

## 1 What's Wrong with Maximum Likelihood?

Suppose we have a data set  $\mathbf{Y} = \{y_i\}_{i=1}^n$  and a probability density model  $f(\cdot | \boldsymbol{\theta})$  where  $\boldsymbol{\theta}$  is the parameter. If we try to fit model  $f$  with the data  $\mathbf{Y}$  and obtain the estimate of the parameter  $\boldsymbol{\theta}$ ,

$$\hat{\boldsymbol{\theta}}_{\mathbf{Y}} := \arg \max_{\boldsymbol{\theta}} \log f(\mathbf{Y} | \boldsymbol{\theta}). \quad (\text{ML})$$

What are we *actually* doing here? We are supposing that *if*  $\mathbf{Y}$  is generated from a probability density  $f(\cdot | \boldsymbol{\theta}_0)$ , then  $\hat{\boldsymbol{\theta}}_{\mathbf{Y}}$  is a good estimate for  $\boldsymbol{\theta}_0$ . This is extensively argued by Ronald Fisher, the inventor of the Maximum Likelihood method.

Yet, this approach poses an obvious problem: *What if*  $\mathbf{Y}$  follows another distribution with density function  $g(\cdot | \phi_0)$ ? We can, of course, also find the ML estimate for  $\phi_0$ :

$$\hat{\phi}_{\mathbf{Y}} := \arg \max_{\phi} \log g(\mathbf{Y} | \phi).$$

In the spirit of ML, we can compare the two log-likelihoods,

$$\log f(\mathbf{Y} | \hat{\boldsymbol{\theta}}_{\mathbf{Y}}) \quad \text{and} \quad \log g(\mathbf{Y} | \hat{\phi}_{\mathbf{Y}}), \quad (1)$$

and see which is larger. However, this poses another problem: since we only have one observation  $\mathbf{Y}$ , we can find some density function  $h(\cdot | \boldsymbol{\psi})$  tailored to fit the data at hand  $\mathbf{Y}$  very well, producing a high likelihood  $h(\mathbf{Y} | \boldsymbol{\psi}_{\mathbf{Y}})$ , but fails to produce a high likelihood  $h(\mathbf{X} | \boldsymbol{\psi}_{\mathbf{Y}})$  when another data set  $\mathbf{X}$  is presented. This is referred to as the problem of **overfitting**.

Luckily, in describing **overfitting**, we are motivated to do **cross-validation**, i.e., to use another data  $\mathbf{X}$  (independent to  $\mathbf{Y}$  but follows the sample distribution) to evaluate a parameter estimated under data  $\mathbf{Y}$ .

## 2 Deriving AIC

Let's switch back to using  $f(\cdot | \boldsymbol{\theta})$  for our density function. Also let  $\boldsymbol{\theta}$  be a  $k$ -dimensional vector of parameters. Instead of trying to estimate compare the log-likelihood like in (1), we try to estimate the **cross-validated** version

$$\log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{Y}}).$$

That is, after we obtained the estimator  $\hat{\boldsymbol{\theta}}_{\mathbf{Y}}$  using the data set  $\mathbf{Y}$ , we evaluate the likelihood using another data set  $\mathbf{X}$ . However, since we do not have another independent data set  $\mathbf{X}$ , we need to do some approximation.

First, we approximate  $\log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{Y}})$  by using the second-order Taylor expansion

---

\*In this short introduction, I shall ignore some technical regularity conditions for clarity. I also assume the reader is familiar ML estimator, it's asymptotic properties, and Fisher information.

around  $\hat{\boldsymbol{\theta}}_{\mathbf{X}}$ :

$$\begin{aligned}\log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{Y}}) &\approx \log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{X}}) && \text{(0-th order)} \\ &+ (\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \hat{\boldsymbol{\theta}}_{\mathbf{X}})^\top \left[ \frac{\partial \log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{X}})}{\partial \boldsymbol{\theta}} \right] && \text{(first order)} \\ &+ \frac{1}{2}(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \hat{\boldsymbol{\theta}}_{\mathbf{X}})^\top \left[ \frac{\partial^2 \log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{X}})}{\partial \boldsymbol{\theta} \partial \boldsymbol{\theta}^\top} \right] (\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \hat{\boldsymbol{\theta}}_{\mathbf{X}}) && \text{(second order)}\end{aligned}$$

Note that the first-order term (the Jacobian) is exactly zero since  $\hat{\boldsymbol{\theta}}_{\mathbf{X}}$  is the **ML** estimator. Thus, we have

$$\begin{aligned}\log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{Y}}) &\approx \log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{X}}) \\ &+ \frac{1}{2}(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \hat{\boldsymbol{\theta}}_{\mathbf{X}})^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \hat{\boldsymbol{\theta}}_{\mathbf{X}})\end{aligned}$$

where

$$\mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}}) = \frac{\partial^2 \log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{X}})}{\partial \boldsymbol{\theta} \partial \boldsymbol{\theta}^\top}.$$

This is the key insight of **AIC**: we can obtain the **cross-validated** log-likelihood by making a “correction” to the estimated likelihood  $f(\mathbf{Y} | \hat{\boldsymbol{\theta}}_{\mathbf{Y}})$ . Now we split the correction term into three parts:

$$(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \hat{\boldsymbol{\theta}}_{\mathbf{X}})^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \hat{\boldsymbol{\theta}}_{\mathbf{X}}) = (\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0)^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0) \quad (\text{a})$$

$$+ (\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0)^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0) \quad (\text{b})$$

$$- 2(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0)^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0) \quad (\text{c})$$

We can easily see that part (c) goes to zero asymptotically ( $n \rightarrow \infty$ ):

$$(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0)^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0) = \underbrace{(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0)^\top}_{\xrightarrow{p} 0} \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}}) \underbrace{(\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0)}_{\xrightarrow{p} 0}.$$

Part (a) and (b) are similar in form:

$$\begin{aligned}(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0)^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0) &= \text{trace} \left( n(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0)(\hat{\boldsymbol{\theta}}_{\mathbf{Y}} - \boldsymbol{\theta}_0)^\top \frac{\mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})}{n} \right) \\ (\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0)^\top \mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})(\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0) &= \text{trace} \left( n(\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0)(\hat{\boldsymbol{\theta}}_{\mathbf{X}} - \boldsymbol{\theta}_0)^\top \frac{\mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})}{n} \right).\end{aligned}$$

Since  $\hat{\boldsymbol{\theta}}_{\mathbf{X}}$  and  $\hat{\boldsymbol{\theta}}_{\mathbf{Y}}$  are both **ML** estimators, the expectation of the blue parts is approximately the inverse of Fisher information (asymptotic variance). By information equality,  $\mathbf{J}(\hat{\boldsymbol{\theta}}_{\mathbf{X}})/n$  also converges to the negative of Fisher information in probability. Hence, we have part (a) and (b) approximated as the trace of identity matrices of dimension  $k \times k$ . That is, we have both parts approximated as  $-k$ .

Therefore, our approximation for the **cross-validated** log-likelihood is

$$\log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{Y}}) \approx \log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{X}}) - k.$$

This is the famous **AIC**. However, **AIC** is often written as

$$\text{AIC} = 2k - 2 \log f(\mathbf{X} | \hat{\boldsymbol{\theta}}_{\mathbf{X}}). \quad (\text{AIC})$$

This is due to its connect with information theory and **Kullback-Leibler Divergence**.

### 3 AIC's Connection with Kullback-Leibler Divergence

**KL** divergence is an information theoretic measure of the discrepancy between two distributions. It is defined as

$$\text{KL}(p \parallel q) := \int_{\mathcal{X}} \log \left[ \frac{p(x)}{q(x)} \right] p(x) dx$$

where  $p$  and  $q$  are two densities on the same support  $\mathcal{X}$ . The two main properties of **KL** are

1.  $\text{KL}(p \parallel q) \geq 0 \forall p, q$ .
2.  $\text{KL}(p \parallel q) = 0$  iff  $p = q$  (almost everywhere).

That is,  $\text{KL}(p \parallel q)$  is small when  $p$  and  $q$  are similar.

In our case, we want to know the discrepancy between the “true” likelihood function  $f(\cdot | \theta_0)$  and the estimated likelihood function  $f(\cdot | \hat{\theta}_Y)$ . Hence, we wish to choose the model with small discrepancy between the two:

$$\begin{aligned} \text{KL}(f(\cdot | \theta_0) \parallel f(\cdot | \hat{\theta}_Y)) &= \int_{\mathcal{X}} \log \left[ \frac{f(\mathbf{X} | \theta_0)}{f(\mathbf{X} | \hat{\theta}_Y)} \right] f(\mathbf{X} | \theta_0) d\mathbf{X} \\ &= \int_{\mathcal{X}} \log f(\mathbf{X} | \theta_0) f(\mathbf{X} | \theta_0) d\mathbf{X} && \text{(entropy)} \\ &+ \int_{\mathcal{X}} -\log f(\mathbf{X} | \hat{\theta}_Y) f(\mathbf{X} | \theta_0) d\mathbf{X} && \text{(cross-entropy)} \\ &= \text{constant} - \mathbf{E}_{\mathbf{X}} \log f(\mathbf{X} | \hat{\theta}_Y) \end{aligned}$$

Thus, we can view (**AIC**) as an approximation of **cross-entropy**. Measuring the discrepancy between  $f(\cdot | \theta_0)$  and  $f(\cdot | \hat{\theta}_Y)$  makes intuitive sense: the problem of **overfitting** can be understood as a large discrepancy between the “true” likelihood and the “estimated” likelihood. In the original paper (Akaike, 1974), **AIC** is motivated by **KL**. Hence, **AIC** is represented as the *negative* of the **cross-validated** likelihood to match the sign of **cross-entropy**. Thus in practice, we want to select the model with *small AIC*.

### 4 Why Times Two?

If we consider a Gaussian model with  $\theta = (\mu, \sigma^2)$ , the log-likelihood is written as

$$\log f(\mathbf{X} | \theta) = -\frac{n}{2} \log(2\pi) - \frac{n}{2} \log \sigma^2 - \frac{1}{2\sigma^2} \sum_{i=1}^n (x_i - \mu)^2.$$

It is a lot nicer to write  $2 \log f(\mathbf{X} | \theta)$  so we can get rid of those  $\frac{1}{2}$ 's. That's why. ■

#### Acronyms

<b>AIC</b>	Akaike Information Criterion. 1–3
<b>KL</b>	Kullback-Leibler Divergence. 2, 3
<b>ML</b>	Maximum Likelihood. 1, 2

#### References

Akaike, H. (1974). A new look at the statistical model identification. *IEEE Transactions on Automatic Control*, 19(6), 716–723. <https://doi.org/10.1109/TAC.1974.1100705>