# 2 Statistical Inference

As mentioned in the last chapter, in econometrics, we interpret data x as draws of a random variable X from probability distribution  $p(X|\theta)$  indexed by a parameter  $\theta$ . This distribution is usually specified based on economic theory or a hypothesized relationship among different variables included in x. In some cases, we only specify certain aspects of this distribution, e.g. the (conditional) mean.

Suppose we observe a particular realization x. What can we say about  $\theta$ ? This is the problem of statistical inference, discussed in this chapter. The discussion here covers point estimation (Section 2.1) and its evaluation in finite samples (Section 2.2), hypothesis testing (Section 2.3) and coverage sets (Section 2.4), both from a frequentist/classical as well as a Bayesian point of view.

To convey the fundamentals in a simple setting, unless otherwise stated, this chapter assumes that  $\theta$  is a scalar. Furthermore, we assume that we have a data sample  $\{x_i\}_{i=1}^n$  of n observations (realizations)  $x_i$  of the same underlying RV  $X_i$  which are drawn independently from some distribution  $p(x|\theta)$  (i.e. the observations are i.i.d.). Thereby,  $X_i$  (and hence  $x_i$ ) is assumed to be a scalar. We take our observations together into the  $n \times 1$  vector  $x = (x_1, ..., x_n)'$ .

The previous chapter explicitly distinguished the cdf  $F_X(x)$ , the pdf for continuous RVs  $f_X(x)$  and the probability function for discrete RVs  $\mathbb{P}[X=x]$  or  $\mathbb{P}[x]$ . In this and subsequent chapters we will predominantly deal with continuous random variables and write p(x) for the pdf (and F(x) or  $\mathbb{P}[X \leq x]$  for the cdf, if needed).

#### 2.1 Point Estimation

**Definition 22.** A point estimator  $\delta(X)$  is a mapping from the sample space of X to the parameter space  $\Theta \colon \delta : \mathcal{X} \to \Theta$ .

In other words, for any realization of X,  $x \in \mathcal{X}$ , the point estimator gives us a value  $\delta(x) \in \Theta$ . Loosely speaking,  $\delta(x)$  is our "best guess" where  $\theta$  lies given that we observed X = x. Typically, we denote the point estimator as  $\hat{\theta}$ , but it is crucial to understand that the point estimator of a parameter  $\theta$  is a function of the data X, hence the definition uses  $\delta(X)$ , where  $\delta$  stands for "decision". While we think of the point estimator  $\delta(X)$  as a function of the RV X, if we evaluate it at the particular realization x that we observe, we call the result an estimate rather than an estimator.

There are different estimators that exploit different aspects of  $p(x|\theta)$ , the probability distribution where the data  $x = (x_1, ..., x_n)'$  (supposedly) came from. As a simple running example, let our RV  $X_i$  and parameter  $\theta$  be related as follows:  $X_i|\theta \sim N(\theta, 1)$ . Our discussion start with the classical case where  $\theta$  is treated as a fixed but unknown parameter. For this reason, we condition the moments and distributions on  $\theta$ .

### 2.1.1 Frequentist/Classical Inference

The Least Squares (LS) estimator minimizes the sum of squared deviations of  $x_i$  from their (conditional) mean  $\mathbb{E}[X_i|\theta] = \theta$ :

$$\hat{\theta}_{LS} = \arg\min_{\theta \in \Theta} \sum_{i=1}^{n} (x_i - \theta)^2.$$

In our simple case, this leads to  $\hat{\theta}_{LS} = \frac{1}{n} \sum_{i=1}^{n} x_i$ . Note that the LS estimator only requires assuming  $\mathbb{E}[X_i|\theta] = \theta$ , i.e. we do not need to know the full distribution of  $X_i|\theta$ .

The Method of Moments (MM) estimator solves for  $\hat{\theta}_{MM}$  by setting empirical moments of  $\{x_i\}_{i=1:n}$  equal to the corresponding population moments in  $p(x|\theta)$ . For example, using the mean, we get the same result as under LS, i.e.

$$\mathbb{E}[X_i|\hat{\theta}_{MM}] = \frac{1}{n} \sum_{i=1}^n x_i \quad \Rightarrow \quad \hat{\theta}_{MM} = \frac{1}{n} \sum_{i=1}^n x_i .$$

<sup>&</sup>lt;sup>1</sup>In other words, the point estimator is a rule how we form our best guess based on a RV X, whereas the point estimate is the best guess we obtain given a particular observation x of X.

Under  $\mathbb{V}[X|\theta] = 1$ , we also have  $\mathbb{E}[X_i^2|\hat{\theta}] = \mathbb{V}[X_i|\theta] + \mathbb{E}[X_i|\hat{\theta}] = 1 + \theta^2$ , and we can define  $\hat{\theta}_{MM}$  also as the quantity that equalizes the second moment:

$$\mathbb{E}[X_i^2 | \hat{\theta}_{MM}] = \frac{1}{n} \sum_{i=1}^n x_i^2 \quad \Rightarrow \quad \hat{\theta}_{MM} = \sqrt{\frac{1}{n} \sum_{i=1}^n x_i^2 - 1} .$$

Similarly to LS, the MM estimator does not require the researcher to specify the whole distribution of  $X_i|\theta$ , but only a couple of moments.<sup>2</sup>

The Maximum Likelihood (ML) estimator maximizes the likelihood of observing our particular sample  $x = (x_1, ..., x_n)'$  out of all the possible draws of  $X = (X_1, ..., X_n)'$ :

$$\hat{\theta}_{ML} = \arg\max_{\theta \in \Theta} \mathcal{L}(\theta|x) = \arg\max_{\theta \in \Theta} \ell(\theta|x)$$
,

where  $\mathcal{L}(\theta|X) = p(X|\theta)$  is the pdf of the RV  $X|\theta$  and  $\ell(\theta|X) = \log \mathcal{L}(\theta|X)$  is its log. In our example, under  $X_i|\theta \stackrel{i.i.d.}{\sim} N(0,1)$ , we get

$$\mathcal{L}(\theta|x) = p(x|\theta) = \prod_{i=1}^{n} p(x_i|\theta) = \prod_{i=1}^{n} (2\pi)^{-\frac{1}{2}} exp\left\{\frac{1}{2} \sum_{i=1}^{n} (x_i - \theta)^2\right\},\,$$

because the observations  $x_i$  are independent. In turn, this leads to

$$\ell(\theta|x) = \frac{n}{2}\ln(2\pi) + \frac{1}{2}\sum_{i=1}^{n}(x_i - \theta)^2,$$

which, once again, gives  $\hat{\theta}_{ML} = \frac{1}{n} \sum_{i=1}^{n} x_i$ .

This notation shows that under ML estimation, the pdf  $p(x|\theta)$ , a function of data/realizations x given the parameter  $\theta$ , is interpreted as a function of  $\theta$  given x, which allows us to find an optimal  $\theta$  – the one that maximizes the likelihood – given the observed data x. A useful property of the ML estimator is invariance: if  $\hat{\theta}$  is the ML estimator of  $\theta$ , then  $f(\hat{\theta})$  is the ML estimator of  $f(\theta)$ .

Finite Sample Properties We can analyze the properties of a point estimator  $\hat{\theta}$  for a given sample size n. Recall that  $\hat{\theta}$  is a function the data X, e.g. we obtained  $\hat{\theta} = \frac{1}{n} \sum_{i=1}^{n} X_i$  above. As a result, we can compute the moments of  $\hat{\theta}$  based on the moments of  $X_i$ , while knowing the whole distribution of  $X_i$  allows us to find the distribution of  $\hat{\theta}$ .

<sup>&</sup>lt;sup>2</sup>We need to specify enough moments so that we can solve for  $\theta$ . If  $\theta$  is a scalar, as here, then one (scalar) moment suffices. If  $\theta$  were two dimensional (and  $X_i$  still a scalar), then we would need to specify two (scalar) moments, e.g.  $\mathbb{E}[X_i|\theta]$  and  $\mathbb{V}[X_i|\theta]$ .

**Definition 23.** A point estimator  $\delta(X)$  of  $\theta$  is <u>unbiased</u> if  $\mathbb{E}[\delta(X)|\theta] = \theta$ .

For example,  $\hat{\theta} = \frac{1}{n} \sum_{i=1}^{n} X_i$  is unbiased:

$$\mathbb{E}[\hat{\theta}|\theta] = \mathbb{E}\left[\frac{1}{n}\sum_{i=1}^{n}X_i\right] = \frac{1}{n}\sum_{i=1}^{n}\mathbb{E}[X_i|\theta] = \frac{1}{n}\sum_{i=1}^{n}\theta = \theta.$$

In contrast, the estimator  $\hat{\theta}_* = \frac{1}{n-1} \sum_{i=1}^n X_i$  is not unbiased:

$$\mathbb{E}[\hat{\theta}_*|\theta] = \mathbb{E}\left[\frac{1}{n-1}\sum_{i=1}^n X_i\right] = \mathbb{E}\left[\frac{n}{n-1}\hat{\theta}\right] = \frac{n}{n-1}\theta \neq \theta.$$

We can also compute

$$\mathbb{V}[\hat{\theta}|\theta] = \mathbb{V}\left[\frac{1}{n}\sum_{i=1}^{n}X_{i}\right] = \frac{1}{n^{2}}\sum_{i=1}^{n}\mathbb{V}[X_{i}|\theta] = \frac{1}{n^{2}}\sum_{i=1}^{n}1 = \frac{1}{n}$$

and

$$\mathbb{V}[\hat{\theta}_*|\theta] = \mathbb{V}\left[\frac{n}{n-1}\hat{\theta}\right] = \frac{n^2}{(n-1)^2}\mathbb{V}\left[\hat{\theta}\right] = \frac{n}{(n-1)^2} \ .$$

The quantity  $\mathbb{E}[\hat{\theta}|\theta]$  tells us the average value we expect to get if we were to randomly draw different samples  $x = (x_1, ..., x_n)'$  of size n and compute the point estimator  $\hat{\theta} = \delta(x)$  for these samples. The quantity  $\mathbb{V}[\hat{\theta}|\theta]$  tells us the dispersion we would get in all of these point estimators. Note that to compute  $\mathbb{E}[\hat{\theta}|\theta]$  we had to know  $\mathbb{E}[X_i|\theta]$ , while to compute  $\mathbb{V}[\hat{\theta}|\theta]$ , we also had to specify  $\mathbb{V}[X_i|\theta]$ .

In our example, it turns out that we not only know the mean and variance of  $\hat{\theta}$ , but its whole distribution. Provided that we  $X_i|\theta$  is Normally distributed, we know that  $\hat{\theta}$  is itself Normally distributed, because it is the average of Normally distributed RVs. We get

$$\hat{\theta} = \frac{1}{n} \sum_{i=1}^{n} x_i \sim N(\theta, n^{-1}).$$

**Asymptotic Properties** We can also analyze the properties of a point estimator  $\hat{\theta}$  as the sample size  $n \to \infty$ . This gives us a sense of what we get under large samples.

**Definition 24.** A point estimator  $\delta(X)$  of  $\theta$  is <u>consistent</u> if  $\delta(X) \stackrel{p}{\to} \theta$ .

Recall the definition of convergence in probability, applied to this case:  $\forall \epsilon > 0$ ,  $\lim_{n \to \infty} \mathbb{P}[|\delta(X) - \theta| > \epsilon] = 0$ , i.e. the probability of obtaining a point estimate  $\delta(x)$  that is further away from

 $\theta$  than the distance  $\epsilon$  converges to zero, no matter how small we make  $\epsilon$ . In other words, consistency tells us that our point estimator "gets it right" in the limit, for a large sample. While we sometimes do not have – or do not want (see Section 2.2) – an unbiased estimator, we typically require an estimator to consistent.

We know that  $\hat{\theta} = \frac{1}{n} \sum_{i=1}^{n} X_i$  is consistent, because  $\frac{1}{n} \sum_{i=1}^{n} X_i \xrightarrow{p} \mathbb{E}[X_i | \theta] = \theta$  by the WLLN. The estimator  $\hat{\theta}_* = \frac{1}{n-1} \sum_{i=1}^{n} X_i$  is also consistent:

$$\hat{\theta}_* = \frac{1}{n-1} \sum_{i=1}^n X_i = \frac{n}{n-1} \hat{\theta} \stackrel{p}{\to} \theta ,$$

because  $\hat{\theta} \stackrel{p}{\to} \theta$  and  $\lim_{n\to\infty} \frac{n}{n-1} = 1$ , and the Slutsky theorem tells us that we can form the limit of their product as the product of the limits.

Based on the CLT, we also know the asymptotic distribution of  $\hat{\theta} = \frac{1}{n} \sum_{i=1}^{n} X_i$ :

$$\sqrt{n}\left(\hat{\theta}-\theta\right)|\theta\stackrel{d}{\to}N(0,1)$$
,

because  $V[X_i|\theta] = 1$ . While for the finite sample distribution we had to know the distribution of the data  $X_i|\theta$ , we found this asymptotic distribution under minimal assumptions: independence and identical distribution of our observations  $\{x_i\}_{i=1}^n$ . Therefore, if we do not know the distribution of the data, we can rely on the CLT to obtain the asymptotic distribution of our estimator. In turn, we can approximate the finite sample distribution based on the asymptotic one, by saying that

$$\sqrt{n} \left( \hat{\theta} - \theta \right) | \theta \stackrel{approx.}{\sim} N(0, 1) \quad \text{or, equivalently,} \quad \hat{\theta} | \theta \stackrel{approx.}{\sim} N(\theta, n^{-1}) ,$$

already for our finite sample of size n.<sup>3</sup>

## 2.1.2 Bayesian Inference

The LS, MM and ML estimators all belong to the classical or frequentist paradigm, under which  $\theta$  is thought to be an unknown, but fixed parameter. As opposed to that, in the Bayesian approach,  $\theta$  is treated as a RV. Given some initial belief about its distribution (or about where the fixed, true value of  $\theta$  lies) – the prior distribution – this belief is updated

<sup>&</sup>lt;sup>3</sup>In our particular example with  $X_i|\theta \sim N(\theta,1)$ , by approximating the finite sample distribution via the asymptotic one, we get it exactly right. However, this does not need to be the case. For example, let  $X_i|\theta \sim Exp(\lambda)$ , with  $\mathbb{E}[X_i|\lambda] = \lambda^{-1}$  and  $\mathbb{V}[X_i|\lambda] = \lambda^{-2}$ , and suppose that – for some reason – we form the estimator  $\hat{\lambda} = \frac{1}{n} \sum_{i=1}^{n} X_i$ . Then  $\hat{\lambda} = \frac{1}{n} \sum_{i=1}^{n} X_i \sim Gamma(n, n\lambda)$ , while  $\sqrt{n}\lambda \left(\hat{\lambda} - \lambda^{-1}\right)|\lambda \stackrel{d}{\to} N(0, 1)$ , i.e.  $\hat{\lambda}|\lambda \stackrel{approx}{\sim} N(\lambda^{-1}, \lambda^{-2}n^{-1})$ .

after observing the sample x to the posterior distribution (the distribution of  $\theta$  or the belief about where the true value of  $\theta$  lies after having observed the data) using Bayes' formula:

$$p(\theta|x) = \frac{p(x|\theta)p(\theta)}{p(x)} \propto p(x|\theta)p(\theta)$$
.

Bayesian inference often exploits the fact that the posterior is proportional to the product of the likelihood (i.e. the conditional distribution of x given  $\theta$ ) and the prior, which means that it (often) suffices to analyze the product of likelihood and prior in order to conclude something about the posterior (see Section 1.2). The term  $p(x) = \int p(x|\theta)p(\theta)d\theta$  is just a normalization constant.

In our example, let our prior be  $\theta \sim N(0, 1/\lambda)$ , where  $\lambda$  is the inverse of the prior variance and specifies how confident we are in our belief that  $\theta$  is equal to the prior mean of zero. Under  $X_i|\theta \stackrel{i.i.d.}{\sim} N(0,1)$ , we then get

$$p(\theta|x) \propto \exp\left\{-\frac{1}{2}\sum_{i=1}^{n}(x_i - \theta)^2\right\} \exp\left\{-\frac{1}{2}\lambda\theta^2\right\}$$
$$\propto \exp\left\{-\frac{1}{2}(-2\theta\sum_{i=1}^{n}x_i + \theta^2(n+\lambda))\right\},$$

whereby we used the expression for  $p(x|\theta)$  derived above. The above mentioned proportionality implies that we can drop all terms which do not depend on  $\theta$  (i.e. all constants and all terms that depend only on x). From this expression, we can deduce that  $\theta|x \sim N(\bar{\theta}, \bar{V})$  with  $\bar{V} = \frac{1}{n+\lambda}$  and  $\bar{\theta} = \frac{1}{n+\lambda} \sum_{i=1}^{n} x_i$ .

Once the posterior  $p(\theta|x)$  is obtained, there are several Bayesian point estimators one might construct (see next section). The most common ones are the posterior mean or median. In our example, both are equal to  $\bar{\theta} = \frac{1}{n+\lambda} \sum_{i=1}^{n} x_i$ . Note that this is a weighted average of the ML estimator and the prior mean, with weights given by the inverses of the ML estimator and prior variances, respectively:

$$\mathbb{E}[\theta|x] = \frac{\sum_{i=1}^{n} x_i}{n+\lambda} = \frac{1}{1/\mathbb{V}[\hat{\theta}_{ML}] + \lambda} \left[ \frac{1}{\mathbb{V}[\hat{\theta}_{ML}]} \cdot \hat{\theta}_{ML} + \lambda \cdot 0 \right] ,$$

whereby  $\mathbb{V}[\hat{\theta}_{ML}] = n^{-1}$ . If  $\hat{\theta}_{ML}$  is precise (which happens if n is large, i.e. we have lots of data to draw inference from), the posterior mean will be closer to the ML estimator. The same happens if  $\lambda$  is low, which means that we are not very confident in our prior belief about  $\theta$ . In contrast, as  $\lambda \to \infty$ , the posterior (mean) comes ever closer to the prior (mean).

Frequentist Properties of Bayesian Estimators Despite the fact that the posterior mean  $\bar{\theta}$  is obtained in the Bayesian paradigm, we could analyze the properties of the point estimator  $\hat{\theta}_B = \bar{\theta} = \frac{1}{n+\lambda} \sum_{i=1}^n X_i$  just like the finite sample and asymptotic properties of the classical/frequentist estimators above. Note that this exercise is not a part of Bayesian inference, as it involves thinking about the properties of our estimator under different randomly drawn samples, whereas Bayesian inference conditions on the particular sample x obtained. However, estimators derived under the Bayesian paradigm are often used in classical/frequentist inference. For our point estimator  $\hat{\theta}_B$ , we can write  $\hat{\theta}_B = \frac{n}{n+\lambda}\hat{\theta}_{ML}$  for  $\hat{\theta}_{ML} = \frac{1}{n}\sum_{i=1}^n X_i$ . In turn, we can see that  $\hat{\theta}_B$  is biased,

$$\mathbb{E}[\hat{\theta}_B] = \frac{n}{n+\lambda} \mathbb{E}\left[\hat{\theta}_{ML}\right] = \frac{n}{n+\lambda} \theta \neq \theta ,$$

but consistent,

$$\hat{\theta}_B = \frac{n}{n+\lambda} \hat{\theta}_{ML} \stackrel{p}{\to} \theta ,$$

because  $\hat{\theta}_{ML} \xrightarrow{p} \theta$  and  $\lim_{n\to\infty} \frac{n}{n+\lambda} = 1$ , and we can combine these two results using Slutsky's theorem. Also, we can show that

$$\mathbb{V}\left[\hat{\theta}_{B}\right] = \frac{n^{2}}{(n+\lambda)^{2}} \mathbb{V}[\hat{\theta}_{ML}] < \mathbb{V}[\hat{\theta}_{ML}] \quad \forall \quad \lambda > 0.$$

Compared to  $\hat{\theta}_{ML}$ ,  $\hat{\theta}_{B}$  introduces a bias but reduces the variance. The intuition is that  $\hat{\theta}_{B}$  is not based on the data alone, but shrinks  $\hat{\theta}_{ML}$  to the prior mean of zero.

# 2.2 Evaluation of Estimators

In finite samples, point estimators  $\delta$  are evaluated using a loss function  $L(\theta, \delta)$ . Common choices are the quadratic/L2 loss function  $L(\theta, \delta) = (\theta - \delta)^2$  or L1 loss function  $L(\theta, \delta) = |\theta - \delta|$ . Based on  $L(\theta, \delta)$ , we can define frequentist risk, Bayesian risk and integrated risk.

**Frequentist Risk** takes the expectation of the loss function treating X as a RV and  $\theta$  as fixed:

$$R(\theta, \delta) = \mathbb{E}[L(\theta, \delta(X))|\theta] = \int_x L(\theta, \delta(X))p(x|\theta)dx$$
.

This means that it determines the behavior of an estimator conditional on a true  $\theta$  and under the assumption that nature provides us repeatedly with draws from X. For example, if  $X|\theta \sim N(\theta, 1)$ , we use a quadratic loss function  $L = (\theta - \delta)^2$ , and  $\delta$  is of the form  $\delta = c \cdot X$ ,

we get

$$R(\theta, \delta) = \mathbb{E}[(\theta - cX)^2 | \theta] = \theta^2 - 2c\theta \mathbb{E}[X|\theta] + c^2 \mathbb{E}[X^2 | \theta] = \theta^2 (1 - c)^2 + c^2.$$

Under the quadratic loss function, we can decompose frequentist risk into a bias- and a variance-term. If  $\theta$  is a scalar, we have  $L(\theta, \delta) = (\theta - \delta)^2$  and

$$\begin{split} R(\theta, \delta) &= \mathbb{E}[(\theta - \delta)^2 | \theta] \\ &= \theta^2 - 2\theta \mathbb{E}[\delta | \theta] + \mathbb{E}[\delta^2 | \theta] \\ &= (\theta - \mathbb{E}[\delta[\theta])^2 + \mathbb{E}[\delta^2[\theta] - \mathbb{E}[\delta | \theta]^2 \\ &= (\theta - \mathbb{E}[\delta[\theta])^2 + \mathbb{V}[\delta | \theta] \;, \end{split}$$

i.e.  $R = \text{bias}^2 + \text{variance}$ . If  $\theta$  is a k-dimensional vector, let W be any positive-defiite weighting matrix and define  $L(\theta, \delta) = (\theta - \delta)'W(\theta - \delta)$ . We get

$$R(\theta, \delta) = \mathbb{E}[(\theta - \delta)'W(\theta - \delta)|\theta]$$

$$= \mathbb{E}[\operatorname{tr}(W(\theta - \delta)(\theta - \delta)')|\theta]$$

$$= \mathbb{E}[\operatorname{tr}(W(\theta - \mathbb{E}[\delta|\theta] + \mathbb{E}[\delta|\theta] - \delta)(\theta - \mathbb{E}[\delta|\theta] + \mathbb{E}[\delta|\theta] - \delta)')|\theta]$$

$$= \operatorname{tr}[W((\theta - \mathbb{E}[\delta|\theta])(\theta - \mathbb{E}[\delta|\theta])' + \mathbb{V}[\delta|\theta])].$$

This shows that unbiasedness is not necessarily a desirable property for a point estimator in the sense that there might be biased but less noisy estimators that yield a lower frequentist risk.

Because in general, the risk function depends on  $\theta$ , which is unknown, there is no unique ranking of estimators independent of  $\theta$ . Possible rankings are based on the minimax risk or inadmissibility. These concepts are discussed in the Appendix.

Bayesian Risk (Expected Posterior Loss) takes the expectation of the loss function treating  $\theta$  as a RV and X = x as fixed:

$$P(\mathbb{P}^{\theta}, \delta(x)) = \mathbb{E}[L(\theta, \delta(x))|x] = \int_{\Theta} L(\theta, \delta(x))p(\theta|x)d\theta$$
.

It does the opposite of frequentist risk; it treats data X (the particular realization x we observe) as fixed and averages the loss function over all possible values of  $\theta$  using the posterior distribution. As opposed to frequentist risk, Bayesian risk does not depend on  $\theta$ , but on x, the realization of X, which we know.  $\mathbb{P}^{\theta}$  denotes the prior distribution  $p(\theta)$ , but is written

differently to avoid the wrong conclusion that P depends on  $\theta$ .

**Definition 25.** A <u>Bayesian estimator</u> (or Bayes estimator) associated with a prior distribution  $\mathbb{P}^{\theta}$  and a loss function  $L(\theta, \delta(X))$  is an estimator that minimizes  $P(\mathbb{P}^{\theta}, \delta(x))$ .

Under the quadratic loss function, the Bayes estimator is given by the posterior mean:

$$\min_{\delta \in \mathcal{D}} \mathbb{E}[(\theta - \delta(x))^2 | x] \quad \Rightarrow \quad 2\mathbb{E}[\theta - \delta(x) | x] = 0 \quad \Rightarrow \quad \delta(x) = \mathbb{E}[\theta | x] .^4$$

Other loss functions can lead to other posterior moments as the Bayes estimator, e.g. the posterior median.

**Integrated Risk** computes the mean of the loss function using the joint distribution of  $\theta$  and X, i.e. it averages over all (a priori) possible parameter values as well as all possible realizations of X:

$$r(\mathbb{P}^{\theta}, \delta) = \mathbb{E}[L(\theta, \delta(X))] = \int_{\mathcal{X}} \int_{\Theta} L(\theta, \delta(x)) p(x, \theta) d\theta dx$$
$$= \int_{\Theta} R(\theta, \delta) p(\theta) d\theta$$
$$= \int_{\mathcal{X}} P(\mathbb{P}^{\theta}, \delta(x)) p(x) dx.$$

It gives a real number and therefore a total ordering of estimators. Since p(x) is hard to obtain, it is easiest to compute integrated risk as  $\mathbb{E}[R(\theta, \delta)]$ , i.e. by first taking the conditional expectation using  $p(X|\theta)$  and then the expectation using  $p(\theta)$ . This calculation shows that minimizing the integrated risk means minimizing expected frequentist risk, whereby the expectation is taken using the prior distribution for  $\theta$ .

## 2.3 Hypothesis Testing

Often, we are not interested in obtaining an estimator for  $\theta$  per se, but in testing whether  $\theta$  lies in some set (interval)  $\Theta_0 \subset \Theta$ . More formally, we test the zero hypothesis  $\mathcal{H}_0 = \theta \in \Theta_0$  against the alternative hypothesis  $\mathcal{H}_1 : \theta \in \Theta_1$ . Often,  $\Theta_1$  is simply the complement of  $\Theta_0$ , i.e. all other regions of the parameter space:  $\Theta_1 = \Theta_0^c = \Theta \setminus \Theta_0$ . If  $\Theta_0$  contains a single value  $\theta_0$ , then  $\mathcal{H}_0 : \theta = \theta_0$  is called a point hypothesis, otherwise it is called a composite hypothesis (and analogously for  $\mathcal{H}_1$ ).

<sup>&</sup>lt;sup>4</sup>Note that  $\mathbb{E}[(\theta - \delta(x))^2 | x]$  must be bounded for this exchange of integration and differentiation to hold, i.e. the posterior variance must be bounded (see result in Appendix to Chapter 1).

**Definition 26.** A <u>hypothesis test</u>  $\varphi \in \{0,1\}$  is a rule that specifies when we reject and when we accept (do not reject)  $\mathcal{H}_0$ , with  $\varphi = 0$  indicating rejection.

A hypothesis test can be viewed as a point estimator of the object  $\mathbf{1}\{\theta \in \Theta_0\}$ . Testing procedures  $\varphi$  can be evaluated using the following loss function:

$$L(\theta, \varphi) = \begin{cases} 1 & \text{if } \varphi \neq \mathbf{1} \{ \theta \in \Theta_0 \} \\ 0 & \text{otherwise} \end{cases} \text{ (i.e. we are wrong)}$$
 (i.e. we are right)

#### 2.3.1 Testing in the Frequentist Paradigm

This loss function leads to the following frequentist risk  $R(\theta, \varphi) = \mathbb{E}[L(\theta, \varphi(X))|\theta]$  which depends on  $\theta$ :

- if  $\theta \in \Theta_0$ , then  $R(\theta, \varphi) = \mathbb{P}\left[\text{reject } \mathcal{H}_0 \mid \mathcal{H}_0 \text{ is true}\right] \equiv \alpha(\theta)$ ,
- if  $\theta \in \Theta_1$ , then  $R(\theta, \varphi) = \mathbb{P}[\text{accept } \mathcal{H}_0 \mid \mathcal{H}_0 \text{ is false}] \equiv 1 \beta(\theta)$ .

Often,  $\alpha(\theta)$  is referred to as the type I error,  $1 - \beta(\theta)$  as the type II error, and  $\beta(\theta) = \mathbb{P}$  [reject  $\mathcal{H}_0|\mathcal{H}_0$  is false] as the power of the test  $\varphi$ . However, note that  $\alpha(\theta)$  and  $\beta(\theta)$  are in fact the same function:

$$\beta(\theta) = \mathbb{P}\left[\text{reject } \mathcal{H}_0 \mid \theta\right] = \mathbb{P}[\varphi = 0 \mid \theta]$$

is the power function, defined on the whole sample space  $\Theta$ . People just call it type I error and write it as  $\alpha(\theta)$  if  $\theta \in \Theta_0$ , while they write it as  $\beta(\theta)$  if  $\theta \in \Theta_1$  and call  $1 - \beta(\theta)$  type II error. To avoid confusion, we will just write  $\beta(\theta)$  for all  $\theta \in \Theta$ .

A good test has a power function near 1 for  $\theta \in \Theta_1$  and near 0 for  $\theta \in \Theta_0$ . Usually, we put a constraint on the type I error by specifying a size (or level) requirement and then search for the test with the highest power.

**Definition 27.** For  $\alpha \in (0,1)$ , a test is a <u>size  $\alpha$  test</u> if  $\sup_{\theta \in \Theta_0} \beta(\theta) = \alpha$ , i.e. if its type I error is equal to  $\alpha$ .<sup>5</sup>

Note that with a point hypothesis, the above expression simplifies to  $\beta(\theta_0) = \alpha$  simply.

<sup>&</sup>lt;sup>5</sup>For discrete RVs, sometimes we cannot set  $\sup_{\theta \in \Theta_0} \beta(\theta)$  exactly to  $\alpha$ . In these cases, we use the definition of a level  $\alpha$  test; a test is a level  $\alpha$  test if  $\sup_{\theta \in \Theta_0} \beta(\theta) \leqslant \alpha$ , i.e. if the type I error is at most  $\alpha \forall \theta \in \Theta_0$ .

There are different tests that use different test-statistics. They are all constructed based on the idea that an event considered unlikely under  $\mathcal{H}_0$  – such as an extreme realization – discredits  $\mathcal{H}_0$ . They all have the generic form

$$\varphi(x;\alpha) = \mathbf{1} \{ T(x) < c_{\alpha} \} ,$$

i.e. we reject if our realized test-statistic T(x) is larger than some critical value  $c_{\alpha}$ , whereby different choices of the size  $\alpha$  translate into different values for  $c_{\alpha}$ . Under this form, the power function becomes  $\beta(\theta) = \mathbb{P}[T(x) \geq c_{\alpha} \mid \theta]$ . As a first example, we will consider the popular t-test.

**Definition 28.** Suppose  $\hat{\theta}|\theta \sim N(\theta, v^2)$  and we are testing a point hypothesis  $\mathcal{H}_0: \theta = \theta_0$ . Under the alternative  $\mathcal{H}_1: \theta \neq \theta_0$ , the (two-sided) <u>t-test</u> is  $\varphi_t(x) = \mathbf{1}\left\{\left|\frac{\hat{\theta}-\theta_0}{v}\right| < c\right\}$ .

Note that the test-statistic  $T(X) = \left| \frac{\hat{\theta} - \theta_0}{v} \right|$  is a function of the data X because our estimator of  $\theta$ ,  $\hat{\theta}$ , is a function of X. Previously, we said that when we apply our estimator to our particular, observed realization of the data, x, we call the result an estimate. Similarly, we then have the realized test-statistic T(x).

For example, let  $X|\theta \sim N(\theta, 1)$ , and assume for simplicity that we have a single observation x s.t.  $\hat{\theta} = x$  has the distribution  $\hat{\theta}|\theta = X|\theta \sim N(\theta, 1)$ . If  $\mathcal{H}_0$  is true, then this distribution is  $N(\theta_0, 1)$  and therefore  $\frac{\hat{\theta} - \theta_0}{1} \sim N(0, 1)$ . The two-sided t-test rejects  $\mathcal{H}_0$  if this statistic is (on either side) too far from its supposed mean of zero (i.e. if x is too far from its supposed mean  $\theta_0$ ) and accepts otherwise.

Gunder the alternative  $\mathcal{H}_1: \theta > \theta_0$ , the (one-sided) t-test is  $\varphi_t(x) = \mathbf{1}\left\{\frac{\hat{\theta} - \theta_0}{v} > c\right\}$ , while under the alternative  $\mathcal{H}_1: \theta < \theta_0$ , the (one-sided) t-test is  $\varphi_t(x) = \mathbf{1}\left\{\frac{\hat{\theta} - \theta_0}{v} < c\right\}$ .

<sup>&</sup>lt;sup>7</sup>In actual econometric models,  $\hat{\theta}$  would be a more complicated function of the data. However, nothing changes conceptually; for example, under a linear regression model (see Chapter 3), we estimate a k-dimensional vector  $\beta$ , which (under certain assumptions) gives us  $\hat{\beta}|\beta \sim N(\beta, V)$  and  $\hat{\beta}_j|\beta_j \sim N(\beta_j, V_{jj})$  for a single element  $\beta_j$  of  $\beta$ .

To find c, we set the type I error equal to te desired type I error  $\alpha$ :

$$\alpha = \beta(\theta_0)$$

$$= \mathbb{P}[\varphi_t = 0 | \theta = \theta_0]$$

$$= 1 - \mathbb{P}[-c \le \hat{\theta} - \theta_0 \le c | \theta = \theta_0]$$

$$= 1 - \mathbb{P}[-c \le Z \le c | \theta = \theta_0]$$

$$= 1 - [\Phi(c) - \Phi(-c)]$$

$$= 2(1 - \Phi(c)),$$

where  $\Phi(c) = \mathbb{P}[Z \leq c]$  is the cdf of a standard Normal-distributed RV Z, and we used the fact that it is symmetric around zero:  $\Phi(-c) = 1 - \Phi(c)$ . To get a test of size 10% ( $\alpha = 0.1$ ), we take c = 1.64. Different values of c yield t-tests of different sizes  $\alpha$ .

To compute the power of this test, we need to think about what happens if  $\mathcal{H}_0$  is false. If the true value of  $\theta$  is some  $\tilde{\theta}$ , then  $\hat{\theta} \sim N(\tilde{\theta}, 1)$  and so  $\frac{\hat{\theta} - \theta_0}{1} \sim N(\tilde{\theta} - \theta_0, 1)$ . As a result,

$$\begin{split} \beta(\tilde{\theta}) &= \mathbb{P}[\varphi_t = 0 | \theta = \tilde{\theta}] \\ &= 1 - \mathbb{P}[-c \leq \hat{\theta} - \theta_0 \leq c | \theta = \tilde{\theta}] \\ &= 1 - \mathbb{P}[-c - (\tilde{\theta} - \theta_0) \leq Z \leq c - (\tilde{\theta} - \theta_0) | \theta = \tilde{\theta}] \\ &= 1 - [\Phi(c - (\tilde{\theta} - \theta_0)) - \Phi(-c - (\tilde{\theta} - \theta_0))] \;. \end{split}$$

This function is plotted in Fig. 2.1. Different sizes of the test – embodied by different choices for c – lead to different powers of the test. The lower the size of the test – i.e. the smaller the probability of wrongly rejecting  $\mathcal{H}_0$  – the lower also the power of the test – i.e. the higher the probability of wrongly accepting (or failing to reject)  $\mathcal{H}_0$ . It is common to be rather conservative (i.e. erring on the side of not rejecting  $\mathcal{H}_0$ ) and report test results for sizes of 10%, 5% and 1%.

**Definition 29.** The likelihood ratio (LR) test is 
$$\varphi_{LR}(x) = 1 \left\{ \sup_{\theta \in \Theta_1} p(x|\theta) / \sup_{\theta \in \Theta_0} p(x|\theta) < c \right\}$$
.

Intuitively, this ratio is small – and therefore the test is likely to accept  $\mathcal{H}_0$  – if there are points in  $\Theta_0$  for which the observed x is much more likely than for points in  $\Theta_1$ . For a point zero-hypothesis  $\mathcal{H}_0: \theta = \theta_0$  with the alternative  $\mathcal{H}_1: \theta \neq \theta_0$ , the LR test simplifies to

$$\varphi_{LR}(x) = \mathbf{1} \left\{ \frac{p(x|\hat{\theta}_{ML})}{p(x|\theta_0)} < c \right\}.$$

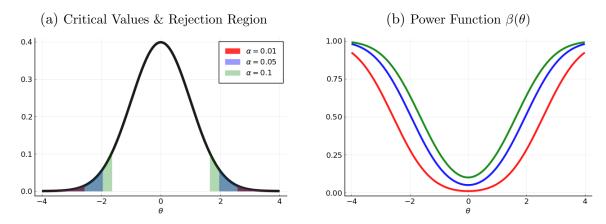


Figure 2.1: Two-Sided t-Tests

Notes: Illustration of the critical values, rejection region and power function for a two-sided t-test with sizes  $\alpha \in \{0.01, 0.05, 0.1\}$ .

Consider our example from before;  $X \sim N(\theta, 1)$ , with single realization x, testing  $\mathcal{H}_0 : \theta = \theta_0$  against  $\mathcal{H}_1 : \theta \neq \theta_0$ . We get

$$T(x) = \frac{p(x|\hat{\theta}_{ML})}{p(x|\theta_0)} = \frac{(2\pi)^{-1/2} exp\left\{-\frac{1}{2}(x-\hat{\theta})^2\right\}}{(2\pi)^{-1/2} exp\left\{-\frac{1}{2}(x-\theta)^2\right\}} = exp\left\{\frac{1}{2}(\hat{\theta}-\theta_0)^2\right\},\,$$

as  $\hat{\theta} = x$ . Instead of working with this test-statistic directly, we can equivalently define it as  $\tilde{T}(x) = 2 \cdot \ln \left( \frac{p(x|\hat{\theta}_{ML})}{p(x|\theta_0)} \right) = (x - \theta_0)^2$  and write our LR test as  $\varphi_{LR}(x) = \mathbf{1} \left\{ \tilde{T}(x) < \tilde{c} \right\}$  with  $\tilde{c} = 2ln(c)$ . In order to get a size  $\alpha$  test, we then find  $\tilde{c}$  so as to set the type I error to  $\alpha$ :

$$\mathbb{P}\left[\tilde{T}(X) \ge \tilde{c} \mid \mathcal{H}_0\right] = \alpha .$$

If  $\mathcal{H}_0$  is true,  $X - \theta_0 \sim N(0, 1)$  and so  $(X - \theta_0)^2 \sim \chi_1^2$  (Chi-squared distribution with one degree of freedom). Therefore,  $\tilde{c}$  is a critical value from that distribution.

**Definition 30.** A test  $\varphi_{\alpha}$  with size  $\alpha$  and power function  $\beta(\theta)$  is a <u>uniformly most powerful</u> size  $\alpha$  test if it maximizes the power uniformly on  $\Theta_1$  among all tests with size  $\alpha$ , i.e. if  $\beta(\theta) \geqslant \beta'(\theta)$  for all  $\theta \in \Theta_1$  and for all power functions  $\beta'(\theta)$  of size  $\alpha$  tests  $\varphi'$ .

The Neyman-Pearson Lemma in the Appendix shows that if both  $\mathcal{H}_0$  and  $\mathcal{H}_1$  are point hypotheses, the LR test is uniformly most powerful.

The above discussion illustrates that to conduct a hypothesis test, we first choose a "test-type" like a t-test or an LR-test, we then find the right critical value for our desired size of the test, and finally we can apply the test to our particular observation. Thereby, finding

<sup>&</sup>lt;sup>8</sup>This presumes that  $\hat{\theta}_{ML} \in \Theta_1$ , i.e.  $\hat{\theta}_{ML} \neq \theta_0$ . Under continuous parameter spaces like  $\mathbb{R}$ , the probability that  $\hat{\theta}_{ML}$  is exactly equal to  $\theta_0$  is zero.

the critical value requires us to know the distribution of our test statistic T(X) under  $\mathcal{H}_0$ , because we need to know the probability of rejection under  $\mathcal{H}_0 - \mathbb{P}[T(X) > c | \mathcal{H}_0]$  – equal to  $\alpha$ . We deduce this distribution based on the distribution of  $X|\theta$  under  $\mathcal{H}_0$ . If we cannot find the distribution of T(x), we can apply a monotonic transformation of both T(x) and c – like we took logs and multiplied by 2 in the previous example – and try to find the distribution of this transformed statistic. Sometimes, however, we cannot proceed analytically. In these cases, we do the equivalent steps numerically: we repeatedly draw  $X|\theta$  from its supposed distribution under  $\mathcal{H}_0$  and simulate the distribution of T(X).

Consider the previous example  $-X|\theta \sim N(\theta, 1)$ , single realization x – but with  $\theta \in \mathbb{R}_+$  restricted to be positive. We get the ML estimator  $\hat{\theta} = \max\{0, x\}$ . We can test  $\mathcal{H}_0 : \theta = \theta_0$  vs.  $\mathcal{H}_1 : \theta \neq \theta_0$  using the LR test with statistic

$$T(x) = 2\left[-\frac{1}{2}(x-\hat{\theta})^2 + \frac{1}{2}(x-\theta_0)^2\right] = (x-\theta_0)^2 - (x-\hat{\theta})^2 = \begin{cases} (x-\tilde{\theta})^2 & \text{if } x > 0\\ \theta_0^2 - 2\theta_0 x & \text{if } x \le 0 \end{cases}.$$

If  $\mathcal{H}_0$  is true, we know  $X|\theta \sim N(\theta_0, 1)$ , but it is very hard to compute the resulting distribution of the test statistic T(X). Instead, we can conduct our hypothesis test numerically as follows.

#### **Algorithm 1** (Numerical Hypothesis Testing).

- 1. For m = 1 : M, draw  $x^m$  from  $N(\theta_0, 1)$ , the distribution of X under  $\mathcal{H}_0$ . For each draw, compute  $T(x^m)$ . This gives you a distribution  $\{T(x^m)\}_{m=1}^M$  that approximates the true distribution of T(X) under  $\mathcal{H}_0$ .
- 2. Sort  $\{T(x^m)\}_{m=1}^M$  in ascending order and take the  $M(1-\alpha)$ th value as  $c_\alpha$ . This empirical quantile approximates the  $100(1-\alpha)$ th quantile of the distribution of T(X) under  $\mathcal{H}_0$ .
- 3. Compute T(x) for your particular, observed realization x. If  $T(x) \leq c_{\alpha}$ , then we accept, otherwise we reject.

**Definition 31.** A <u>p-value</u> is the largest type I error (i.e. largest size, probability of wrong rejection) of a test at which  $\mathcal{H}_0$  is accepted (cannot be rejected).

More formally, take a test  $\varphi(x;\alpha)$  based on a test statistic T(x) and let  $\alpha$  be its size. Then  $p\text{-value} = \sup \alpha \ s.t. \ \varphi(x;\alpha) = 1$ , i.e. we accept  $\mathcal{H}_0$ .

Recall the general structure of a test:  $\varphi(x;\alpha) = \mathbf{1}\{T(x) < c_{\alpha}\}$ , where we now emphasize the dependence of our test  $\varphi$  on the size  $\alpha$ . Our examples illustrated that the size of a test is

embodied in the critical value c based on which we judge our test-statistic T(x), with larger sizes corresponding to smaller critical values. Of course, if we are conservative and take a very small size, i.e. small probability of wrong rejection, we have a large critical value, which means that we are likely to accept  $\mathcal{H}_0$ . In fact, as  $\alpha \to 0$ , we have  $c \to \infty$ , and we always accept! You can think of the p-value as the size obtained when we take the smallest c s.t. we accept  $\mathcal{H}_0$ . If the p-value is large, this means that we can even take a small c and still accept  $\mathcal{H}_0$ , which means that it is likely to be true. In contrast, if the p-value is small, this means that we indeed need a very large critical value to accept  $\mathcal{H}_0$ , which means that  $\mathcal{H}_0$  is likely to be wrong (essentially we only accept it if we are very conservative, specifying a very small probability of wrong rejection). See illustration in Fig. 2.2.

Consider the two-sided t-test example from above:  $X \sim N(\theta, 1)$ , and we test  $\mathcal{H}_0 : \theta = \theta_0$  vs.  $\mathcal{H}_1 : \theta \neq \theta_0$  using the test  $\varphi(X; \alpha) = \mathbf{1} \{ T(X) < c_{\alpha} \}$  based on the statistic  $T(X) = \left| \hat{\theta} - \theta_0 \right|$ , with  $\hat{\theta} = X$ . Then

$$p = \sup_{\alpha} \mathbb{P} \left[ \varphi(X; \alpha) = 0 | \mathcal{H}_0 \right]$$

$$= \sup_{\alpha} \mathbb{P} \left[ |X - \theta_0| > c_{\alpha} | \mathcal{H}_0 \right]$$

$$= \sup_{\alpha} 2 \cdot \mathbb{P} \left[ X - \theta_0 > c_{\alpha} | \mathcal{H}_0 \right]$$

$$= \sup_{\alpha} 2(1 - \Phi(c_{\alpha})).$$

This expression decreases with  $c_{\alpha}$ . Therefore, the smallest  $c_{\alpha}$  we can take s.t. we still accept  $\mathcal{H}_0$  is  $c_{\alpha} = T(x) = |x - \theta_0|$ . We obtain  $p = 2(1 - \Phi(|x - \theta_0|))$ .

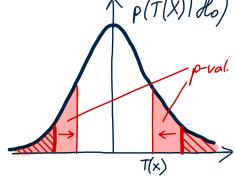
The p-value reports the results of a test on a continuous (0,1)-scale rather than a discrete  $\{0,1\}$ -scale, i.e. accept/reject. It can be (and often is) used as a decision rule; we reject if the p-value is smaller than a desired size  $\alpha$ , and accept otherwise.

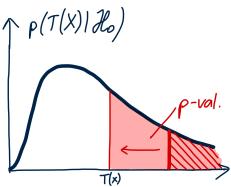
### 2.3.2 Testing in the Bayesian Paradigm

As mentioned at the beginning of this section, a hypothesis test can be viewed as a point estimator of the object  $\mathbf{1} \{ \theta \in \Theta_0 \}$ . The frequentist approach considers  $\theta$  to be a fixed number. Therefore,  $\mathbf{1} \{ \theta \in \Theta_0 \}$  is an unknown but fixed parameter. In other words, a hypothesis  $\mathcal{H}_0 : \theta \in \Theta_0$  can either be true or false;  $\theta$  either is in  $\Theta_0$  or it is not. In contrast, under the Bayesian approach,  $\theta$  is a RV and we obtain probabilities for a hypothesis  $\mathcal{H}_0$  (and  $\mathcal{H}_1$ ) being true or false. The prior probability of  $\mathcal{H}_0$  being true is  $\mathbb{P}[\theta \in \Theta_0]$ , while the

#### (a) Two-sided test with $T(X) \in \mathbb{R}$







(b) Test with  $T(X) \in \mathbb{R}_{++}$ 

Figure 2.2: p-Values

Notes: Illustration of the critical values with corresponding rejection region (thick red line) and p-values (red area) for two different tests; left the two-sided t.test and right a test with positive support of the test-statistic (e.g. the LR test in the example in the text, where  $T(X) \sim \chi^2$  under  $\mathcal{H}_0$ ).

posterior probability is  $\mathbb{P}[\theta \in \Theta_0|X]$ . Hypotheses are then evaluated using

posterior odds 
$$=\frac{p(\theta \in \Theta_0|X)}{p(\theta \in \Theta_1|X)}$$
,

or

Bayes factors = 
$$\frac{\text{posterior odds}}{\text{prior odds}} = \frac{p(\theta \in \Theta_0|X)/p(\theta \in \Theta_1|X)}{p(\theta \in \Theta_0)/p(\theta \in \Theta_1)}$$
,

The difference in testing philosophy between the two approaches is best illustrated using a simple example. Let  $\Theta = \{0, 1\}$  and  $\mathcal{X} = \{0, 1, 2, 3, 4\}$  with the following probabilities.

	0	1	2	3	4
$\mathbb{P}[X \theta=0]$ $\mathbb{P}[X \theta=1]$					

Suppose we observe x=2. As  $P[X\geqslant 2|\ \theta=0]=0.11$  and  $\mathbb{P}[X\geqslant 2|\theta=1]=0.049$ , from the frequentist point of view, we would reject  $\mathcal{H}_0:\theta=0$  at any  $\alpha>0.11$  and we would reject  $\mathcal{H}_0:\theta=1$  at any  $\alpha>0.049$  (testing based on p-value). From the Bayesian point of view, if we consider  $\theta=1$  and  $\theta=0$  as equally likely a-priori, then  $\mathbb{P}(\theta=j|x=2)=\frac{\mathbb{P}(x=2|\theta=j)\mathbb{P}(\theta=j)}{\mathbb{P}(x=2)}=\frac{0.04\cdot0.5}{\mathbb{P}(x=2)}$  is the same for both j=0,1. We get the posterior odds  $\frac{p(\theta=0|x=2)}{P(\theta=1|x=2)}=1$  and so the observation x=2 does not favor one model against the other. This illustrates that the Bayesian approach picks the better model (i.e. better value of  $\theta$ ) among all considered ones, while the Frequentist approach picks the conjectured model (value of  $\theta$ ) if it is "good" in the sense that the realization obtained is not too extreme/unlikely under the supposed model  $\mathcal{H}_0$ .

Two comments are in order. First, for Bayesian testing, there is no type I error; we cannot condition on  $\mathcal{H}_0$  being true, because  $\theta$  is not a parameter, but a RV in this perspective. Second, under continuous (prior and posterior) distributions for  $\theta$ , non-trivial adjustments need to be made to test point hypotheses, because the probability of any single point is zero.

# 2.4 Coverage Sets

#### 2.4.1 Frequentist Confidence Sets

In the frequentist paradigm, a confidence set  $C(X) \subseteq \Theta$  is a (random) set that should cover the true  $\theta$  with a prespecified probability:

$$\inf_{\theta \in \Theta} \mathbb{P}[\theta \in C(X)|\theta] = 1 - \alpha .$$

For a scalar  $\theta$ , we speak of confidence intervals. (The infimum operator appears because generally, this probability can depend on the true value of  $\theta$ , and because we do not know the true  $\theta$ , the coverage probability has to be guaranteed  $\forall \theta \in \Theta$ ). The probability is taken with respect to the RV X, with  $\theta$  fixed. Hence, the definition of C(X) above says that, if we look at many different, random realizations of X, the set C(X) should contain the true  $\theta$  in  $100(1-\alpha)\%$  of cases (even if the true  $\theta$  is such that this probability is lowest).

We can construct C(X) based on a test  $\varphi$ , testing the point-hypothesis  $\mathcal{H}_0: \theta = \theta_0$ . Let C(x) contain all the values for  $\theta_0$  that we would accept given our realization x:

$$C(x) = \{\theta_0 \in \Theta : \varphi(x; \theta_0) = 1\}.$$

Then

$$\inf_{\theta \in \Theta} \mathbb{P}[\theta \in C(X) | \theta] = \inf_{\theta \in \Theta} \mathbb{P}[\varphi(x; \theta) = 1 \mid \theta] = 1 - \underbrace{\sup_{\theta \in \Theta} \left( \mathbb{P}[\varphi(x; \theta) = 0 \mid \theta] \right)}_{\text{type I error; reject } \theta \text{ given } \theta \text{ is true}} = 1 - \alpha ,$$

i.e. we get the desired coverage probability.<sup>9</sup>

Consider the example from above:  $X \sim N(\theta, 1)$ , and we test  $\mathcal{H}_0: \theta = \theta_0$  vs.  $\mathcal{H}_1: \theta \neq \theta_0$ . Using the two-sided t-test  $\varphi_t(x; \theta_0) = \mathbf{1}\{|x - \theta_0| < c_{\alpha}^t\}$ , we accept  $\mathcal{H}_0$  if  $-c_{\alpha}^t \leqslant x - \theta_0 \leqslant c_{\alpha}^t$ ,

<sup>&</sup>lt;sup>9</sup>Just as for some discrete distributions of X, we cannot construct a test with size  $\alpha$ , but with size at most  $\alpha$  (i.e.  $\leq \alpha$ ), for such distributions we cannot construct a C(X) for which the coverage probability is exactly  $1-\alpha$ . The actual, general definition of C(X) in the expression above uses a larger-or-equal sign:  $\inf_{\theta \in \Theta} \mathbb{P}[\theta \in C(X)|\theta] \geq 1-\alpha.$ 

which shows that the set of all  $\theta_0$  we would accept is  $C_{\alpha}^t(x) = [x - c_{\alpha}^t, x + c_{\alpha}^t]$ . In the case of a size  $\alpha = 0.05$  test, we get the 95% confidence interval  $C_{0.05,t}(x) = [x - 1.96, x + 1.96]$ . Under the LR test,  $\varphi_{LR}(x;\theta_0) = \mathbf{1}\{(x-\theta_0)^2 \le c_{\alpha}^{LR}\}$ , which means we accept  $\mathcal{H}_0$  if  $-\sqrt{c_{\alpha}^{LR}} \le (x-\theta_0) \le \sqrt{c_{\alpha}^{LR}}$  and yields  $C_{\alpha}^{LR}(x) = [x - \sqrt{c_{\alpha}^{LR}}, x + \sqrt{c_{\alpha}^{LR}}]$ .

For the same test statistic, there are (infinitely) many confidence sets one could construct. Typically, we want the confidence set to be as small (short) as possible (conditioning on a given coverage probability). Essentially, constructing C(X) from the acceptance region of a point-hypothesis test leads to the smallest C(X).

Just as sometimes, we cannot set up a test analytically, in these cases we cannot find an analytical expression for the confidence set, but have to construct it numerically. Consider the example from the previous section:  $X|\theta \sim N(\theta,1)$ , single realization x, with parameter space  $\theta \in \mathbb{R}_+$ . The previous section showed how to test  $\mathcal{H}_0: \theta = \theta_0$  vs.  $\mathcal{H}_1: \theta \neq \theta_0$  numerically using the LR test  $\varphi(x;\theta_0) = \mathbf{1}\{T(x;\theta_0) < c\}$ . Based on this test, we can construct C(x) numerically as follows.

#### Algorithm 2 (Numerical Confidence Set Construction).

- 1. Choose a grid  $\mathcal{T}$  of values for  $\theta_0$ .
- 2. For each  $\theta_0 \in \mathcal{T}$ ,
  - (a) for m = 1: M, draw  $x^m$  from  $N(\theta_0, 1)$ , the distribution of X under  $\mathcal{H}_0: \theta = \theta_0$ . For each draw, compute  $T(x^m; \theta_0)$ . This gives you a distribution  $\{T(x^m; \theta_0)\}_{m=1}^M$  that approximates the true distribution of  $T(X; \theta_0)$  under  $\mathcal{H}_0: \theta = \theta_0$ .
  - (b) get the critical value for a size  $\alpha$  test,  $c_{\alpha}(\theta_0)$ , as the (empirical)  $100(1-\alpha)th$  quantile of the distribution of  $\{T(x^m;\theta_0)\}_{m=1}^M$ .
  - (c) compute  $T(x; \theta_0)$  for your particular, observed realization x. If  $T(x; \theta_0) \leq c_{\alpha}(\theta_0)$ , then  $\theta_0 \in C(x)$ , otherwise  $\theta_0 \notin C(x)$ .

## 2.4.2 Bayesian Credible Sets

Bayesian credible sets C(X) are defined by

$$\mathbb{P}[\theta \in C(X)|X] = 1 - \alpha .^{10}$$

In contrast to the frequentist paradigm, here X is fixed while  $\theta$  is random. In line with the frequentist paradigm, here too there are many different sets C(X) one could construct. The

<sup>&</sup>lt;sup>10</sup>Again, more formally, this definition uses a greater-or-equal sign.

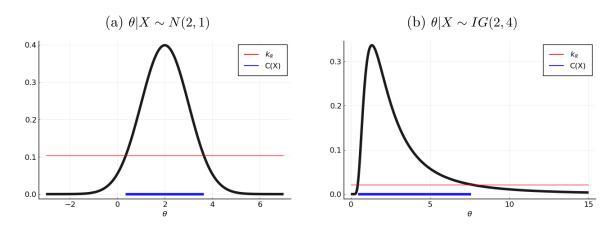


Figure 2.3: Bayesian Highest Posterior Density (HPD) Sets

Notes: Illustration of the HPD sets with coverage probabilities  $1 - \alpha \in \{0.9, 0.95\}$  under different posteriors.

smallest is the highest posterior density (HPD) set

$$C(X) = \{\theta \in \Theta : p(\theta|X) \ge k_{\alpha}\}\$$
, where  $k_{\alpha}$  is s.t.  $\mathbb{P}[\theta \in C(X)|\theta] \ge 1 - \alpha$ .

See illustration in Fig. 2.3. Essentially, we use a threshold value  $k_{\alpha}$  to slice the posterior horizontally and look at the part of the x-axis (the values for  $\theta$ ) that correspond to the highest values of the posterior density.

# **Appendix**

**Definition 32.** The <u>minimax risk</u> is  $\bar{R} = \inf_{\delta \in \mathcal{D}} \sup_{\theta \in \Theta} R(\theta, \delta)$ , where  $\mathcal{D}$  is the set of all possible point estimators. The minimax estimator is  $\delta_0 = \arg\min_{\delta \in \mathcal{D}_0} (\max_{\theta \in \Theta} R(\theta, \delta))$ .

The minimax estimator leads to the best among all possible worst cases. Intuitively, nature (by choosing  $\theta$ ) maximizes, the econometrician (by choosing  $\delta$ ) minimizes the risk.

Consider the example with  $L=(\theta-\delta)^2$ ,  $X|\theta\sim N(0,1)$  and  $\delta=c\cdot X$ . Let  $\Theta=\mathbb{R}$ . Taking c=1 leads to the minimax estimator, since for any  $c\neq 1$ , nature could choose  $\theta=\pm\infty$ , which results in  $R(\theta,\delta)=\pm\infty$  for any  $\delta=cx,\,c\neq 1$ .

**Definition 33.** A point estimator  $\delta_1$  is <u>inadmissible</u> if  $\exists \ \delta_2 \ s.t. \ R(\theta, \delta_1) \ge R(\theta, \delta_2) \ \forall \ \theta$  and  $\exists \ \theta' \ s.t. \ R(\theta', \delta_1) > R(\theta', \delta_2).$ 

In other words,  $\delta_1$  is inadmissible if there is another point estimator which never has a higher frequentist risk and for one value of  $\theta$  even leads to a lower frequentist risk than  $\delta_1$ . In our example,  $\forall c > 1$ ,  $R(\theta, cX) = \theta^2(1-c)^2 + c^2 > 1 = R(\theta, x)$  and so any c > 1 is inadmissible.

Proposition 18 (Neyman-Pearson Lemma).

Let  $\mathcal{H}_0: \theta = \theta_0$  and  $\mathcal{H}_1: \theta = \theta_1$ . Then  $\exists$  a UMP test  $\forall \alpha \in (0,1)$  and it is of the form  $\varphi(x) = \mathbf{1}\left\{\frac{p(x|\theta_1)}{p(x|\theta_0)} < k\right\}$ , with k determined so as to achieve the required type I error:  $\mathbb{P}\left[\frac{p(X|\theta_1)}{p(X|\theta_0)} > k|\theta_0\right] = \alpha$ . In other words, when both hypotheses are point hypotheses, the LR test is the UMP test.

**Proof:** Take the above  $\varphi(x)$ . Suppose  $\exists \varphi^*$  that satisfies the size constraint:  $R(\theta_0, \varphi^*) = \int (1 - \varphi^*) p(x|\theta_0) dx \leqslant \alpha$ . If  $x \in S^+$ ,  $\varphi(x) = 1$  (we reject) and so  $p(x|\theta_1) > kp(x|\theta_0)$ . If  $x \in S^-$ ,  $\varphi(x) = 0$  (we do not reject) and so  $p(x|\theta_1) < kp(x|\theta_0)$ . Therefore,

$$\int (\varphi^r - \varphi) \left[ p(x|\theta_1) - kp(x|\theta_0) \right] dx$$

$$= \int_{S^+} (\varphi^* - \varphi) \left[ p(x|\theta_1) - kp(x|\theta_0) \right] dx + \int_{S^-} (\varphi^* - \varphi) \left[ p(x|\theta_1) - kp(x|\theta_0) \right] dx \geqslant 0.$$

This means that

$$\beta(\theta_1) - \beta^*(\theta_1) = \int [(1 - \varphi) - (1 - \varphi^*)] \rho(x|\theta_1) dx$$

$$= \int (\varphi^* - \varphi) p(x|\theta_0) dx$$

$$\geqslant \int (\varphi^* - \varphi) k \rho(x|\theta_0) dx$$

$$= k[1 - \alpha^* - (1 - \alpha)]$$

$$= k(\alpha - \alpha^*) \geqslant 0$$

as  $\alpha^* \leqslant \alpha$ .