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Bayesian inference for neutrino physics



Probability of the hypothesis H given the data D = "posterior probability on H"



Probability of the hypothesis H given the data D = "posterior probability on H"



= "posterior probability on H"



• Derivation from conditional probabilities

• Probability to observe A and B:

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Probability to observe B

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• Physical interpretation

• In physics, we often have A as the hypothesis H, and B as the data D:

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

• The posterior probability is the prior probability when it is weighted by the likelihood of observing the data, normalised by the probability of observing any data

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- Also referred to as "updating belief on H "

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• Example from neutrino physics

- We do not know $\delta_{CP} \rightarrow$ our prior probability on the parameter is flat
- We compute the likelihood of observing the ν_e and $\bar{\nu_e}$ spectra according to several δ_{CP} hypothetical values
- We measure the posterior probability for each hypothesis
- We estimate which value of δ_{CP} corresponds to the highest posterior probability

Posterior probability sampling

• Simple 1 parameter case:

- Chose a grid, and evaluate the posterior probability at each point e.g. $P(\delta_{CP,i}|D)$ for $i \in [-\pi, \pi]$ with steps of $\pi/100$
- Easy to extend to 4 oscillation parameters

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• But: systematics!

- At each step, one needs to evaluate the posterior probability varying the systematical parameters as well (throws)
- Neutrino physics → Ø(100) systematical parameters (flux, interaction model, detector response...)
- The real posterior probability is: $P(\delta_{CP,i}, \overrightarrow{\varsigma_j} | D)$ where $\overrightarrow{\varsigma_i}$ is the vector of systematical uncertainties
- Grid searches can become computationally very expensive (many points to evaluate)

Markov Chain Monte-Carlo

• Markov Chain Monte-Carlo (MCMC)

- Alternate procedure to grid to sample the space of oscillation parameters $\vec{\vartheta}$ and systematics parameters $\vec{\zeta}$
- Semi-random walk in the parameter space
- Stochastic model: randomness of throws
- Sequential process: the state of a throw only depends on the throw before

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• Metropolis-Hastings algorithm

- Most generic implementation of Markov Chain Monte-Carlo (MCMC)
- The semi-random walks is *proportional* to the target distribution
- The collection of steps are *samples* from the posterior distribution
- Ensure 2 specific properties of the MCMC: *Aperiodicity*: do not oscillate between same values *Ergodicity*: can converge to the stationary distribution

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$$r < u \rightarrow$$
 reject step $i + 1$

count again step i

- $r \ge 1 \rightarrow \text{accept step } i+1$
- Iterate process until obtaining enough step to analyse the distribution



• The detailed balance equation ensure that the samples follow the target distribution

• The acceptance function is: $A(x_{i+1}, x_i) = min(1, r)$

$$r \ge 1 \rightarrow A(x_{i+1}, x_i) = 1$$
$$r < 1 \rightarrow A(x_{i+1}, x_i) = r$$

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- Defining the probability to transition to the step x_{i+1} , i.e. the transition probability: $T(x_{i+1} | x_i) = J(x_{i+1} | x_i) A(x_{i+1}, x_i)$

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• Interpretation: if we propose a step with $G(x_{i+1}) > G(x_i)$

The acceptance function is: $A(x_{i+1}, x_i) = 1$

The transition probability is: $T(x_i | x_{i+1}) = \frac{G(x_{i+1})}{G(x_i)}$

→ The probability to jump back on the previous step is proportional to the ratio of G(x) value

Application for neutrino physics

$\circ~$ In the case of neutrino physics

- The target distribution is the posterior probability on the oscillation parameters $\vec{\vartheta}$ and systematics parameters $\vec{\zeta}$
- All parameters are treated the same, they just have different prior probabilities
- All parameters are inferred at the same time
 - → joint analysis of near and far detector values



• Jump function parameter

• The jump function can be symmetrical \rightarrow Metropolis algorithm

or asymmetrical \rightarrow Hastings addition

- The jump function has a width parameter:
 - \rightarrow this is referred to as the step size
 - \rightarrow its value is heuristic, although literature exist about its optimisation
 - → strongly impacts the convergence rate of the chain



Burn-in

• The Markov chain takes time to reach equilibrium

- The chain can start far from the target distribution
 → creates a bias towards initial values
- The first values must be discarded: "burn-in"
- The burn-in size can be determined from the trace of the chain



Autocorrelation

• The steps are correlated between them

- Independent samples can be selected by subsampling the chain
- Value of subsampling order can be determined from the autocorrelation function

$$\mathscr{A}(k) = \frac{\varrho(k)}{\varrho(0)}$$

where:

$$\varrho(k) = \mathbb{E}(x_i - \bar{x}) \mathbb{E}(x_{i+k} - \bar{x})$$
$$= \frac{1}{N-k} \sum_{i=1}^{N-k} (x_i - \bar{x})(x_{i+k} - \bar{x})$$

E = expectation value



(b) Δm_{32}^2

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Convergence tests

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- Test: comparison of independent chains



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- Test: comparison of independent chains

• Geweke diagnostic

- Compare the beginning and the end of a Markov chain
- Select 5% of the chain from its beginning and increment of 5% e.g. [0-5%], [5-10%], ..., [45-50%] and compare with remaining 50% of the chain: [50-100%]
- Useful to determine burn-in value and spot issues

$$G = \frac{\bar{x}_{ini} - \bar{x}_{fin}}{\sqrt{\sigma(x)_{ini}^2 + \sigma(x)_{fin}^2}}$$

Note: 5% is not a hard rule, other binning can be chosen



Credible intervals

- In Bayesian probabilities, results are given as credible intervals
 - Area where there is the highest probability that the true value lies in
 - E.g. there is a 90% probability that the true value of $\sin^2 \theta_{23}$ is in [0.42, 0.59]



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- The posterior probabilities are automatically marginalised
 - When projecting in lower dimensions than the Markov chain, the shape of the posterior probabilities of the other parameters is included in the integral



 θ_{13}

sin²

marg. post. proba. density

80.0

0.0

0.06



 $\sin^2 \theta_{23}$

350

300

Bayes factor

• Comparison of 2 hypotheses

• If we have 2 hypotheses H_1 and H_2 , we can compare them with the Bayes factor, i.e. the ratio of marginalised likelihood

• Bayes factor:
$$B_F = \frac{P(D \mid H_1)}{P(D \mid H_2)}$$

- If the prior probabilities are the same, this is equivalent to the ratio of posterior probabilities
- Example: the Bayes factor for normal ordering is $B_F = 3.72$ on this plot



Changing the prior

• The posterior probability can be evaluated for a different definition of the prior

- Equivalent to a variable change of the distribution: prior in $x \rightarrow \text{prior}$ in y = f(x)
- Need to evaluate the Jacobian of the transformation:

$$P(H(x)) \rightarrow P(H(y)) = P(H(x)) |J(y)|$$
$$= P(H(x)) |\frac{\partial x}{\partial y}|$$

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• A useful way to:

- Check the robustness of the prior
- Answer a different question e.g. what is the probability of CP-violation (instead of what is the δ_{CP} value)



Comparison with frequentist stat.

	Frequentist	Bayesian
probability	frequency of occurence	degree of belief
parameters	fixed (once chosen)	uncertain
observation	fluctuates	certain (once observed)

The two approaches in a nutshell:

- frequenstist \rightarrow probability of observation, given a model
- bayesian \rightarrow probability of a model, given an observation

Methodologies

- frequenstist: estimates frequencies, by emulating repetitions of the experiment (toys) for a given parameter, using the likelihood as PDF
- bayesian: exploits the Bayes theorem to compute the posterior
 P(para|obs), using the prior P(para) and P(obs|para) the likelihood
- **3.** Both approaches get unifed when
 - there is an infinite number of measurements
 - the prior is uniform: $P(par|obs) = A \times \mathcal{L}(par; obs)$

(same equation, but its meaning and the question it addresses are different)

Slide by Romain Madar, School of Statistics 2022

• Bayesian inference consist in computing a posterior probability density

- Update the probability of a hypothesis according to the information on the data
- Markov Chain Monte-Carlo is a useful tool to sample high dimensional cases
- Can infer any shape of posterior probabilities

• The process requires careful tuning

- Asymptotically, MCMC properties ensure that it will converge to the target distribution
- We do not have infinite time, neither an infinite number of processors
- Ensuring convergence is key to the process
 - \rightarrow convincing ourselves that the output is the needed one is not easy!
- Extensive literature about it, but no « one-solution-fit-all »
- Does not mean it should no be used! But not blindly