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Programming in Python¹

Mattia Monga

Dip. di Informatica Università degli Studi di Milano, Italia mattia.monga@unimi.it

Academic year 2024/25, I semester

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How science works

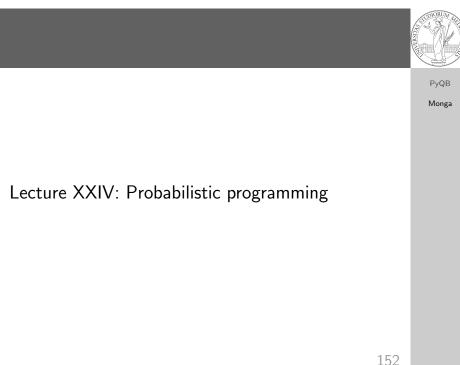


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Describing one single "scientific method" is problematic, but a schema many will accept is:

- Imagine a hypothesis
- ② Design (mathematical/convenient) models consistent with the hypothesis
- ③ Collect experimental data
- ④ Discuss the fitness of data given the models

It is worth noting that the falsification of models is not *automatically* a rejection of hypotheses (and, more obviously, neither a validation).



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The role of Bayes Theorem

In this discussion, a useful relationship between data and models is Bayes Theorem.

$$P(M,D) = P(M|D) \cdot P(D) = P(D|M) \cdot P(M)$$

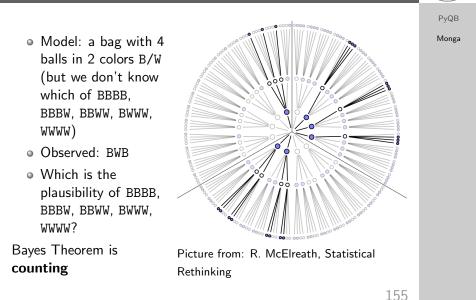
Therefore:

$$P(M|D) = \frac{P(D|M) \cdot P(M)}{P(D)}$$

The plausibility of the model given some observed data, is proportional to the number of ways data can be *produced* by the model and the prior plausibility of the model itself.

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Simple example



Classical binomial example

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- Which is the proportion p of water covering Earth? The models are indexed by the float 0
- Given *p*, the probability of observing some W,L in a series of **independent random observations** is:
- $P(W, L|p) = \frac{(W+L)!}{W! \cdot L!} p^W \cdot (1-p)^L$ (binomial distribution).
- Do we have an initial (prior) idea?
- Make observations, apply Bayes, update prior!





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This Bayesian strategy is (conceptually) easy to transform in a computational process.

- Code the models
- ② Run the models
- 3 Compute the plausibility of the models based on observed data

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A conventional way of expressing the model



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 $W \sim Binomial(W + L, p)$ $p \sim Uniform(0, 1)$

Probabilistic programming is systematic way of coding this kind of models, combining predefined statistical distributions and Monte Carlo methods for computing the posterior plausibility of parameters.

In principle you can do it by hand



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import pymc as pm

PyMC

W, L = 7, 3earth = pm.Model() with earth: p = pm.Uniform("p", 0, 1) # uniform prior w = pm.Binomial("w", n=W+L, p=p, observed=W) posterior = pm.sample(2000)

posterior['p']

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def dbinom(success: int, size: int, prob: float) -> float: fail = size - success return math.factorial(size)/(math.factorial(success)*math.factorial(fail))*prob**succ | \hookrightarrow ess*(1-prob)**(fail) Then, W, L = 7, 3 # for example 'WWWLLWWLWW' p_grid = np.linspace(start=0, stop=1, num=20) prior = np.ones(20)/20likelihood = dbinom(W, n=W+L, p=p_grid) unstd_posterior = likelihood * prior posterior = unstd_posterior / unstd_posterior.sum() Unfeasible with many variables!

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