

Today's topic is mainly about **simple linear regression models**.

Suppose lots of data in a population with random variables  $X$  and  $Y$ . (There is an obvious trend going through the data points.) The size of the population is  $N$ , which is hard to know, because we always sample the population to inference.

We have  $E[Y|X] = \beta_0 + \beta_1 X_i$  as the mean of the model.

For data point  $(X_i, Y_i)$ , what we really care for is that

$$Y_i = \beta_0 + \beta_1 X_i + \epsilon_i, i = 1, 2, \dots, N$$

where for  $i = 1, 2, \dots, N$ ,  $\epsilon_i$  is some random variation, the vertical distance from  $(X_i, Y_i)$  to the line. And in the population we call it the **error**.

But in practice, we never have the chance to draw a line like this because that is the population. In fact, hopefully after a representative sampling process, we can draw a sample picture instead.

What's changed? Axes becomes  $x, y$  and the number of data points decreases (because of sample). Then we are gonna fit a model into the data we have as estimation. We wish it could reflect the true model in population.

Note that  $\beta_0$  was the intercept of linear line to x-axis,  $\beta_1$  was the slope of the line. And we are gonna estimate those two:

$$\hat{\beta}_0 = b_0, \hat{\beta}_1 = b_1$$

where  $b_0$  is the intercept of the sample line,  $b_1$  is the slope of the sample line.

And for each data point in the sample  $(x_i, y_i), i = 1, 2, \dots, n$ ,  $n$  is the sample size, there is a corresponding point on the linear line  $(x_i, \hat{Y}_i)$ , and the vertical difference between them is  $e_i = y_i - \hat{Y}_i$  which is called the **residual**.

So the line should be:

$$\hat{Y}_i = \hat{\beta}_0 + \hat{\beta}_1 x_i, i = 1, 2, \dots, n$$

Ok, we've all set up. Back to the big question: **How do we estimate  $\hat{\beta}_0 = b_0, \hat{\beta}_1 = b_1$ ?**

We use Gauss' method of least squares (check textbook): find  $b_0, b_1$  that minimize the sum of squares of the errors.

$$\begin{aligned} \text{population } \sum_{i=1}^N \epsilon_i^2 &= \sum_{i=1}^N (Y_i - \hat{Y}_i)^2 \\ \text{or sample } \sum_{i=1}^n e_i^2 &= \sum_{i=1}^n (y_i - (\hat{\beta}_0 + \hat{\beta}_1 x_i))^2, \\ \hat{\beta}_0 = b_0, \hat{\beta}_1 = b_1 &\text{ are the estimates that minimize this!} \end{aligned}$$

To calculate  $b_0, b_1$  in practice we need means and variances of the  $x, y$  sample variables and we also need the covariance of  $X, Y$ :

To estimate this in the sample we use sum of products of the deviation from  $x$  to its mean and the deviation of  $y$  to its mean (over degrees of freedom):

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

then

$$\begin{aligned} \hat{\beta}_1 = b_1 &= \frac{\frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})(y_i - \bar{y})}{\frac{1}{n-1} \sum_{i=1}^n (x_i - \bar{x})^2} = \frac{S_{xy}}{S_{xx}} \\ \hat{\beta}_0 = b_0 &= \bar{y} - b_1 \bar{x} \end{aligned}$$

Two competing models

model 1: population is  $Y = \beta_0 + \epsilon$ , the estimated version (sample) is  $\hat{Y} = \bar{y}$

model 2: population  $Y = \beta_0 + \beta_1 X + \epsilon$ , sample  $\hat{Y} = \hat{\beta}_0 + \hat{\beta}_1 x = b_0 + b_1 x$

What's the difference between the two models? The term  $\beta_1 X$ . The first does not assume that  $X$  has any effect.

Do we need this term?

- if we don't then  $\beta_1 = 0$  and we have model 1.
- if we are convinced that a positive linear trend is a better fit then  $\beta_1 > 0$  and we have model 2.

We should do hypothesis test of  $H_0 : \beta_1 = 0$  v.s.  $H_A : \beta_1 > 0$ .

By definition, the standard error of  $\beta_1$  is the standard deviation of the sampling distribution of  $\beta_1$ . So how do we work this out?

Assumptions underlying a simple linear regression (SLR) model

1. General assumptions (applicable to most statistical models)
  - (a) that the sample is representative of the population of interest
  - (b) that the explanatory ( $X$ ) variables are measured without error (or at least minimal error of  $Y$ )  $\rightarrow$  all the error is in the  $Y$  direction (vertical on the earlier plots)
  - (c) that a model of the proposed form (e.g. a linear model) is appropriate
2. Model-specific assumptions (most regression-type models including SLR)
  - (a) (population)  $Y_i = \beta_0 + \beta_1 X_i + \epsilon_i, i = 1, 2, \dots, N$  where  $\beta_0 + \beta_1 X_i$  is the deterministic model for the mean  $E[Y_i|X] = \beta_0 + \beta_1 X_i$ , and  $\epsilon_i$  is the stochastic model for the variance. The assumptions, specific to this model, are about  $\epsilon_i$

$$\epsilon_i \stackrel{iid}{\sim} N(0, \sigma^2)$$

That errors ( $\epsilon_i$ ) are independent and identically (normally) distributed with mean 0 and constant variance  $\sigma^2$ . This in a nutshell is the variance model.