



**UNIVERSITY  
OF TURKU**



**Turku PET  
CENTRE**



# Bayesian Statistical Inference

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PhD project: The human dopamine system, funded by Päivikki and Sakari Sohlberg Foundation

Licensed psychologist, undergraduate student in Statistics

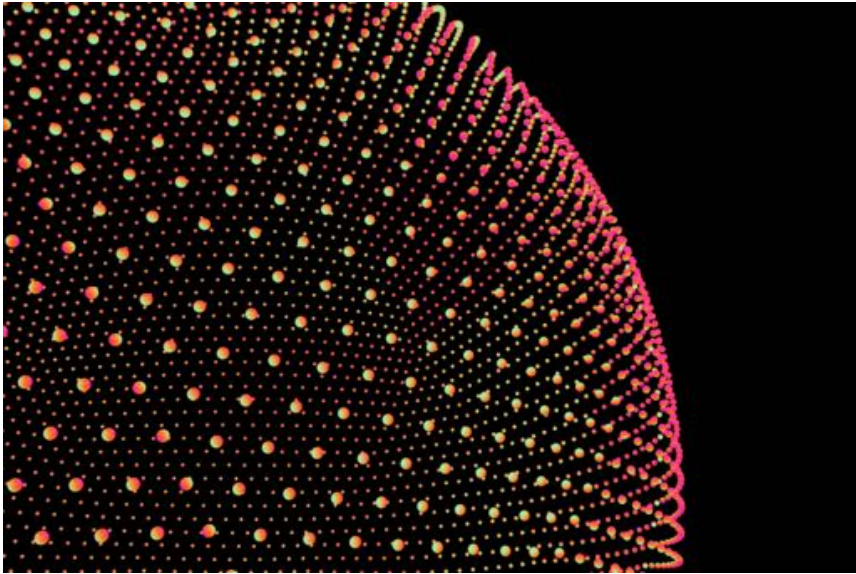
# Contents

- 1. Theory and philosophy**
2. How, when, why to apply?

# Statistical inference

*“Statistical inference is the process of drawing conclusions about an **underlying population** based on a **sample** or subset of the data.”*

Statistical inference = Translating data into understanding

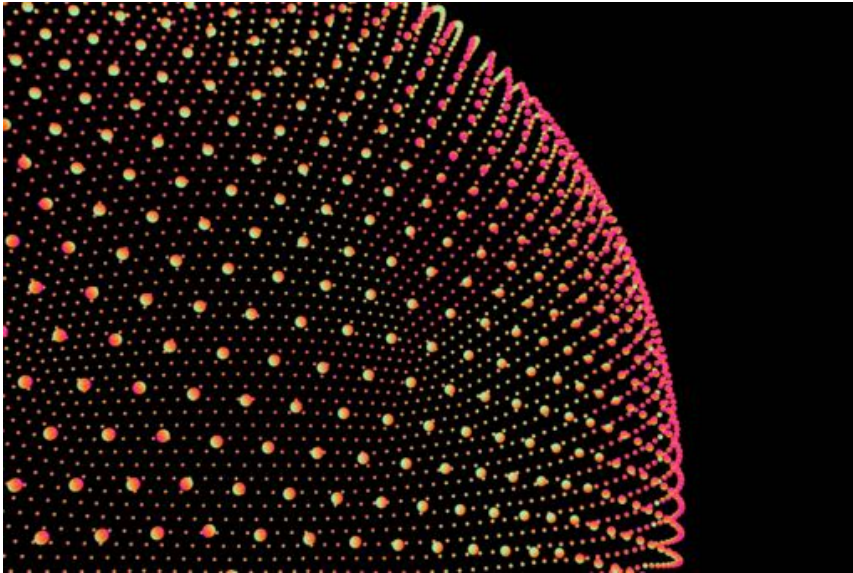


Data



Underlying population

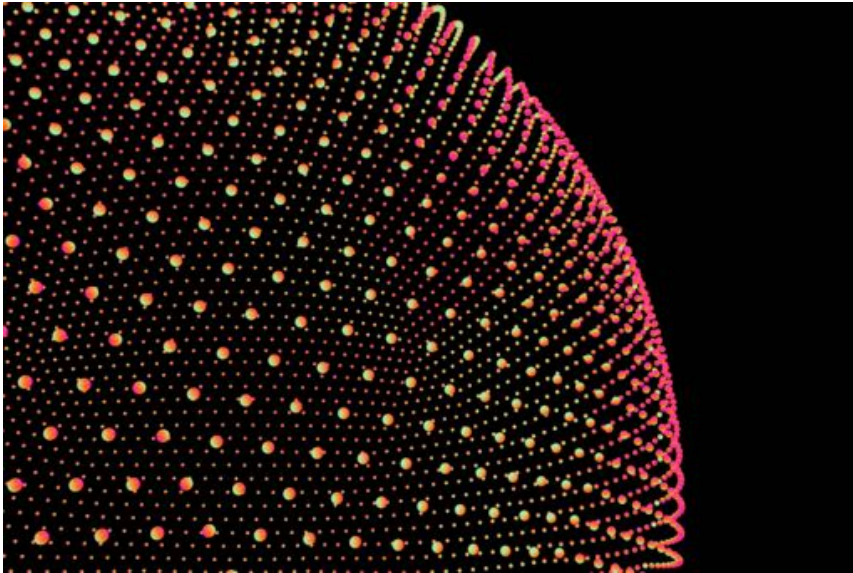
# Statistical inference = Translating data into understanding



Example: How much people love statistics?

- Subset: not practical to ask everyone
- Sample bias: poll at stats dept. – generalize to others?
- Measuring variation: bad day?

# Statistical inference



$\neq$



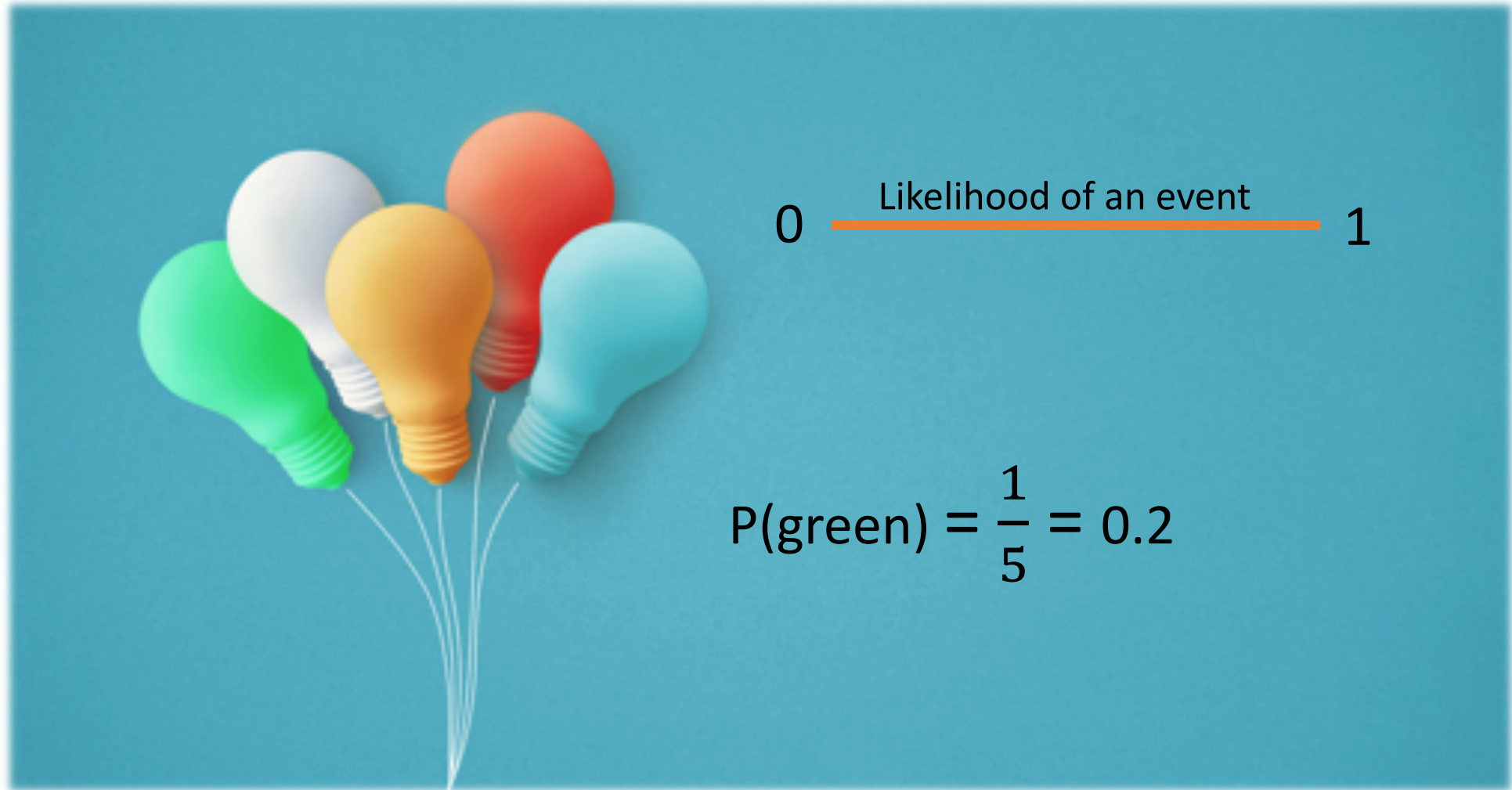
→ Dealing with **uncertainty**:

Gap


The heart and soul of statistical inference



# Describing uncertainty: Probability



# Describing uncertainty: Probability



0 Likelihood of an event 1

$$P(\text{green}) = \frac{1}{5} = 0.2$$

Rarely that simple in research!  
But a key concept that can be applied in various ways

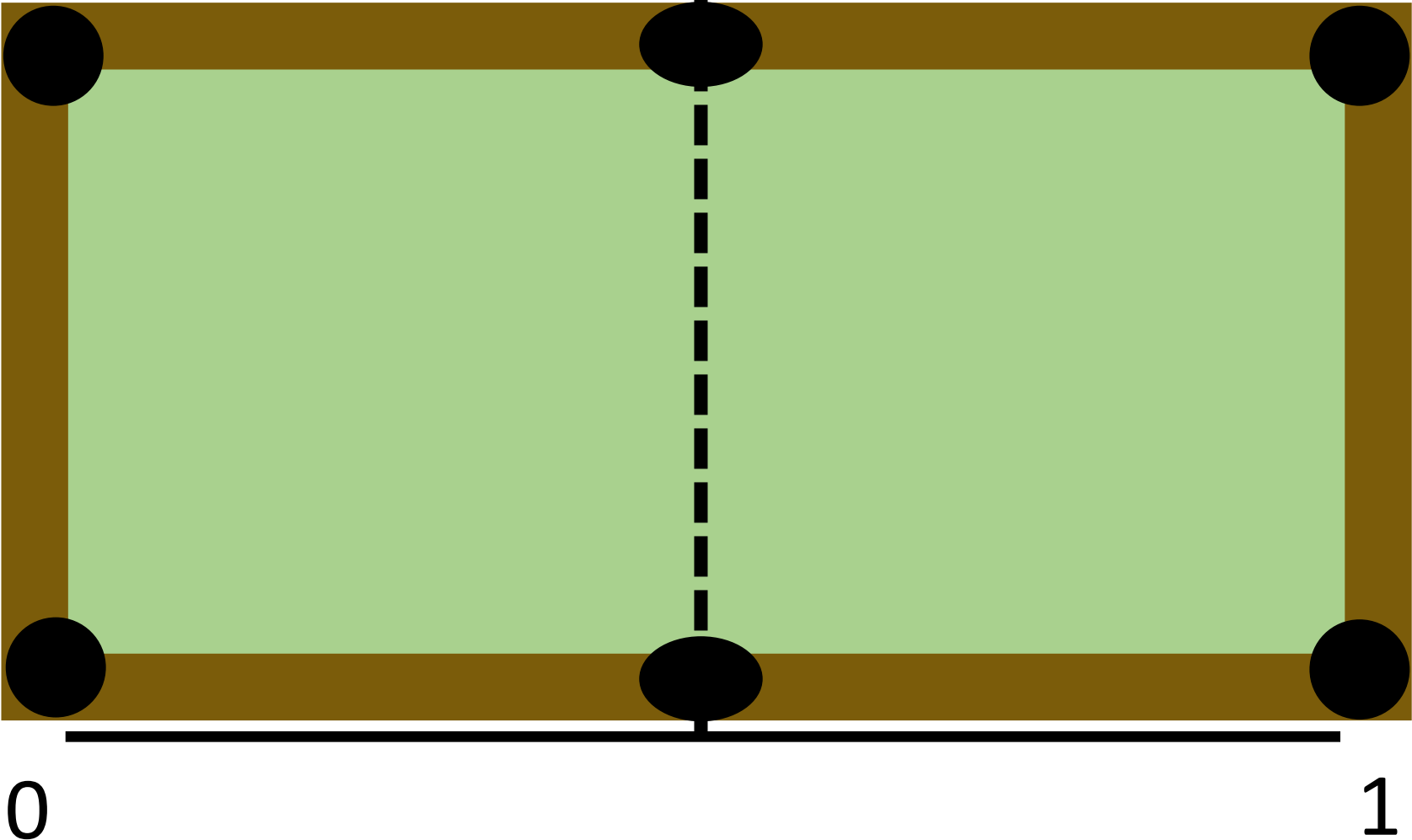


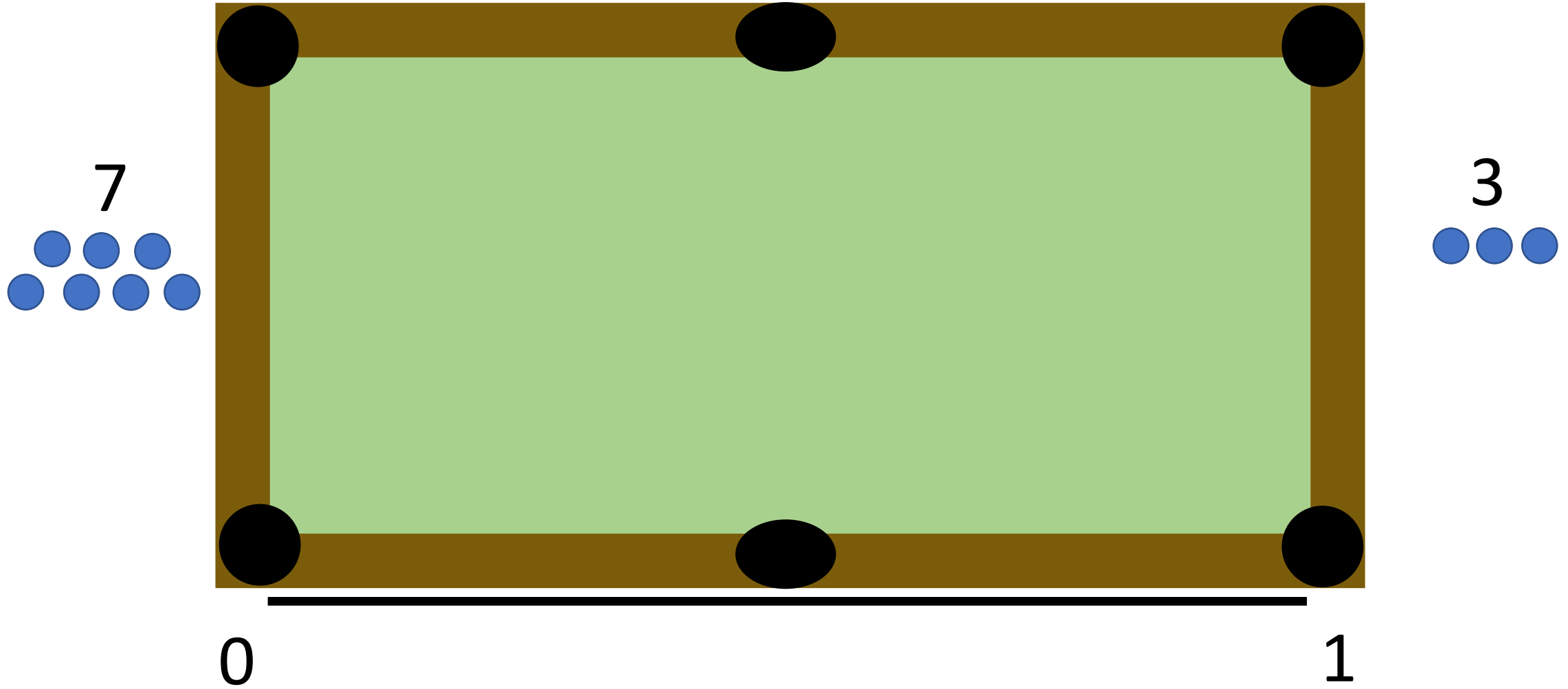
# Two approaches to statistical inference

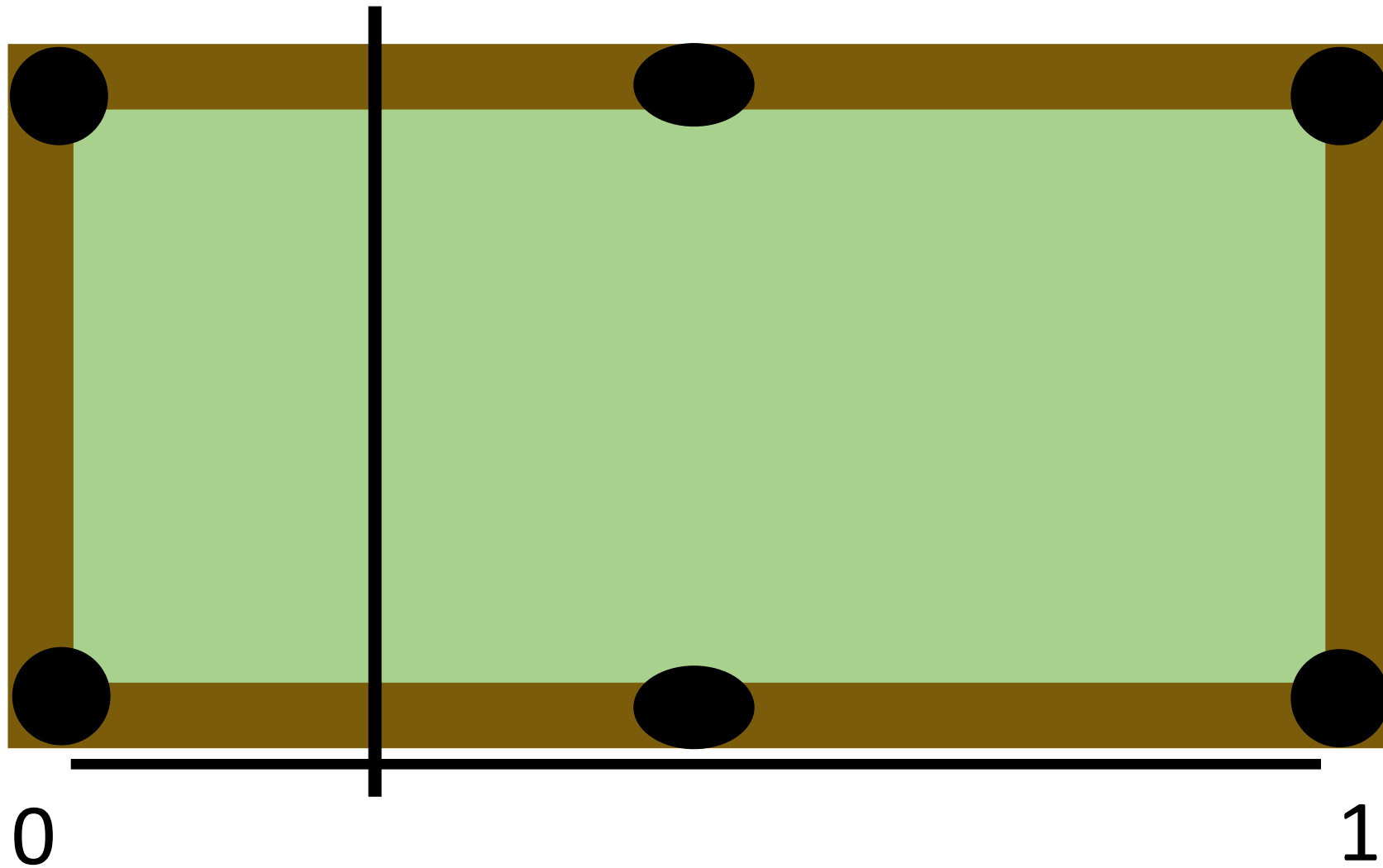
1. Bayesian
2. Frequentist

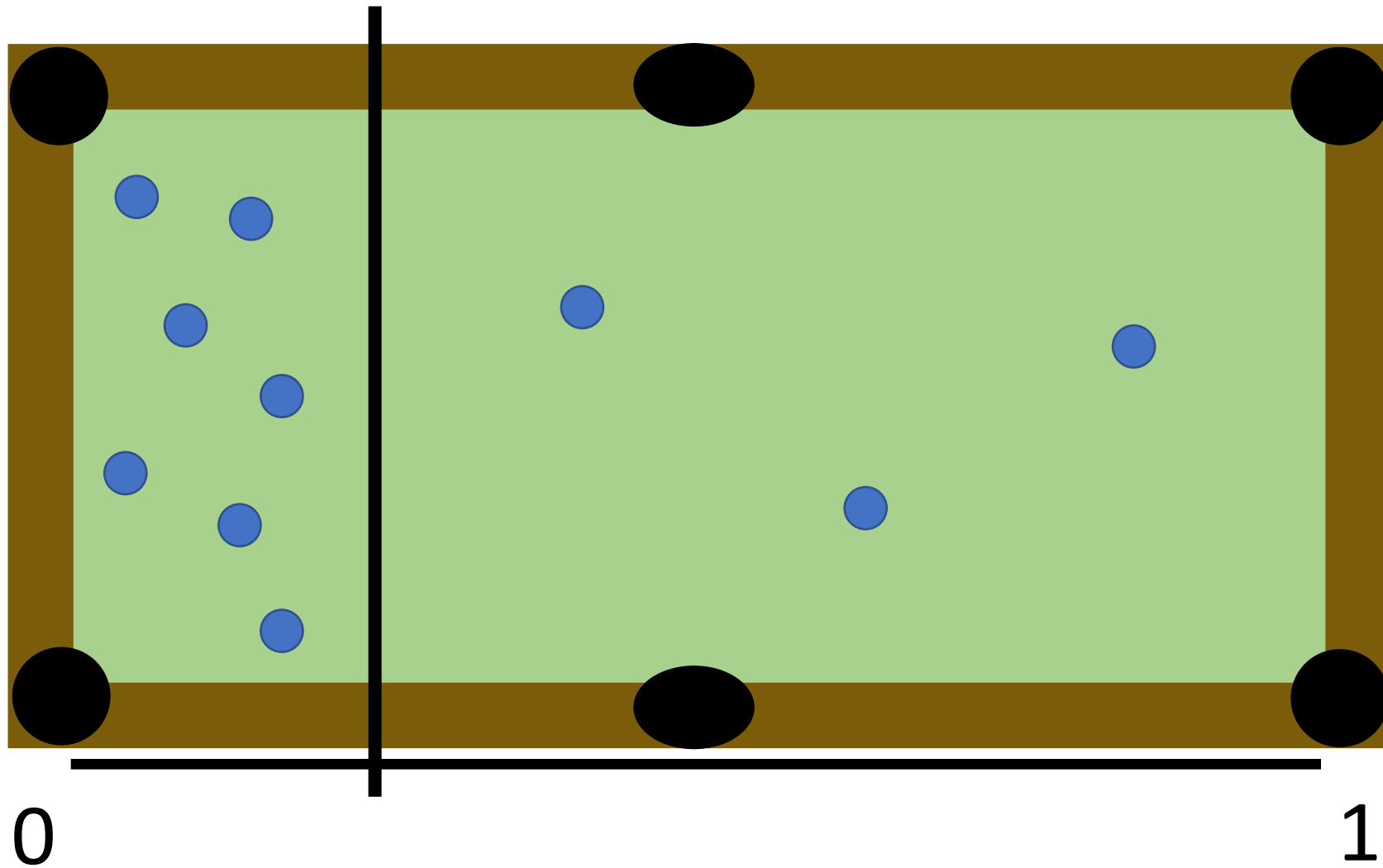


# Thomas Bayes

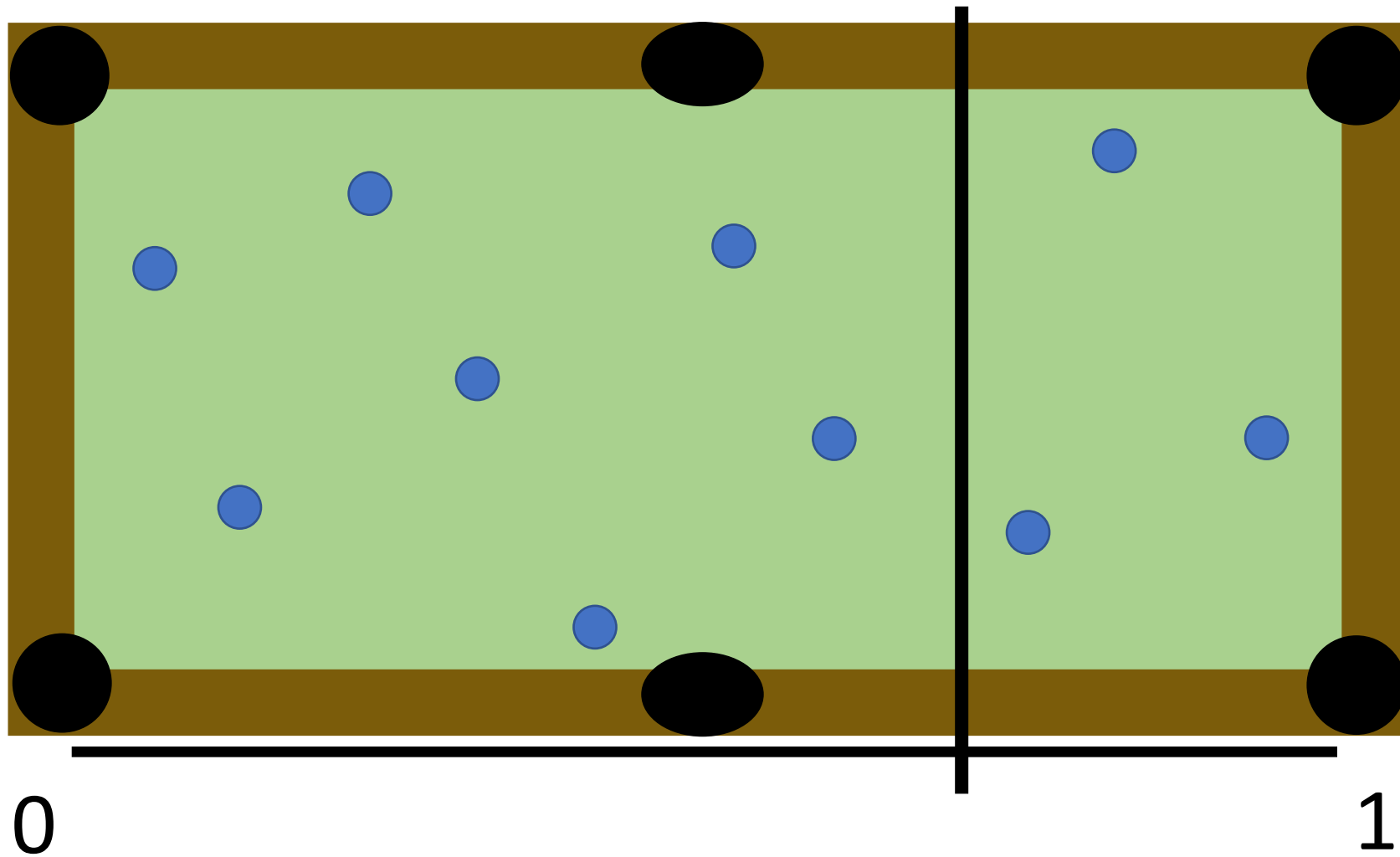












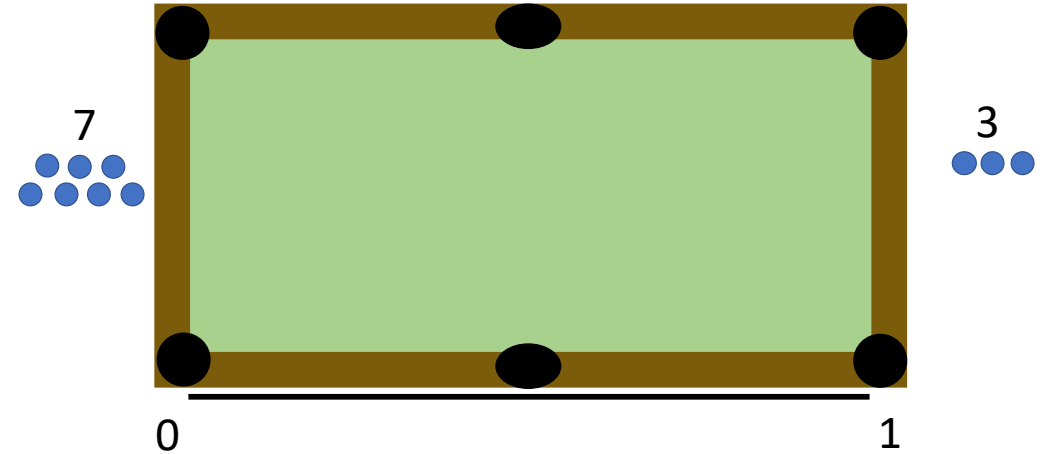
# Where is the line?

1) What do we know prior to data?

- There is a line, somewhere from 0 to 1

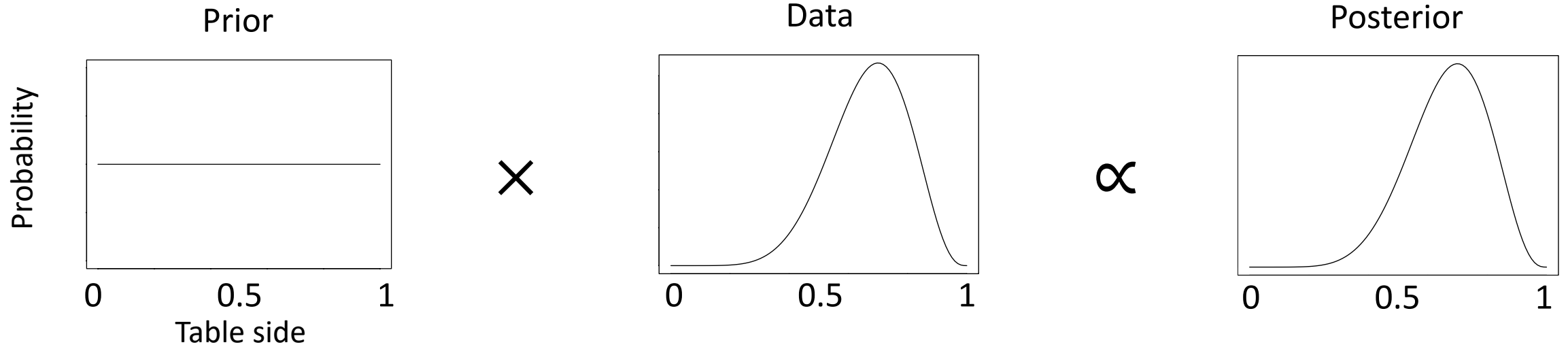
2) What do we observe?

- 7 on the left
- 3 on the right

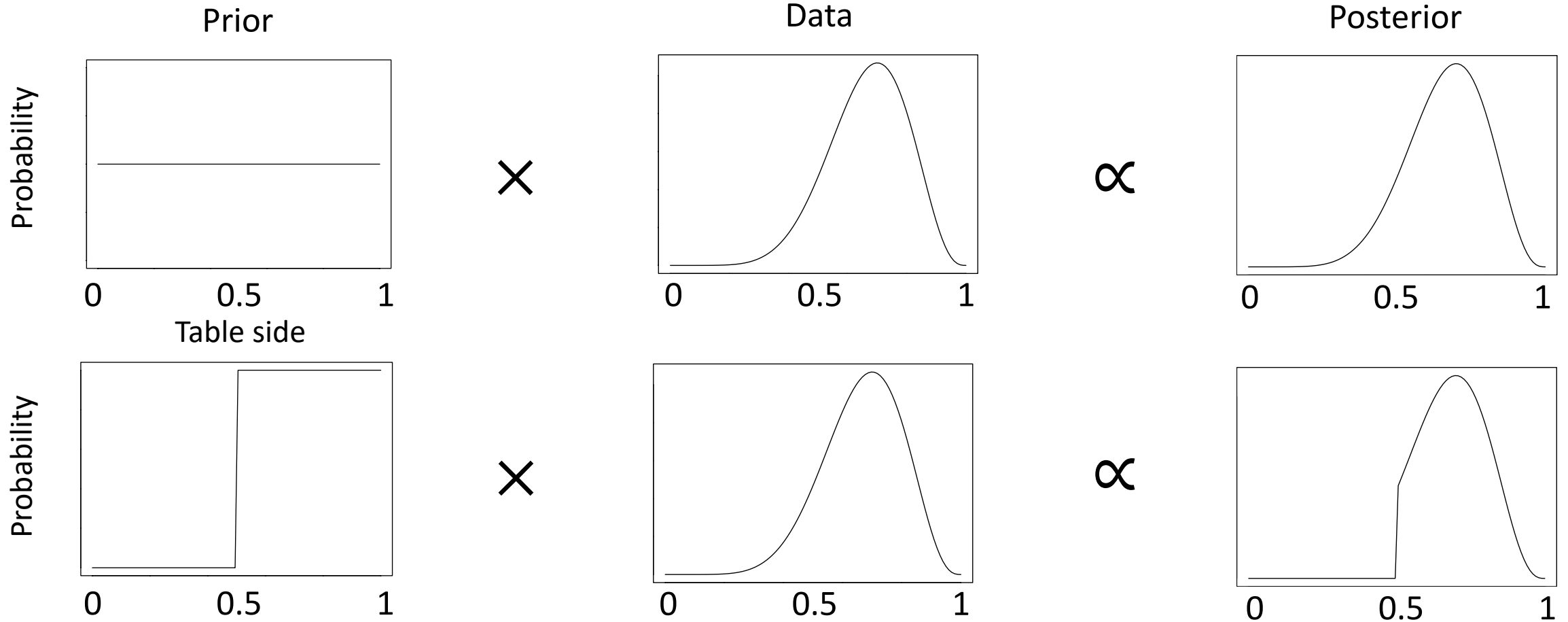


Both the **prior** knowledge and the **data** (observations) affect our **posterior** understanding

# Where is the line?

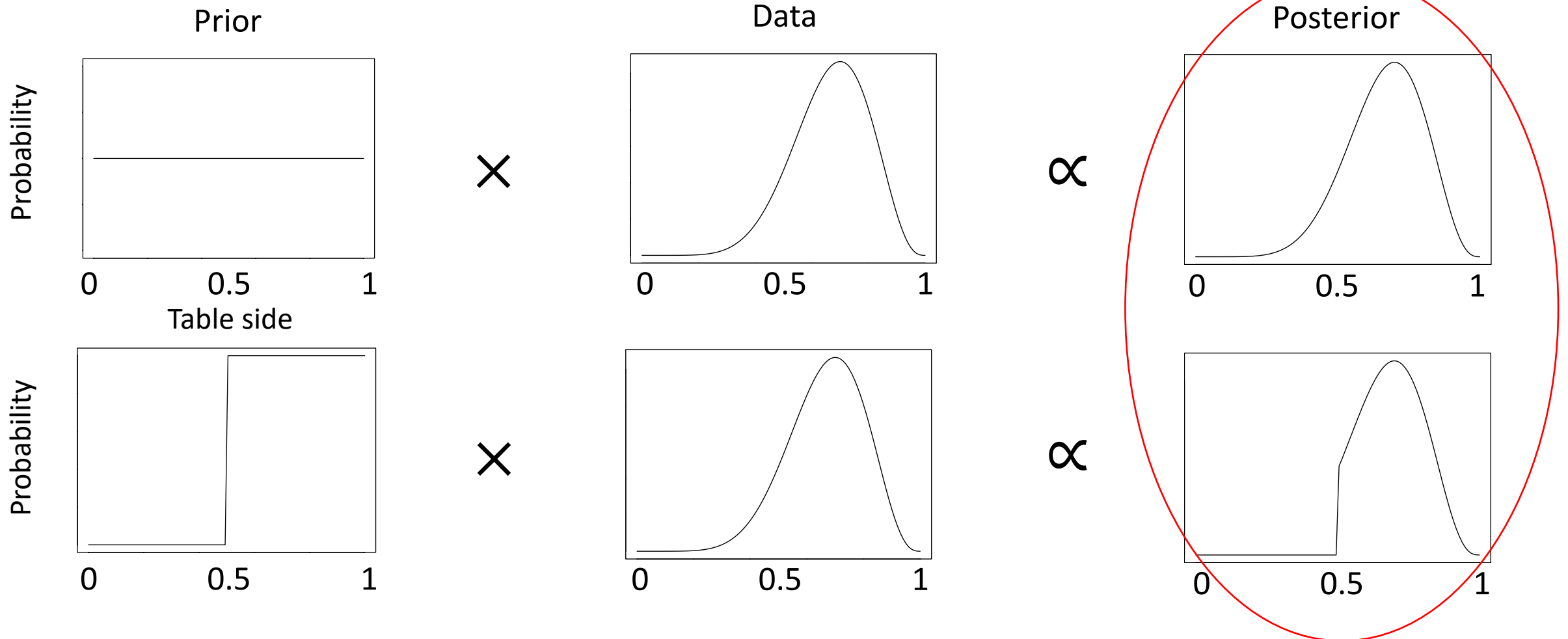


# Where is the line?



# Where is the line?

Can be an iterative process:  
If we got more data,  
we could use the posterior as the new prior





# The result is a probability distribution for the parameter

Parameter= the unknown ground truth we try to solve

Probability for each possible location of the line  
But the line is somewhere, we just don't know it

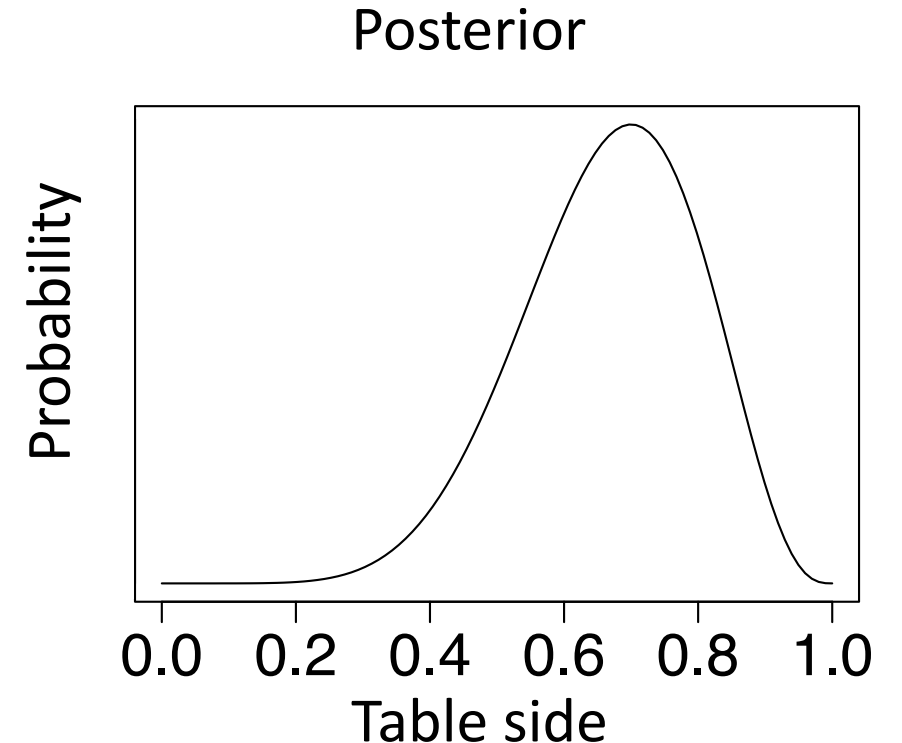
(50% probability one has a virus, is pregnant...)

What does it mean to be 50% likely pregnant?

One is or not

## Subjective probability:

- Give parameter probabilities
- Get posterior probabilities back (updated with data)



# Where is the probability?

## Data

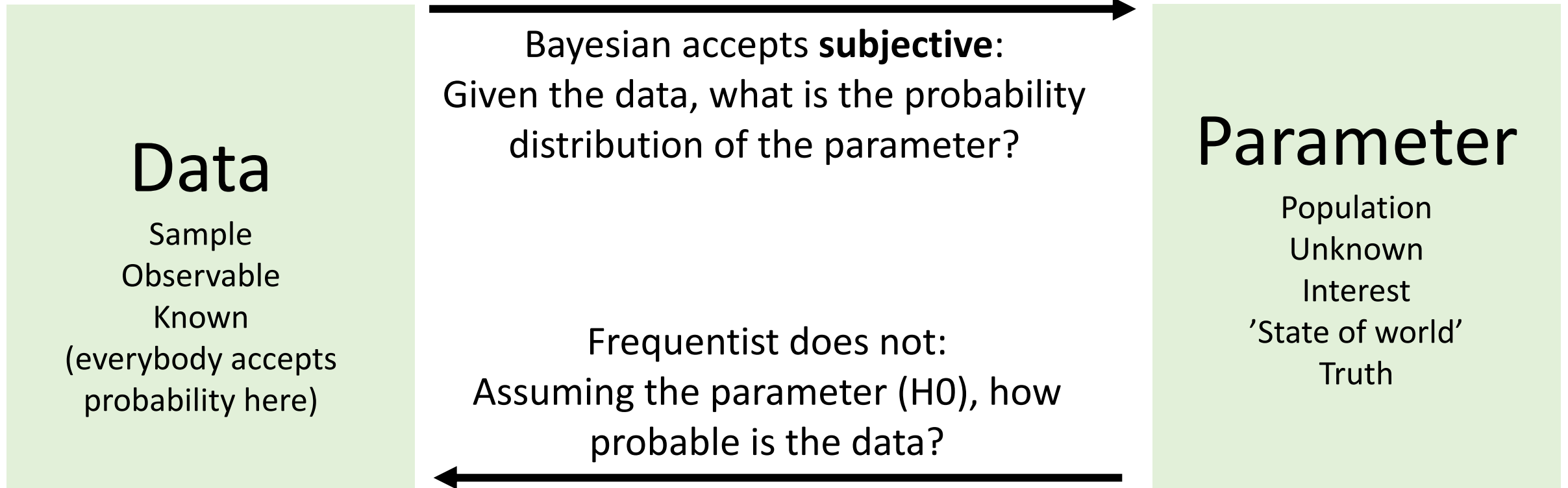
Sample  
Observable  
Known  
(everybody accepts  
probability here)

Bayesian accepts **subjective**:  
Given the data, what is the probability  
distribution of the parameter?

## Parameter

Population  
Unknown  
Interest  
'State of world'  
Truth

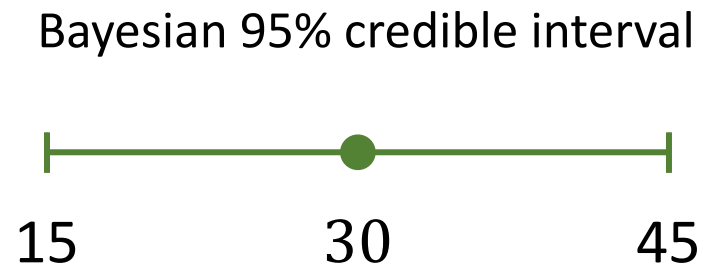
# Where is the probability?



# Example: Laugh

- How much people laugh?
- Sample  $n = 100$  people
- Laughtermeter: Measure how long a subject laughs during a day
- Parameter: true mean laughing time in the population

# Example: Laugh

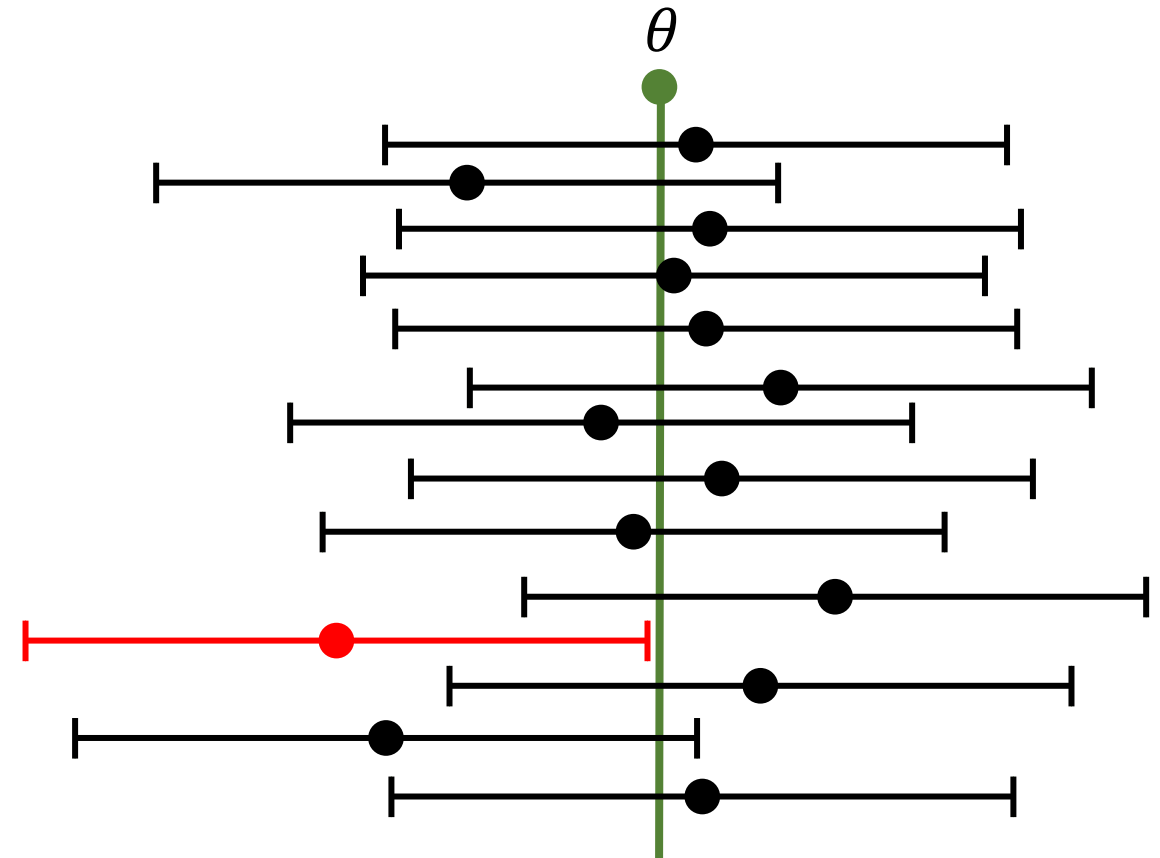


95% probability that people laugh 15-45 min a day  
Most likely 30 min

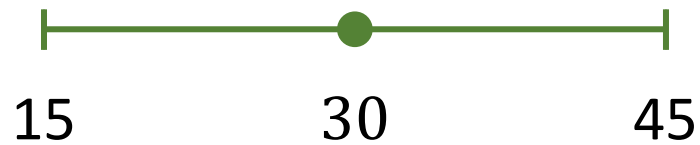


# Example: Laugh

## Frequentist 95% confidence interval



## Bayesian 95% credible interval



95% probability that people laugh 15-45 min a day  
Most likely 30 min

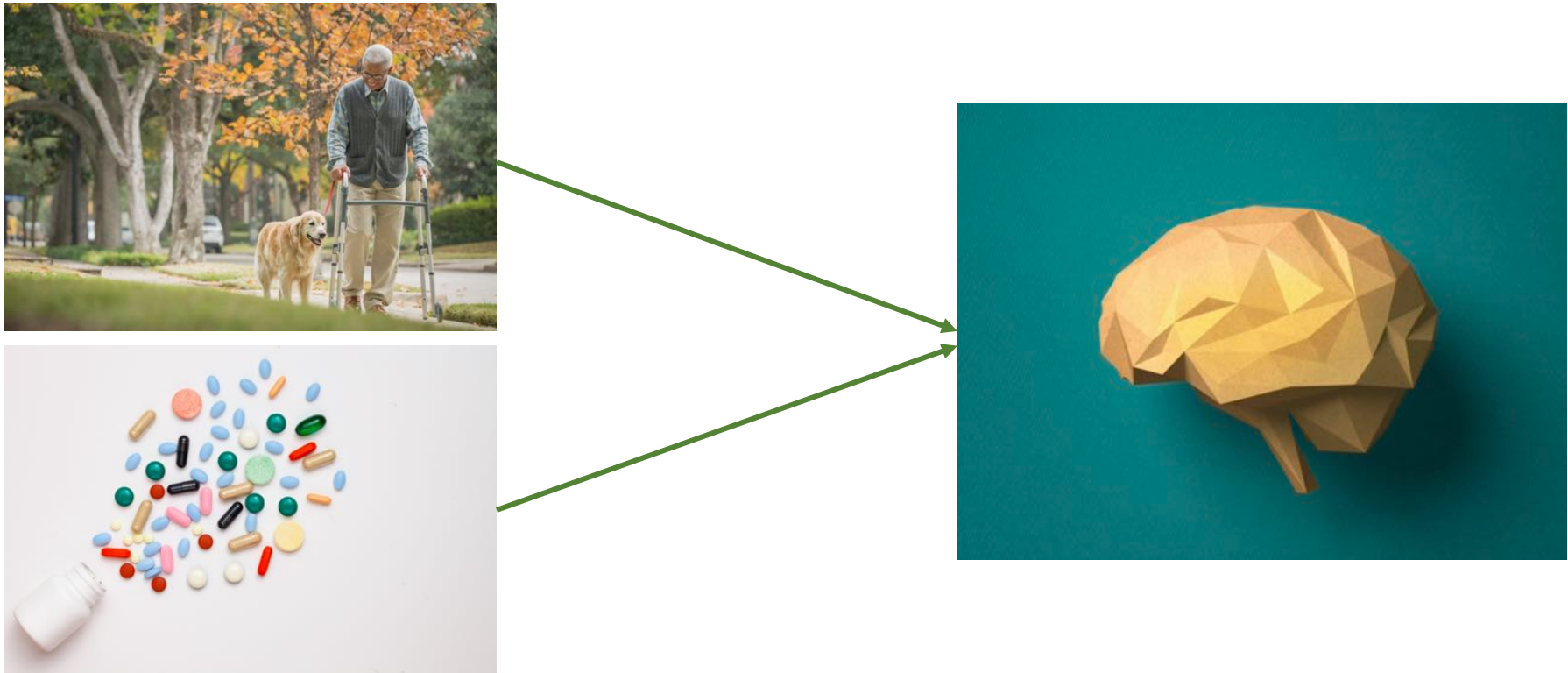
When we take many many samples and calculate confidence interval for each, 95% of the times the true mean laughing time is within the interval

# Contents

1. Theory and philosophy
2. **How, when, why to apply?**

# Usual interest

- How some variable is related to another

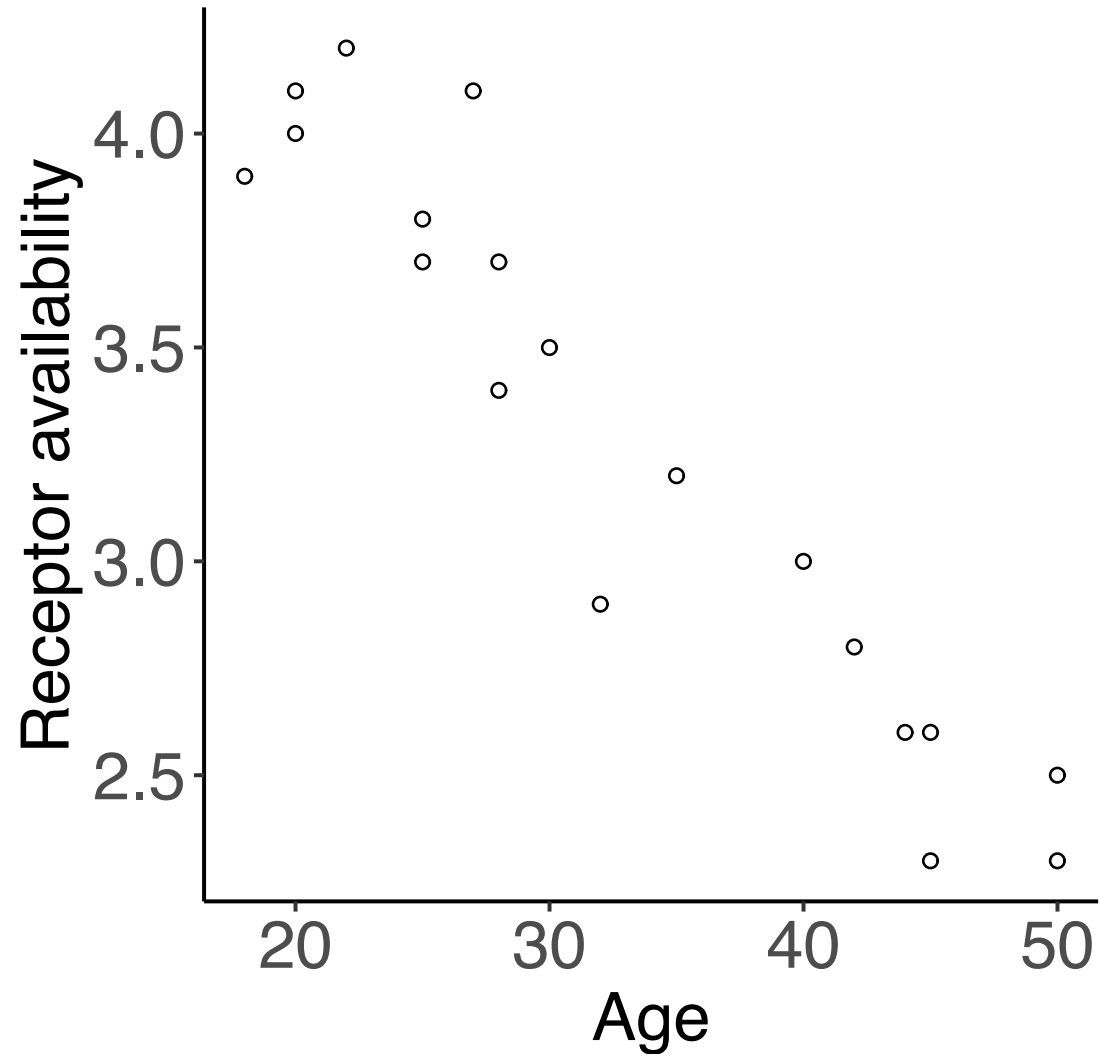


# Example

How is **age** associated with (dopamine) **receptor availability** in a region of interest (ROI)?

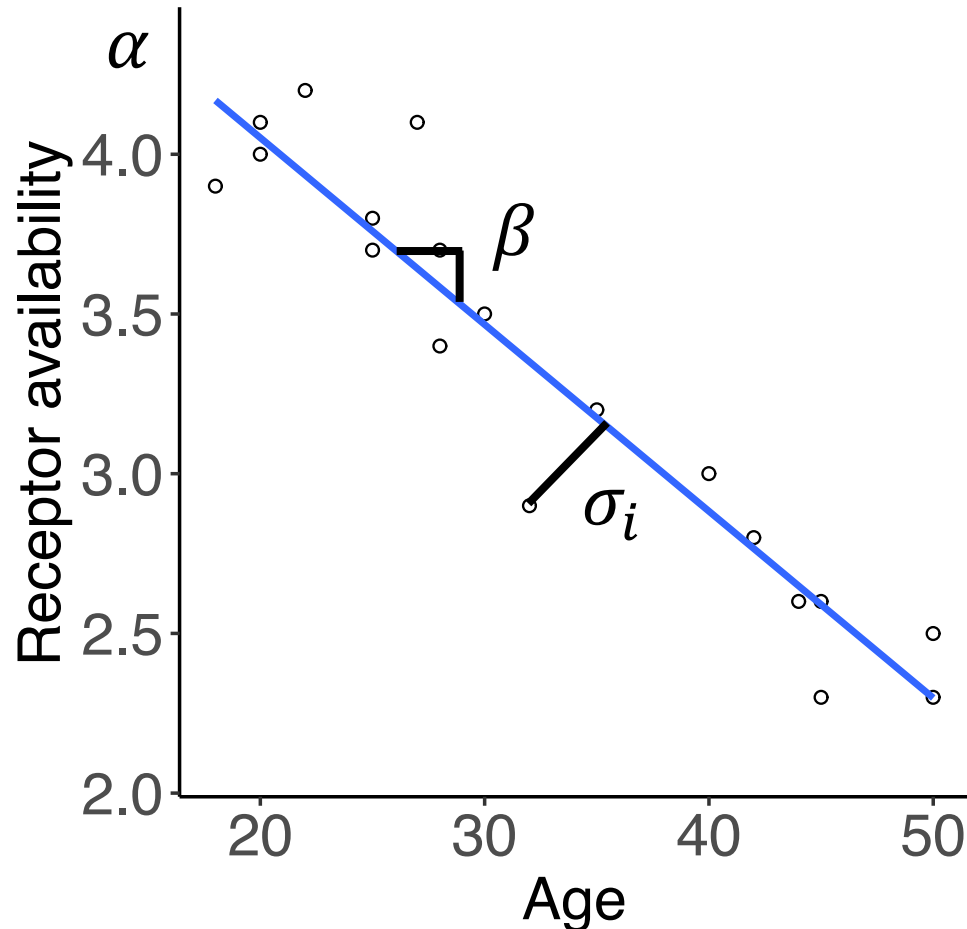
Receptor availability (binding potential = bp)

- Always positive value acquired from a PET image (1 estimate/ ROI for each subject)
- The density (and affinity) of available (unoccupied) dopamine receptors in a specific region  
→ information about dopamine function
- Normal( $\mu, \sigma$ )



- n= 20
- Age
- Receptor availability (bp) from PET
- 1 brain region

# Linear regression: linear relation



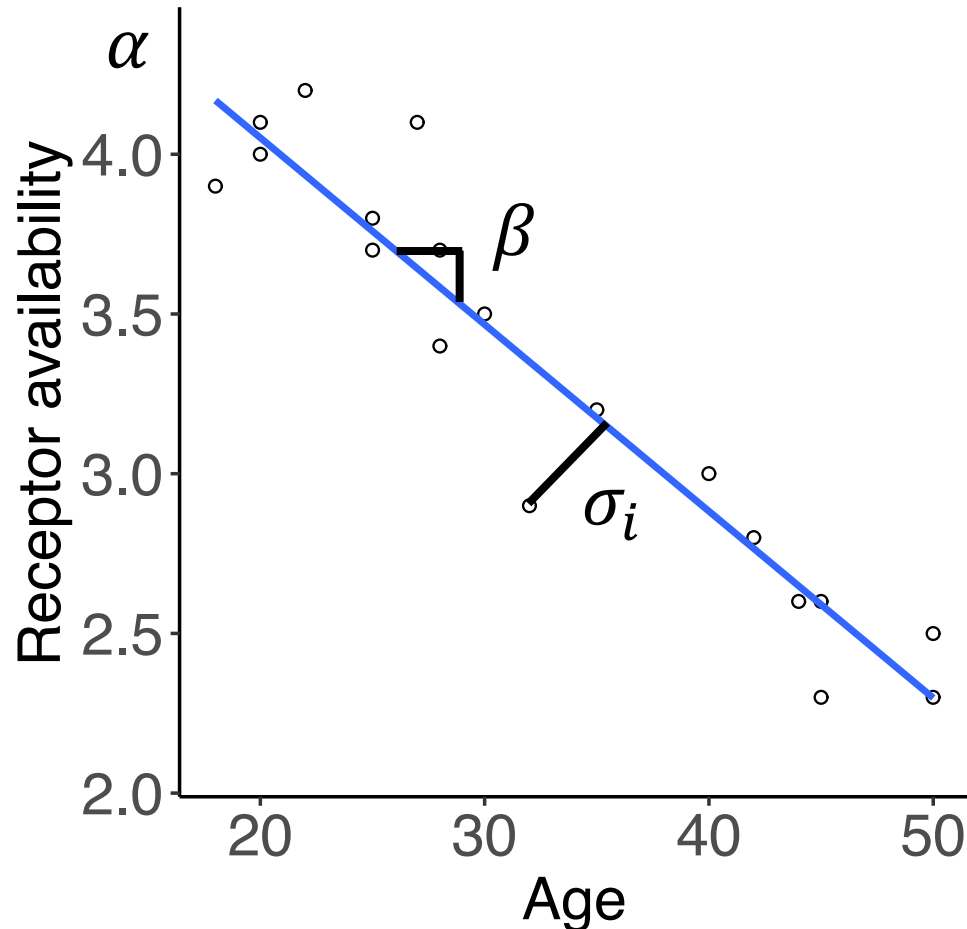
Parameters:

$\alpha$  = expected availability, when age = 0

$\beta$  = regression coefficient (slope), steepness of the line, the change in availability when age increases 1 unit

$\sigma_i$  = distance between an observation and regression line

# Linear regression: linear relation



Parameters:

$\alpha$  = expected availability, when age = 0

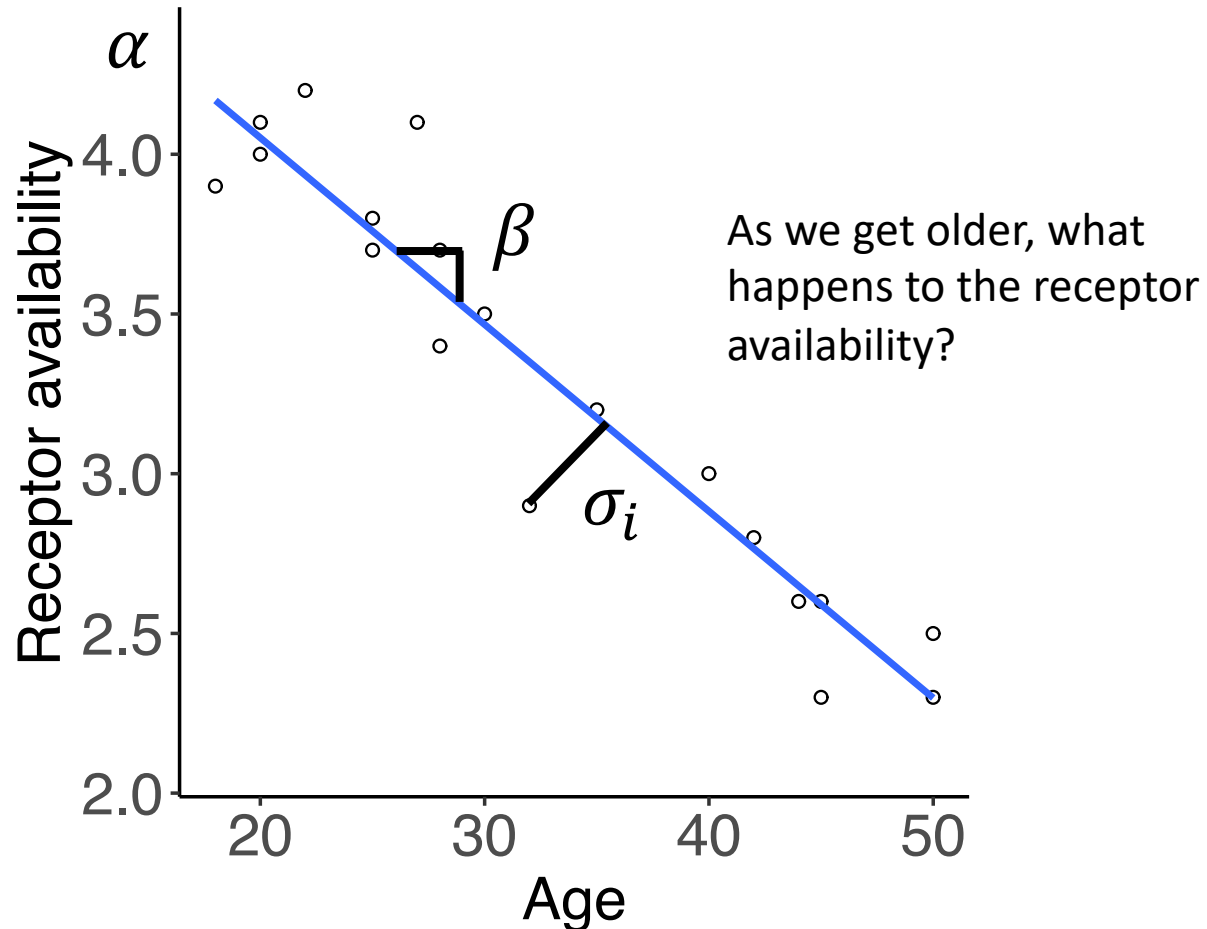
$\beta$  = regression coefficient (slope), steepness of the line, the change in availability when age increases 1 unit

$\sigma_i$  = distance between an observation and regression line

Linear model:

$$\mu = \alpha + \beta \times \text{age}$$

# Linear regression: linear relation



Parameters:

$\alpha$  = expected availability, when age = 0

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Linear model:

$$\mu = \alpha + \beta \times \text{age}$$

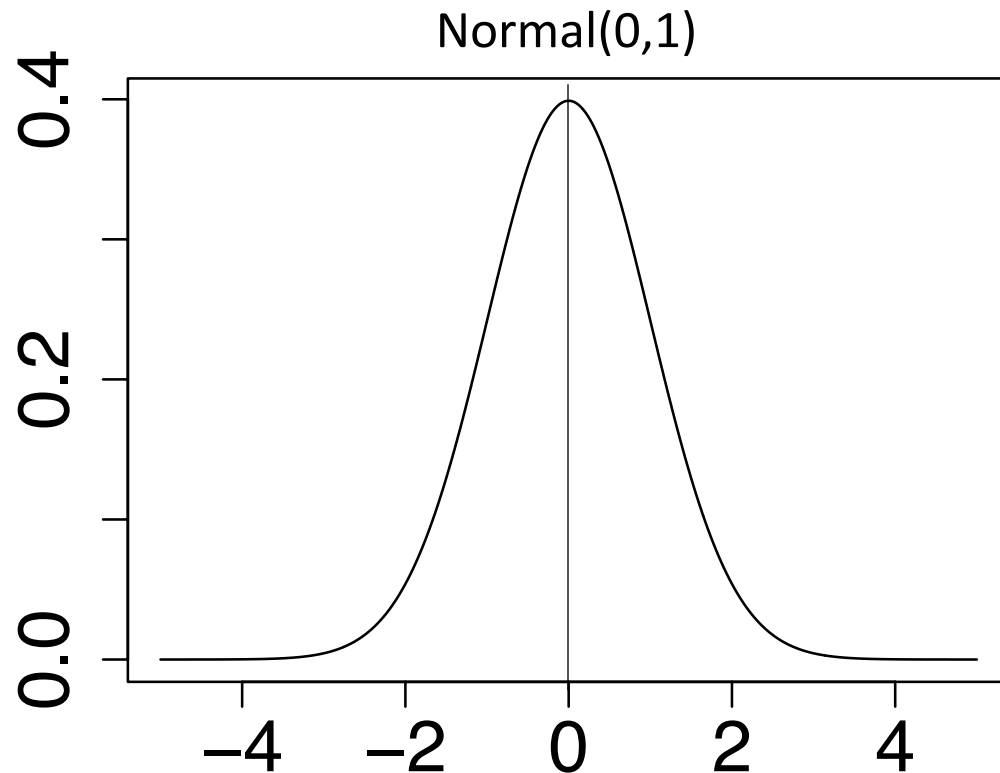
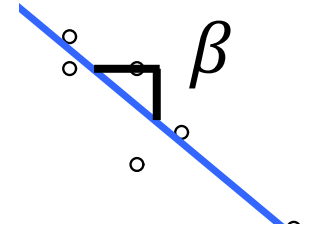


Let's solve  $\beta$  the Bayesian way  
(= what happens to the receptor availability when we age)

# Tools

- brms-package in R

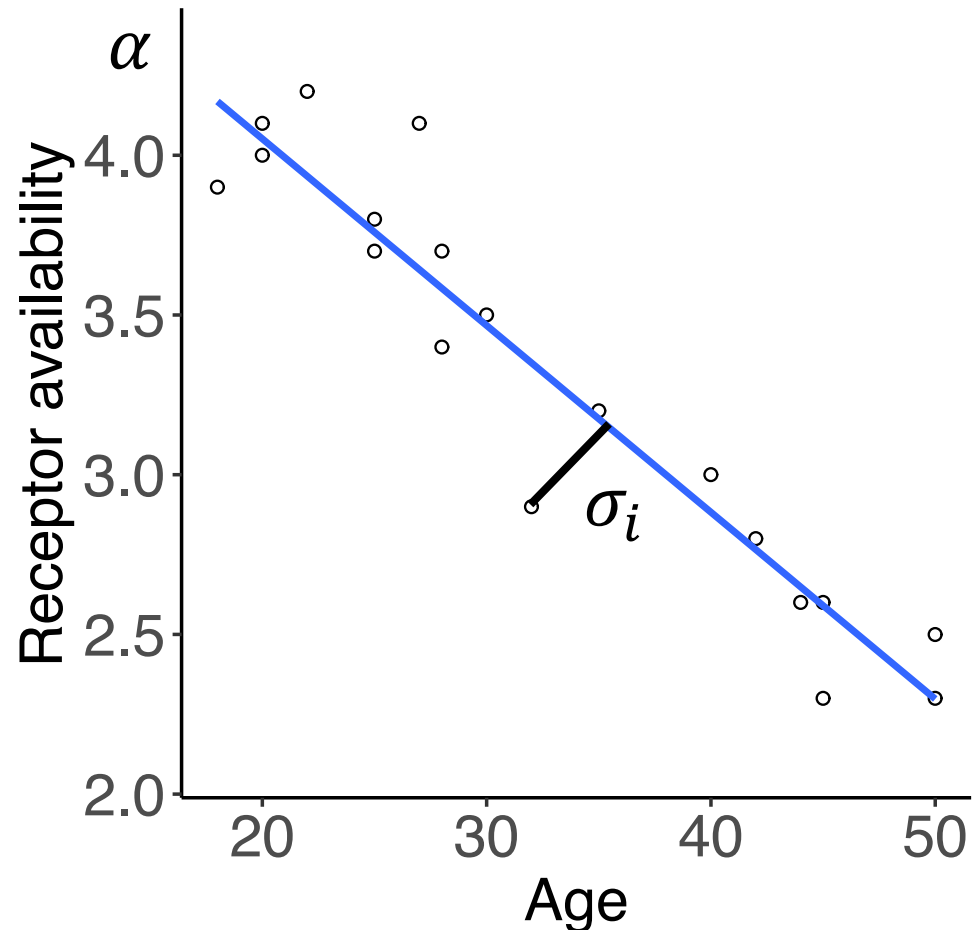
# Prior for each parameter



$\beta$  = regression coefficient (slope)

- Mean is at "zero-effect" (no association)
- The extremely high associations (steep line) are suggested less probable than weaker associations: does not push toward strong associations  $\rightarrow$  if one found the evidence from data
- No values for  $\beta$  are excluded (no 0 probability given)
- Symmetrical: neither positive nor negative slope is given higher probability than the other

# Prior for each parameter



Also  $\alpha$  and  $\sigma$  are parameters of the model: need priors

For now brms package default priors (weakly informative)

Good to be aware of all priors used and their fit in your data and population

```

#### REQUIRED PACKAGES
library(rstan)
library(brms)

### STATISTICAL MODELING

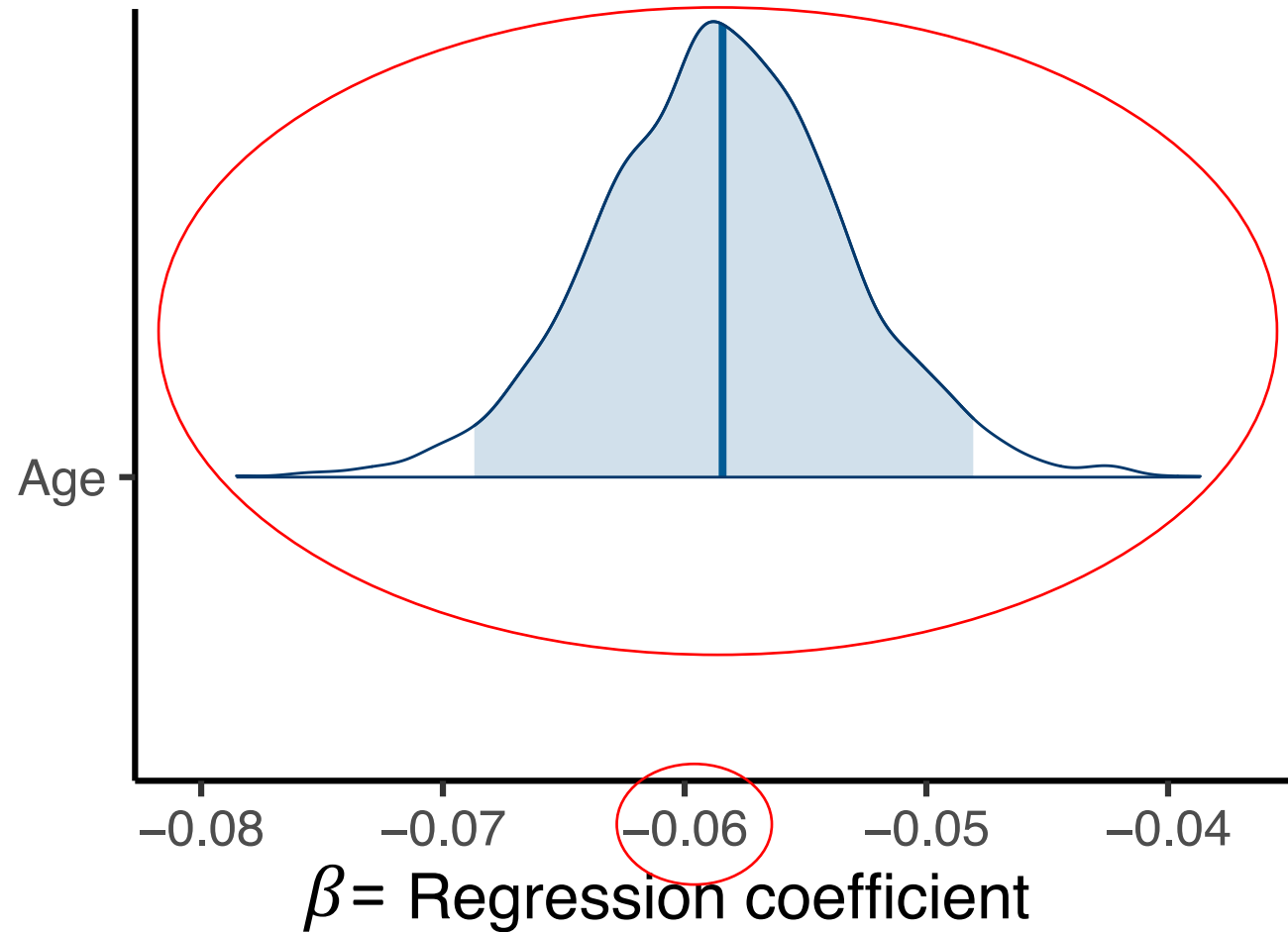
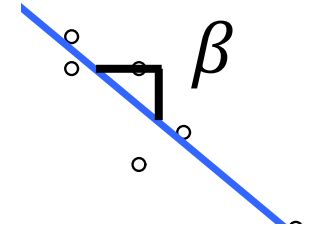
prior <- c(set_prior("normal(0,1)", class = "b")) # prior for beta

model <- bp ~ age # bp = receptor availability

fit <- brm(formula = model,
           data = data, # mock-up
           family = gaussian(), # receptor availability normally distributed
           prior = prior,
           warmup = 1000, iter = 2000, chains = 4, # sampling settings, see brms
           control = list(adapt_delta = 0.95)) # sampling settings, see brms

```

# Result for $\beta$



Ageing 1 year,  
The availability decreases 0.06 units  
=  
Ageing 10 years  
The availability decreases 0.6 units

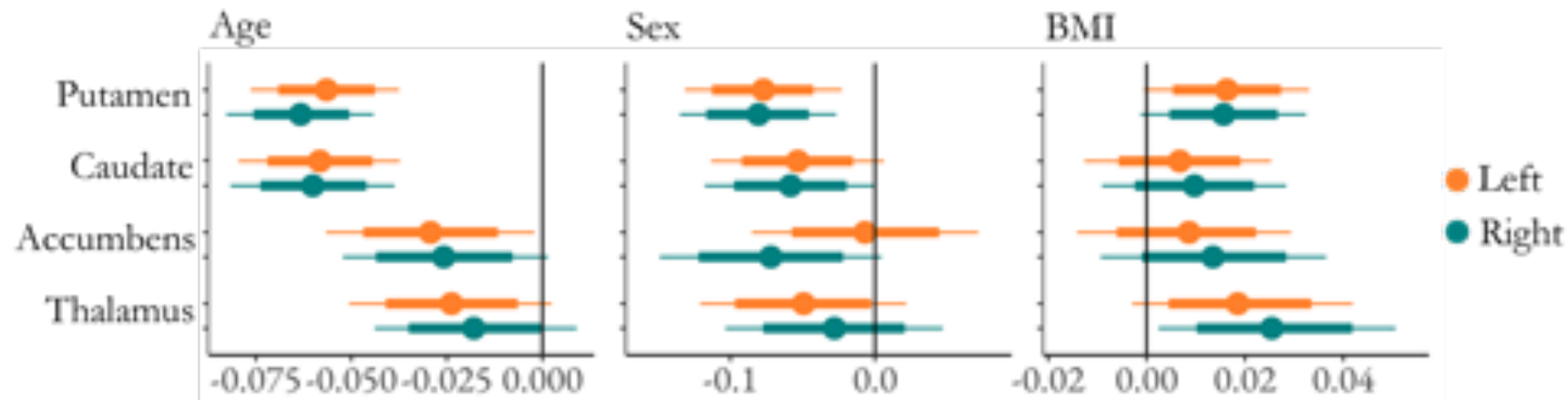
# When?

- Simple enough setting
  - Computational methods (sampling from the posterior distribution)
  - Regional rather than voxel-level
  - Some predictors (e.g. age, sex...) but not too many

# When?

Age, sex, BMI → dopamine receptor availability (preprint)

- Ageing 10 years and 5% decrease in the availability (putamen and caudate)
- Females had higher availability than males
- BMI: weak positive association with the availability
- Please see the preprint for more information about the scales to use (standardizing, log-transformation)



Bayesian methods not limited to linear regression



# Why?

## Interpretation intuitive

- Probabilities for parameter values (what we want)
  - Instead of having probability for the data in the light of only 1 assumed parameter value (null-hypothesis)
- One sample enough -> imaginary resampling not needed

# To take home

- Prior and data form posterior
- If interested in how your prior affects the result, you can try different ones and compare your posteriors
- We have information about the world before we see the data and we can use it (as long as we accept the subjective probability)!

# Further Learning Material

- Bayesian Data Analysis (Gelman, Carlin, Stern Rubin), Chapman Hall, 1995, 2003, 2013
  - <http://www.stat.columbia.edu/~gelman/books>
- McElreath, R. (2018). *Statistical rethinking: A Bayesian course with examples in R and Stan*. Chapman and Hall/CRC.
  - <https://www.youtube.com/watch?v=4WVeICswXo4>
  - Scripts and datasets

# Acknowledgements

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- **Severi Santavirta**, MD, Doctoral candidate, University of Turku

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## R

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- Bürkner, P. (2021). Rdocumentation. brms. Retrieved from: <https://www.rdocumentation.org/packages/brms/versions/2.15.0>. Cited: 17.8.2021.
- Gabry J, Mahr T (2021). "bayesplot: Plotting for Bayesian Models." R package version 1.8.0, <URL:https://mc-stan.org/bayesplot/>.

## Images

- PowerPoint image bank