# Session 2.1: Bayesian inference

Bayesian modelling for Spatial and Spatio-temporal data, Imperial College

### Learning Objectives

After this session you should be able to:

- Understand how Bayes Theorem can be applied to random variables
- Describe what conjugacy means
- Obtain posterior distribution for the Beta-Binomial and Gamma-Poisson families

The topics treated in this lecture are presented in Chapter 3 of the book Blangiardo and Cameletti (2015) and in Chapter 2.3, 3.1-3.4 and 5.2-5.3 of the book Johnson, Ott, and Dogucu (2022).

#### Outline

- 1. Bayes Theorem for random variables
- 2. A quick recap
- 3. What is conjugacy?
- 4. Some conjugacy models: Beta-Binomial
- 5. Some conjugacy models: Gamma-Poisson

# Bayes Theorem for random variables

### An example

- A clinical trial is carried out to assess the efficacy of a preventive treatment for migraine
- A group of 20 patients are offered the new drug and they have to report if they have a migraine episode in the next week after taking the medication.
- ullet Our aim is to estimate the probability of success of this new drug ( heta)
- ullet Our data here is Y: the number of patients reporting no migraine episodes.

Note that Y is a **random variable** (look at recording 3 for a more detailed description of this concept)

## Prior probability model

As a first step we need to assign a prior on  $\theta$ .

As a simplification let's assume that  $\theta$  is discrete and can only get thevalues 0.1,0.4,0.8 with the following probability function, which specifies the prior probability of each possible  $\theta$  value:

$$\frac{\theta}{0.1} \frac{p(\theta)}{0.2}$$
0.4 0.5
0.8 0.3

Note that this prior reflects some sort of information from a previous study on a similar compound and put 50% probability on the event that 40% of the patients will report a reduction in migraine.

#### The Binomial data model

- ullet Our random variable Y can take any discrete value between 0 and 20 (total number of patients)
- ullet It will depend on the probability heta
- ullet For our formal Bayesian analysis, we must model this dependence of Y on heta through a **conditional probability** model
- We make two assumptions about the trials: (1) the outcome of any one patient doesn't influence the outcome of another; and (2) the probability of success does not change among patients
- We can use the Binomial model

 $Y \sim \operatorname{Binomial}(n, \theta)$ 

with conditional probability function

$$p(y\mid heta) = rac{n!}{y!(n-y)!} heta^y(1- heta)^{n-y}$$

- ullet Mean of this distribution is E(Y)=n heta
- ullet Variance is V(Y)=n heta(1- heta)

Check out recording 4 for a recap on the Binomial distribution

#### The Binomial data model

This model allows to calculate ANY conditional probability. For instance, conditioning on heta=0.4 let's see the difference in the probability of getting 12 or 15 successes



$$p(y=12 \mid heta=0.4) = rac{20!}{12!8!} 0.4^{12} (1-0.4)^8 = 0.035$$



$$p(y=15 \mid heta=0.4) = rac{20!}{15!5!} 0.4^{15} (1-0.4)^5 = 0.0013$$

--

Note you can get these results in R using

```
> #1
> dbinom(12,20,0.4)
```

[1] 0.03549744

```
> #2
> dbinom(15,20,0.4)
```

[1] 0.001294494

#### Binomial likelihood function

- The Binomial provides a theoretical model of the data we might observe.
- ullet In the end we observe 10 successes out of the 20 patients (y=10).
- ullet The next step in our Bayesian analysis is to determine how compatible this particular data is with the various possible heta

We need to evaluate the *likelihood* of getting 10 successes in the trial under each possible value of heta

Similarly to last week with events, the likelihood function follows from evaluating the conditional probability function  $p(y=10\mid\theta)$  for all the possible  $\theta$  values:

$$p(y=10 \mid heta=0.1) = rac{20!}{10!10!} 0.1^{10} (1-0.1)^{10} = 0.00000644$$

2

$$p(y=10 \mid heta=0.4) = rac{20!}{10!10!} 0.4^{10} (1-0.4)^{10} = 0.117$$

3

$$p(y=10 \mid heta=0.8) = rac{20!}{10!10!} 0.8^{10} (1-0.8)^{10} = 0.002$$

## Likelihood vs probability function

Let's recall the fundamental difference between probability function and likelihood function

When  $\theta$  is known, the **conditional probability** function  $p(\cdot \mid \theta)$  allows us to compare the probabilities of different values of the data Y occurring with  $\theta$ :

$$p(y_1 \mid \theta) \text{ vs } p(y_2 \mid \theta)$$

When Y is known, the **likelihood function**  $L(\cdot \mid y) = p(y \mid \cdot)$  allows us to compare the relative likelihood of the data y under different possible values of  $\theta$ :

$$L( heta_1 \mid y) = p(y \mid heta_1) ext{ vs } L( heta_2 \mid y) = p(y \mid heta_2)$$

So  $L(\cdot \mid y)$  provides the tool we need to evaluate the relative compatibility of data Y=y with various heta values.

## Normalising constant

- Now we have a prior for  $\theta$  and a likelihood and as Bayesian we want to **balance** these two pieces of information to obtain the posterior.
- Something is missing...

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This is the total probability of having y=10 successes. How do we get this?

Law of total probability:

$$P(A) = \sum_{j} P(A \mid B_{j}) P(B_{j}) ext{ (for events)}$$

$$P(Y=y) = \sum_{j} p(Y \mid heta_{j}) p( heta_{j}) ext{ (for discrete probability functions)}$$

So we get

$$P(Y=10) = 0.00000644 \times 0.2 + 0.117 \times 0.5 + 0.002 \times 0.3 = 0.059$$

Interpretation: across all the possible heta there is only around 6% chance that there are 10 successes.

#### Posterior distribition

Finally we are ready to apply Bayes theorem and get the posterior distribution

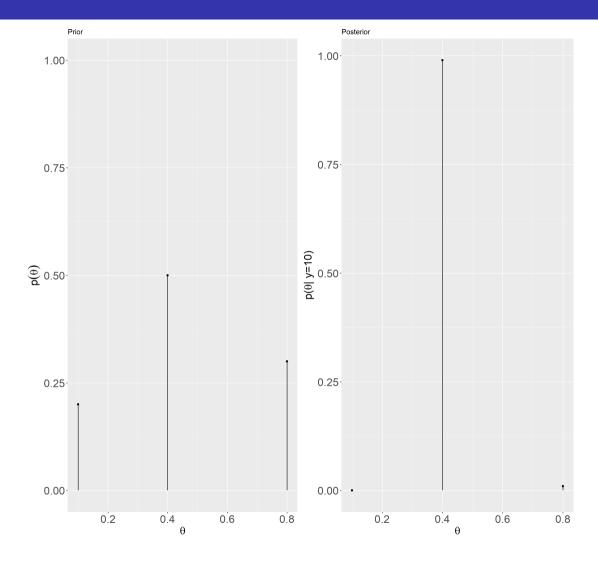
$$p( heta \mid y = 10) = rac{p( heta)L( heta \mid y = 10)}{p(y = 10)} ext{ for } heta \in \{0.1, 0.4, 0.8\}$$

which gives us:

$$p( heta=0.1 \mid y=10) = rac{0.2 imes 0.0000064}{0.059} = 0$$
 $p( heta=0.4 \mid y=10) = rac{0.5 imes 0.117}{0.059} = 0.99$ 
 $p( heta=0.8 \mid y=10) = rac{0.3 imes 0.002}{0.059} = 0.01$ 

# Comparing prior and posterior

$\theta$	$p(\theta)$	$p(\theta \mid y = 10)$
0.1	0.2	0.00
0.4	0.5	0.99
0.8	0.3	0.01



#### Posterior shortcut

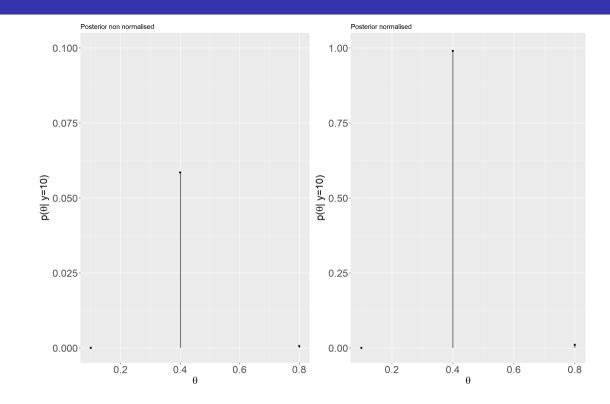
- Moving forward we can actually forget about calculating the normalising constant
- ullet Note that on slide 12 the p(y=10) appears at the denominator of all the  $p( heta\mid Y)$

It normalises the posterior probabilities so they sum to 1

ullet So we can simply acknowledge that p(y=10)=1/c and replace the posterior as:

$$p(\theta \mid y = 10) \propto p(\theta) \times L(\theta \mid y)$$

• The proportionality means that if we compare the normalised and unnormalised posterior they preserve their relative relationship



# A quick recap

## So a quick general recap: Bayesian inference

#### Makes fundamental distinction between

- Observable quantities y, i.e.~the data
- Unknown quantities heta
- $\theta$  can be statistical parameters, missing data, mismeasured data...
  - $\rightarrow$  parameters are treated as random variables
  - $\rightarrow$  in the Bayesian framework, we make probability statements about model parameters

### So a quick general recap: Bayesian inference

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Note that in the Frequentist framework, parameters are fixed non-random quantities and the probability statements concern the data

### Bayesian inference

- ullet As with any statistical analysis, we start building a model which specifies  $p(Y=y\mid heta)$
- This is the data distribution, which relates all variables into a full probability model
- The choice of data distribution depends on the nature of the data:
- e.g. are we analysing continuous or discrete data, are the data symmetric or skewed, etc.
  - As we observe the data, we can use descriptive tools (e.g. plots) to visualise the data and choose the best likelihood

### Bayesian inference

#### From a Bayesian point of view

- ullet is unknown so should have a probability distribution reflecting our uncertainty about it before seeing the data
- ightarrow need to specify a prior distribution p( heta)
  - *y* is known so we should condition on it
    - $\rightarrow$  use Bayes theorem to obtain conditional probability distribution for unobserved quantities of interest given the data:

$$p( heta \mid y) = rac{p( heta)\,p(y \mid heta)}{\int p( heta)\,p(y \mid heta)\,d heta} \propto p( heta)\,p(y \mid heta)$$

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#### This is the posterior distribution

- ullet The prior distribution p( heta), expresses our uncertainty about heta before seeing the data
- The posterior distribution  $p(\theta \mid y)$ , expresses our uncertainty about  $\theta$  after seeing the data

#### The posterior distribution

Posterior distribution forms basis for all inference --- can be summarised to provide

- point and interval estimates of Quantities of Interest (QOI), e.g. treatment effect, small area estimates, ...
- point and interval estimates of any function of the parameters
- probability that QOI (e.g. treatment effect) exceeds a critical threshold
- prediction of QOI in a new unit
- prior information for future experiments, trials, surveys, ...
- inputs for decision making

• ...

What is conjugacy?

### How to select a prior

- Selecting the prior is crucial for a Bayesian analysis
  - There is no right way to select a prior
  - The choices often depend on the objective of the study and the nature of the data
- There are other criteria to consider when choosing a prior model:

#### Computational ease

Especially if we don't have access to computing power, it is helpful if the posterior model is easy to build.

#### Interpretability

The posterior balance (is a compromise) the data and the prior. A posterior model is interpretable, and thus more useful, when you can look at its formulation and identify the contribution of the data relative to that of the prior.

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#### Conjugate prior

Let the prior model for a parameter  $\theta$  with  $p(\theta)$  and the model of data Y conditioned on  $\theta$  have likelihood function  $L(\theta \mid y)$ .

If the resulting posterior model  $p(\theta \mid y) \propto p(\theta) \times p(y \mid \theta)$  is of the same family as the prior, then we say it is a **conjugate prior**.

# Some conjugacy models: Beta-Binomial

#### Beta-Binomial model for proportions: example

- We consider an early investigation of a new drug
- Experience with similar compounds has suggested that response rates between 0.2 and 0.6 could be feasible
- We interpret this as a distribution with mean = 0.4 and standard deviation 0.1
- A Beta(9.2,13.8) distribution has these properties (Check recording 7 to see how to go from the mean and sd to the a and b parameters of a Beta distribution)
- ullet Suppose we now treat n=20 volunteers with the compound and observe y=15 positive responses

## Identifying the different model components

• Assuming patients are independent, with common unknown response rate heta, leads to a binomial data distribution

$$p(y\mid n, heta) = \left(rac{n}{y}
ight) heta^y(1- heta)^{n-y} \; \propto \; heta^y(1- heta)^{n-y}$$

•  $\theta$  needs a continuous prior distribution:

$$egin{aligned} heta &\sim \mathrm{Beta}(a,b) \ p( heta) &= rac{\Gamma(a+b)}{\Gamma(a)\Gamma(b)} \ heta^{a-1} (1- heta)^{b-1} \end{aligned}$$

## Combining prior and data

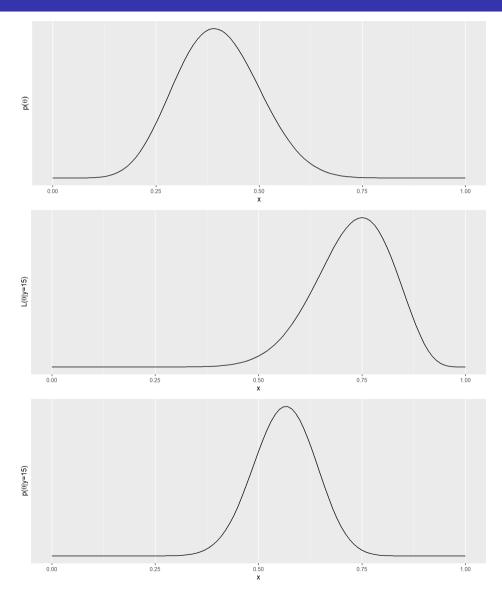
Combining the Binomial data and the Beta prior gives the following posterior distribution

$$egin{split} p( heta \mid y,n) &\propto p(y \mid heta,n) p( heta) \ &\propto heta^y (1- heta)^{n-y} heta^{a-1} (1- heta)^{b-1} \ &= heta^{y+a-1} (1- heta)^{n-y+b-1} \end{split}$$

The posterior is still a Beta distribution (with different parameters):

$$p(\theta \mid y, n) \propto \mathrm{Beta}(y + a, \ n - y + b)$$

# Comparing prior, likelihood and posterior



#### Gamma-Poisson model for count data: example

- For a recap on the Poisson distribution see recording 5
- In epidemiology we are often interested in estimating the rate or relative risk rather than the mean for Poisson data:
- Suppose we observe y=7 cases of leukaemia in one region;
- ullet The expected number of cases is E=4
- Data distribution: Poisson with mean  $\theta = \lambda \times E$ , where  $\lambda$  is the unknown incidence ratio:

$$p(y \mid \lambda, E) = rac{(\lambda E)^y e^{-\lambda E}}{y!}$$

• **Prior**: Gamma(a, b) on the the risk  $\lambda$ :

$$p(\lambda) = rac{b^a}{\Gamma(a)} \lambda^{a-1} e^{-b\lambda}$$

Check recording 8 for a recap on the Gamma distribution

## Combining likelihood and prior

This implies the following posterior

$$egin{aligned} p(\lambda \mid y) &\propto p(\lambda) \, p(y \mid \lambda) \ &= rac{b^a}{\Gamma(a)} \lambda^{a-1} e^{-b(\lambda)} e^{-(\lambda E)} rac{(\lambda E)^y}{y!} \ &\propto \lambda^{a+y-1} e^{-(b+E)\lambda} \ &= \operatorname{Gamma}(a+y,\,b+E) \end{aligned}$$

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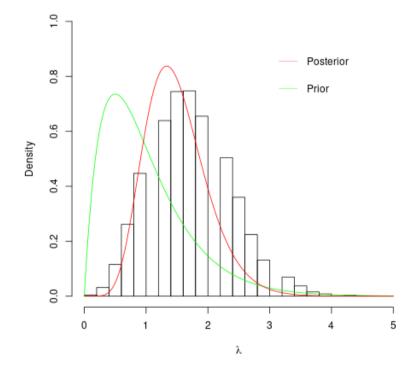
The posterior is another (different) Gamma distribution

$$E( heta \mid y) = rac{a+y}{b+E}$$

So posterior mean depends on the prior (a,b) and on the data (y,E)

## Prior, likelihood, posterior for the Leukaemia example

- ullet Assuming a prior  $\lambda \sim \mathrm{Gamma}(2,2)$
- ullet Considering the data y=7, E=4
- ullet We obtain a posterior  $\lambda \mid y \sim \mathrm{Gamma}(9,6)$  centered around 1.5



The posterior becomes a compromise between the prior and the data

#### References

Blangiardo, M. and M. Cameletti (2015). *Spatial and spatio-temporal Bayesian models with R-INLA*. John Wiley & Sons. Johnson, A. A., M. Q. Ott, and M. Dogucu (2022). *Bayes Rules!: An Introduction to Applied Bayesian Modeling*. CRC Press.