

Linear regression cheatsheet

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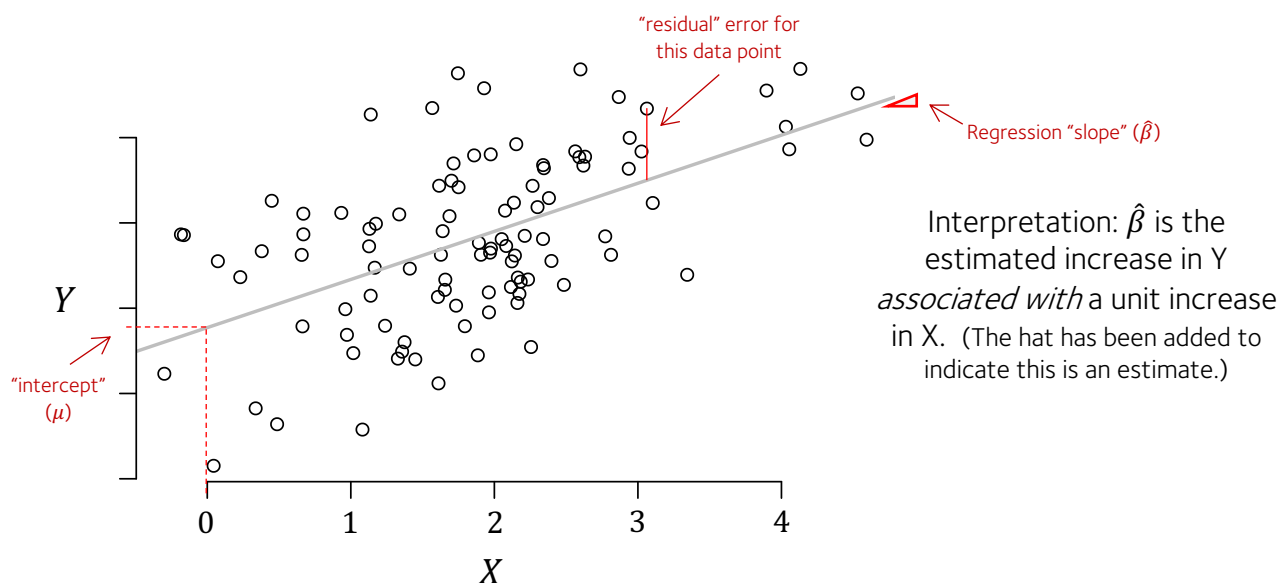
Linear regression models an outcome variable (Y) in terms of one or more predictor variables (X). The model asserts that Y is a linear combination of columns of X plus some noise. The noise is assumed to be Gaussian with some variance σ^2 . The residual variance is assume to be the same for all data points).

$$Y = \mu + X_1\beta_1 + X_2\beta_2 + \dots + X_d\beta_d + \epsilon \quad \epsilon \sim N(0, \sigma^2)$$

Or using matrix notation:

$$Y = \mu + X\beta + \epsilon \quad \epsilon \sim N(0, \sigma^2)$$

Matrix multiplication of the d -dimensional *row* vector of predictors X and the d -dimensional *column* vector of parameters β



The likelihood function. The regression likelihood composes the above into a single formula – the likelihood of Y given X and the parameters. (It is simplest to write this if we instead imagine μ to be the first entry of β . This works out if we add a single 1 as the first entry of X :

For a single sample:
$$P(Y|X, \beta) = \frac{1}{\sqrt{2\pi\sigma^2}} \cdot e^{-\frac{1}{2} \frac{(Y-X\beta)^2}{\sigma^2}}$$
← Squared residual (distance) from regression line

The outcome values are assumed independent of each other (probabilities multiply). So for multiple samples the likelihood is:

For multiple samples:
$$P(Y|X, \beta) = \frac{1}{\sqrt{2\pi\sigma^2}} \cdot e^{-\frac{1}{2} \frac{\sum_n (Y_n - X_n\beta)^2}{\sigma^2}}$$
← "sum of squared errors"
← The exponent is negative. Maximising the likelihood is therefore the same as minimizing the sum of squared errors – it finds the 'best-fitting line'.

Basic linear regression (maximum likelihood) in R:

```
> fit = lm( Y ~ X, data = D )

> coefficients(fit)
(Intercept)          X
0.0007606242 0.3135072376

> logLik(fit)
'log Lik.' -132.981 (df=3)

> residuals(fit)
      1      2      3      4      . . .
-0.6115976 -0.3239313  0.7034511 -0.2934937 . . .

> summary(fit)$coefficients
              Estimate Std. Error  t value    Pr(>|t|)
(Intercept) 0.0007606242 0.09412669  0.008080856 0.9935689071
X            0.3135072376 0.08512788  3.682780013 0.0003778035
```

This turns out to have an analytic solution:

$$\hat{\beta} = (X^t X)^{-1} X^t Y$$
← Maximum likelihood estimate (MLE)

$$\text{variance}(\hat{\beta}) = \sigma^2 (X^t X)^{-1}$$
← Variance of MLE

$$\text{se}(\hat{\beta}_j) = \sqrt{\sigma^2 (X^t X)^{-1}_{jj}}$$
← Standard error of MLE

But what if you want to fit with prior information included? Use **brms** package:

```
> library( brms )

> fit = brm(
  Y ~ X,
  data = data,
  prior = set_prior( "normal(0,1)" )
)

> fit$fit
Inference for Stan model: ca2436c230608c2ca38ebc402110120d.
4 chains, each with iter=2000; warmup=1000; thin=1;
post-warmup draws per chain=1000, total post-warmup draws=4000.

              mean se_mean  sd  2.5%  25%   50%   75%  97.5% n_eff Rhat
b_Intercept  0.25   0.00 0.05  0.16  0.22  0.25  0.28  0.34 3549  1
b_X          -0.05   0.00 0.04 -0.13 -0.08 -0.05 -0.02  0.03 4293  1
sigma        0.45   0.00 0.03  0.39  0.42  0.44  0.47  0.51 3729  1
lp__        -65.24   0.03 1.25 -68.44 -65.82 -64.91 -64.32 -63.81 1972  1
```

Samples were drawn using NUTS(diag_e) at Thu Nov 11 17:56:07 2021. For each parameter, n_{eff} is a crude measure of effective sample size, and R_{hat} is the potential scale reduction factor on split chains (at convergence, $R_{\text{hat}}=1$).