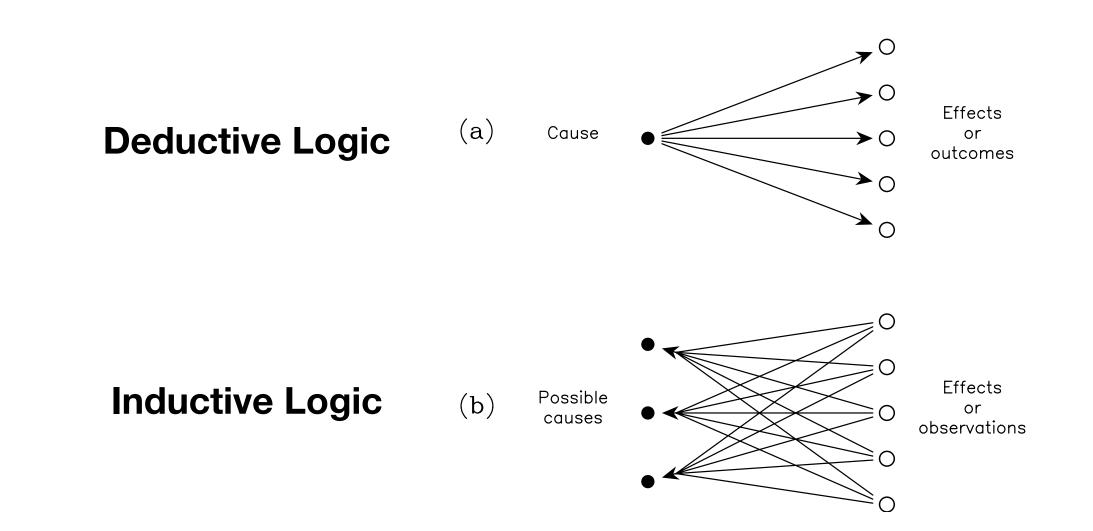
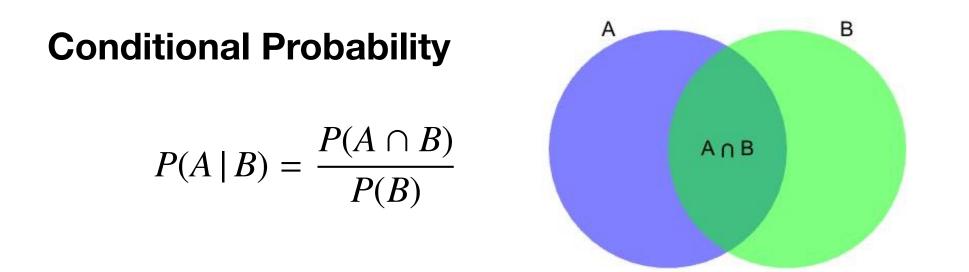
Teeraparb Chantavat IF, Naresuan University

Probability Theory

Approach	Probability definition
Frequentist Statistical Inference	 P(A) = long-run relative frequency with which A occurs in identical repeats of an experiment. "A" restricted to propositions about random variables.
Bayesian Inference	P(A B) = a real number measure of the plausibility of a proposition/hypothesis A, given (conditional on) the truth of the in- formation represented by proposition B. "A" can be any logical proposition, <u>not</u> restricted to propositions about random variables.





- P(A)Observing the data.P(B)The theory is true.
- P(A | B) The data is observed given that the theory is true

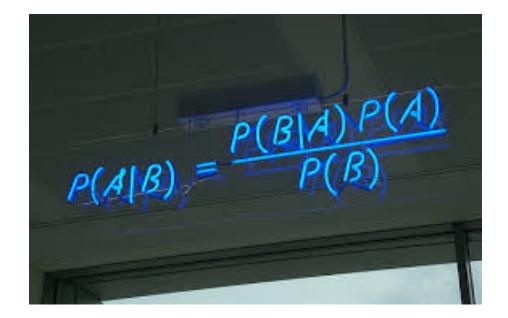
Symmetry Rule

 $P(B \cap A) = P(A \cap B)$

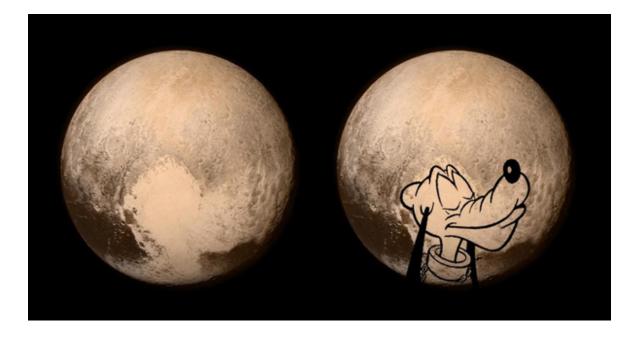
P(B | A)P(A) = P(A | B)P(B)

Bayesian Rule

$$P(B | A) = \frac{P(A | B)P(B)}{P(A)}$$



"There are no problems left in statistics except the assessment of probability"



$P(A \mid B) \neq P(B \mid A)$

$$P(H \mid D) = \frac{P(D \mid H)P(H)}{P(D)}$$

- P(H) Probability that the hypothesis is true. **Prior**
- P(D|H) Probability that the data is observed **Likelihood** given that the hypothesis is true.
- P(D) Probability that the collections of data **Evidence** is liable.
- P(H|D) Probability that the hypothesis is true **Posterior** given that the data is true.

• A theory usually have many parameters, for example, a two-parameter model

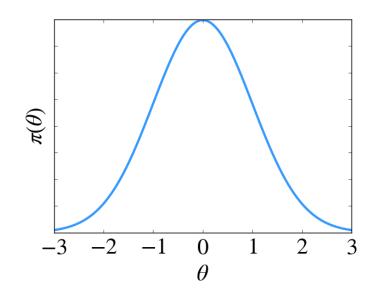
 $\boldsymbol{\Theta} = \{\boldsymbol{\Theta}_1, \boldsymbol{\Theta}_2\}$

 The hypothesis is the assumption that the parameter have a particular value for example

$$H_1: \theta_1 = 1.0 \text{ and } \theta_2 = 2.0$$

$$\mathbf{H}_1 \equiv \boldsymbol{\theta}_1 = (\theta_1, \theta_2)$$

- The prior probability is the distribution of the parameters we know before the experiment (degree of believe).
- We can have a uniform distribution for total ignorance or a normal distribution if mean and standard deviation are given.



• The evidence is usually considered as a normalization constants — nothing to do with parameter estimations.

 $P(\theta | \mathbf{x}) \propto \mathcal{L}(\mathbf{x} | \theta) \pi(\theta)$

• However, the **evidence** is important **model comparison**.

 In most cases, we are working the logarithm of the likelihood function called log-likelihood

$$L(\boldsymbol{x} | \boldsymbol{\theta}) = \log_{e} \mathcal{L}(\boldsymbol{x} | \boldsymbol{\theta})$$

• Expanding around the maximum of the loglikelihood at θ_0 i.e.

$$\left. \frac{\partial L}{\partial \theta_{\alpha}} \right|_{\theta = \theta_0} = 0$$

$$L(\boldsymbol{x} \mid \boldsymbol{\theta}) = L(\boldsymbol{x} \mid \boldsymbol{\theta}_{0}) + \frac{1}{2} \sum_{\alpha, \beta} \frac{\partial L}{\partial \theta_{\alpha}} \frac{\partial L}{\partial \theta_{\beta}} \Big|_{\boldsymbol{\theta} = \boldsymbol{\theta}_{0}} (\theta_{\alpha} - \theta_{\alpha 0}) (\theta_{\beta} - \theta_{\beta 0})$$

• We define the **precision matrix P** as

$$L(\boldsymbol{x} | \boldsymbol{\theta}) = L(\boldsymbol{x} | \boldsymbol{\theta}_0) - \frac{1}{2} (\boldsymbol{\theta} - \boldsymbol{\theta}_0)^{\mathrm{T}} \cdot \mathbf{P} \cdot (\boldsymbol{\theta} - \boldsymbol{\theta}_0)$$

where
$$P_{\alpha\beta} \equiv \frac{\partial^2 L}{\partial \theta_{\alpha} \partial \theta_{\beta}} \Big|_{\theta=\theta_0}$$

• The log likelihood is given by

$$\mathcal{L}(\boldsymbol{x} | \boldsymbol{\theta}) \propto \exp\left(-\frac{1}{2} \left(\boldsymbol{\theta} - \boldsymbol{\theta}_0\right)^{\mathrm{T}} \cdot \mathbf{P} \cdot \left(\boldsymbol{\theta} - \boldsymbol{\theta}_0\right)\right)$$

 The inverse of the precision matrix is called covariance matrix

$$\mathbf{P} \equiv \mathbf{C}^{-1}$$

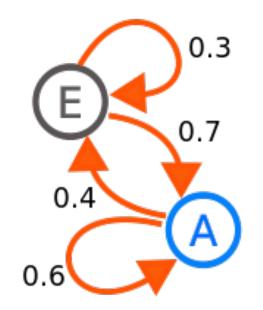
$$\mathcal{L}(\boldsymbol{x} | \boldsymbol{\theta}) \propto \exp\left(-\frac{1}{2} \left(\boldsymbol{\theta} - \boldsymbol{\theta}_0\right)^{\mathrm{T}} \cdot \mathbf{C}^{-1} \cdot \left(\boldsymbol{\theta} - \boldsymbol{\theta}_0\right)\right)$$

The variance of the parameter can be estimated as

$$\operatorname{Var}\left(\theta_{\alpha}\right) = \mathbf{C}_{\alpha\,\alpha}$$

Markov Chain Monte Carlo (MCMC)

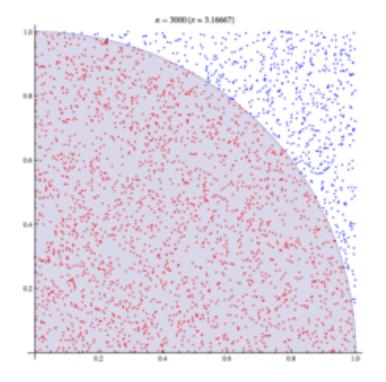
A Markov chain is a chain of states in a parameter space that is "memoryless" (Markov property).

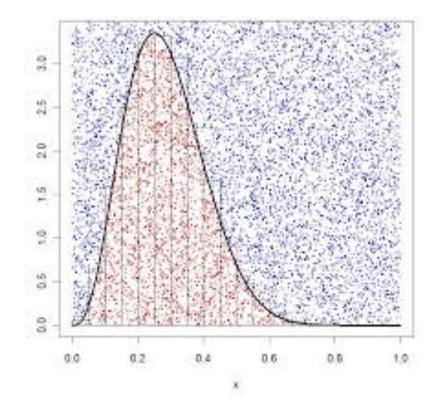


How the state change depends only on the current state.

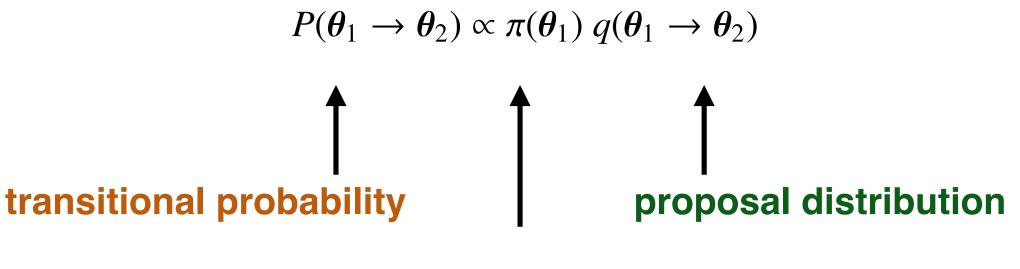
Markov Chain Monte Carlo (MCMC)

A **Monte Carlo** is a method using random walk to generate the output.





The **Metropolis-Hastings algorithm** is an algorithm for random walks that will eventually converge to a true distribution of the parameter space.



prior probability

The change of state from θ_1 to θ_2 is governed by the **acceptance rate**

$$\alpha(\boldsymbol{\theta}_1 \to \boldsymbol{\theta}_2) = \min\left\{1, \frac{\pi(\boldsymbol{\theta}_2) \ q(\boldsymbol{\theta}_2 \to \boldsymbol{\theta}_1)}{\pi(\boldsymbol{\theta}_1) \ q(\boldsymbol{\theta}_1 \to \boldsymbol{\theta}_2)}\right\}$$

We are assumed an equilibrium state; hence,

$$q(\theta_1 \rightarrow \theta_2) = q(\theta_2 \rightarrow \theta_1)$$

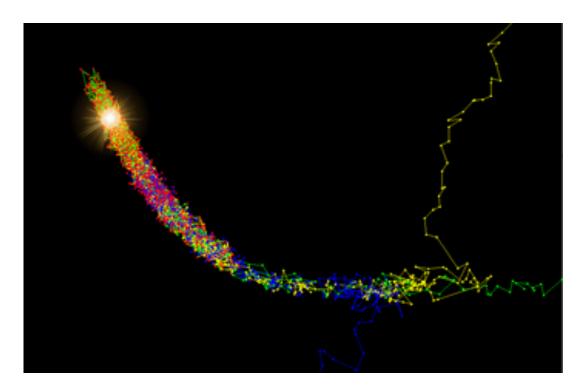
Therefore,

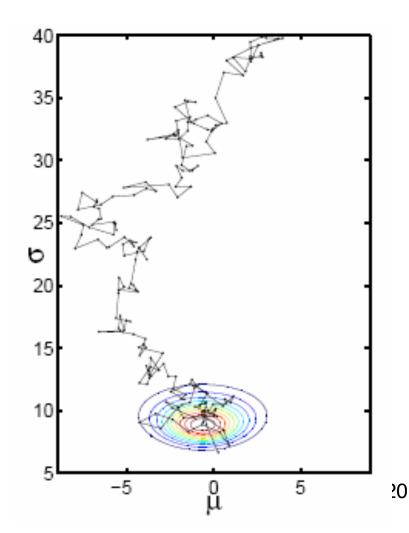
$$\alpha(\boldsymbol{\theta}_1 \to \boldsymbol{\theta}_2) = \min\left\{1, \frac{\pi(\boldsymbol{\theta}_2)}{\pi(\boldsymbol{\theta}_1)}\right\}$$

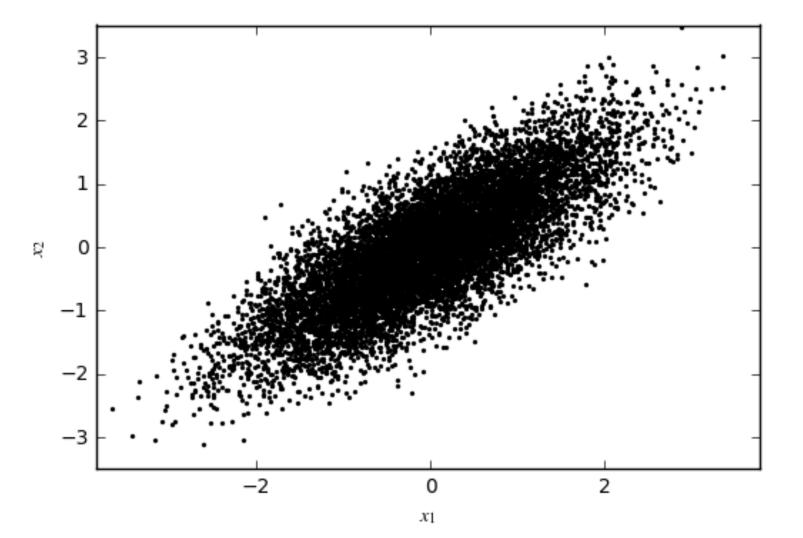
Pseudo code for Metropolis-Hastings Algorithm

```
alpha = likelihood2 / likelihood1;
if alpha > 1:
   jump to the new state;
else:
   if alpha > rand();
    jump to the new state;
   else:
      remain in the same state;
```

The chain will take some time to stabilize this is called the **burn-in phase**



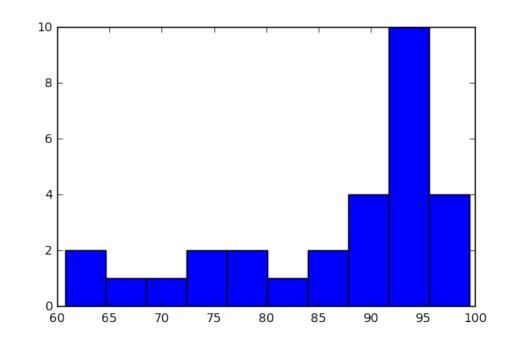




 Histogram is a common way to make sense of discrete data.

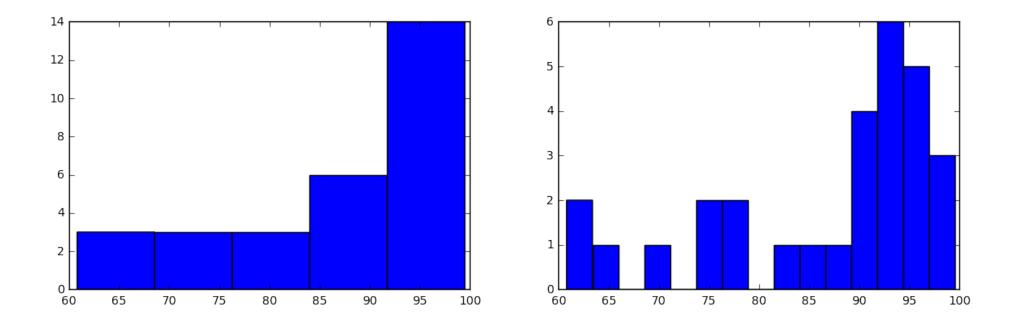
Data

93.5, 93, 60.8, 94.5,
82, 87.5, 91.5, 99.5,
86, 93.5, 92.5, 78,
76, 69, 94.5, 89.5,
92.8, 78, 65.5, 98,
98.5, 92.3, 95.5, 76,
91, 95, 61.4, 96, 90

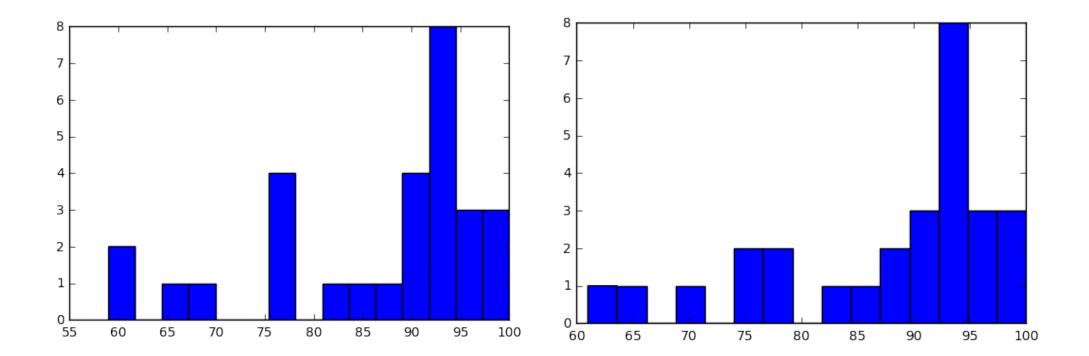


Histogram

 The same data could generate different histograms with different number of bins.



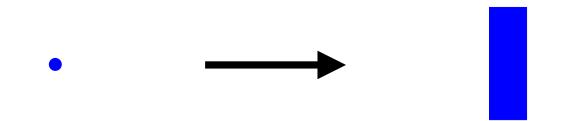
• The same data could generate different histograms with different starts of left-edge of bins.



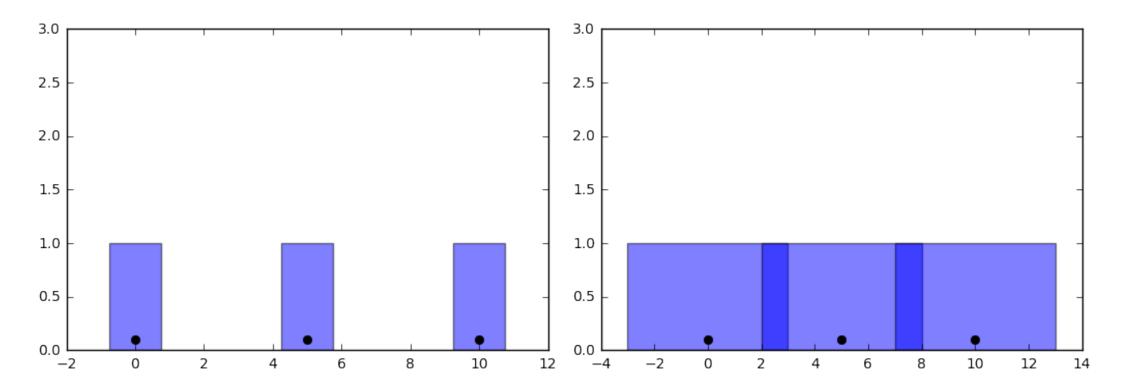
Drawbacks of Histogram

- Not smooth
- Depend on width of bins
- Depends on end points of bins

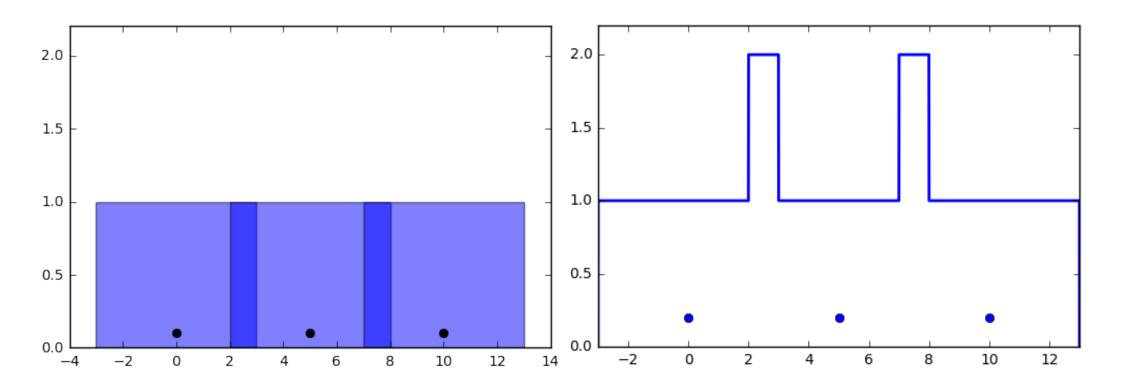
• If we instead replace the data point by a kernel function



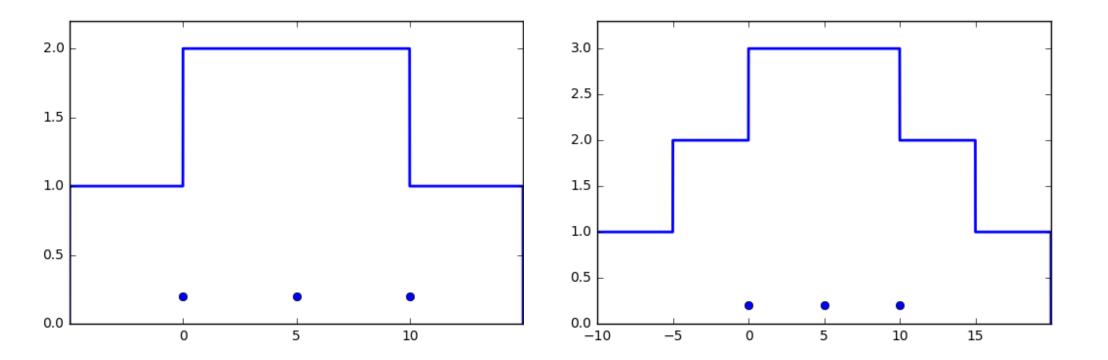
• For, simplicity suppose we have only three data points **0**, **5**, **10**



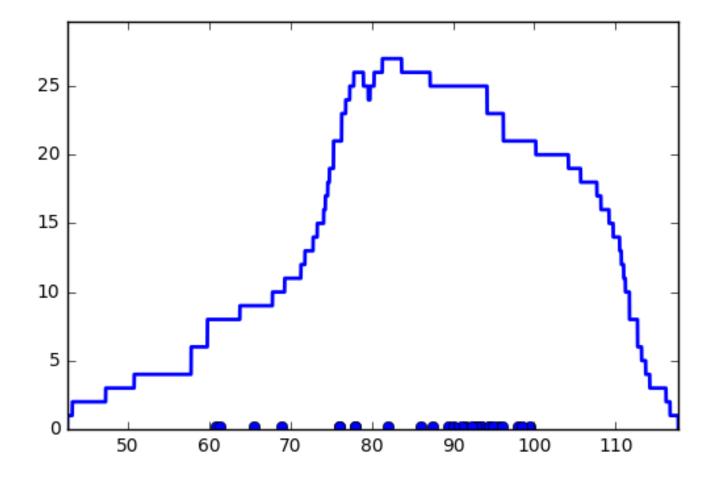
 For, simplicity suppose we have only three data points 0, 5, 10



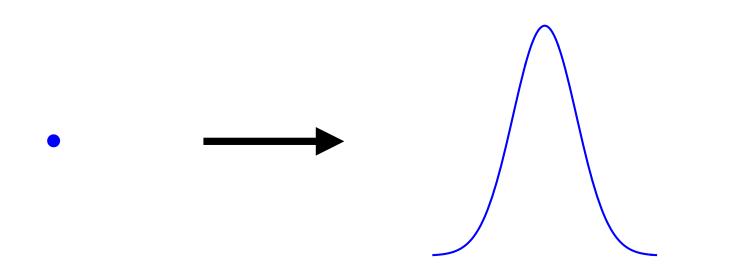
 For, simplicity suppose we have only three data points 0, 5, 10



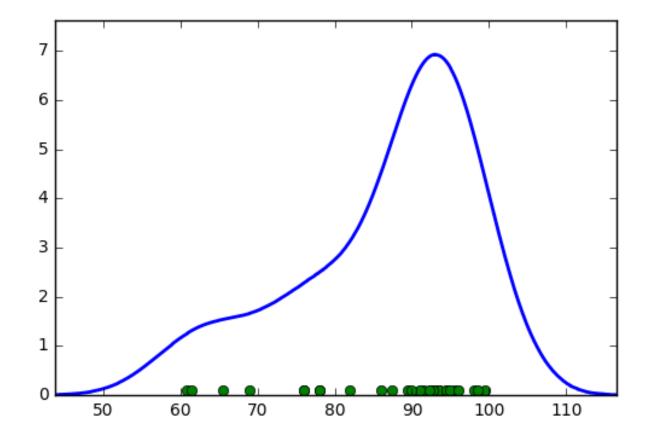
• With our previous data



- The plot is not smooth because we have a nonsmooth kernel function
- We can use a smooth kernel function for example a Gaussian function

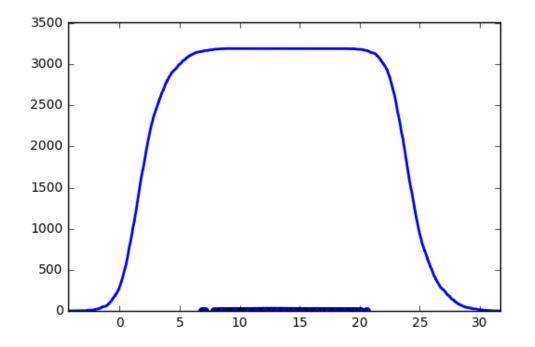


• We have a smooth distribution.



• We oversmooth the distribution.

Oversmoothed



- The optimal bandwidth has to be estimated.
- A normal way to estimated the optimal bandwidth is to minimize the Asymptotic Mean Integrated Squared Error (AMISE)

$$\int (f(x) - f_n(x))^2 \mathrm{d}x$$

