Rain2	-0.229364	0.088635	-2.59	0.0140

Polynomial Fit degree =2:

YIELD = 21.66 + 1.057 **RAINFALL** - 0.229 (**RAINFALL** - 10.784)² (**from Fit Y by X Platform**) $R^2 = 0.26$ **Parameter Estimates**

lerm	Estimate	Std Error	t Ratio	Prob> t
Intercept	21.660175	3.094868	7.00	<.0001
RAINFALL	1.0572654	0.293956	3.60	0.0010
(RAINFALL-10.7842)^2	-0.229364	0.088635	-2.59	0.0140





Interpretation: The linear model seems inadequate. The quadratic model describes the curvature in the relationship between rainfall and corn yield that was evident is the scatterplot and patterns of the residuals.

With a significant quadratic term in the model, *corn yield depends on the specific value taken by rainfall*. The mean yield increases up to about 14 inches of rain per summer, then corn yield declines with additional amount of rain. It seems that there is an *optimum* amount of rain.

In fact, because β_2 is *negative* (i.e. the relationship is concave-down), the value of X that maximizes $\mu\{Y \mid X\} = \beta_0 + \beta_1 X + \beta_2 X^2$ is:

 $X_{\text{max}} = -\beta_1/2\beta_2$ (use β_1 and β_2 from the Fit *Model platform* to solve for the optimum).

For a *positive* β_2 (relationship concave-up), the value above would minimize the mean response (see Sleuth p. 290).

Interpretation depends on other explanatory variables included in the model

Example: Determine the association between weight (Y), height (X_1) and age (X_2) in 12 children.

Term	Estimate	Std Err	t Ratio	Prob> t
Intercept	6.55	10.94	0.60	0.564
Height	0.72	0.26	2.77	0.021
Age	2.05	0.94	2.19	0.056

Multiple Regression: weight = $\beta_0 + \beta_1$ height + β_2 age

A one-unit increase in height resulted in an estimated 0.72 unit increase in mean weight (t = 2.77, P= 0.021), independently of the effect of age. There is also evidence that a one-unit increase in age was associated with an increase in weight of 2.05 units (t = 2.19, P = 0.056), after accounting for the effect of height.

There is no evidence that β_0 (intercept) is different from zero (as expected, the mean weight of children is zero when both height and age are equal to zero).

Let's re-analyze the data but this time only considering the relationship between height and weight.

Term	Estimate	Std Err	t Ratio	Prob> t
Intercept	6.19	12.84	0.48	0.64
Height	1.07	0.24	4.44	0.0013

Simple linear regression: weight = $\beta_0 + \beta_1$ height

The coefficient associated with height is *larger* than before. This is because taller children also tend to be older. The coefficient of height in this *reduced model* contains the effect of height on weight and in addition reflects the association between age and weight.

In the model Weight = $\beta_0 + \beta_1$ height + β_2 age, the coefficient of *height* reflects the association between height and weight *after accounting for the effect of age*. In other words, the coefficient of *height* describes the association between height and weight *for children of the same age*.

A strategy for data analysis (with few explanatory variables)

Because values of the regression coefficients change when different effects are included in the regression model, one should not think that a single model will be better than all others in describing the data (especially with complex models).

There is a general strategy that can help developing an *inferential model*.

After considering the questions of interests and choosing a design:

- 1. *Explore* the data with graphical tools: consider transformations; check outliers.
- 2. Formulate an inferential model by wording questions in term of model parameters.
- 3. *Check* the model fit. a) if appropriate, fit a richer model, i.e. one with interactions or curvature
 - b) examine residuals and check for nonconstant variance and outliersc) drop unnecessary extra terms
- 4. *Infer* answers to the questions of interest using appropriate tools, such as confidence intervals and hypothesis tests.
- 5. Communicate results in language of ecology, not statistics.

Example. What lifestyles allow evolution of large brains in mammals?

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- a) Assuming that brains are energetically costly to produce, one hypothesis is that brain size in mammals is limited by the rate at which mothers can provide nutrition to their offspring. If this is true, we would expect a *negative* association between brain weight and litter size.
- b) Another reasonable assumption is that producing larger brains require more time than smaller ones. Thus one would expect brain weight to be associated *positively* with gestation period.
- c) Because body size is related to brain weight, metabolic rate, and gestation length through allometric relationships, the two hypotheses above should be investigated after accounting for variation in body weight.

Finding the expected relationships would provide support for the hypotheses that production of large brains:

1) requires energy efficiency from mothers (could it be related to diet quality?)

2) may require time (could it be disfavored in "r-selected" species and favored in "K-selected" species?).

response variable:	brain weight (g)
explanatory variables:	body weight (kg), gestation length (days), litter size

Which, if any, variables are associated with brain size, after accounting for variation in body size?

Not transformed

ln(litter)

In(gest)

In(body)

- 0. 310071

0.4179421

0.5750714

0. 115927

0. 140782

0. 032588



0.0089

0.0038

<. 0001

- 1. 0

- 1

0

1

2 3

5

6

4

In(brain) Predicted

8 9

- 2.67

17.65

2.97

Log transformed

There is evidence that brain weight is positively associated with gestation length (two-sided p-value = 0.0038 for a test that the slope is zero) after accounting for the effect of body weight and litter size. Brain weight is also negatively associated with litter size (two-sided p-value = 0.0089) after accounting for body weight and gestation period.

Although there are likely *cluster effects* due to taxonomic relatedness that may strongly influence statistical inferences, this analysis provides some evidence in favor of the ideas that rate at which mothers provide nutrition to their offspring or gestation length could limit the evolution of large brains in mammals.