

Generalized linear regression

Outline

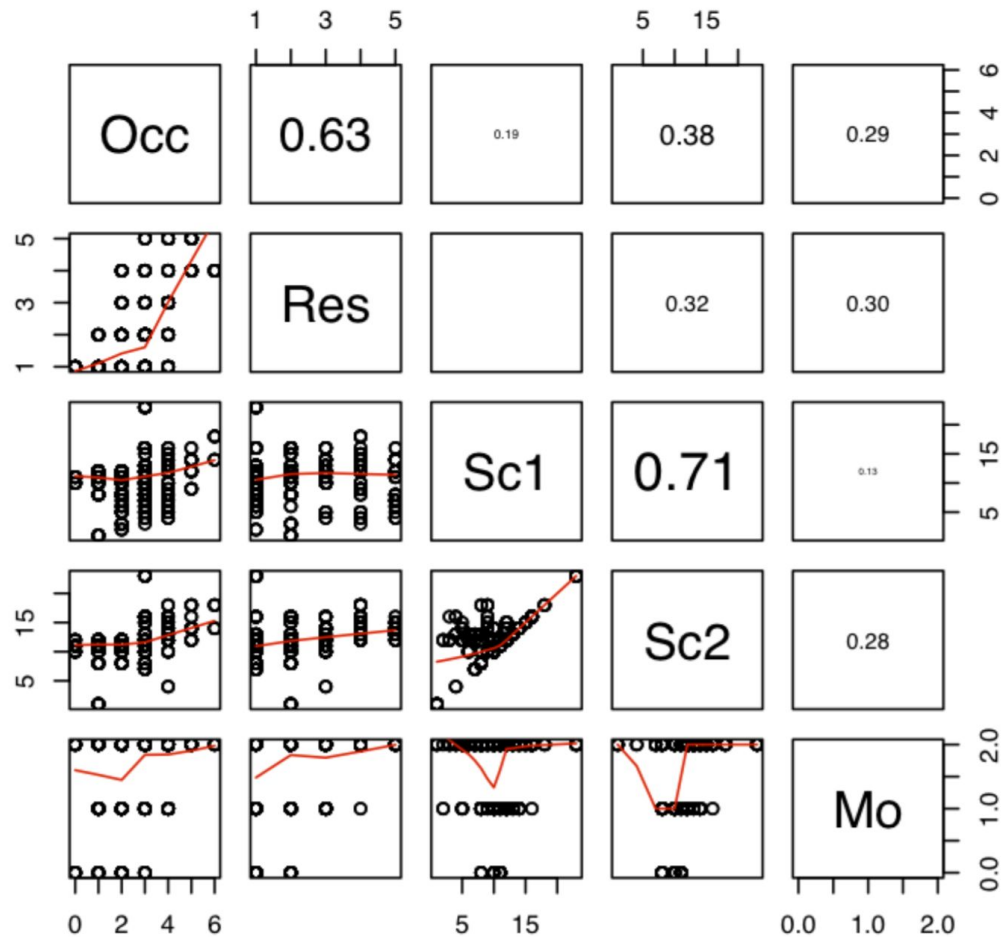
- Homework tips
- Interactions between independent variables
- *Generalized linear regression*
 - In particular, *logistic regression*, used for binomial dependent variables
- For home consumption: isotonic regression

Homework 05 tips (1/)

- If you're using `stopifnot` to verify properties, and the condition fails, try `printing` out the relevant values just before you run `stopifnot`.
 - Unfortunately R doesn't let you specify a logging statement with `stopifnot`, though Python does with `assert`.
- The command `all.equal` can be used to check if two values are very close, and plays nicely with `stopifnot`.

Homework 05 tips (2/)

- A few people were confused by the notation $Y \sim 1$:
 - If you have $Y \sim X$ and you want to "drop out" X , R doesn't let you write $Y \sim$.
 - The 1 here symbolizes the intercept.
 - If for some reason you want a model *without* an intercept you can write $Y \sim -1$.
- Generalizing a bit, if you have $Y \sim X_1 + X_2$:
 - If you want to "drop out" X_1 , you write $Y \sim X_2$.
 - If you want to "drop out" X_2 , you write $Y \sim X_1$.
- The function `lrtest` from the `lmtest` package can compute the log-likelihood ratio test (both the test statistic and the p -value) for two models so long as one nests the other; you still have to fit the "dropped out" models though.



Questions?

Interactions

Interaction terms

In some cases we are interested not just in the effect of a given independent variable (IV) but its *interaction* with another IV.

The nature and interpretation of the interaction depends on the the types of IVs involved.

Types of interaction

- Interaction of two binomial IVs: is there an change in Y when both X_1 and X_2 are active (i.e., true) beyond that associated with X_1 and X_2 ?
 - Are the effects of X_1 and X_2 on Y independent?
- Interaction of a binomial IV X_b and a continuous IV X_c : is the change in Y associated with X_c different when X_b is active?
 - Is the slope of X_c with respect to Y different when X_b is active?
- Interaction of two continuous IVs: is Y also sensitive to the product of X_1 and X_2 ?

The interaction of two multinomial IVs is best understood by decomposing them into binomial IVs.

Specifying interactions in R formulae

Long form:

$$Y \sim X1 + X2 + X1 : X2$$

Short form:

$$Y \sim X1 * X2$$

You *can* specify an interaction without a main term (e.g., $Y \sim X1 : X2$) but it is rarely needed.

Example (1/)

Is petal length influenced by the *interaction* between sepal length and sepal width?

Q: What kind of interaction is this?

A: An interaction between continuous IVs.

```
> r.simple <- lm(Petal.Length ~ Petal.Width +
+               Sepal.Length + Sepal.Width,
+               data = iris)
> summary(r.simple)
```

...

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	-0.26271	0.29741	-0.883	0.379	
Petal.Width	1.44679	0.06761	21.399	<2e-16	***
Sepal.Length	0.72914	0.05832	12.502	<2e-16	***
Sepal.Width	-0.64601	0.06850	-9.431	<2e-16	***

...

```
> r.intrct <- lm(Petal.Length ~ Petal.Width +
+               Sepal.Length * Sepal.Width,
+               data = iris)
> summary(r.intrct)
```

```
...
              Estimate Std. Error t value Pr(>|t|)
(Intercept)    0.71482    1.56623    0.456  0.6488
Petal.Width    1.43584    0.06991   20.539 <2e-16 ***
Sepal.Length   0.56175    0.26970    2.083  0.0390 *
Sepal.Width   -0.97041    0.51486   -1.885  0.0615 .
Sepal.Length:Sepa...  0.05642    0.08874    0.636  0.5259
...
```

Example (2/)

The three-way interaction model

```
Petal.Length ~ Petal.Width + Sepal.Length + Sepal.Width +  
Petal.Width:Sepal.Length +  
Petal.Width:Sepal.Width +  
Sepal.Length:Sepal.Width +  
Petal.Width:Sepal.Length:Sepal.Width
```

can also be fit (if you have enough data), but is basically uninterpretable.

Example (3/)

Is there an interaction between age and gender in the ANAE low back merger data (from homework 05)?

Q: What kind of interaction is this?

A: An interaction between a continuous IV (age) and a binomial IV (gender...uh... at least as it is coded in that data).

```

> r.simple <- lm(distance ~ age + gender + dialect,
+               data = anae)
> summary(r.simple)
...

```

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	176.8756	7.8732	22.465	< 2e-16	***
age	0.7735	0.2954	2.618	0.009177	**
gender1	4.4896	4.4425	1.011	0.312821	
dialect1	-103.6565	30.3734	-3.413	0.000708	***
dialect2	-138.7272	34.7547	-3.992	7.80e-05	***
dialect3	-132.0029	18.7318	-7.047	8.00e-12	***
dialect5	1.5156	48.6868	0.031	0.975181	

```

...

```



```
> r.intrct <- lm(distance ~ age + gender + age:gender +  
+ dialect, data = anae)
```

```
> summary(r.intrct)
```

...

	Estimate	Std. Error	t value	Pr(> t)	
(Intercept)	177.0400	7.8752	22.481	< 2e-16	***
age	0.6877	0.3080	2.233	0.026093	*
gender1	4.2793	4.4478	0.962	0.336575	
age:gender1	0.2984	0.3024	0.987	0.324383	
dialect1	-102.7246	30.3890	-3.380	0.000795	***
dialect2	-141.1245	34.8407	-4.051	6.14e-05	***
dialect3	-132.4366	18.7376	-7.068	7.02e-12	***

...

Example (4/)

There is no consensus whether it is sensible to test for interactions (e.g., of age and gender) when one or the other non-interaction term is non-significant (e.g., above, where the standard error for age was larger than the coefficient).

Nota bene

- Caution is necessary when "dropping" (i.e., doing likelihood ratio tests on) models with interaction terms:
 - If the model is $Y \sim X1 * X2$, to drop the interaction you write $Y \sim X1 + X2$.
 - To make this even clearer, you can specify the full model as $Y \sim X1 + X2 + X1:X2$.
 - If you drop a (non-interaction) independent variable, you should also remove its interactions.
- `drop1` does not understand interactions and gives nonsensical results if they are present; stepwise fitting functions may similarly be confused.
- Three- and four-way interactions are hard to interpret, burn up degrees of freedom quickly, and tend to give short shrift to main effects:

"Analysts usually steer clear of higher-order interactions".

Generalized linear regression

Generalizing linear regression

Two key assumptions of linear regression (as well as ANOVA) is that the dependent variable (DV) is

- normally distributed (or the central limit theorem applies), and
- a linear sum of the coefficients and their IVs.

This assumption is *flagrantly* violated when the DV is a proportion or probability, as

- probabilities violate the assumption of homogeneity of variance, and
- for probability DVs a linear model can predict $p < 0$ or $p > 1$.

Problems with percentages

A change of a percentage $\hat{p} = .5$ is "less" (according to the binomial distribution) of a change than a change for probability close to 0 or 1, so

- effects close to 0 or 1 are underestimated and
- effects close to .5 are overestimated.

"In what space can we capture these intuitions?"

Desiderata:

- smooth, continuous, differentiable transformation function
- domain $[0, 1]$, range $(-\infty, +\infty)$

The arcsine transformation

One traditional answer to these questions is the *arcsine* (or *angular*) transformation, defined by the inverse sine of the square root of the proportion, or

$$\arcsin \sqrt{p}$$

which has a range of $[0, 2\pi]$. Or in R:

```
> asin(sqrt(p))  
[1] 0.4636476
```

From probabilities to odds

The odds of a probability p is simply:

$$O = p / (1 - p)$$

This has the range $[0, +\infty]$.

For instance, for $p = .9$, $O = 9$, and for $p = .1$, $O = 0.1111$.

From odds to log-odds

Because of the strange range...

e.g., $p < .5$ implies $0 < O < 1$, whereas $p > .5$ implies $1 < O < \infty$,

it is often preferable to work in log-space, where the range is $[-\infty, +\infty]$.

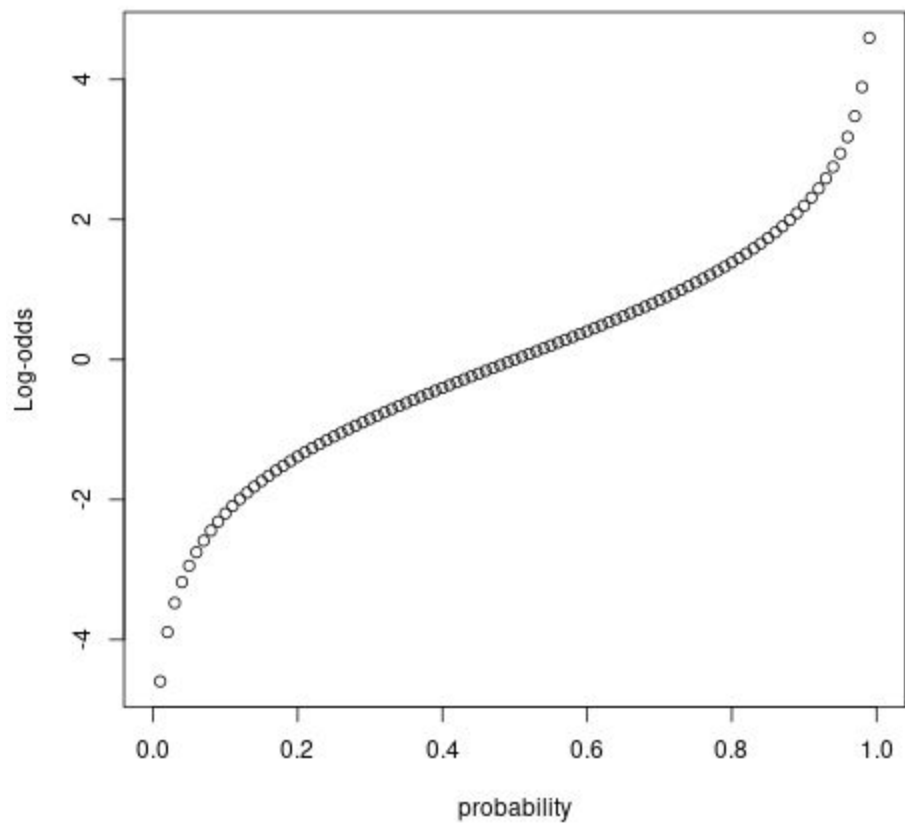
$$\begin{aligned}\log O &= \log p - \log(1 - p) \\ &= \log c - \log(N - c)\end{aligned}$$

For instance, for $p = .9$, $\log(O) = 2.197$, and for $p = .1$, $\log(O) = -2.197$.

Introducing `qlogis`

In R, the transformation from probabilities to log-odds, the *logit*, is performed by `qlogis`.

```
> qlogis(seq(0, 1, .1))  
[1]          -Inf -2.1972246 -1.3862944 -0.8472979 -0.4054651  
[6] 0.0000000 0.4054651 0.8472979 1.3862944 2.1972246  
[7] Inf
```



Introducing logistic regression

The framework of *generalized linear models* are linear models augmented with *link functions* (such as the logit) which map arbitrary types of DVs onto a linear function.

With some magic (not covered in this class), we can then estimate the parameters of such models.

A generalized linear models with logit link functions are known as *logistic regression* models.

FYI: the *logistic* is the inverse of the logit function, mapping from log-odds to probabilities.

Logistic regression in R

To specify a logistic regression, we use the function `glm` (which fits generalized linear models) and specify `family = binomial` (which enables a logit link function, giving us *logistic* regression) in particular.

Nearly all other linear model functions work the same; we can call `summary`, compute `residuals`, perform the likelihood ratio test with `drop1`, etc.

An example from the [ʃ]treets of Columbus (1/)

The envelope of variation is pronunciation of word-initial *str-* as [str] vs. [ʃtr].

Data collected using a *rapid anonymous* design: ask for directions to a nearby bank so as to elicit tokens of *street* (cf. Labov 1966 on post-vocalic *r* in New York via *fourth floor*, Prichard 2010 on /ay/-monophthongization in Atlanta via *five o'five*).

An example from the [j]treets of Columbus (2/)

Predictors include:

- Gender
- Emphasis (normal vs. a second rendition after "what did you say?")
- Age (coded as "young", "middle", or "old")
- Social class (coded as "working class", "lower middle class", and "upper middle class")

```
> xtabs(~ str + emphatic, data = cbus)
```

```
  emphatic
```

```
str      Less More
```

```
shtr     43   14
```

```
str      77  106
```



```
> r <- glm(str ~ emphatic, data = cbus, family = binomial)
> summary(r)
```

Call:

```
glm(formula = str ~ emphatic, family = binomial, data = cbus)
...
```

Coefficients:

	Estimate	Std. Error	z value	Pr(> z)	
(Intercept)	0.5826	0.1904	3.060	0.00221	**
emphaticMore	1.4418	0.3422	4.213	2.52e-05	***
...					

Interpretation notes

Using the standard procedure of computing estimates for Y by adding the intercept and the product of coefficients and IVs, we obtain a number in log-odds space. We can convert back to an estimated probability using the R function `plogis`, the inverse of `qlogis`.

E.g., to estimate $P(\text{str} \mid \text{"more emphatic"})$, we have:

```
> intercept <- 0.5826
> moreEmphatic <- 1.4418
> plogis(intercept + moreEmphatic)
[1] 0.8833352
```

Nota bene

There is a strong connection between (binomial) logistic regression for statistical inference and so-called *multinomial logistic regression* (or *maxent models*) used for classification in speech and language processing, though

- they often use different representations of the IVs (e.g., dense continuously-valued vs. sparse booleans), and
- they often use different learning algorithms (e.g., iteratively-reweighted least squares vs. stochastic gradient descent).

Questions? Please take
them to email, or Slack.