

6/10/19 Bayes2019 TALENT Lecture M1b

• Notebooks:

- (A) • From M1a - topics/basics-of-bayesian-statistics/Exploring-pdfs.ipynb
 - topics/bayesian-parameter-estimation/
 - (B) parameter-estimation-in-bayesTALENT-intro.ipynb
 - (C) parameter-estimation-fitting-straight-line-I.ipynb

• Topic: Parameter Estimation I (of 3)

• Overview comments

- In general terms, "parameter estimation" in physics means obtaining values for parameters (constants) that appear in a theoretical model that describes data. (Exceptions exist, of course)
- Conventionally this process is known as "fitting the parameters" and the goal is to find the "best fit".
- We will make particular interpretations of these phrases from our Bayesian point of view.
- Today we'll set up the problem and look at how familiar ideas like "least-squares fitting" show up from a Bayesian perspective.

• There are many examples of where parameter estimation is needed in low-energy nuclear physics (as examples) and every other subfield of physics.

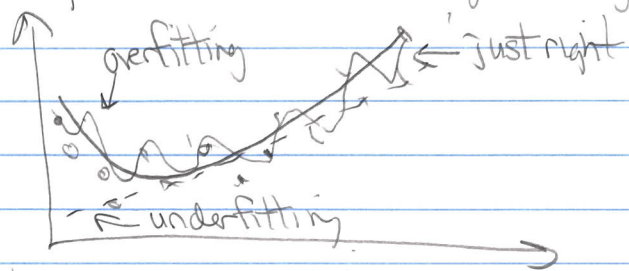
we'll do a toy model of this to explore issues

- the parameters in an effective field theory Hamiltonian (chiral, pions, halo, deformed nuclei) - usually called low-energy constants or LECs
- parameters that define an energy density functional (eg. Skyrme type)
- parameters in an optical potential used in reactions calculations
- parameters in a model for extrapolating to an infinite model space (eg. no-core shell model)
- and so on.

• As we proceed, we will make the case that a Bayesian approach is the way to go.

6/10/19

As a teaser, let's ask: what can go wrong in a fit?



Bayesian methods can prevent/identify both underfitting (model is not complex enough to describe the data) or overfitting (model tunes to data fluctuations or terms are underdetermined, leading to them playing off each other).

- We'll see how this plays out!

Let's step through part of the notebook you were sent last month — with some supplementary material.

- Load notebook (B). We'll run using RISE. But you just use it normally. ;)
- We'll include "footnotes" here on Python, Jupyter, Bayesian statistics, physics

• Import of modules

- Note "cell magic" %matplotlib inline (alternative %matplotlib notebook, has interactive figures)
- use of seaborn here is just to make the graphs look good.
- We'll use emcee (cf. MC) to do "sampling" later. corner is used to make a particular type of plot.

• Example from Sivia's book: Gaussian noise and averages.
 • This is an excellent book!

• $p(x | \mu, \sigma) = \frac{1}{\sqrt{2\pi\sigma^2}} \cdot e^{-\frac{(x-\mu)^2}{2\sigma^2}}$ ← μ, σ are given. Probability density of x given μ, σ . Normalized

dimensions of $1/x$ ⇒ $\int_{-\infty}^{\infty} \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(x-\mu)^2}{2\sigma^2}} dx = 1$

• Justification as theoretical model — in Bayesian circles from maximum entropy. Usually justified from "central limit theorem". How many know about that?

6/10/19

• M measurements $D \equiv \{X_k\} = (x_1, \dots, x_M)$ eg. $M=100$
distributed according to $p(x|\mu, \sigma)$.

• How do we get Num here? As in Exploring-pdfs.ipynb.
• "Sample" from $N(\mu, \sigma^2) \Rightarrow$ we'll see this.

• Goal: find approximate μ, σ
Frequentist: maximum likelihood method
Bayesian: compute posterior pdf $p(\mu, \sigma | x, I)$ ← other information

• Random seed of 1 runs same series of random numbers. If you put 2 or 42, then different from 1, but still the same with every run.

• stats.norm, rvs as in Exploring-pdfs.ipynb
• size=M is a "keyword argument" (often kw \equiv keyword)
 \Rightarrow optional and there is a default value (here 1).

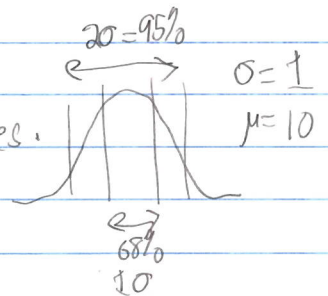
• shift-tab-tab after evaluating cell.
• eg. place on "norm" or "rvs"

• Output D is a numpy array. ← everything in Python is an object, so more than just a datatype \Rightarrow extra methods.

• Put cursor after D then shift-tab-tab
• $[\dots]$ when printed.

• Discuss about number of entries in tails amongst selves.

• Hint: "tail" of Gaussian, say beyond 20
 $\Rightarrow x > 12$ or $x < 8$.



• How many do you expect on average?
 $20 \Rightarrow 95\%$ so about 5/100.

• Here 4 in that range. If there were zero is there a bug?
No, there is a chance that will happen.

• Note the pattern (or lack) and repeat to get different numbers. How?
Change the random seed from 1. (You are invited to try.) Always play.

6/10/19

• Questions about plotting?

- We'll repeatedly use constructions like this, so get used to it!
- ; means we put on same line. Not necessary.
- alpha=0.5 just makes the (default) color lighter.
- try color='red' on your own in scatter plot (as in vlines)
- might prefer side-by-side => alternative code.
- An "axis" in Matplotlib means an entire subfigure, not just the x-axis or y-axis.
- If you want to know about a plotting command already there, shift-tab-tab (usually, sometimes not).
- To find vlines (vertical lines), google "matplotlib vertical line" (try it).
- fig.tight_layout() for good spacing with subplots.

- Ask questions this afternoon and throughout if you are confused by code!

• Observations on graphs?

- scatter plot shows tail => in this case there are 5, but rerun and it will be more or less => everything is a pdf
- histogram is imperfect. Problem? of Exploring pdfs at end (sampling ID pdfs)
- tails fluctuate

• Frequentist approach

- true value for params μ, σ , not a pdf
- Use of \mathcal{L} is notation commonly used
- * Why the product? Assumed independent. Reasonable?
- $\log \mathcal{L}$ for several reasons (note: "log" always means ln. If we want base 10, then \log_{10})

$\mathcal{L} = (\text{const}) e^{-\chi^2}$ so maximizing $\mathcal{L} = \text{maximizing } \log \mathcal{L} = \text{minimizing } \chi^2$

• You can all carry out the maximization

eg. $\frac{\partial \log \mathcal{L}}{\partial \mu} = -\frac{1}{\sigma^2} \sum_{i=1}^M \frac{\partial (x_i - \mu)}{\partial \mu} \cdot -1 = \frac{1}{\sigma^2} \sum_{i=1}^M (x_i - \mu) = \frac{1}{\sigma^2} \left(\sum_{i=1}^M x_i - M\mu \right)$
 => set to zero => $M\mu_0 = \sum_{i=1}^M x_i$ or $\mu_0 = \frac{1}{M} \sum_{i=1}^M x_i$ You do σ_0^2 (easier than σ_0 is $\frac{1}{\sigma_0^2}$)

6/10/19

* Do these make sense?

- μ_0 is mean of data \rightarrow estimator for "true mean"
- σ_0 gives spread about μ_0 .

- Note use of `.sum` to add up D array elements
- Printing with f strings
`f'...' or f'''...'''` \leftarrow multiline string
`.2f` means float with 2 decimal points

- Note comment on "unbiased estimator"
 - an accurate statistic
 - Here compare μ_0 from $\frac{1}{M}$ and $\frac{1}{M-1}$
 - If you do this many times, you'll find $\frac{1}{M}$ doesn't quite give μ_{true} correctly (take mean of μ_0 's from many trials), but $\frac{1}{M-1}$ does. (Try it!)
 - The difference is $O(\frac{1}{M})$, so small for large M.

• Compare estimates to true. Are they good estimates?
 How can you tell? Eg. should they be within .1, .01, or what?
 \Rightarrow more as we go!

- Bayesian approach
 $p(\mu, \sigma | D, I)$ is posterior: probability (density) of finding some μ, σ given data D and what else we know (I).
 "I" could be that $\sigma > 0$ or μ should be near zero.

Frequentist probability: long-run frequency of (real or imagined) trials.
 \Rightarrow data is probabilistic (repeat experiment and get different result)
 but model parameters are not (universe stays the same with more observations)

Bayesian probability: quantification of information (what you know, often said "what you believe"). Data are fixed (it's what you found) but knowledge of true model parameters is fuzzy (and gets update with more trials - coin flipping).

6/10/19

Bayes' Theorem

class: you label each term

likelihood

$$\text{posterior} \Rightarrow P(\mu, \sigma | D, I) = \frac{P(D | \mu, \sigma, I) P(\mu, \sigma | I)}{P(D | I)}$$

← prior

← data probability (or "fully marginalized likelihood" or "evidence or ...")

It will become intuitive!

• tells you how to flip $P(\mu, \sigma | D, I) \leftrightarrow P(D | \mu, \sigma, I)$

hard ← → easy

Aside on denominator

general vector of parameters

$$P(D | I) = \int P(D | \theta, I) P(\theta) d\theta$$

so integrate ("marginalize") over all values of θ . Numerically costly \Rightarrow more later on how to do it.
(parameter)

• For model fitting, we don't need $P(D | I)$ calculated. Find the posterior and just normalize that function (or we might only need relative probabilities).

• If $P(\mu, \sigma | I) \propto 1 \Rightarrow$ "flat prior" (more later)
then

$$P(\mu, \sigma | D, I) \propto \mathcal{L}(D | \mu, \sigma)$$

then F and B get same answer for most likely values μ_0, σ_0 (called "point estimates" as opposed to a full pdf)

• Back to the prior, \Rightarrow include additional information. What you know before a measurement.

- We will talk much more.
- F says it is nonsense; subjective, individual

\Rightarrow discuss this amongst yourselves.

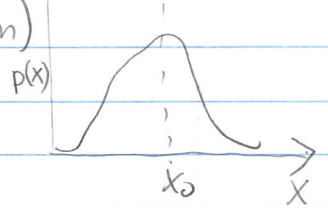
• How to compute $P(\mu, \sigma | D, I)$ in practice? Often with MCMC. Just look now and return Wednesday

6/10/19

General reason why Gaussians may show up:

- Given $p(x|D, I)$, then our "best estimate" from

$$\frac{dp}{dx} \Big|_{x_0} = 0 \text{ with } \frac{d^2p}{dx^2} \Big|_{x_0} < 0 \text{ (maximum)}$$

Look nearby to characterize posterior $p(x)$.

$p(x)$ varies too fast, so characterize $\log p$

$$\Rightarrow L(x) := \log p(x|D, I) = L(x_0) + \frac{dL}{dx} \Big|_{x_0} (x-x_0) + \frac{1}{2} \frac{d^2L}{dx^2} \Big|_{x_0} (x-x_0)^2 + \dots$$

- If we can neglect higher order terms, then

$$p(x|D, I) \approx A e^{\frac{1}{2} \frac{d^2L}{dx^2} \Big|_{x_0} (x-x_0)^2}$$

← normalization

\Rightarrow very generally looks like Gaussian.

$$p(x|D, I) = \frac{1}{\sqrt{2\pi}\sigma} e^{-\frac{(x-\mu)^2}{2\sigma^2}} \Rightarrow \mu = x_0, \sigma = \left(-\frac{d^2L}{dx^2} \Big|_{x_0} \right)^{-1/2}$$

- we usually quote $x = x_0 \pm \sigma$, because if Gaussian, this is sufficient to tell us the entire distribution.

- For Bayesian: full posterior $p(x|D, I)$ for $\forall x$ is general result, and $x = x_0 \pm \sigma$ may be an approximate characterization.

- What if asymmetric $p(x|D, I)$? Multimodal?

6/10/19

95%

Bayesian vs. Frequentist confidence interval

- Bayesian is easy: a credible interval or Bayesian confidence interval or degree-of-belief (DOB) interval is: given some data, 95% chance (probability) that the interval contains the true parameter.

- Frequentist 95% confidence interval

- If large # of repeat samples, 95% of these intervals include the true value of the parameter

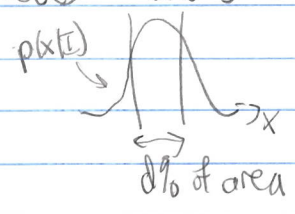
- So the parameter is fixed (no pdf) and the confidence interval depends on data, (random sampling)

"There is a 95% probability that when I compute a confidence interval from data of this sort that the true value of θ will fall within the (hypothetical) space of observations."

- What?

- One key difference: Bayesian includes prior.

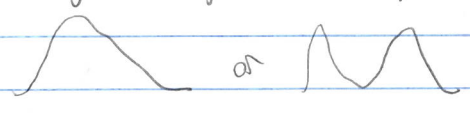
- Issues: One-D \rightarrow if symmetric pdf, then clear how to define ^{d%} confidence interval,



Algorithm: start from center, step outward adding area, stop at d%

Two-D: need a way to integrate from top.

- What if asymmetric or multimodal?



Two of the possible choices:

- Equal-tailed interval (central interval): area above and below interval are equal

- Highest posterior density (HPD) region: Posterior density for every point is higher than the posterior density for any point outside the interval.