

COSMOLOGICAL DATA ANALYSIS

中山大学
王鑫



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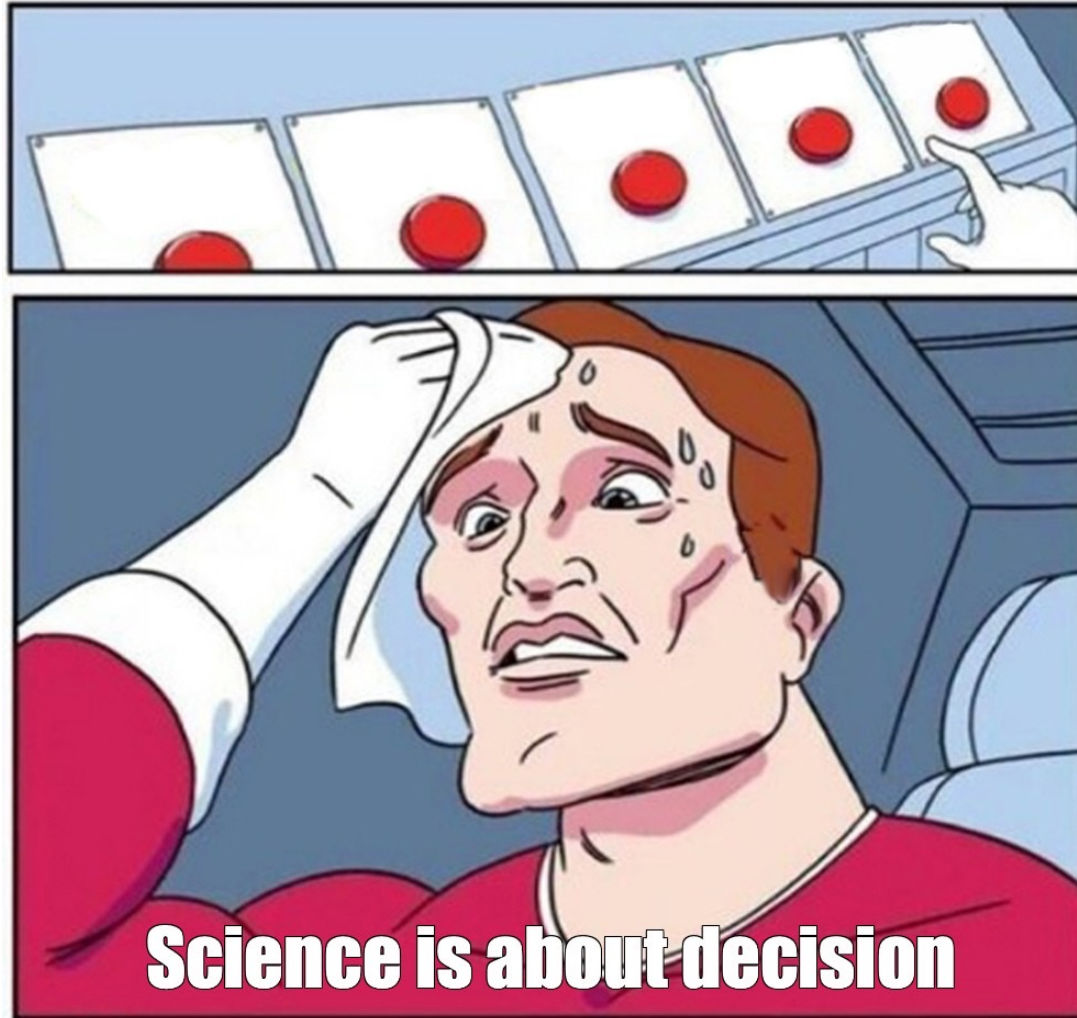


BASED ON ...

- ***Practical statistics for Astronomer***, Cambridge University Press, J. V. Wall and C. R. Jenkins
- ***Modern Cosmology***, Second Edition, Academic Press, Scott Dodelson, Fabian Schmidt
 - Chapter 14. “Analysis and inference”
- ***Information Theory, Inference, and Learning Algorithms***, Cambridge University Press, David J.C. MacKay



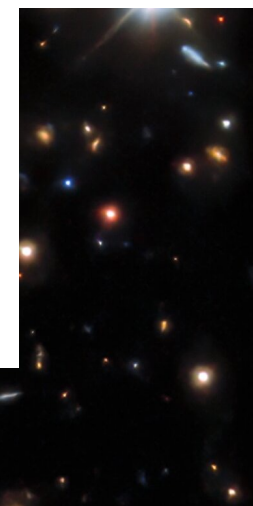
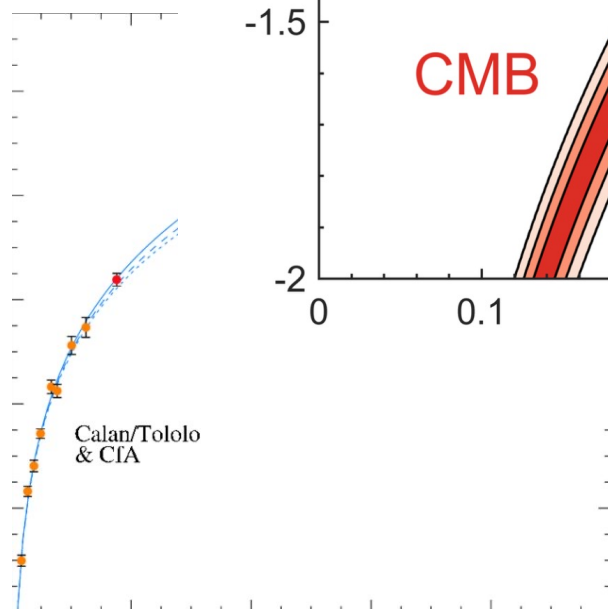
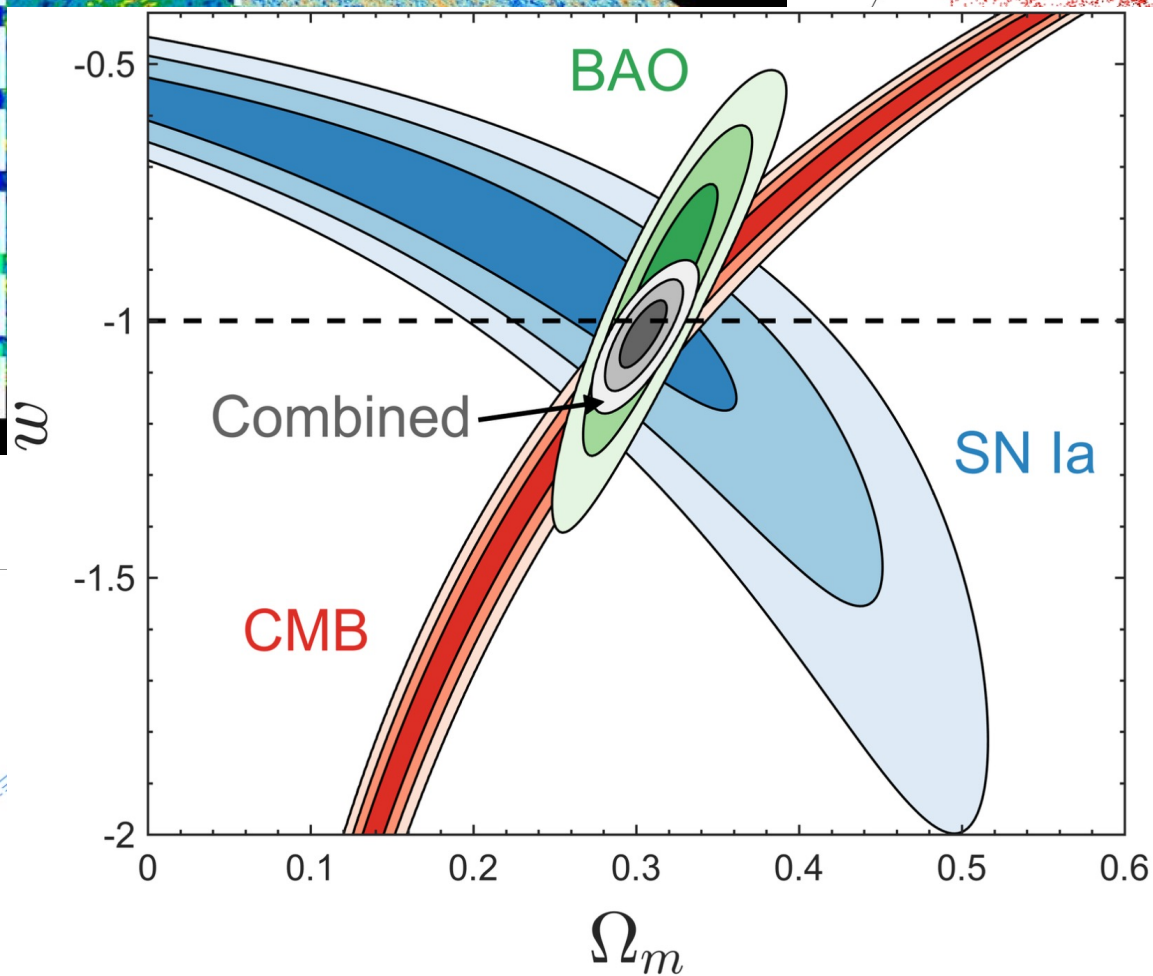
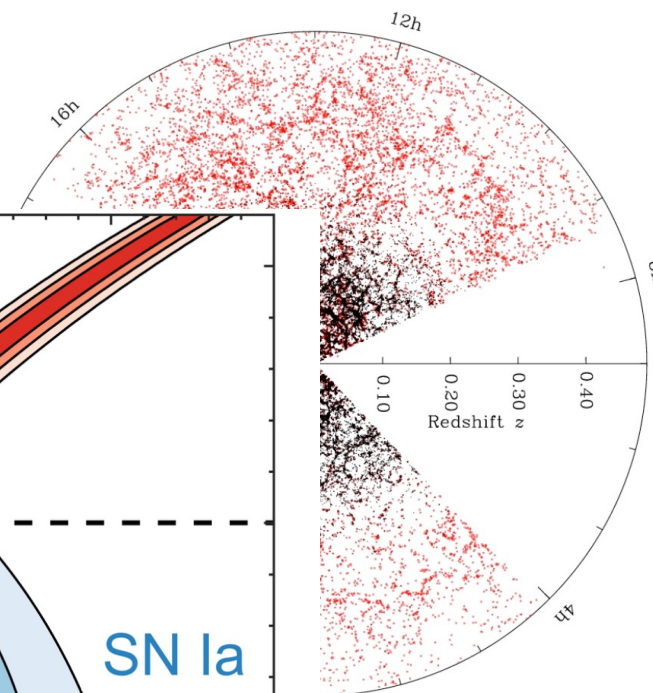
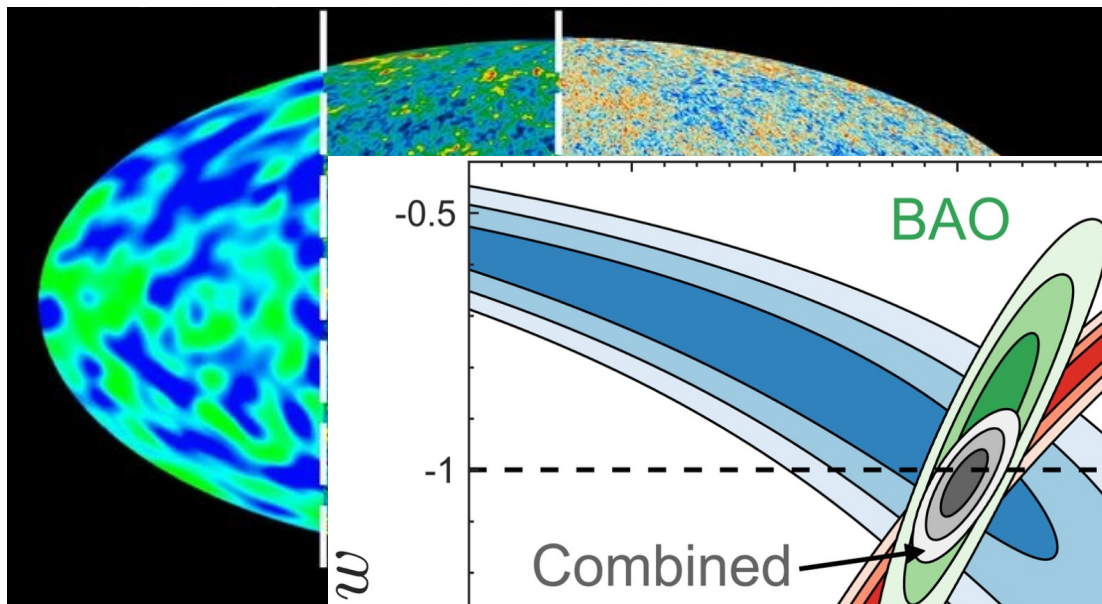
SCIENCE IS ABOUT DECISION



SCIENCE IS ABOUT DECISION

- *star vs galaxy ?*





ASTRONOMERS CANNOT AVOID STATISTICS

- **Data Error** (range):
 - *How can data be used best? Or at all?*
 - *Correlation, testing the hypothesis, model fitting; how do we proceed?*
- **Incomplete samples:**
 - *samples from an experiment cannot be re-run (cosmology)*
 - *upper limits*
- **We must decide:**
 - *The decision process need some methodology*
 - *no matter how good the experiment.*



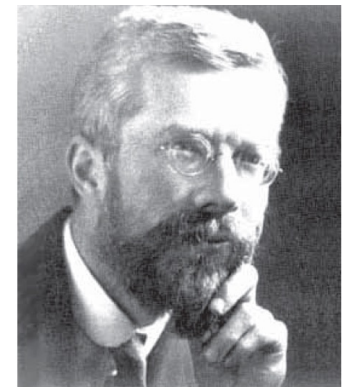
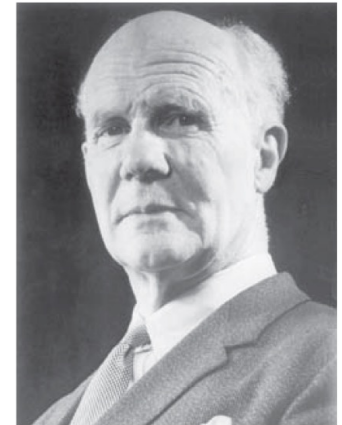
BAYESIAN VS. FREQUENTIST



Thomas Bayes
(1702–1761)



What's **probability**?
How to handle **uncertainty**?
How to incorporate **prior knowledge**?



Jerzy Neyman, Egon
Pearson and Ronald
Fischer.

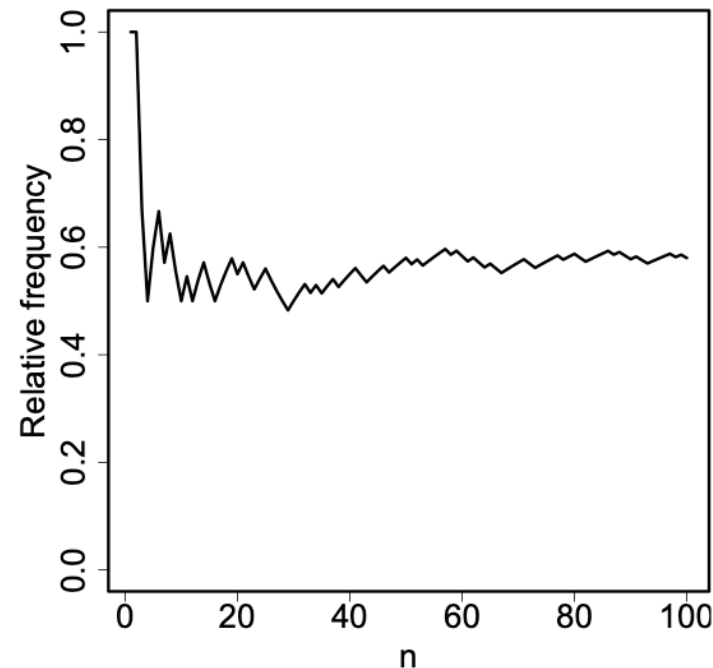


FREQUENTISTS

- Consider experiments with a random component.
- **Probabilities:** the relative frequency of an event, A , is defined as

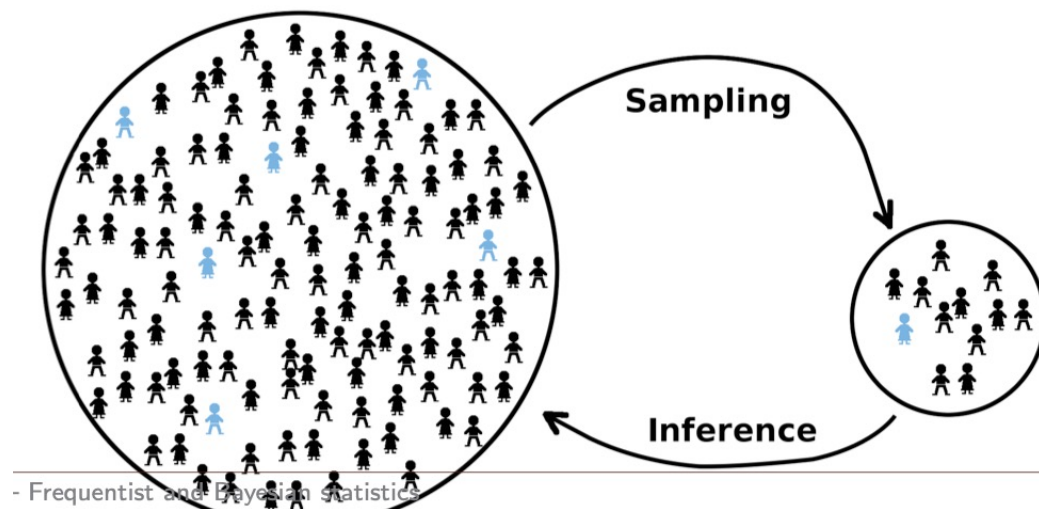
$$P(A) = \frac{\text{\# of outcomes consistent with } A}{\text{\# of experiments}}$$

- The **probability** of event A is the **limiting** relative frequency



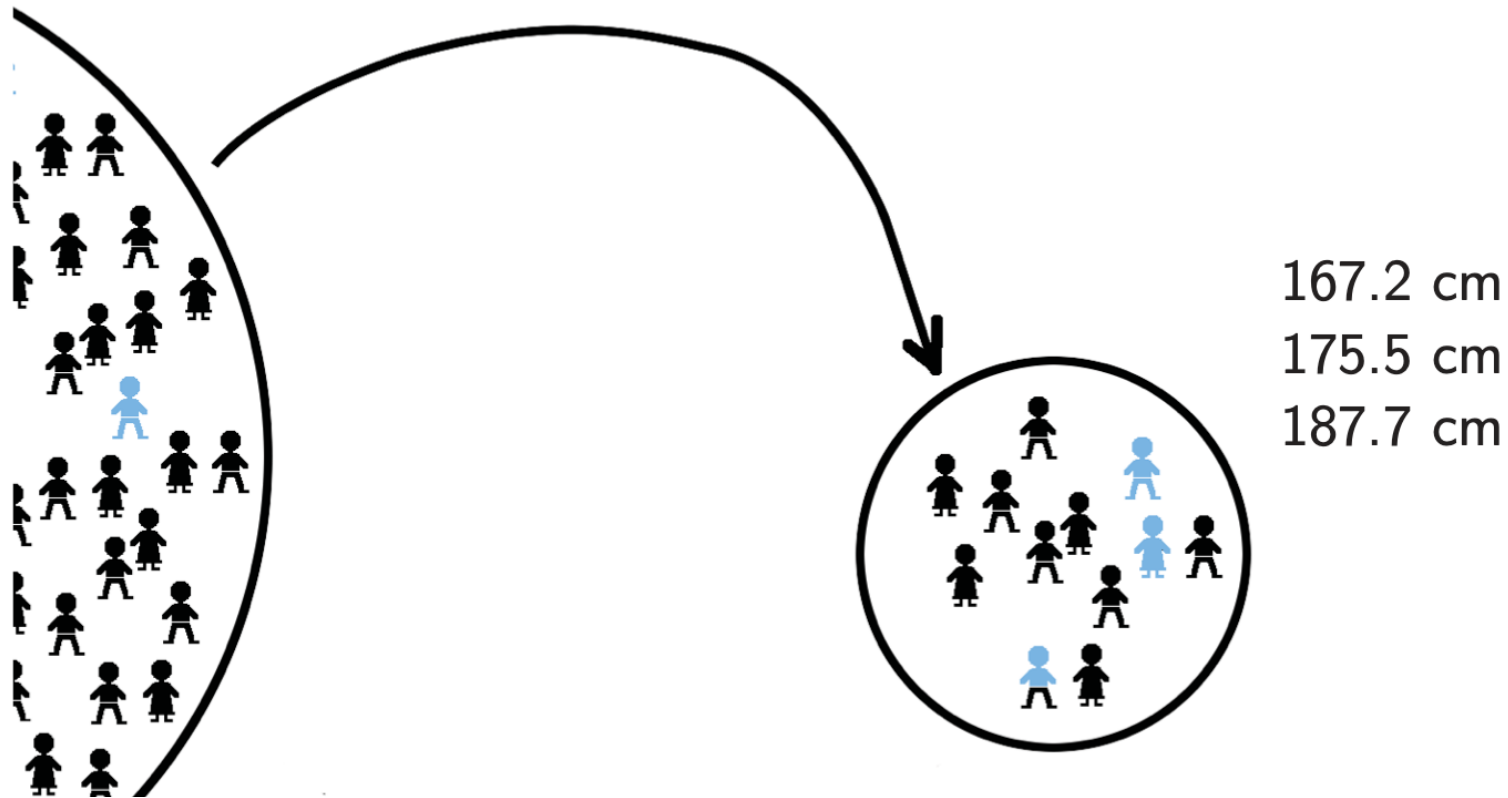
FREQUENTISTS

- This definition **restricts** the things we can add probabilities to:
 - What is the probability of there being life on Mars 100 billion years ago?
 - not a repeatable experiment
- Assumption: there is an **unknown but fixed underlying parameter**, θ , for a population (e.g., the mean height of men).
- Random variation:
 - environmental factors, measurement errors, ...



THE META-EXPERIMENT IDEA

- *meta-experiments*: consider the current dataset as *a single realization* from all possible datasets.



THE META-EXPERIMENT IDEA

- For example: a population mean is *real* but *unknown*, and *unknowable*
 - can only be estimated from the data.
- From the distribution for the sample mean, constructs a confidence interval, centered at the *sample mean*.
 - Regardless of the location of true mean;
 - *Can't say* there's a 95% probability that the true mean is in this interval, because it's either already in, or it's not.
 - the true mean is a fixed value, which doesn't have a distribution.
 - The sample mean does have a distribution!
 - "95% of similar intervals would contain the true mean if each interval were constructed from a different random sample like this one."



BAYESIANS

- A numerical formalization of our degree of belief:
 - personal belief → the **prior**:
 - No fixed values for parameters but a distribution.
 - All distributions are subjective: Yours is as good as mine
- Treat the parameters (e.g. mean) as random var.
 - mean : the mean of my distribution
- Only the data are real:
 - The population mean is an abstraction
 - Exist only conceptually
 - some values are more believable than others based on the data and their prior beliefs.



BAYESIANS

- **Credibility interval:**
 - Posterior: centered near the sample mean, tempered by “prior”.
- Bayesian can say what the frequentist cannot:
 - “There is a 95% probability (*degree of believability*) that this interval contains the mean.”

	Advantages	Disadvantages
Frequentist	Objective Calculations	Confidence intervals (not quite the desired)
Bayesian	Credibility intervals (usually the desired) Complex models	Subjective Calculations



IN SUMMARY

- *A frequentist is a person whose long-run ambition is to be wrong 5% of the time.*
- *A Bayesian is one who, vaguely expecting a horse, and catching a glimpse of a donkey, strongly believes he has seen a mule.*

A frequentist uses impeccable logic to answer the wrong question, while a Bayesian answers the right question by making assumptions that nobody can fully believe in.

P. G. Hamer



EXAMPLE

- Before 1987, 4 supernovae had been recorded in 10 centuries. What, before 1987, was the probability of a bright supernova happening in the 20th century?
- Three possible answers:
 - Probability is meaningless in this context.
 - Supernovae are physically determined events, not random.
 - From this God's-eye viewpoint, probability is indeed meaningless; 'God does not play dice...'
 - Frequentist: our best estimate of the probability is $4/10$, although it is obviously not very well determined.
 - Bayesian: a-priori assignment:
 - Modeling: stellar mass function, stellar birth rate, metallicity etc.



PROBABILITY

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PROBABILITY

- *Measure of belief (Cox, 1946):*
 - *A, B and C are three events*
 - *wish to have some measure of how strongly we think each is likely to happen*
 - *apply the rule: if A is more likely than B, and B is more likely than C, then A is more likely than C.*
- *Axioms of probability (Kolmogorov)*
 - *any random event A has a probability $P(A)$ between 0 and 1;*
 - *the sure event has $P(A) = 1$;*
 - *If A and B are exclusive events, then $P(A \text{ or } B) = P(A) + P(B)$;*



CONDITIONALITY AND INDEPENDENCE

- Two events A and B are said to be independent:

$$P(A \text{ and } B) = P(A)P(B)$$

- Probability of one is unaffected by what we may know about the other.

- conditional probability:

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

- Marginalization:

$$P(A) = \sum_i P(A|B_i)P(B_i)$$

- To get rid of these 'nuisance parameters' B_i .



BAYES' THEOREM

- Bayes' theorem:

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

- Particularly, θ is theory, d is data:

$$P(\theta|d) = \frac{P(d|\theta)P(\theta)}{P(d)}$$

- *posterior* probability: $P(\theta|d)$
- *prior* probability: $P(\theta)$, state of belief before the experiment
- Likelihood function: $P(d|\theta) = \mathcal{L}(\theta), \mathcal{L}(d|\theta)$
- normalizing factor: $P(d)$



EXAMPLE

- There are N **red balls** and M **blue balls** in a jar, and $N + M = 10$. We draw $T = 3$ times (putting the balls back after drawing them) and $R = 2$ **red balls**. How many red balls are there in the jar?

- Model (hypothesis) probability of red ball: $N / (N + M)$

- The likelihood (probability of getting R **red balls**):

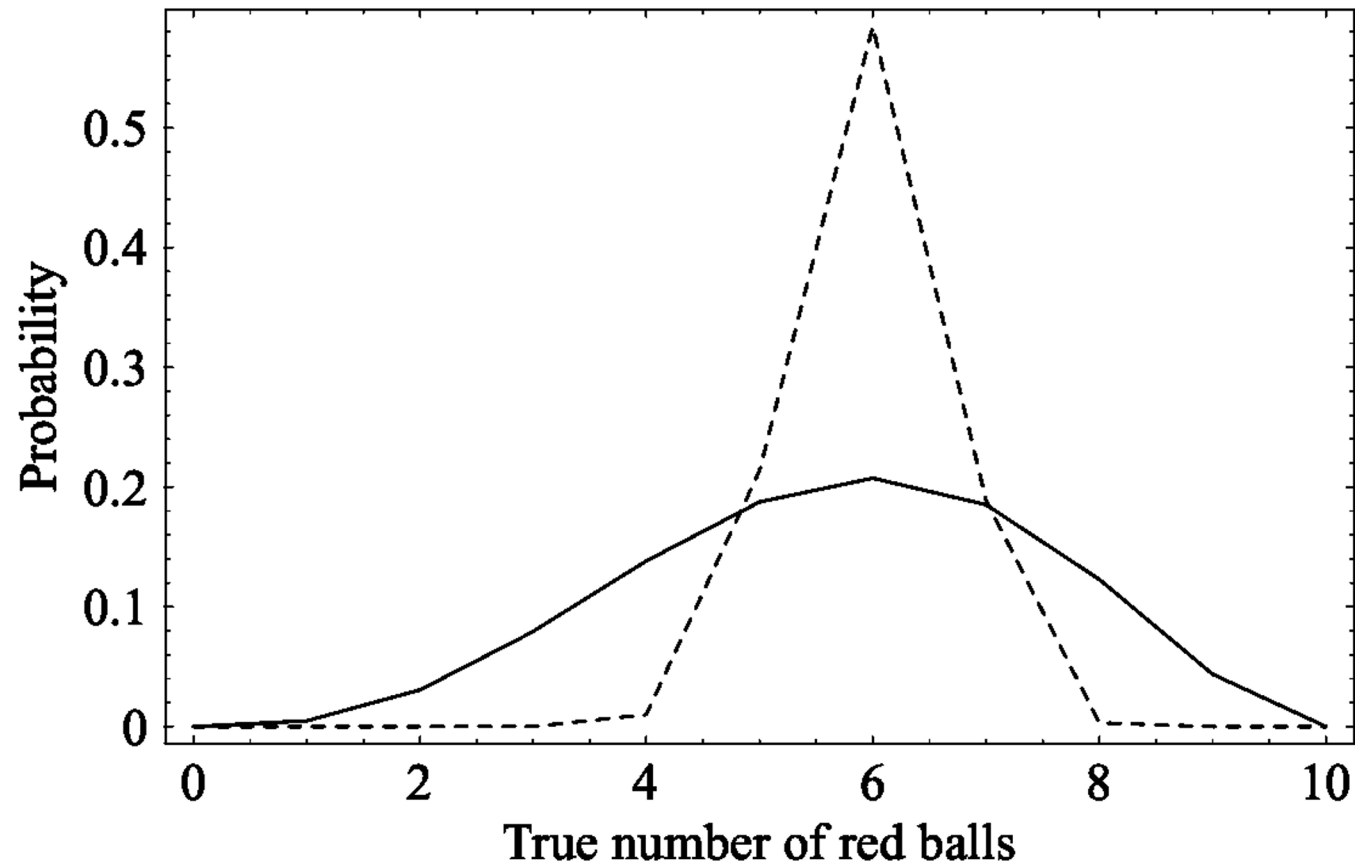
$$\binom{T}{R} \left(\frac{N}{N + M} \right)^R \left(\frac{M}{N + M} \right)^{T-R}$$

- Prior (uncertain):

- uniformly likely between 0 and $N + M$



EXAMPLE



The probability distribution of the number of red balls, for five drawings (solid curve) and 50 drawings (dashed curve).



BINOMIAL DISTRIBUTION

- 2 outcomes: 'success' or 'failure':
 - each successive trials are independent, with probability ρ
- the chance of n successes in N trials:

$$P(n) = \binom{N}{n} \rho^n (1 - \rho)^{N-n}$$

- The mean:

$$\sum_{n=0}^N n P(n) = N\rho$$

- The variance (mean square value):

$$\sum_{n=0}^N (n - N\rho)^2 P(n) = N\rho(1 - \rho)$$



EXAMPLE

- Before 1987, 4 supernovae had been recorded in 10 centuries. What, before 1987, was the probability of a bright supernova happening in the 20th century?

- Define supernova rate per century ρ , posterior:

$$P(\rho|data) \propto \binom{10}{4} \rho^4 (1 - \rho)^6 \times (\text{prior on } \rho)$$

- Take prior to be uniform, and normalize:

$$\int_0^1 P(\rho|data) d\rho = 1$$

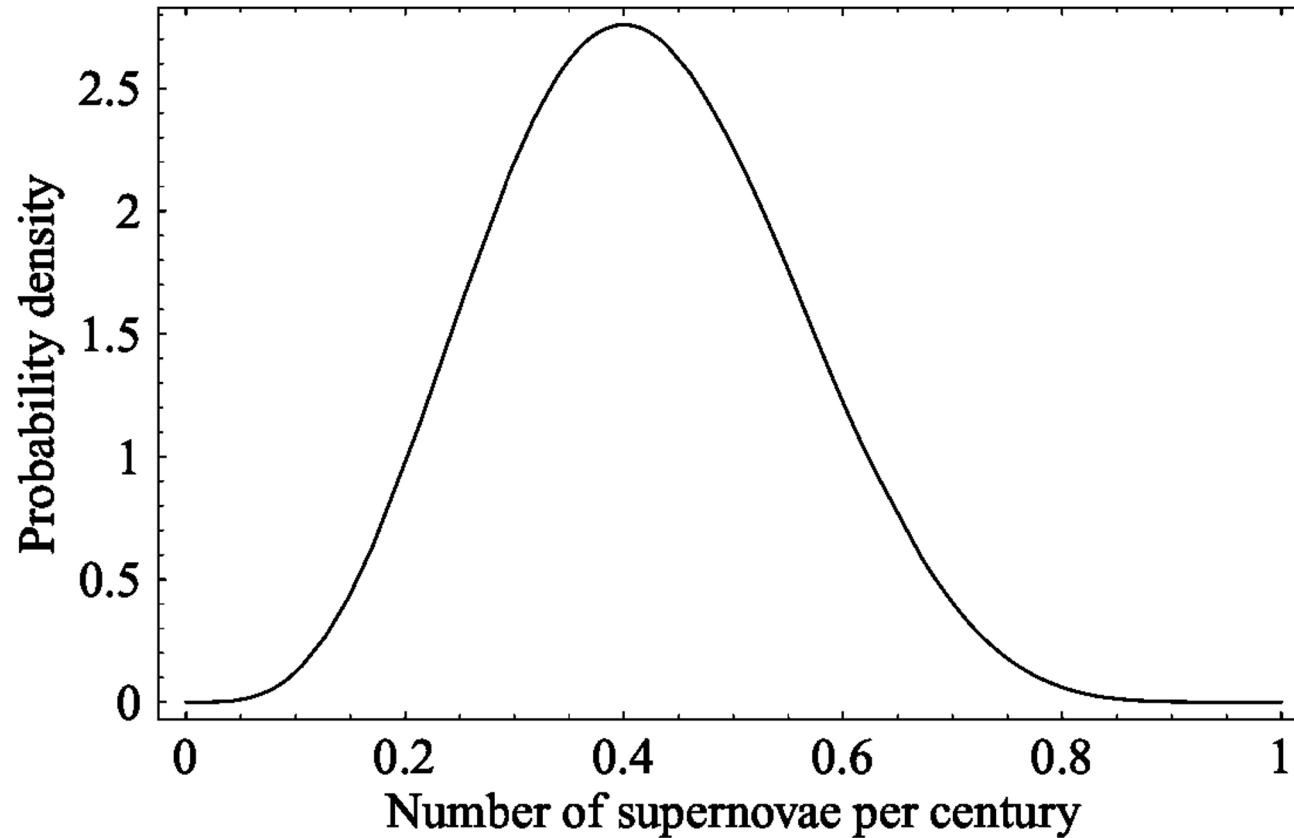
- for n supernovae in m centuries, the distribution is:

$$P(\rho|data) = \rho^n (1 - \rho)^{m-n} / B[n + 1, m - n + 1]$$

- Here $B[n, m]$ is beta function.



EXAMPLE



The posterior probability distribution for ρ , given that we have four supernovae in 10 centuries.

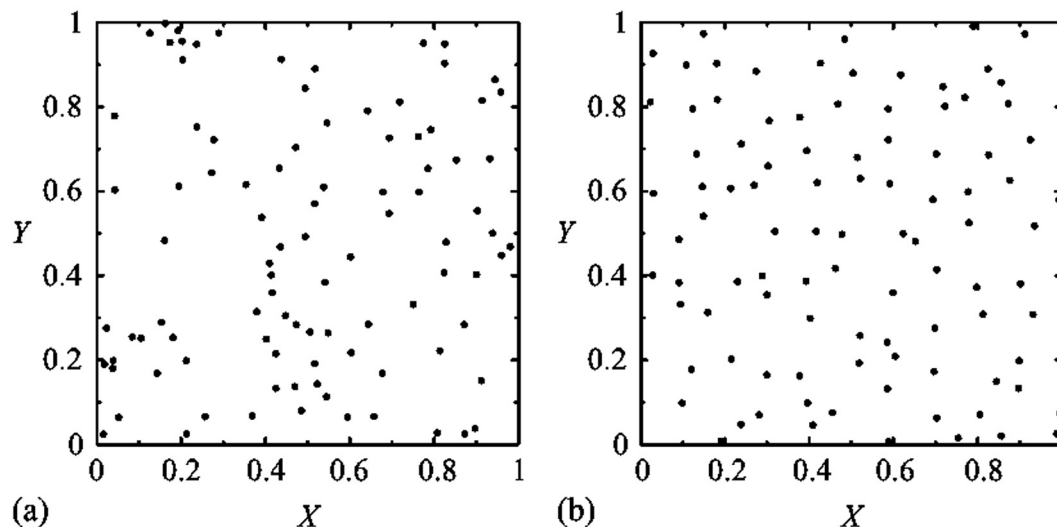


POISSON DISTRIBUTION

- derives from the binomial in the limiting case of very rare (independent) events and a large number of trials:
- Binomial: $\rho \rightarrow 0$, $N\rho \rightarrow \mu = \text{finite value}$.
- The probability of n events in a given interval, with the expectation of μ events in the same interval

$$P(n) = \frac{\mu^n}{n!} e^{-\mu}$$

- The variance of the Poisson distribution is also μ .



EXAMPLE

- Poisson statistics govern the number of photons arriving during an integration.
- The probability of a photon arriving in a fixed interval of time is (often) small. The arrivals of successive photons are independent.
- Integration over time t of photons arriving at a rate λ .
- Mean photons:

$$\mu = \lambda t$$

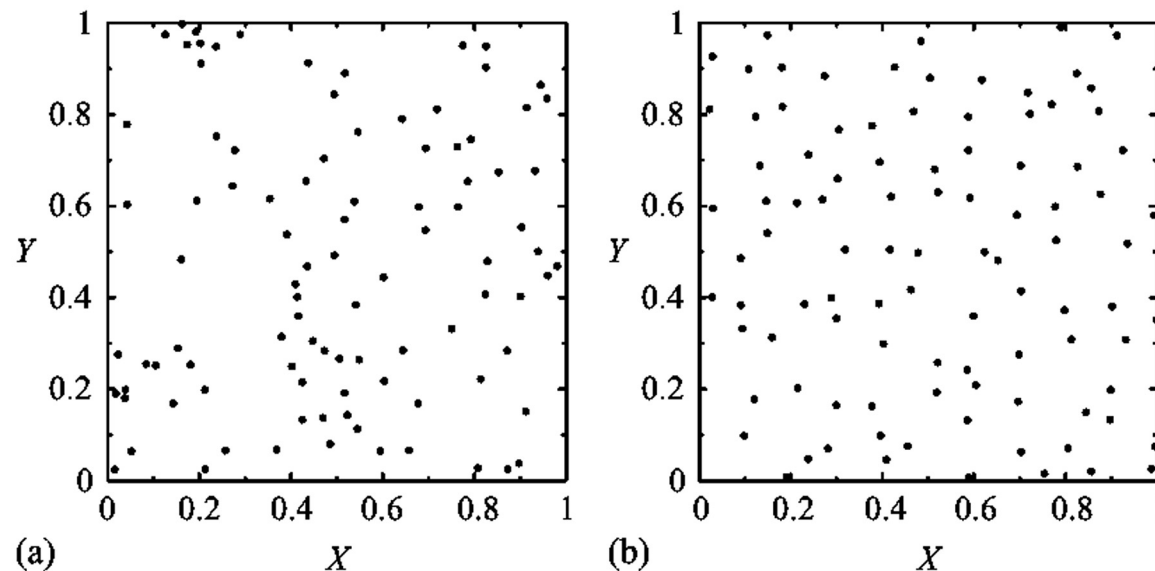
- the fluctuation on this number will be

$$\sigma = \sqrt{\mu}$$



EXAMPLE

- In large-scale structure, the galaxy distribution is discrete;
- In a voxel of volume V , $\langle N \rangle$ average galaxies;
- The variance $\sigma_N^2 = \langle N \rangle = nV$;
- "shot noise" contribution of the galaxy distribution: $1/n$



PROBABILITY DENSITY FUNCTION

- if x is a continuous random variable, then $f(x)$ is its probability density function (PDF), when:
 - $P(a < x < b) = \int_a^b f(x)dx$
 - $\int_{-\infty}^{\infty} f(x)dx = 1$
 - $f(x)$ is a single-valued non-negative number for all real x
- Cumulative probability distribution function:

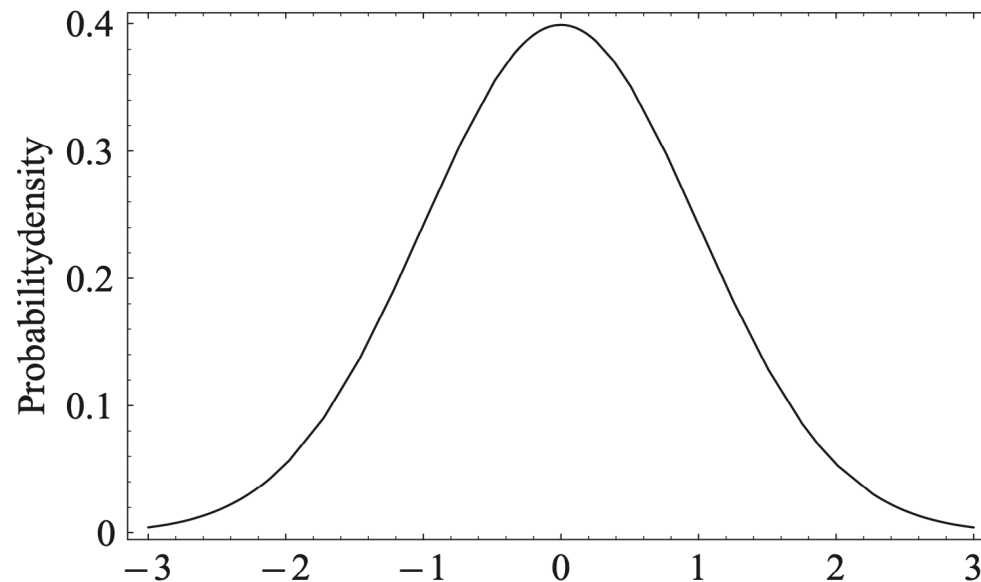
$$F(x) = \int_{-\infty}^x f(y)dy$$



GAUSSIAN DISTRIBUTION

- *Large-N limit of both binomial and Poisson distributions:*

$$P(x) = \frac{1}{\sigma\sqrt{2\pi}} \exp \left[-\frac{(x - \mu)^2}{2\sigma^2} \right]$$



the area between 1σ is 0.68; between 2σ is 0.95; and between 3σ is 0.997.



CENTRAL LIMIT THEOREM:

- Form averages M_n

$$M_n = \frac{1}{n} \sum_{i=1}^n x_i$$

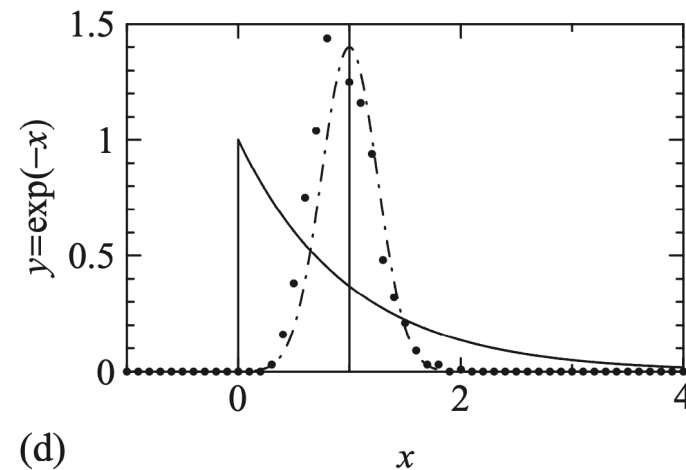
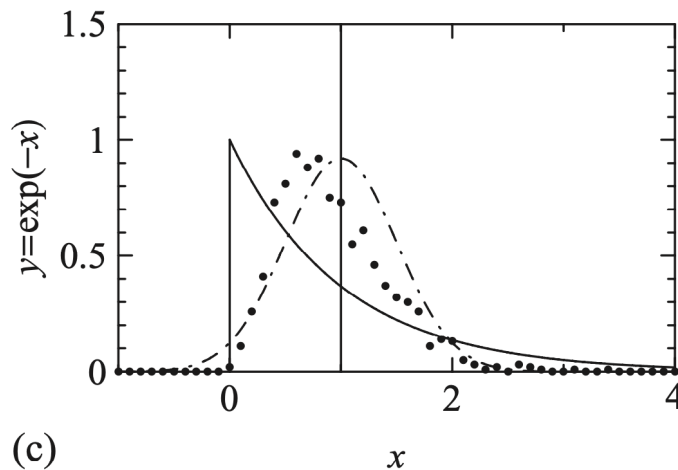
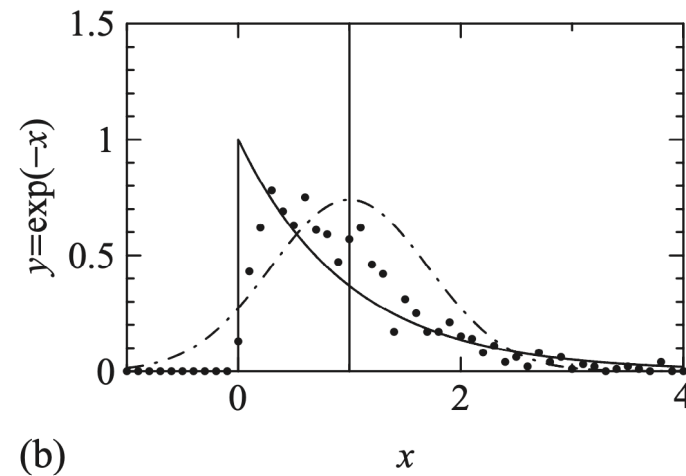
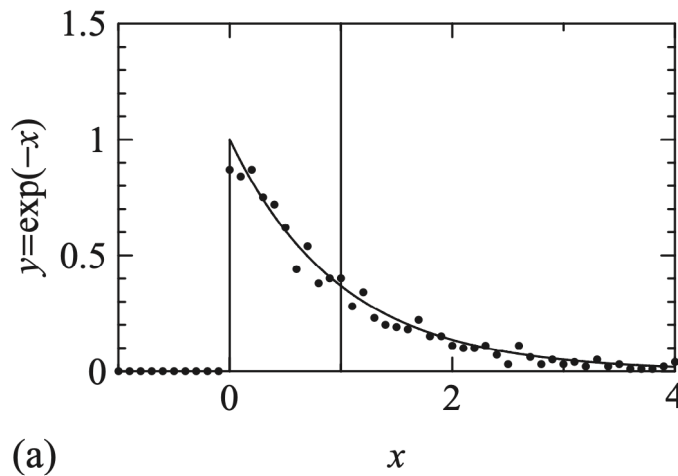
- from repeatedly drawing n samples from a population x_i with finite mean μ , variance σ^2 , then

$$\lim_{n \rightarrow \infty} \left[\frac{M_n - \mu}{\sigma / \sqrt{n}} \right] \rightarrow N(0,1)$$

- Averaging of large number of samples will produce Gaussian, **no matter** what the shape of the distribution from which the sample is drawn.
 - errors on averaged samples will always look ‘Gaussian’



CENTRAL LIMIT THEOREM:



An indication of the power of the central limit theorem. The panels show successive amounts of 'integration': in (a), a single value has been drawn; in (b), 200 values have been taken from an average of two values; (c), 200 values from an average of four; (d), 200 values from an average of 16.



STATISTICS



STATISTICS

- **Statistic:** some function of the data alone
- **Examples:**
 - *the location of the data:*
 - **Average:** $\bar{X} = 1/N \sum_{i=1}^N X_i$
 - **Median:** $X_{med} = X_j$ ($j = N/2 + 0.5$, N is odd; $N/2$ N is even)
 - **Mode:** $X_{mode} =$ value of X_i occurring most frequently, peak location in the histogram
 - *the scale or amount of scatter in the data*
 - **Mean deviation:** $\overline{\Delta X} = 1/N \sum_{i=1}^N |X_i - X_{med}|$
 - **Mean square deviation:** $S^2 = 1/N \sum_{i=1}^N |X_i - \bar{X}|^2$
 - **Root-mean-square deviation:** $rms = S$



STATISTICS

- some function f of a random variable x , with distribution function g , the **expectation value**:

$$E[f(x)] = \langle f(x) \rangle = \int f(x) g(x) dx$$

- With large # of experiments, the average of $\bar{X}$ will converge to the true mean value:

$$E[x] = \langle x \rangle = \int x g(x) dx$$

- Similarly, the statistics S^2 will converge to true variance

$$\text{var}[(x - \mu)^2] = E[(x - \mu)^2] = \int (x - \mu)^2 g(x) dx$$

- n -th central moments

$$\mu_n = \int (x - \mu)^n g(x) dx$$



REQUIREMENTS FOR STATISTICS

- unbiased

- e.g. Gaussian distribution:

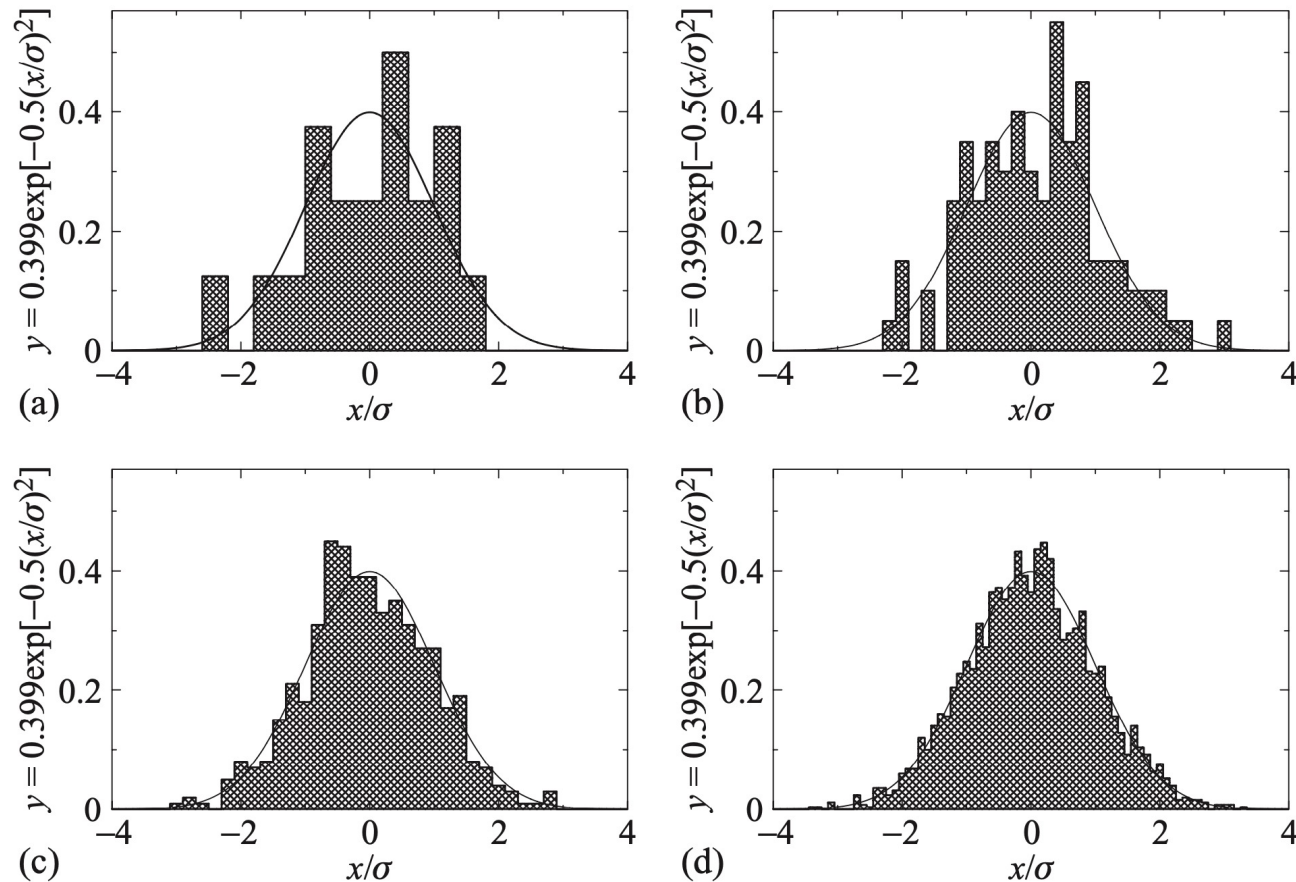
- $\bar{X} = 1/N \sum_{i=1}^N X_i$ is indeed an unbiased estimate of mean μ ;
- unbiased estimation of the population variance σ^2 (sample variance):

$$\sigma_s^2 = \frac{1}{N-1} \sum_{i=1}^N (X_i - \bar{X})^2$$

- differs from the expectation value of S^2 by the factor $N/(N-1)$
- $\bar{X}$: yields a minimum value from the sum of the squares of the deviations, thus a low estimate of the variance.



REQUIREMENTS FOR STATISTICS



x_i drawn at random from a Gaussian distribution of $\sigma = 1$: (a) 20 values, (b) 100 values, (c) 500 values, (d) 2500 values. The average values of x_i are 0.003, 0.080, 0.032 and 0.005; the median values 0.121, 0.058, 0.069 and 0.003; and the rms values 0.968, 1.017, 0.986 and 1.001. Solid curves represent Gaussians of unit area and standard deviation.



REQUIREMENTS FOR STATISTICS

- consistent:

$\lim_{n \rightarrow \infty} \text{estimator} \rightarrow \text{true value}$

- *rms is a consistent measure of the standard deviation of a Gaussian distribution for large N;*
- *but biased for small N.*

- closeness

- *smallest possible deviation from the truth*

- robust

- *robust against outliers*
- *e.g. median is far more robust than average*



RANDOM VS. SYSTEMATIC

- Variance on the average:

$$S_m^2 = E \left[\left(\frac{1}{N} \sum_{i=1}^N X_i - \mu \right)^2 \right]$$
$$\rightarrow S_m^2 = \frac{\sigma^2}{N} + \frac{1}{N^2} \sum_{i \neq j} E[(X_i - \mu)(X_j - \mu)]$$

- Neglecting the second term, the error on the average scale with $1/\sqrt{N}$.
- Second term contains the covariance:
$$\text{cov}[X_i, X_j] = E[(X_i - \mu)(X_j - \mu)]$$
- Describes the correlation between x_i and x_j .



RANDOM VS. SYSTEMATIC

- For *independent* measurements (e.g. photometric measurements of some objects):

$$\text{cov}[X_i, X_j] = \int (x_i - \mu_i)g(x_i|\cdots)dx_i \int (x_j - \mu_j)g(x_j|\cdots)dx_j = 0$$

- Here $g(x|\text{some parameters})$ describe the PDF of X .
- Random vs. systematics:
 - *random errors* decrease with larger N ($1/\sqrt{N}$ or slower);
 - *Systematic errors persist* no matter how much data are collected
 - can only be reduced by better understanding the experiments



ERROR PROPAGATION

- *measure variables $x, y, z \dots$ with independent errors $\delta X, \delta Y, \delta Z \dots$*

- *interested in some function $f(x, y, z, \dots)$, the error on f*

$$\delta f = \left. \frac{\partial f}{\partial x} \right|_{x=X} \delta X + \left. \frac{\partial f}{\partial y} \right|_{y=Y} \delta Y + \left. \frac{\partial f}{\partial z} \right|_{z=Z} \delta Z + \dots$$

- *Assuming independent error:*

$$\begin{aligned} & \text{var}[f] \\ &= \left(\left. \frac{\partial f}{\partial x} \right|_{x=X} \right)^2 \sigma_x^2 + \left(\left. \frac{\partial f}{\partial y} \right|_{y=Y} \right)^2 \sigma_y^2 + \left(\left. \frac{\partial f}{\partial z} \right|_{z=Z} \right)^2 \sigma_z^2 + \dots \end{aligned}$$



ERROR PROPAGATION

- So we can have variance inverse weighting:
 - Data with larger variance are down-weighted.
- e.g. the weighted mean

$$\bar{X}_w = \sum_{j=1}^n w_j \bar{X}_j / \sum_{j=1}^n w_j$$

- The weight $w_j = 1/\sigma_j^2$
- The variance of $\bar{X}_w$ is then

$$\sigma_w^2 = 1 / \sum_{j=1}^n 1/\sigma_j^2$$



COMBINING DISTRIBUTIONS

- Assuming we have the measured x , with PDF g , and function $f(x)$
- The PDF $h(f)$ of f could be derived because the conservation of the probability:

$$h(f)df = g(x)dx$$

- Example:
- $g(x) = \exp(-x)$ for positive x , and $f(x) = \log x$.
- It's easy to see that the PDF of f :

$$h(f) = \exp[-\exp(f)] \exp(f)$$



COMBINING DISTRIBUTIONS

- Assuming x, y follow PDF of $g(x), g(y)$, the PDF $h(z)$ of their sum $z = x + y$

$$h(z) = \int g(z - x)g(x)dx$$

- Is therefore a convolution
- Similarly, for $z = xy$:

$$h(z) = \int \frac{1}{|x|} g(x)g\left(\frac{z}{x}\right) dx$$

- And for $z = x/y$:

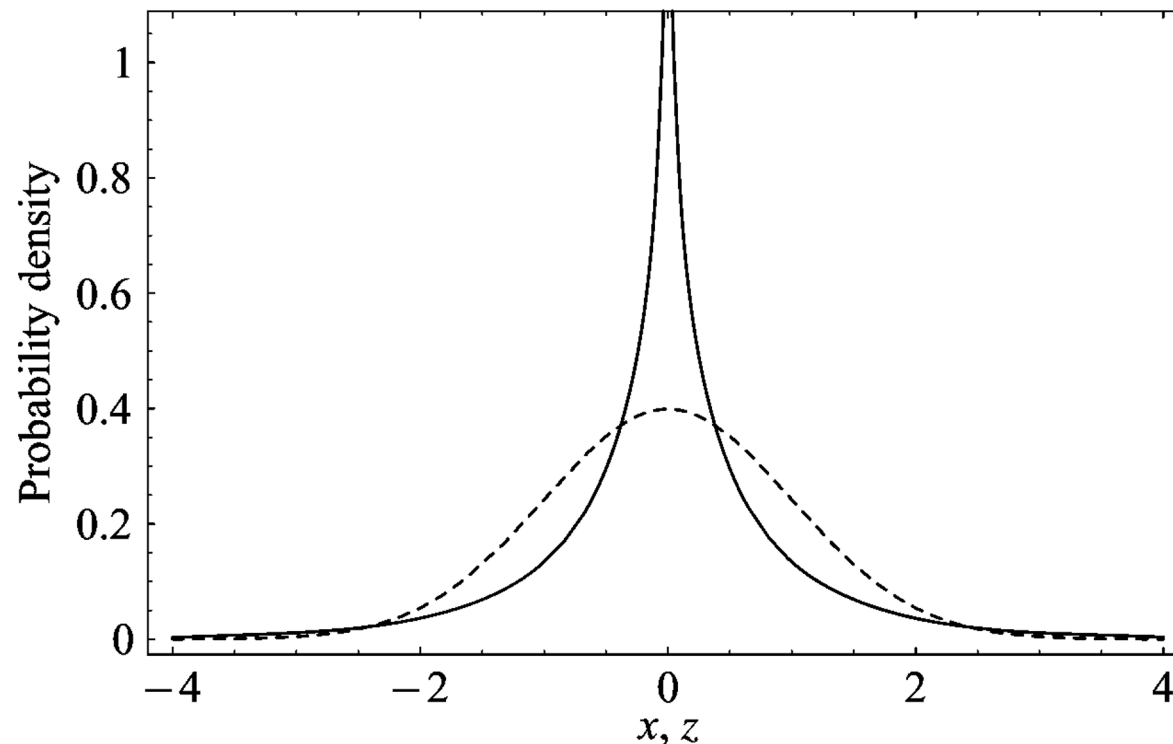
$$h(z) = \int |x|g(x)g(zx)dx$$



EXAMPLE

- *the product of two Gaussian variables of zero mean (e.g. in radio astronomy, visibility).*
- *The distribution of the product is (modified Bessel function)*

$$h(z) = \frac{2}{\pi \sigma^2} K_0 \left[\frac{|z|}{\sigma^2} \right]$$

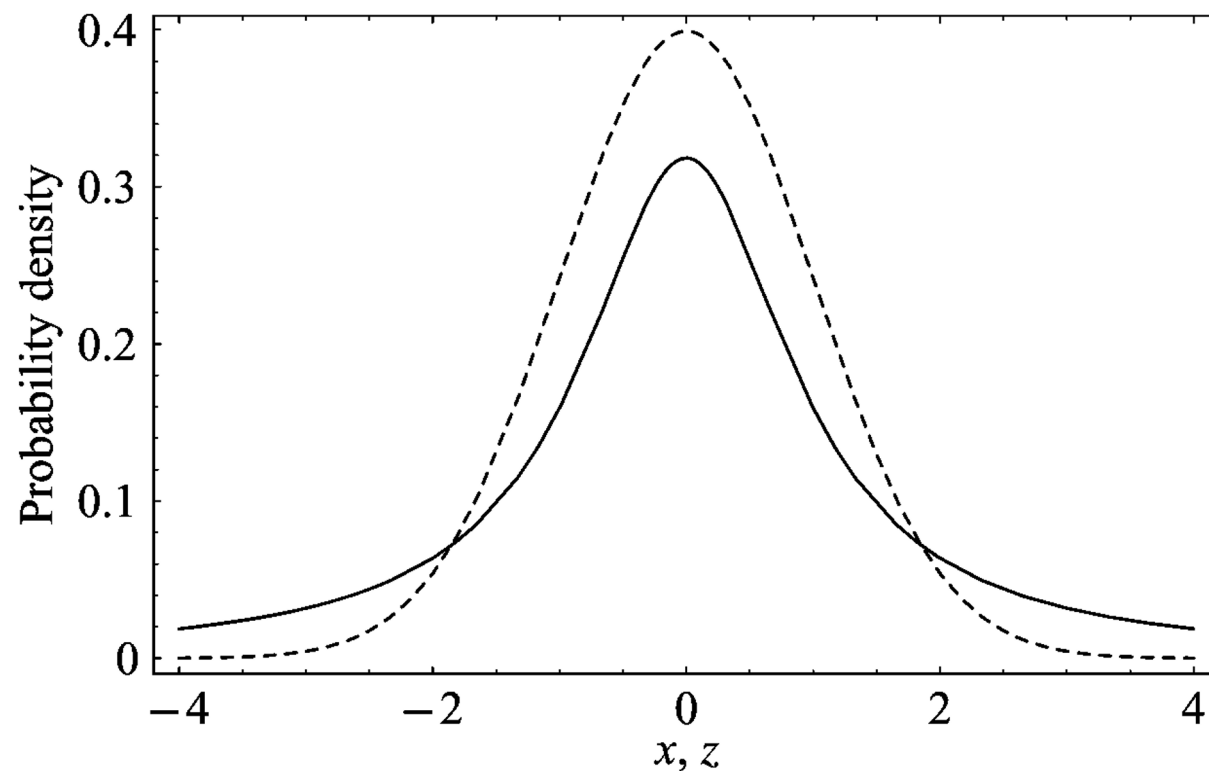


EXAMPLE

- *the ratio of two Gaussian variables of zero mean*

$$h(z) = \frac{1}{\pi} \frac{1}{1 + z^2}$$

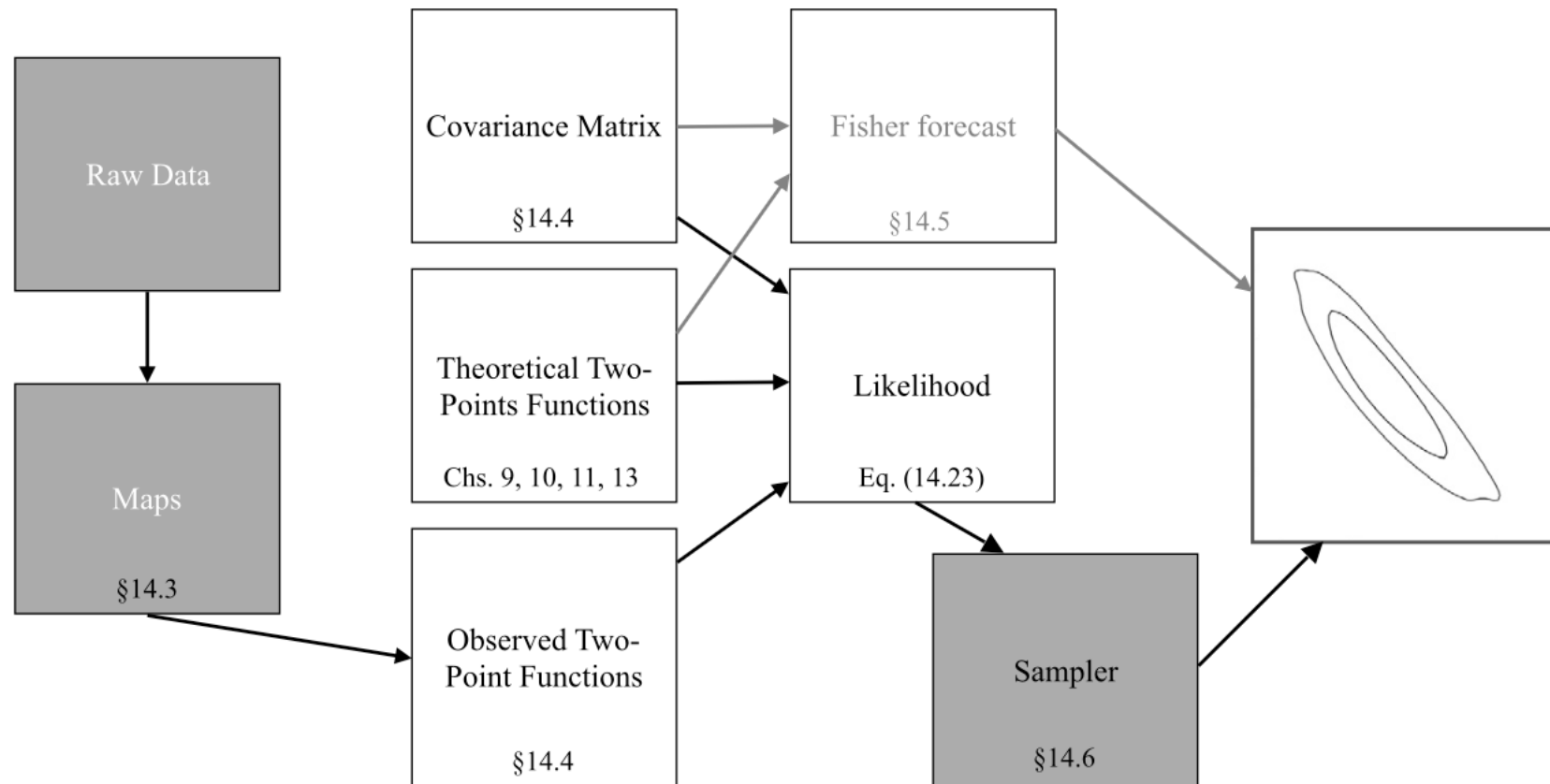
- *! Independent of σ*



LIKELIHOOD & INFERENCE



FROM RAW DATA TO PARAMETER



LIKELIHOOD



- Suppose you want to weigh *somebody*:

$$X_i = \mu(\vec{\alpha}) + n_i$$



LIKELIHOOD

- Consider N data X_i , we estimate the statistics $(1/N) \sum_i X_i$
- It's a good estimator for $\mu(\vec{\alpha})$, where $\vec{\alpha} = (\alpha_1, \dots, \alpha_n, \dots)$ are unknown parameters (slopes, intercepts etc.)
- We believe that the measurement is scattered with Gaussian error around

$$X_i = \mu(\vec{\alpha}) + n_i$$

- with Gaussian distribution

$$P(x|\vec{\alpha}) = \frac{1}{\sigma\sqrt{2\pi}} \exp \left[-\frac{(x - \mu(\vec{\alpha}))^2}{2\sigma^2} \right]$$



LIKELIHOOD

- From Bayes' theorem, the posterior probability distribution for the parameters $\vec{\alpha}$

$$P(\vec{\alpha}|X_i) \propto \prod_i \frac{1}{\sigma\sqrt{2\pi}} \exp \left[-\frac{(X_i - \mu(\vec{\alpha}))^2}{2\sigma^2} \right] P(\vec{\alpha})$$

- prior information $P(\vec{\alpha})$.
- Here μ is “given”, assuming everything depends on it being the correct model
- Leads to the maximum likelihood method and method of least squares.
- Easy to update models (posterior $\rightarrow$ prior)



MAXIMUM LIKELIHOOD

- Maximum likelihood (ML)
 - derived by Bernoulli in 1776 and Gauss around 1821
 - worked out in detail by Fisher in 1922
- From the probability density function $f(x, \alpha)$, we'd like to estimate parameter α .
- Data $X_1, X_2, \dots, X_N$ independently drawn from f , the likelihood function is

$$\mathcal{L}(X_1, X_2, \dots, X_N) = \prod_{i=1}^N f(X_i | \alpha)$$

- assuming that the priors are 'diffuse', meaning that they change little over the peaked region of the likelihood function.



MAXIMUM LIKELIHOOD

- Construct the 'best' estimate of α : peak of $\mathcal{L}$
- Define maximum likelihood estimator (MLE) $\hat{\alpha}$:

$$\left. \frac{\partial}{\partial \alpha} \ln \mathcal{L}(\alpha) \right|_{\alpha=\hat{\alpha}} = 0$$

- MLE is a statistic:
 - It depends only on the data, not on parameters.
- This estimator would have some trouble if the priors are not diffuse
 - meaning they are having as strong an effect on our conclusions as the data
- minimum variance compared to any other estimate
- not always unbiased



EXAMPLE

- Measured Y_i at given independent variables X_i , with model

$$y(a, b) = ax + b$$

- Assuming Y_i have a Gaussian scatter, each term of likelihood:

$$\mathcal{L}_i(y|(a, b)) = \exp \left[-\frac{(Y_i - (aX_i + b))^2}{2\sigma^2} \right]$$

- Maximizing the log-likelihood gives:

$$\frac{\partial \ln \mathcal{L}}{\partial a} = -2 \sum_i X_i (Y_i - aX_i - b) = 0$$

$$\frac{\partial \ln \mathcal{L}}{\partial b} = -2 \sum_i (Y_i - aX_i - b) = 0;$$

- This produce the ordinary least squares estimate:

$$\hat{a} = \frac{\overline{XY} - \bar{X}\bar{Y}}{\overline{X^2} - (\bar{X})^2}, \quad \hat{b} = \bar{Y} - \hat{a}\bar{X}$$



EXAMPLE

- *The source count of extragalactic radio sources:*

$$N(> S) = kS^{-\gamma}$$

- *N : the number of sources on a particular patch of sky with flux density greater than S .*

- *probability distribution:*

$$P(S|\gamma) = dN/dS = \gamma k S^{-(\gamma+1)}$$

- *Assuming observed M sources with flux densities S brighter than S_0 , the normalization k :*

$$\int_{S_0}^{\infty} P(S|\gamma) dS = 1 \rightarrow k = S_0^{\gamma}$$



EXAMPLE

- So the likelihood function:

$$\mathcal{L}(\gamma) = \prod_i^M P(S_i|\gamma) = \gamma^M S_0^{M\gamma} \prod_i^M S_i^{-(\gamma+1)}$$

- So the log-likelihood:

$$\ln \mathcal{L}(\gamma) = M \ln \gamma - \gamma \sum_i \ln \frac{S_i}{S_0} - \sum_i \ln S_i$$

- Assuming observed M sources with flux densities S brighter than S_0

$$\hat{\gamma} = M / \sum_i \ln \frac{S_i}{S_0}$$



EXAMPLE

- The intensity of neutrino burst after supernova decays exponentially after the core collapse. N neutrinos were detected with arrival times (in order) $T_1, T_2, \dots$

- The probability of a neutrino arriving at time t is:

$$P(t) = \exp[-(t - t_0)]$$

- for $t > t_0$ and zero otherwise. To estimate **parameter t_0** ,

$$\ln \mathcal{L}(t_0) = Nt_0 - \sum_i T_i$$

- does not appear to have a maximum by derivative;
- But clearly $t_0 < T_1$, within the range, best estimator: $\hat{t}_0 = T_1$.



DEVIATIONS & FISHER MATRIX

- MLE $\hat{\alpha}$ is distributed around true value;
- The covariance matrix of this distribution involves the curvature (Hessian matrix) of $\mathcal{L}$:

$$\mathcal{H} = \begin{bmatrix} \frac{\partial^2 \ln \mathcal{L}}{\partial \alpha_1^2} & \frac{\partial^2 \ln \mathcal{L}}{\partial \alpha_1 \partial \alpha_2} & \dots \\ \frac{\partial^2 \ln \mathcal{L}}{\partial \alpha_2 \partial \alpha_1} & \frac{\partial^2 \ln \mathcal{L}}{\partial \alpha_2^2} & \dots \\ \vdots & \vdots & \ddots \end{bmatrix}$$

- This matrix depends on the data, taking the expectation value, we have the Fisher information matrix:

$$F = E[\mathcal{H}]$$



DEVIATIONS & FISHER MATRIX

- the covariance matrix of the MLEs of the parameters:

$$C = F^{-1}$$

- Fisher matrix describes the width of the likelihood function, the scatter in the maximum-likelihood estimators

- The probability distribution of our N MLEs $\hat{\alpha}$ is then

$$P(\hat{\alpha}_1, \hat{\alpha}_2, \dots) = \frac{1}{\sqrt{(2\pi)^N |\det C|}} \exp \left[-\frac{1}{2} (\hat{\vec{\alpha}} - \vec{\alpha}) C^{-1} (\hat{\vec{\alpha}} - \vec{\alpha})^T \right]$$

- Taking the expectation value is important, as otherwise the matrix would be different for each set of data



SIMPLE EXAMPLE

- *A Gaussian of true mean μ and variance σ^2 , if we have N data X_i , the log likelihood is*

$$\ln \mathcal{L} = -\frac{1}{2\sigma^2} \sum_i (X_i - \mu)^2 - N \ln \sigma$$

- *And Fisher ‘matrix’:*

$$F = -E \left(\frac{\partial^2 \ln \mathcal{L}}{\partial \mu^2} \right) = \frac{N}{\sigma^2}$$

- *Therefore, the variance on the estimate of the mean is*

$$\frac{\sigma^2}{N}$$

- *As expected~*



EXAMPLE

- In the source-count example, we have just one parameter, the variance on $\hat{\gamma}$ is then:

$$\frac{1}{E[\partial^2 \mathcal{L}(\gamma) / \partial \gamma^2]}$$

- Which is

$$\frac{\gamma^2}{M}$$

- As long as the errors are the small, we can approximate

$$\frac{\hat{\gamma}^2}{M}$$



BAYESIAN LIKELIHOOD ANALYSIS

- *From Bayes' theorem*

$$P(\vec{\alpha}|X_i) \propto \mathcal{L}(\vec{\alpha}|X_i)P(\vec{\alpha})$$

- *Two great strengths of the Bayesian approach:*
 - *deal with nuisance parameters via marginalization*
 - *the evidence or Bayes factor to choose between models*
- *asymptotic distribution of the likelihood function:*

$$\mathcal{L}(\vec{\alpha}|X_i) = \mathcal{L}(\hat{\vec{\alpha}}|X_i) \exp\left(-\frac{1}{2}(\hat{\vec{\alpha}} - \vec{\alpha})F(\hat{\vec{\alpha}} - \vec{\alpha})^T\right)$$

- *Here F is the Fisher information matrix.*
- *This is called the Laplace approximation.*



BAYES FACTOR

- Need to check the 'fit' of the two models:
 - choice of Model A or Model B
- Using the Bayes factor, to calculate the posterior odds on Model A, compared to Model B,

$$\mathcal{P} = \frac{\int_{\alpha} p_A \mathcal{L}(X_i | \alpha, A) P(\alpha | A)}{\int_{\alpha} p_B \mathcal{L}(X_i | \alpha, B) P(\alpha | B)}$$

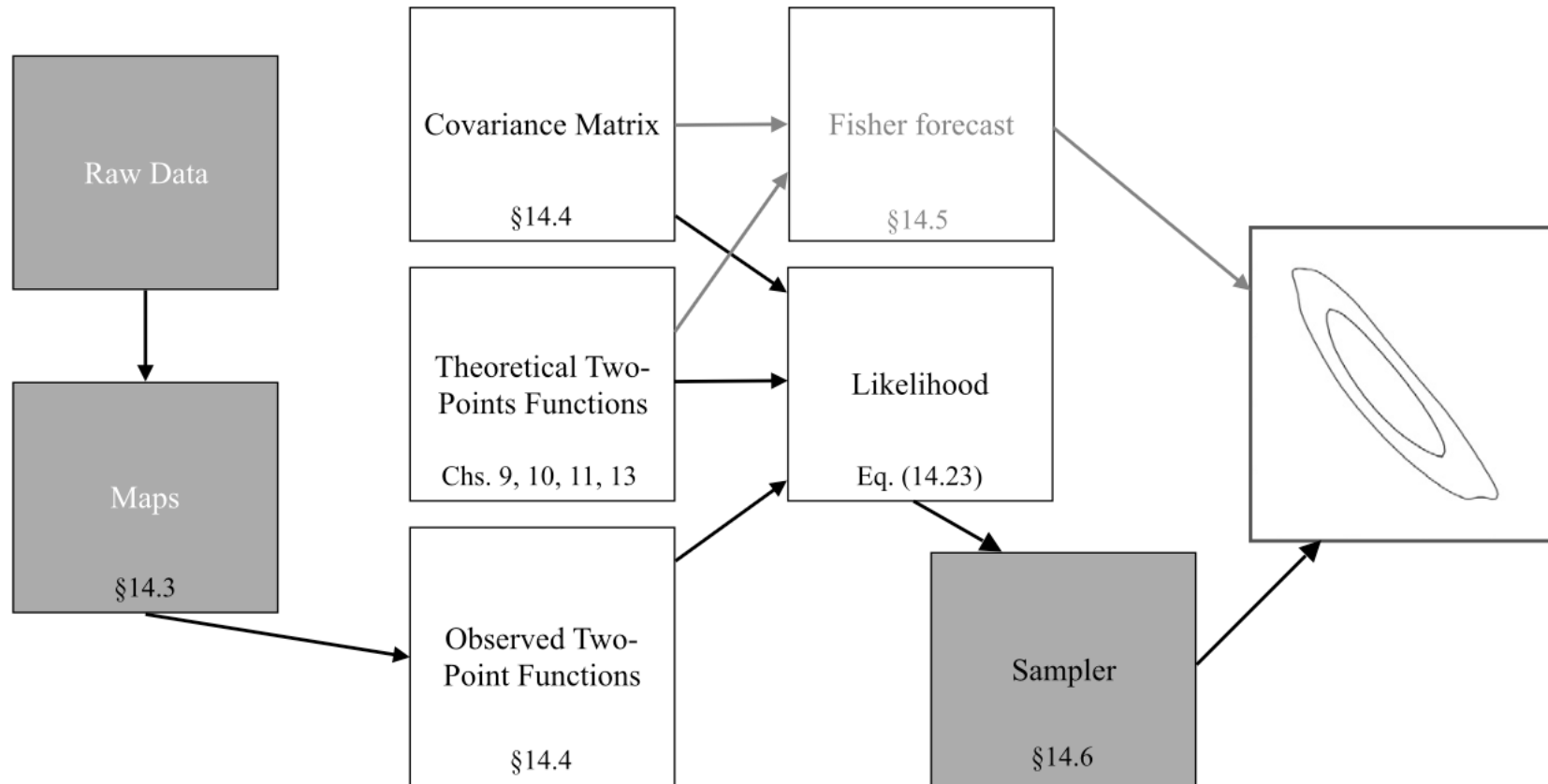
- The Integration might be cumbersome, but it's worth the effort.



COSMOLOGICAL DATA ANALYSIS



FROM RAW DATA TO PARAMETER CONSTRAINTS



$$\ln \mathcal{L}(\lambda_\alpha) = -\frac{1}{2} \sum_{ll'} \left(\hat{C}(l) - C^{\text{theory}}(l, \lambda_\alpha) \right) \left(\text{Cov}^{-1} \right)_{ll'} \left(\hat{C}(l') - C^{\text{theory}}(l', \lambda_\alpha) \right)$$



MAPMAKING

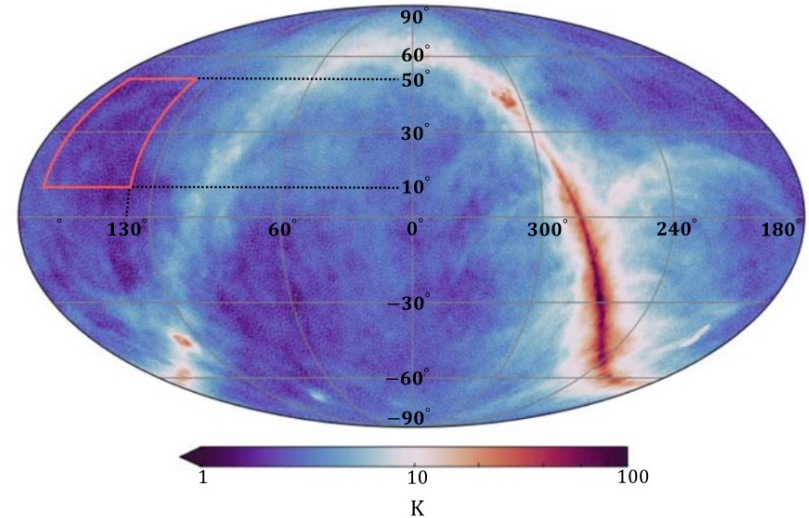


MAPMAKING

- Assume the observed time ordered data

$$d_t = P_{ti}s_i + \eta_t$$

- s_i is sky map with pixel index of i , η_t is noise, and P_{ti} is the pointing matrix.



- To extract the signal from data, consider the likelihood

$$\begin{aligned}\chi^2 &= -2 \ln \mathcal{L}(\{d_t\}|\{s_k\}) \\ &= \sum_{tt'kl} (d_t - P_{tk}s_k)(N^{-1})_{tt'}(d_{t'} - P_{t'l}s_l)\end{aligned}$$



MAPMAKING

- So the MLE is

$$\frac{\partial \chi^2}{\partial s_i} = -2 \sum_{tt'j} P_{ti} (N^{-1})_{tt'} (d_{t'} - P_{t'l} s_l) = 0$$

- Which leads to

$$\begin{aligned} \hat{s}_i &= \sum_{tt'j} (C_N)_{ij} P_{tj} (N^{-1})_{tt'} d_{t'} \\ (C_N^{-1})_{ij} &= \sum_{tt'} P_{ti} (N^{-1})_{tt'} P_{t'j} \\ &\rightarrow \hat{s} = C_N P^T N^{-1} d \end{aligned}$$



TWO-PIONT FUNCTIONS



CMB POWER SPECTRUM

- Transfer the observed map into spherical harmonics

$$a_{lm}^{\text{obs}} = \int d\Omega Y_{lm}^*(\hat{\mathbf{n}}) \Delta(\hat{\mathbf{n}})$$

- the fractional temperature fluctuation $\Delta = (T - T_0)/T_0$

$$\Delta(\hat{\mathbf{n}}) = \int d\Omega' \Theta(\hat{\mathbf{n}}') B(\hat{\mathbf{n}}, \hat{\mathbf{n}}') + \eta(\hat{\mathbf{n}})$$

- Combine two definitions, one has

$$a_{lm}^{\text{obs}} = \sum_{l'm'} B_{lm,l'm'} a_{l'm'} + \eta_{lm}$$

- isotropic beam $\rightarrow a_{lm}^{\text{obs}} = a_{lm} B_l + \eta_{lm}$



CMB POWER SPECTRUM

- The likelihood function of

$$\langle a_{lm} \rangle = 0; \quad \langle a_{lm} a_{l'm'}^* \rangle = \delta_{ll'} \delta_{mm'} C(l)$$

$$\langle \eta_{lm} \eta_{l'm'}^* \rangle = N(l) \delta_{ll'} \delta_{mm'}$$

$$P(\{a_{lm}^{\text{obs}}\} | C(l)) = \prod_{m=-l}^l \int da_{lm} P(a_{lm}^{\text{obs}} | a_{lm}) P(a_{lm} | C(l))$$

$$P(a_{lm}^{\text{obs}} | a_{lm}) = \frac{1}{\sqrt{2\pi N(l)}} \exp \left[-\frac{1}{2N(l)} |a_{lm}^{\text{obs}} - B_l a_{lm}|^2 \right]$$

- The probability $P(a_{lm} | C(l))$ is Gaussian with mean zero and variance $C(l)$;
- Carry out the integral, we have

$$\begin{aligned} \mathcal{L} \equiv P(\{a_{lm}^{\text{obs}}\} | C(l)) &= \\ &= \left(2\pi [C(l) B_l^2 + N(l)] \right)^{-(2l+1)/2} \exp \left\{ -\frac{1}{2} \sum_{m=-l}^l \frac{|a_{lm}^{\text{obs}}|^2}{C(l) B_l^2 + N(l)} \right\} \end{aligned}$$



CMB POWER SPECTRUM

- *The first derivative of the log of the likelihood*

$$\frac{d \ln \mathcal{L}}{dC(l)} = -\frac{(2l+1)B_l^2/2}{C(l)B_l^2 + N(l)} + \frac{1}{2} \sum_{m=-l}^l \frac{|a_{lm}^{\text{obs}}|^2 B_l^2}{[C(l)B_l^2 + N(l)]^2}$$

- *Setting to zero, obtain the estimator for $C(l)$ is:*

$$\hat{C}(l) = B_l^{-2} \left(\frac{1}{2l+1} \sum_{m=-l}^l |a_{lm}^{\text{obs}}|^2 - N(l) \right)$$

- *The error of this estimator:*

$$\text{Var}[\hat{C}(l)] = \langle \hat{C}(l)^2 \rangle - C(l)^2$$



CMB POWER SPECTRUM

- Expand the first term and use $\langle |a_{lm}^{\text{obs}}|^2 \rangle = C(l)B_l^2 + N(l)$

$$\begin{aligned} & \left\langle B_l^{-4} \left(\frac{1}{2l+1} \sum_{m=-l}^l |a_{lm}^{\text{obs}}|^2 - N(l) \right)^2 \right\rangle - C(l)^2 = \\ & = \left\langle B_l^{-4} \left(\frac{1}{2l+1} \sum_{m=-l}^l |a_{lm}^{\text{obs}}|^2 \right)^2 \right\rangle \longrightarrow = \frac{2l+3}{2l+1} [C(l) + N(l)B_l^{-2}]^2 \\ & - 2B_l^{-4}N(l) \left(C(l)B_l^2 + N(l) \right) + B_l^{-4}N(l)^2 - C(l)^2 \end{aligned}$$

- So the error on estimator

$$\begin{aligned} \sqrt{\text{Var} [\hat{C}(l)]} &= \sqrt{\frac{2}{2l+1} [C(l) + N(l)B_l^{-2}]} \\ \text{Cov}_{ll'} &= \frac{2}{2l+1} [C(l) + N(l)B_l^{-2}]^2 \delta_{ll'} \end{aligned}$$



GALAXY POWER SPECTRUM

- *discrete Fourier transform of the galaxy density field:*

$$\delta_g(\mathbf{k}) = L^{3/2} \sum_i^{K_{\text{grid}}^3} \delta_g(\mathbf{x}_i) e^{-i\mathbf{k} \cdot \mathbf{x}_i}, \quad \text{where } \mathbf{k} \in (n_x, n_y, n_z) k_F$$

- *The fundamental frequency $k_F = 2\pi/L$*
- *Similar to CMB $C(l)$, and setting $B_l = 1$, we have estimator*

$$\hat{P}_g(k_\alpha) = \frac{1}{m_{k,\alpha}} \sum_{\mathbf{k}}^{||\mathbf{k}| - k_\alpha| < \Delta k/2} |\delta_g(\mathbf{k})|^2 - P_N$$

- *The number of modes in each shell:*

$$m_{k,\alpha} = \frac{1}{2} \frac{4\pi k_\alpha^2 \Delta k}{k_F^3} = \frac{1}{4\pi^2} V k_\alpha^2 \Delta k.$$



GALAXY POWER SPECTRUM

- *The covariance matrix:*

$$\begin{aligned}\text{Cov}_{\alpha\beta} &\equiv \left\langle \hat{P}_g(k_\alpha) \hat{P}_g(k_\beta) \right\rangle - \left\langle \hat{P}_g(k_\alpha) \right\rangle \left\langle \hat{P}_g(k_\beta) \right\rangle \\ &= \frac{1}{m_{k,\alpha}} \sum_{\mathbf{k}}^{||\mathbf{k}| - k_\alpha| < \Delta k/2} \frac{1}{m_{k,\beta}} \sum_{\mathbf{k}'}^{||\mathbf{k}'| - k_\beta| < \Delta k/2} \left[\left\langle |\delta_g(\mathbf{k})|^2 |\delta_g(\mathbf{k}')|^2 \right\rangle - \left\langle |\delta_g(\mathbf{k})|^2 \right\rangle \left\langle |\delta_g(\mathbf{k}')|^2 \right\rangle \right]\end{aligned}$$

- *Under Gaussian assumption, this eventually leads to*

$$\text{Cov}_{\alpha\beta} = \frac{2}{m_{k,\alpha}} \left[P_g(k_\alpha) + P_N \right]^2 \delta_{\alpha\beta}$$



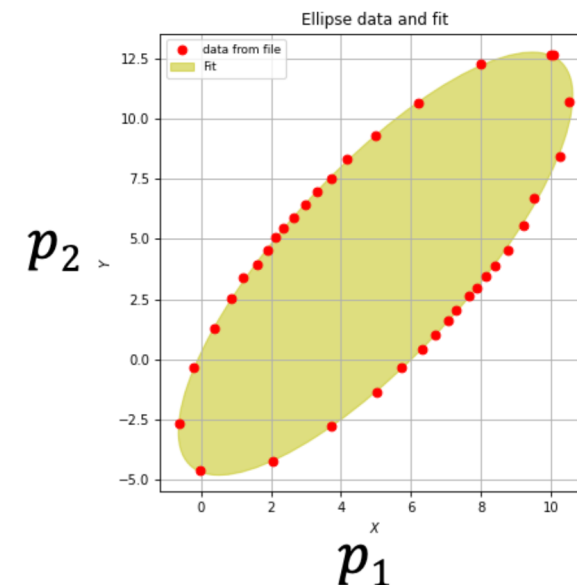
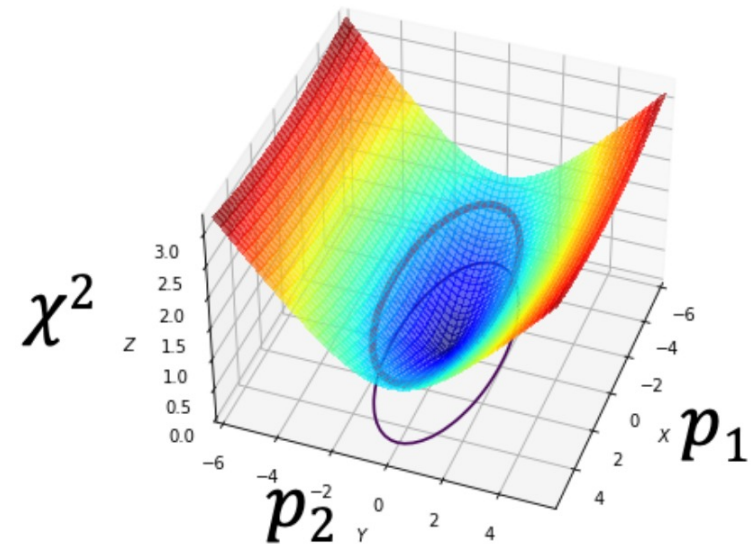
SAMPLING THE LIKELIHOOD FUNCTION



A DIRECT METHOD: GRID SCAN

- Divide parameter space into small grids
- Find the minimum chi-square $\chi^2_{\min}$
- Find confidence levels by
$$\Delta\chi^2 = \chi^2 - \chi^2_{\min}$$

- Limitation:
- Time-consuming: $t \sim n_{\text{grid}}^N$, N is the number of parameters
- Marginalization is complicated!



SAMPLING

- *Purposes:*
- *To generate samples $\{\mathbf{x}^{(r)}\}_{r=1}^R$ from a given probability distribution $P(\mathbf{x})$*
- *To estimate expectation of a function $\phi(\mathbf{x})$ under this distribution $P(\mathbf{x})$*

$$\Phi = \langle \phi(\mathbf{x}) \rangle = \int d^N x P(\mathbf{x}) \phi(\mathbf{x})$$

- *Therefore,*

$$\Phi = \frac{1}{R} \sum_r \phi(\mathbf{x}^{(r)})$$



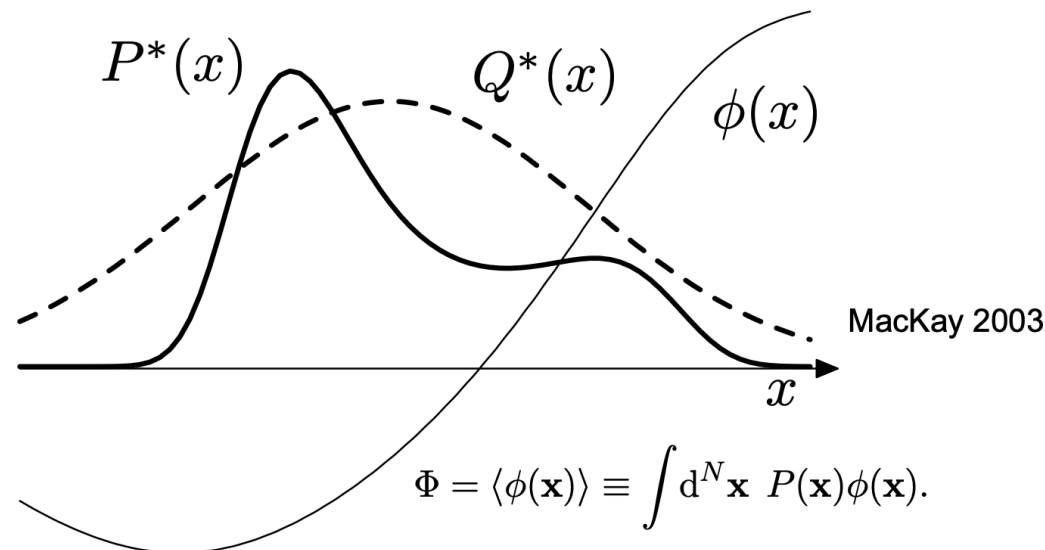
SAMPLING METHODS

- Monte Carlo (MC): *Random sampling*
 - Uniform sampling
 - Importance sampling
 - Rejection sampling
- Markov chain Monte Carlo (MCMC)
 - Metropolis-Hastings method
 - Gibbs sampling
 - Slice sampling
 - Hamiltonian Monte Carlo
 -



IMPORTANCE SAMPLING

- Generate samples from a simpler (wrong) distribution $Q(x)$
- Assign high probability to “important” values
 - adjusting the “importance” of each point by introducing a weight $w_r = P^*(\mathbf{x}^{(r)}) / Q^*(\mathbf{x}^{(r)})$
- The expectation is $\Phi = \sum_r w_r \phi(\mathbf{x}^{(r)}) / \sum_r w_r$

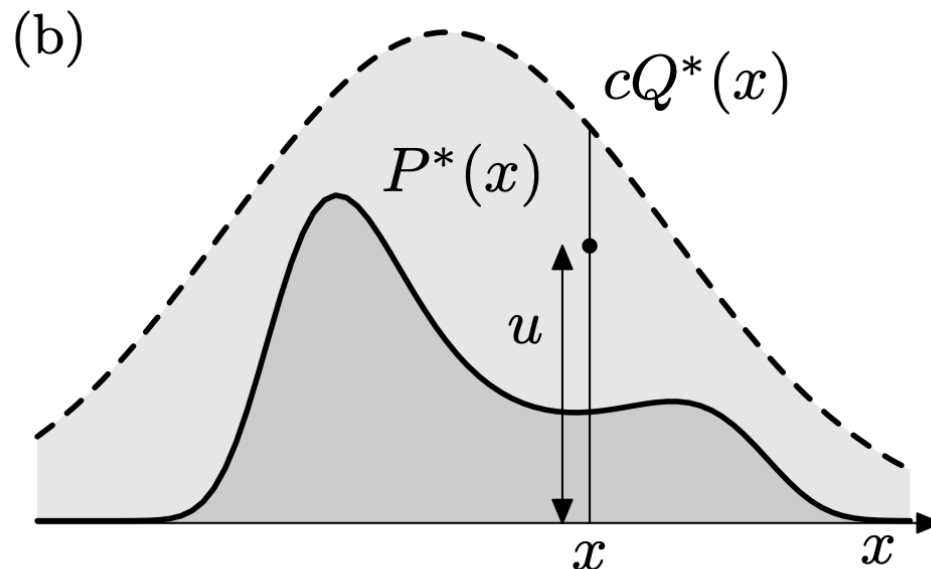
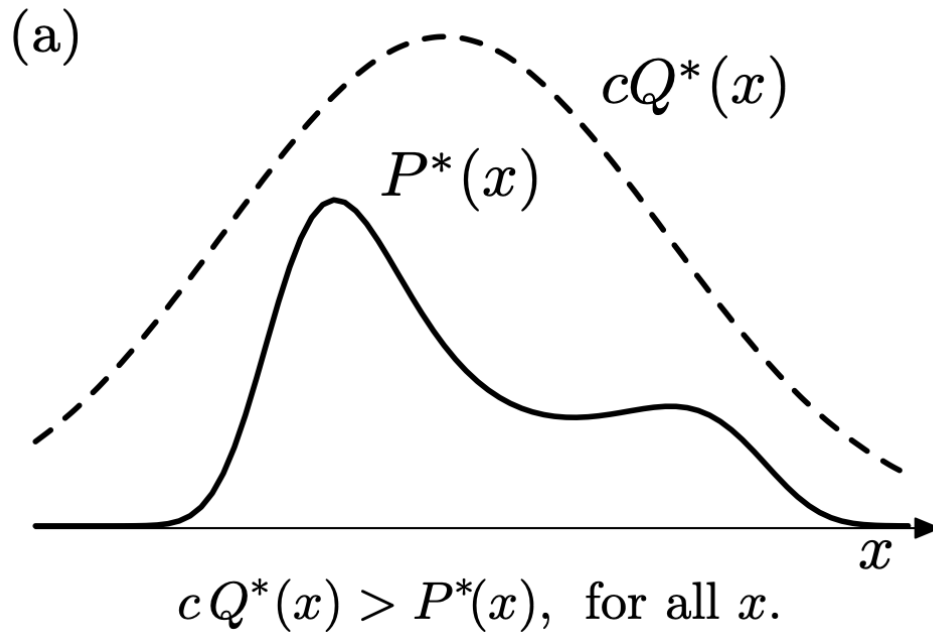


REJECTION SAMPLING

- *Selecting a simpler proposal distribution $Q(x)$, and find a constant c such that $cQ(x) > P(x)$ for all x ;*
- *Generating two random numbers:*
 - *Sample x from $Q(x)$;*
 - *Generate a uniformly distributed random variable u from the interval $[0, cQ(x)]$;*
- *If $u > P(x)$, x is rejected;*
- *else, it is accepted;*



REJECTION SAMPLING



- Work best if Q is a good approximation to P ;
- otherwise acceptance rate will be too low;
- Not suitable in high-dimension (N parameters) case

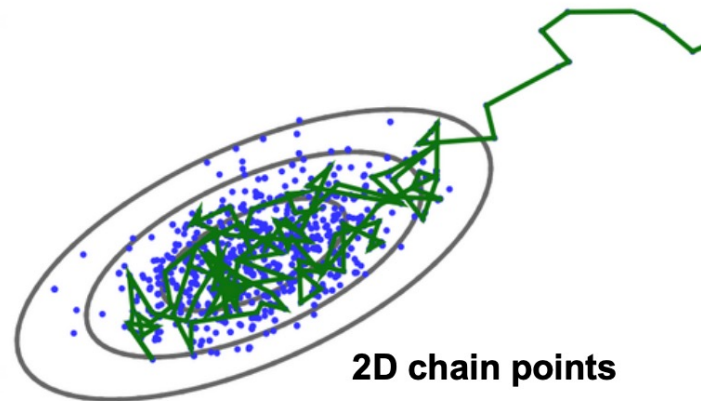


MARKOV CHAIN MONTE CARLO

- *Based on simulation;*
- *Computational time cost is approximately linearly with the number of parameters $t \sim N$;*
- *Generating samples from the full posterior distribution, easy marginalization;*
- *Algorithm:*
 - *Metropolis-Hastings, Gibbs, Slice sampling, Hamiltonian MC, ...*



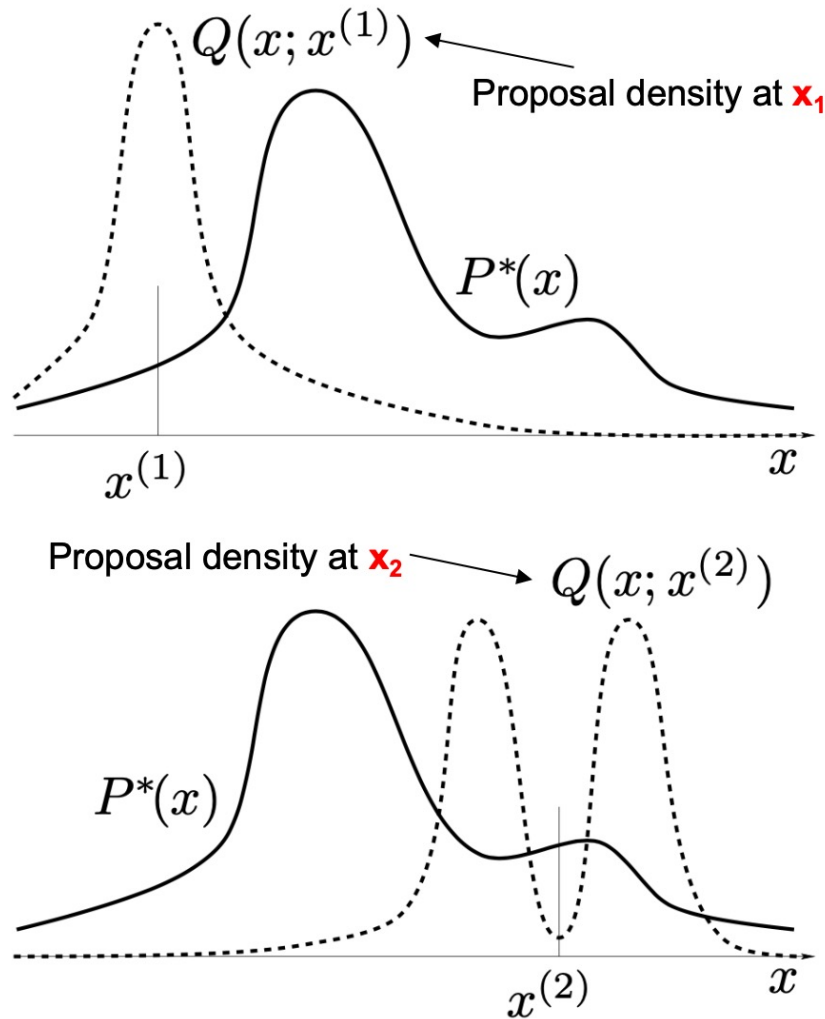
MARKOV CHAIN MONTE CARLO



- **Markov Chain:** a chain composed by a sequence of steps (or chain points), that the next 'step' in the sequence depends **only upon** the **previous one**;
- **Monte Carlo:** a computational algorithms which rely on **random sampling**, with the algorithm being guided by some rules designed to give the desired outcome



METROPOLIS-HASTINGS METHOD



- It makes use of a **proposal density Q** which depends on the current state x . Proposal density at x . Q can be **any fixed density** from which we can draw samples;
- First a new sample is proposed **based on the previous sample**, then the proposed sample is either added to the sequence or rejected depending on **the value of the probability distribution at that point**;
- It generally used for sampling from **multi dimensional distributions**, especially when the number of dimensions is high.



METROPOLIS-HASTINGS METHOD

- MH method obtains the MCMC chain points by applying the **acceptance probability**, which is defined as:

$$a(\theta_{n+1}|\theta_n) = \min \left\{ \frac{p(\theta_{n+1}|\mathbf{d}) q(\theta_n|\theta_{n+1})}{p(\theta_n|\mathbf{d}) q(\theta_{n+1}|\theta_n)}, 1 \right\}$$

- θ_n is the n th chain point (or a parameter set), $\mathbf{d}$ is the observational data, $p(\theta|\mathbf{d})$ is the **posterior distribution**, $q(\theta_{n+1}|\theta_n)$ is the **proposal density**.



METROPOLIS-HASTINGS METHOD

- Based on Bayes' theorem, *posterior distribution* can be estimated by

$$p(\theta|\mathbf{d}) = \frac{\mathcal{L}(\mathbf{d}|\theta) p(\theta)}{\int \mathcal{L}(\mathbf{d}|\theta) p(\theta) d\theta}$$

- $p(\theta)$ is the *prior probability*, and $\mathcal{L}(\mathbf{d}|\theta_n)$ is the *likelihood function*.
- When assuming an uniform prior distribution, we have $p(\theta|\mathbf{d}) \sim \mathcal{L}(\mathbf{d}|\theta_n)$
- Besides, if assume that the proposal density follows the same Gaussian distribution for every chain point, we have $q(\theta_{n+1}|\theta_n) = q(\theta_n|\theta_{n+1})$, then we find:

$$a(\theta_{n+1}|\theta_n) = \min \left\{ \frac{\mathcal{L}(\mathbf{d}|\theta_{n+1})}{\mathcal{L}(\mathbf{d}|\theta_n)}, 1 \right\}$$



METROPOLIS-HASTINGS METHOD

- 0. *Select* an initial starting point, randomly jump a few steps with a *step size*;
- 1. Based on θ_n , find θ_{n+1} by evaluating *proposal density* $q(\theta_{n+1}|\theta_n)$;
- 2. Calculate $a(\theta_{n+1}|\theta_n)$;
- 3. When $a = 1$, accept θ_{n+1} ; otherwise, accept θ_{n+1} by a probability a ;
- 4. If θ_{n+1} is accepted, $\theta_{n+1} = \theta_{n+1}$; if not, $\theta_{n+1} = \theta_n$;
- 5. repeat the first step.



```

import numpy as np
import matplotlib.pyplot as plt

# Target distribution: Univariate Gaussian (mean = 5, standard deviation = 2)
def target_distribution(x):
    return np.exp(-0.5 * ((x - 5) / 2)**2) / (2 * np.pi * 2**2)**0.5

# Metropolis-Hastings algorithm
def metropolis_hastings(iterations, proposal_std):
    samples = []
    current_sample = np.random.randn() # Initial sample from a standard normal distribution

    for _ in range(iterations):
        # Propose a new sample from a Gaussian distribution centered at the current sample
        proposed_sample = current_sample + np.random.normal(scale=proposal_std)

        # Calculate acceptance ratio
        acceptance_ratio = min(1, target_distribution(proposed_sample) / target_distribution(current_sample))

        # Accept or reject the proposed sample based on the acceptance ratio
        if np.random.rand() < acceptance_ratio:
            current_sample = proposed_sample

        samples.append(current_sample)

    return samples

# Parameters
iterations = 10000
proposal_std = 1.0

# Generate samples using Metropolis-Hastings algorithm
samples = metropolis_hastings(iterations, proposal_std)

# Plot the histogram of generated samples and the target distribution
plt.hist(samples, bins=50, density=True, alpha=0.6, label='Metropolis-Hastings Samples')
x_range = np.linspace(min(samples), max(samples), 100)
plt.plot(x_range, target_distribution(x_range), 'r', label='Target Distribution')
plt.legend()
plt.xlabel('Sample Value')
plt.ylabel('Density')
plt.title('Metropolis-Hastings Sampling')
plt.show()

```

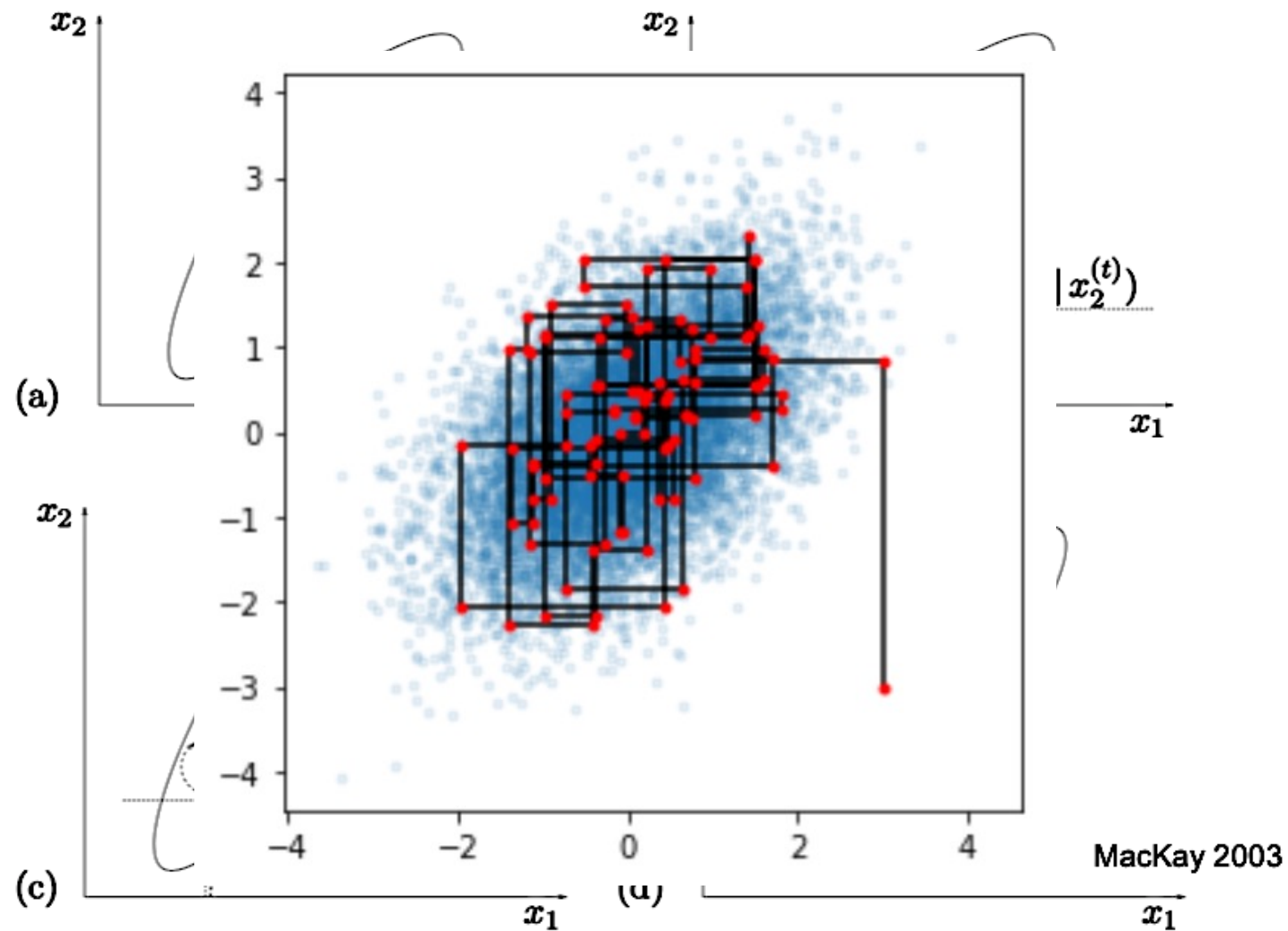


GIBBS SAMPLING

- A method for sampling from distributions over *at least two dimensions*;
- It can be viewed as a *special case of the Metropolis method*, in which a sequence of proposal distributions Q are defined in terms of the *conditional distributions* of the joint distribution $P(x)$;
- It assumed that, although $P(x)$ is *too complex* to draw samples from directly, its *conditional distribution* $P(x_i | \{x_j\}_{j \neq i})$ *are tractable* to work with.



GIBBS SAMPLING



SLICE SAMPLING

- *It can be applied when the target density $P(x)$ can be evaluated at any point x ;*
- *Step-size is less important than Metropolis method;*
- *No requirement that the one-dimensional conditional distributions be easy to sample from, like Gibbs sampling;*
- *Similar to rejection sampling, but no requirement for an upper-bounding function.*



CONVERGENCE CRITERION

- Consider M **parallel chains** ($j = 1 \cdots M$), each chain contains N points ($i = 1 \cdots N$), then the chain element is y_i^j (a point in parameter space);

- Define the mean of **the chain**:

$$\bar{y}^j = \frac{1}{N} \sum_{i=1}^N y_i^j$$

- Define the mean of **all chains**:

$$\bar{y}^j = \frac{1}{NM} \sum_{i,j=1}^{N,M} y_i^j$$



CONVERGENCE CRITERION

- *Then the variance between chains:*

$$B_n = \frac{1}{M-1} \sum_{j=1}^M (\bar{y}^j - \bar{y})^2$$

- *The variance within a chain:*

$$W = \frac{1}{M(N-1)} \sum_{ij} (y_i^j - \bar{y}^j)^2$$

- *Define the quantity:*

$$\hat{R} = \frac{[N(N-1)]W + B_n(1 + 1/M)}{W}$$

- *The MCMC chains converge when:*

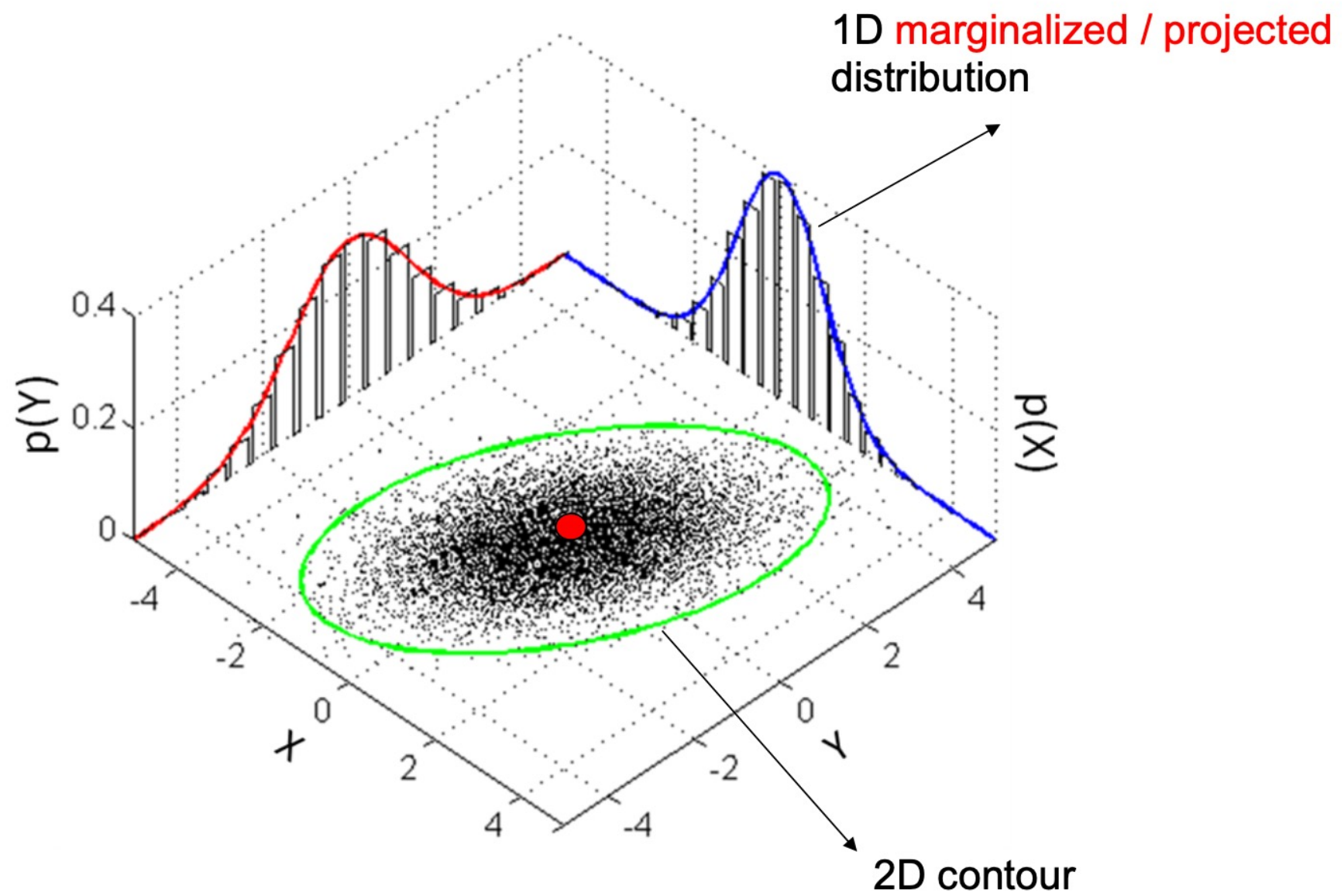
$$\hat{R} < 1.1$$



AFTER OBTAINING THE MCMC CHAINS

- **Burn-in period:** although the Markov chain eventually converges to the desired distribution, the **initial samples** may follow a very different distribution, especially if the starting point is in a region of low density. As a result, a burn-in period is typically necessary, where **an initial number of samples are thrown away (usually the first 100-1000 samples)**.
- **Thinning process:** after burn-in, in order to obtain independent samples, we should **only keep every n th sample in the chains (usually $n = 10 - 100$)**.
- Finally, about **$10^4 - 10^5$ samples** or chain points are needed to illustrate the posterior distribution.





THANK YOU

