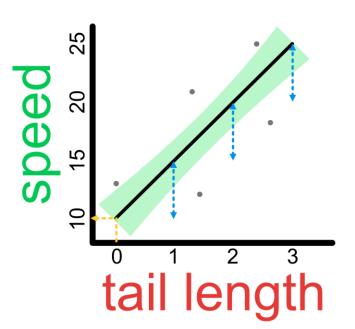
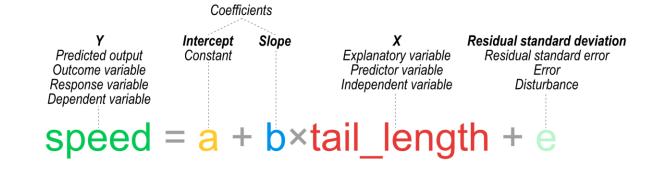
Modelling normal distribution

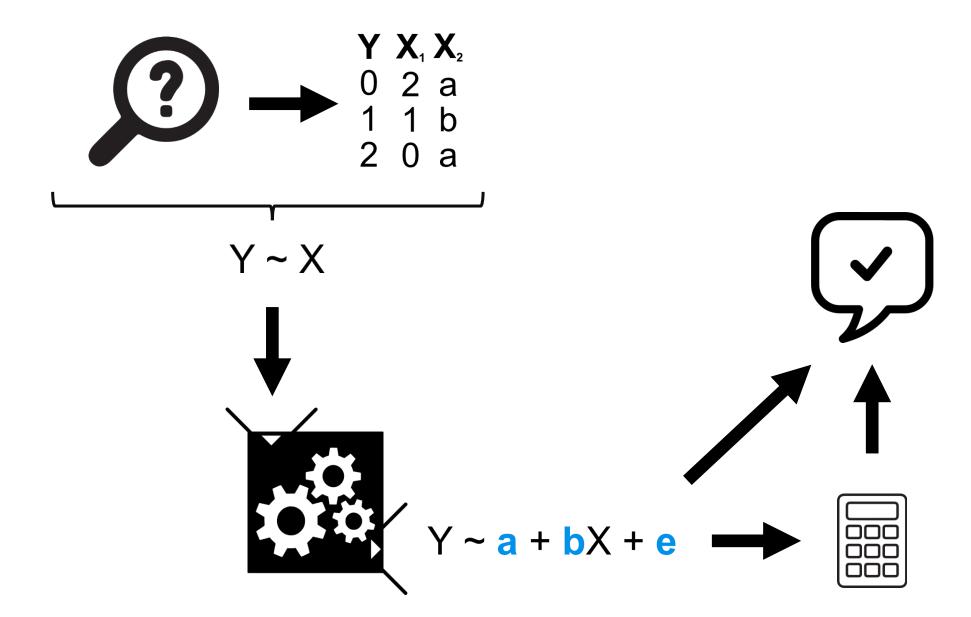
Pärt Prommik, PhD Ülo Maiväli, PhD



Recap

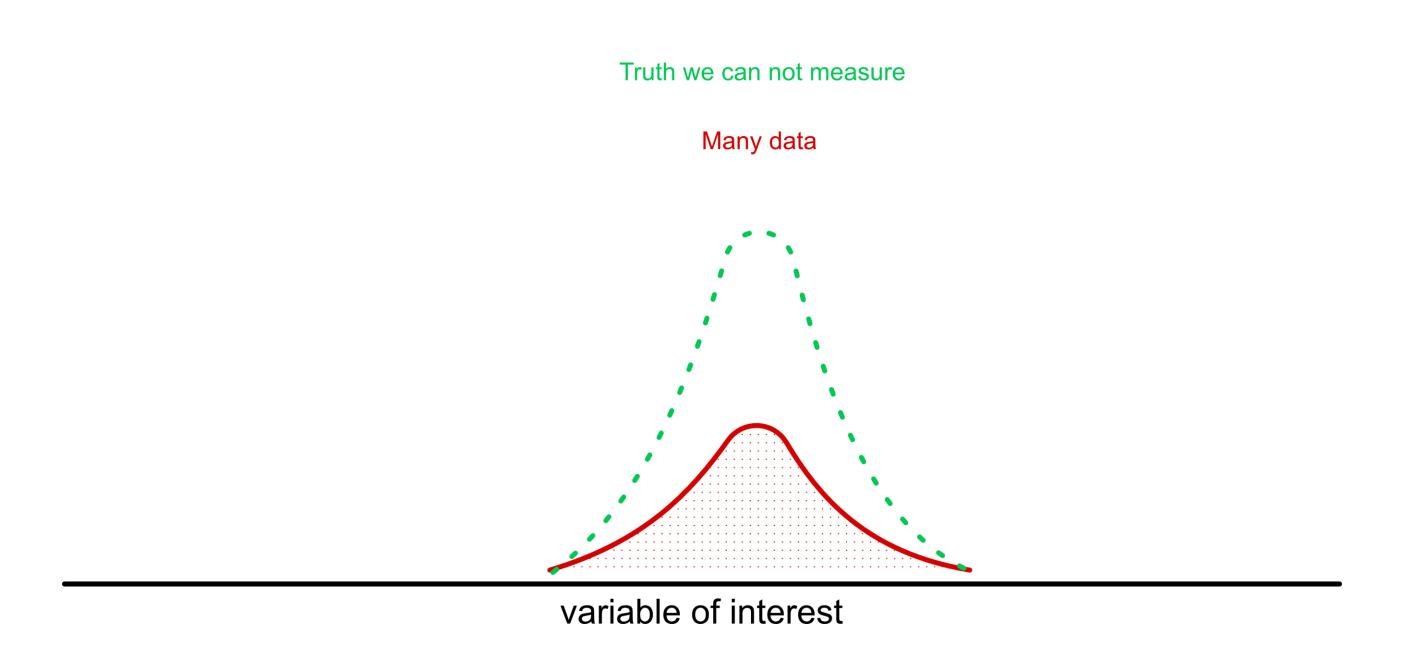






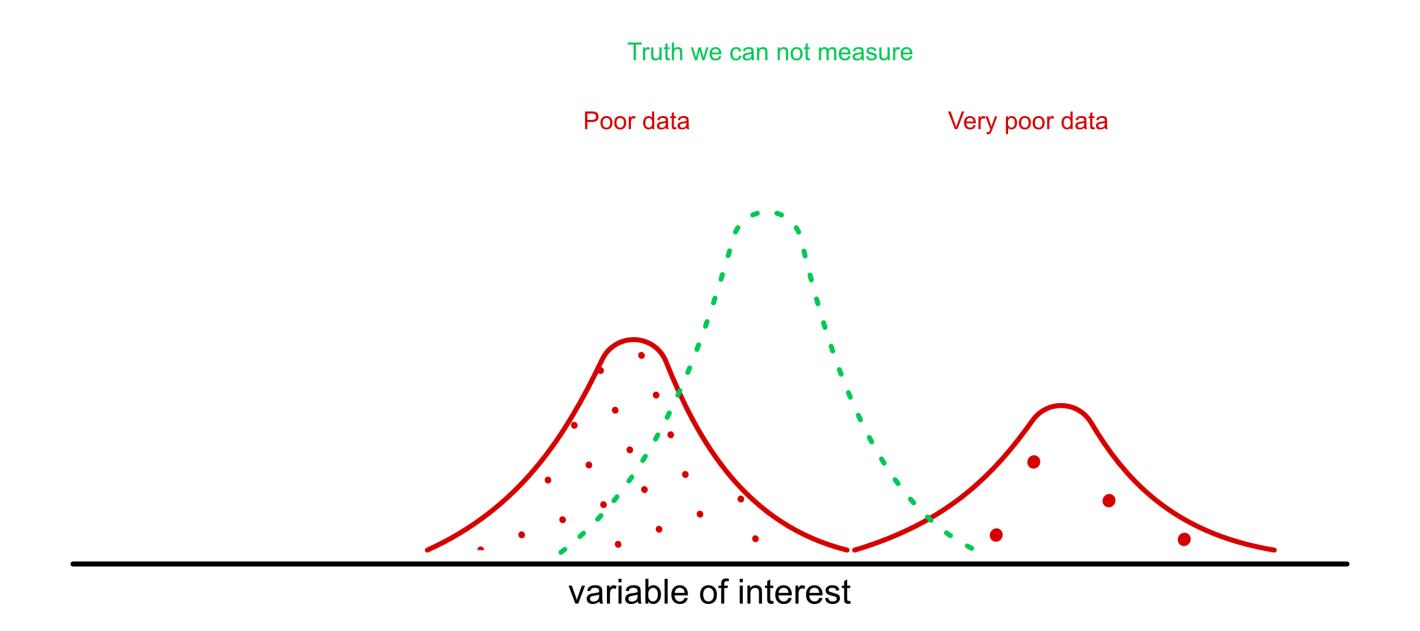


Classical regression is data-driven



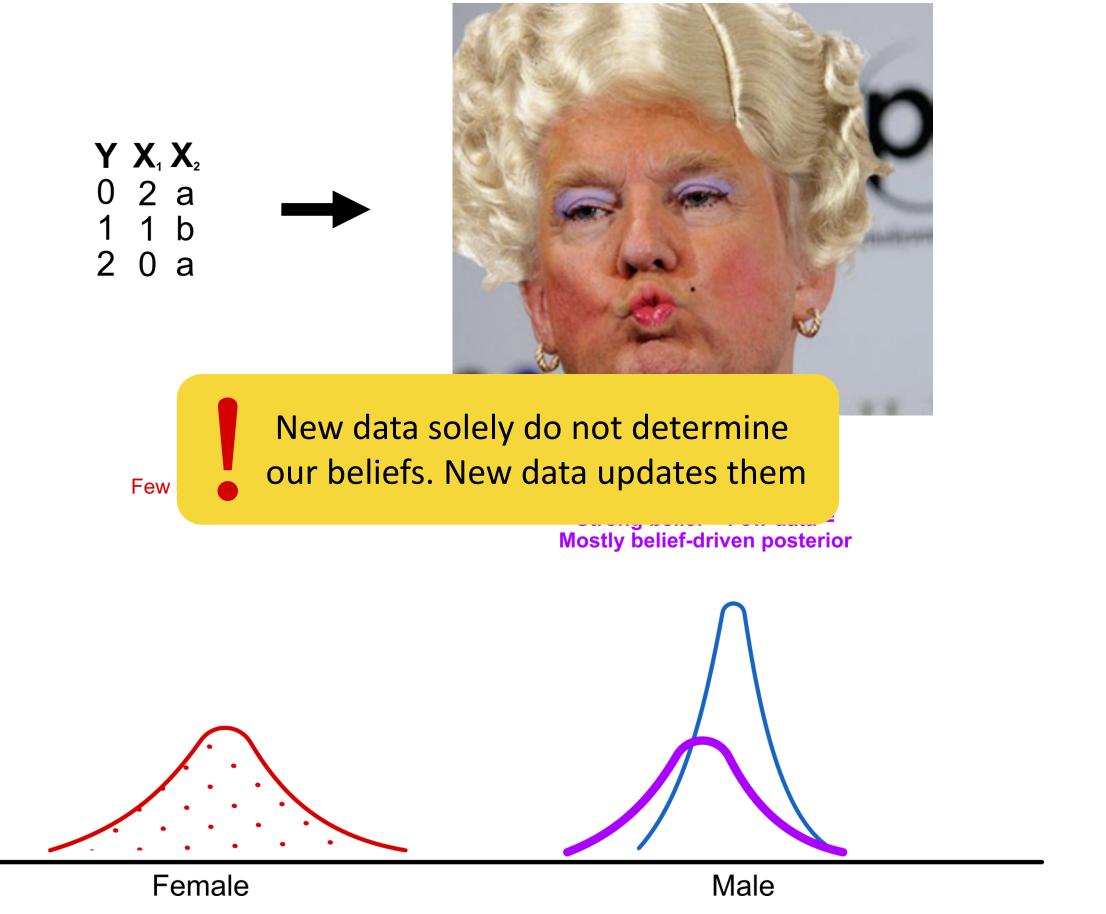


Classical regression is data-driven



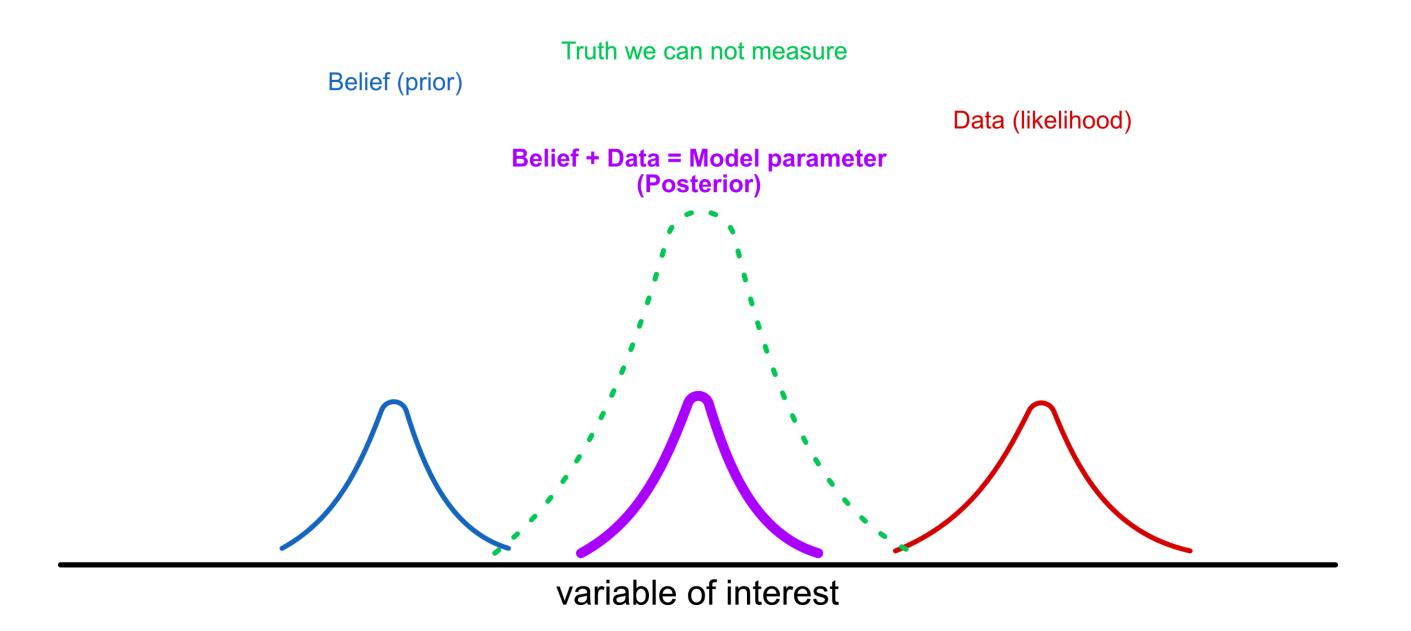


Bayesian thinking is different



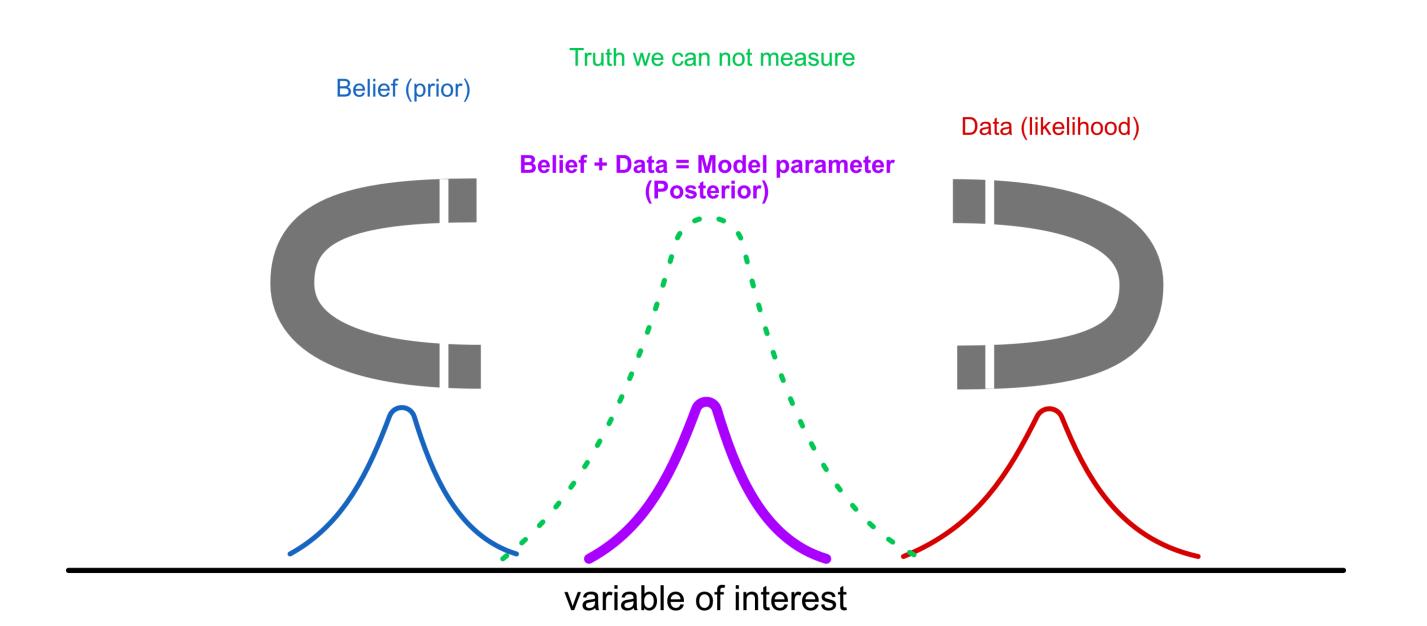


Inputting domain knowledge to an analysis





Inputting domain knowledge to an analysis

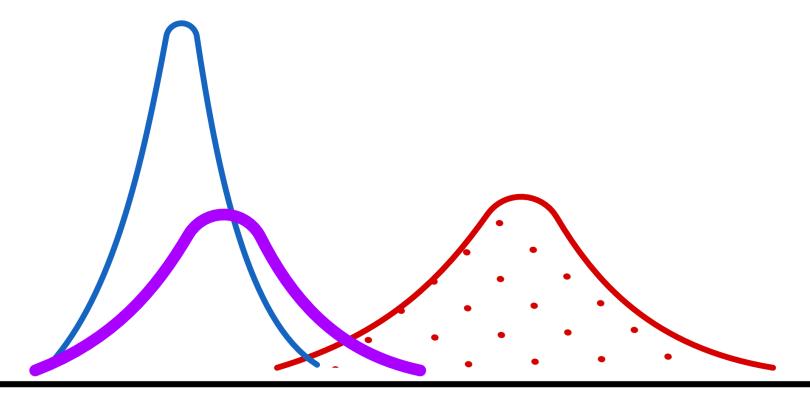




Strong belief (prior)

Few data (likelihood)

Strong belief + Few data = Mostly belief-driven posterior



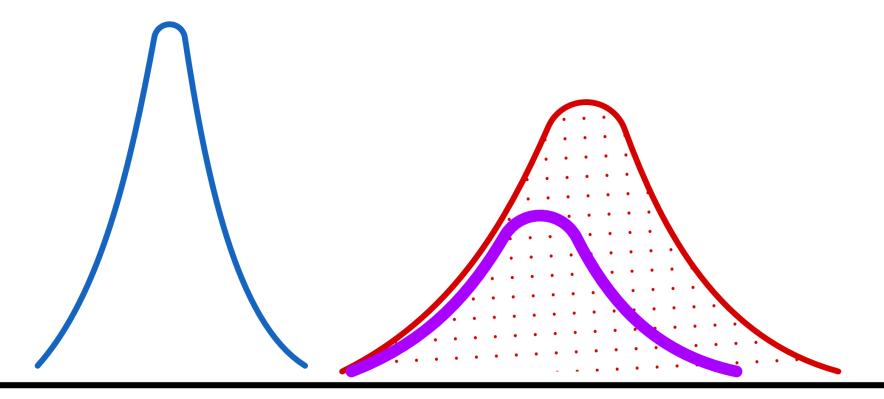
variable of interest





Many data (likelihood)

Strong belief + Few data = Mostly belief-driven posterior



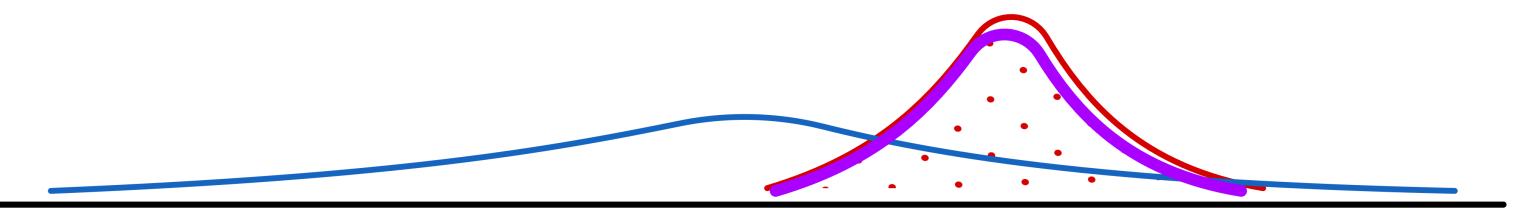
variable of interest



Weak belief (prior)

Few data (likelihood)

Weak belief + Data = Mostly data-driven posterior



variable of interest

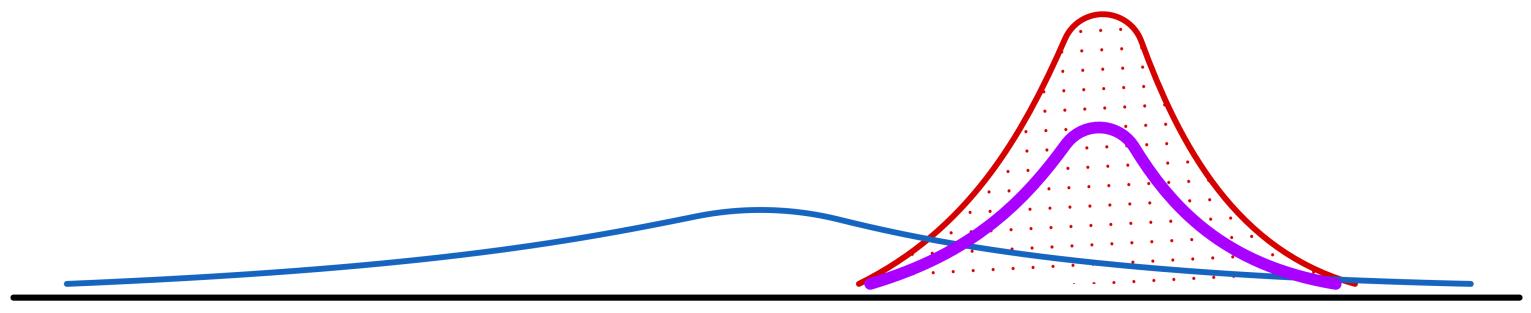




Many data (likelihood)

Weak belief + Data =

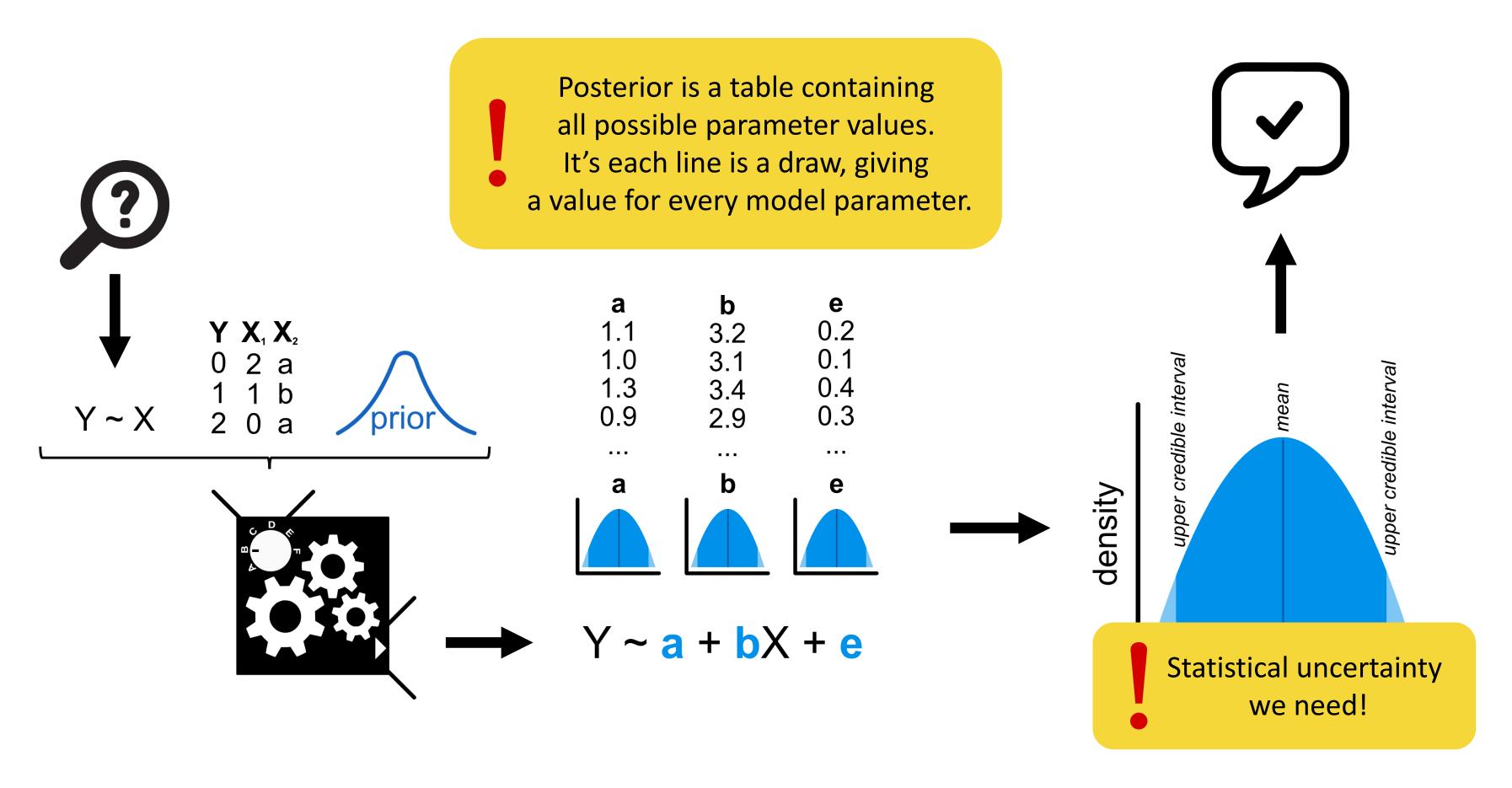
Mostly data-driven posterior



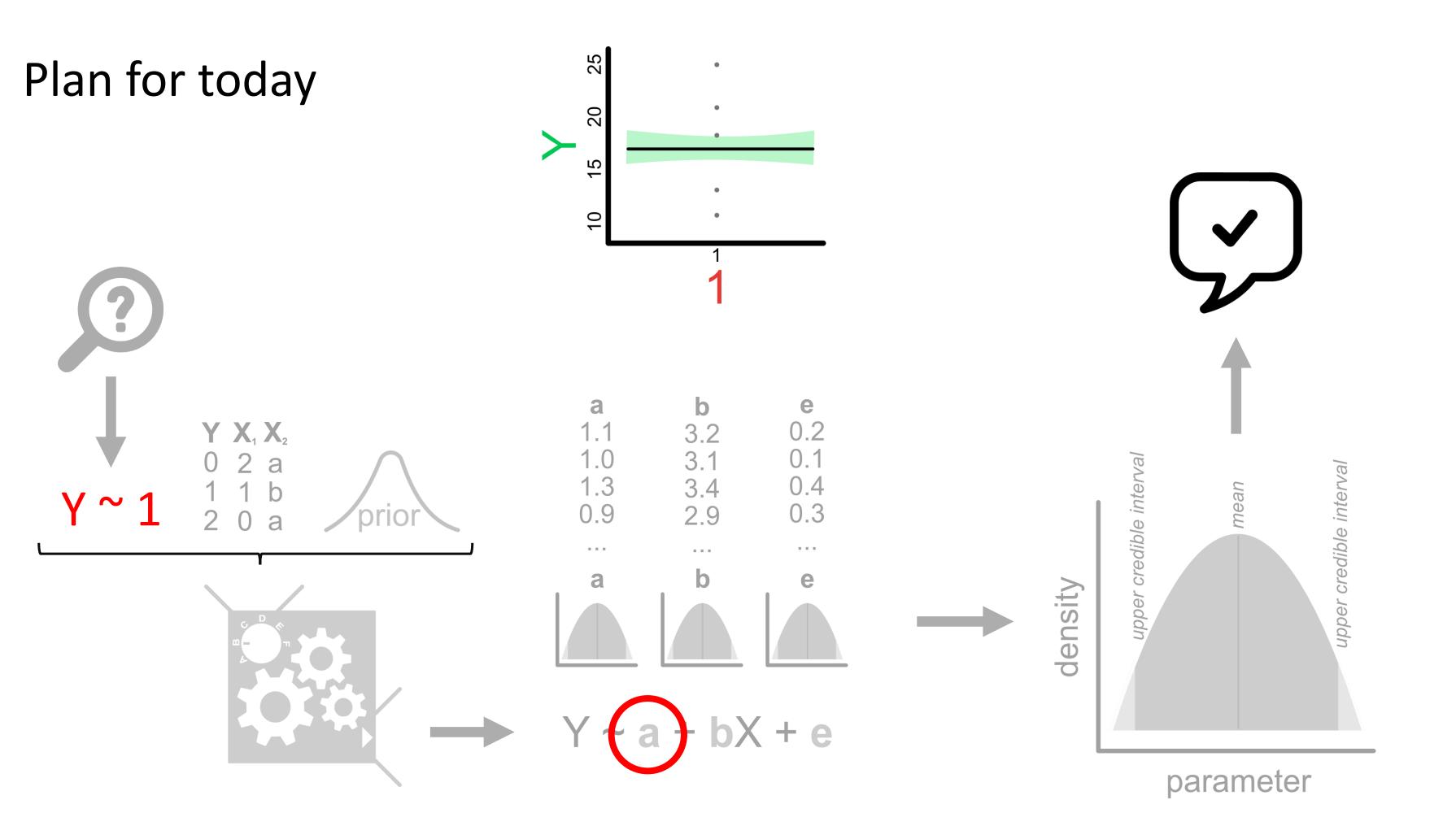
variable of interest



Bayesian regression = working with distributions

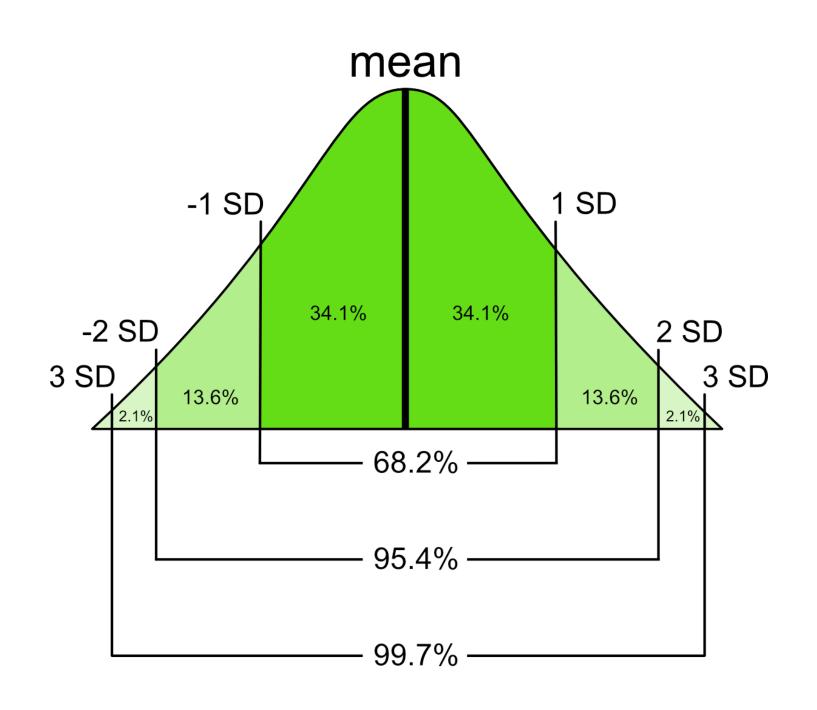


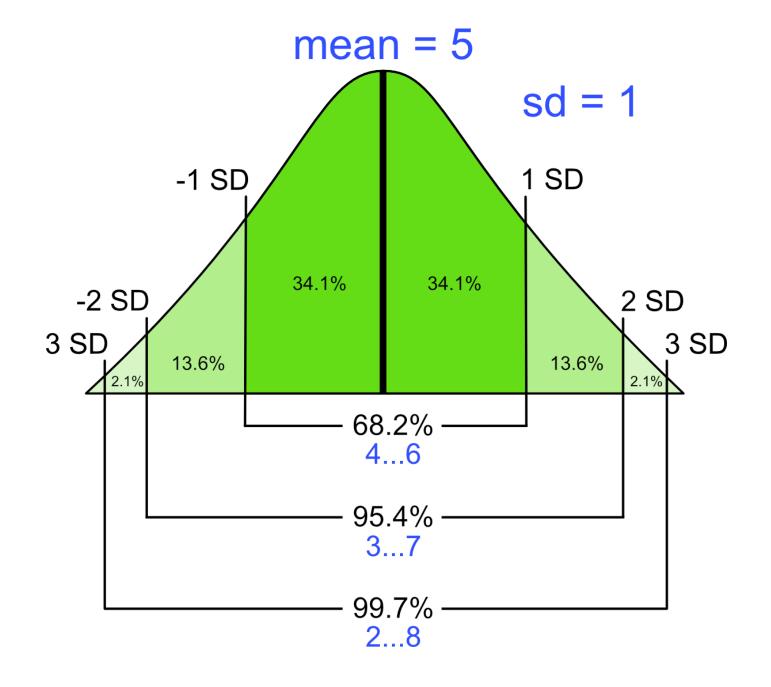






Prior is specified as a distribution

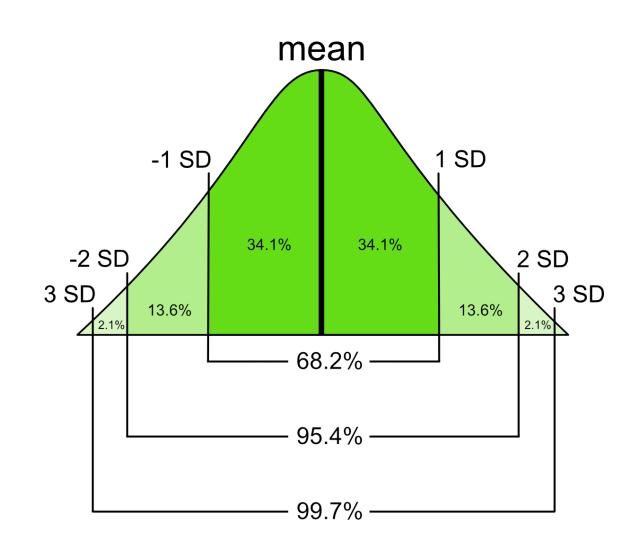


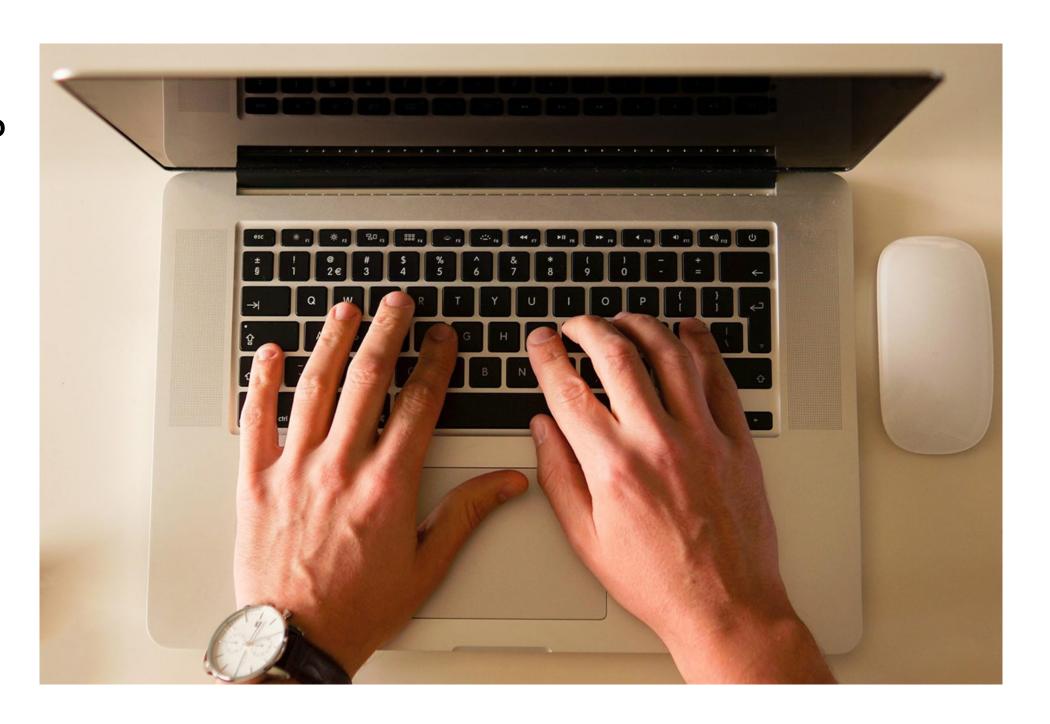




Define your first prior for writing speed in general population

- 1. Mean typed words per minute
- 2. How much the mean may vary (SD)?







Let's do some modelling



```
Family: gaussian
  Links: mu = identity; sigma = identity
Formula: words ~ 1
  Data: data (Number of observations: 17)
  Draws: 4 chains, each with iter = 2000; warmup = 1000; thin = 1;
         total post-warmup draws = 4000
Population-Level Effects:
          Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
             52.73
                                42.06
                                         63.42 1.00
                                                                 2300
                        5.45
                                                        2509
Intercept
Family Specific Parameters:
      Estimate Est.Error 1-95% CI u-95% CI Rhat Bulk_ESS Tail_ESS
                    3.26
                            16.54
                                     29.34 1.00
                                                             2411
         22.24
                                                    2635
sigma
Draws were sampled using sampling(NUTS). For each parameter, Bulk_ESS
and Tail_ESS are effective sample size measures, and Rhat is the potential
scale reduction factor on split chains (at convergence, Rhat = 1).
```



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                                                       2509
                                                                2300
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```



Family: gaussian

```
Y<sub>1</sub> Y<sub>2</sub> Y<sub>3</sub>
1 F 0
2 T 1
3 F 0
```

```
Links: mu = identity; sigma = identity
Formula: words ~ 1
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                   3.26
                         16.54 29.34 1.00
        22.24
                                                   2635
                                                            2411
sigma
```

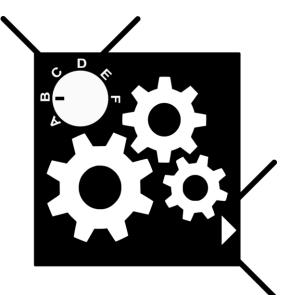
Draws were sampled using sampling(NUTS). For each parameter, Bulk_ESS

scale reduction factor on split chains (at convergence, Rhat = 1).

and Tail_ESS are effective sample size measures, and Rhat is the potential

Y₁ Y₂ Y₃ 1 F 0 2 T 1 3 F 0







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```

Y₁ Y₂ Y₃
1 F 0
2 T 1
3 F 0

Y ~ X

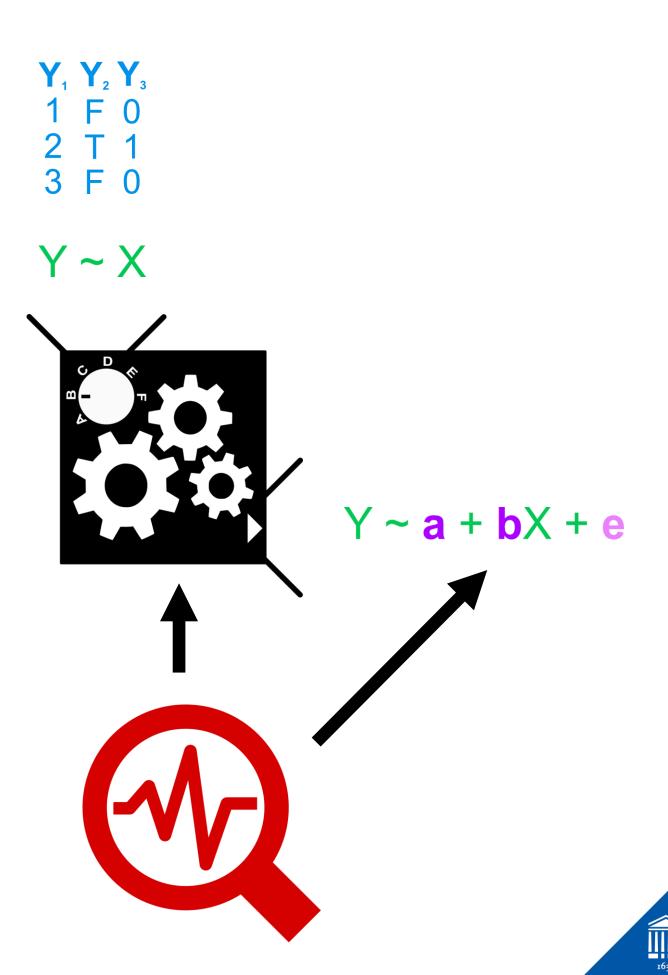


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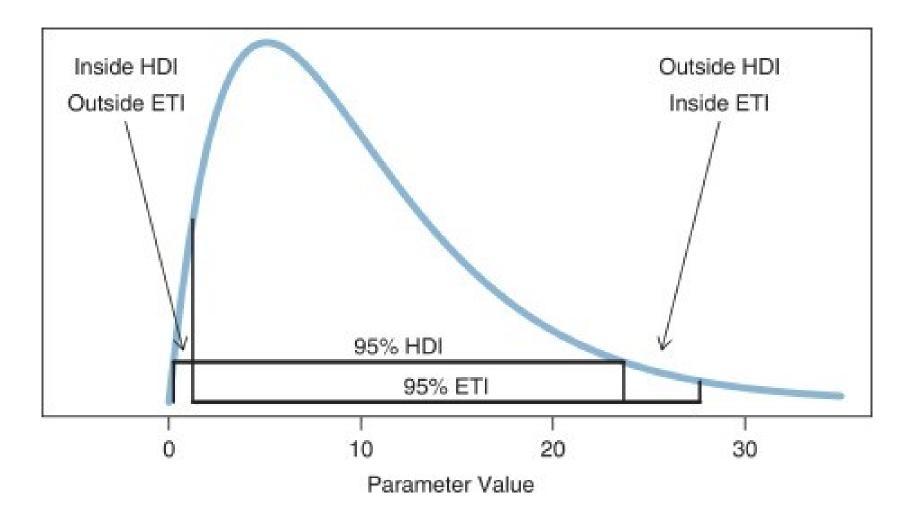
Y, Y, Y, 1 1 F 0 2 T 1 3 F 0 Y ~ X Y ~ a + bX +



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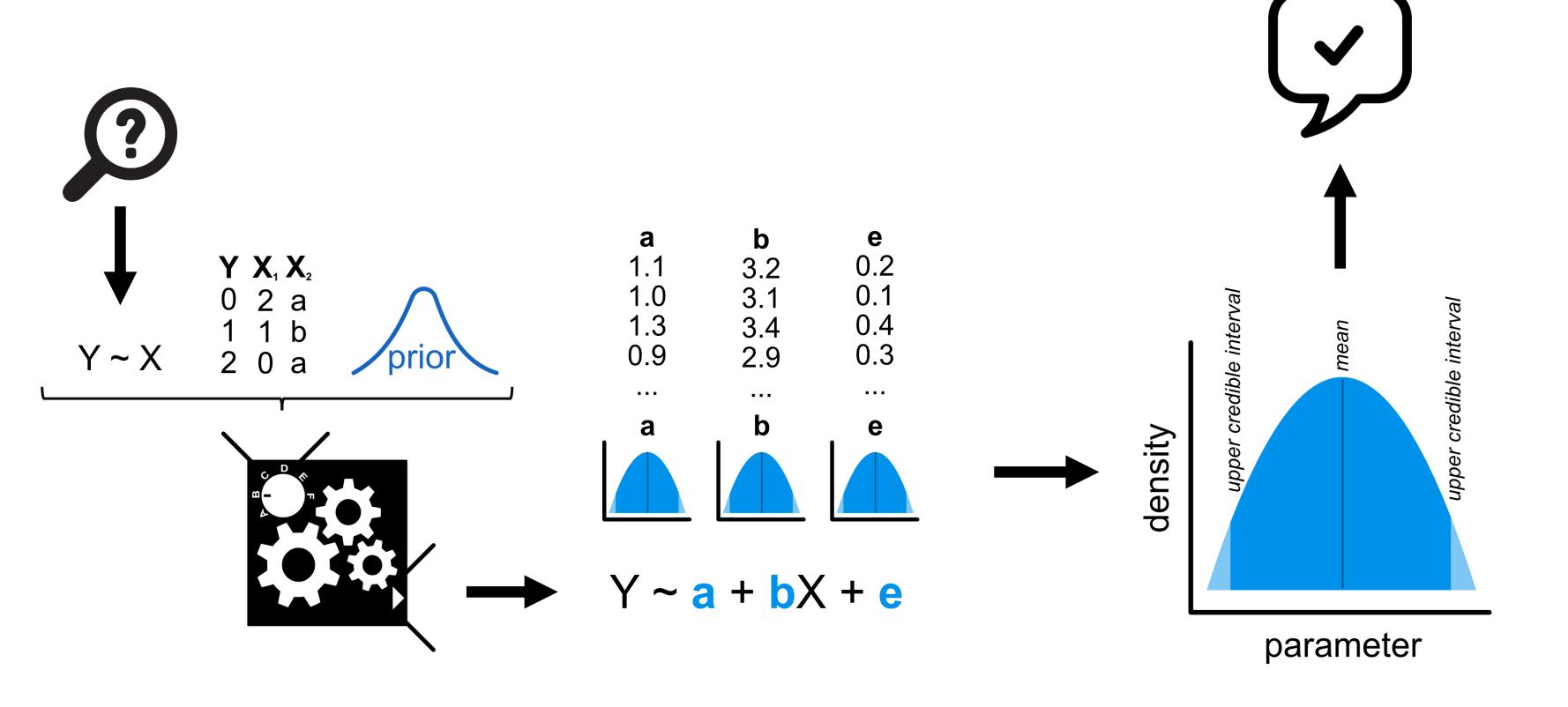


HDI vs ETI





Summary







Need help?



