

# Bayesian statistics

## 3/4

### *Hypothesis testing*

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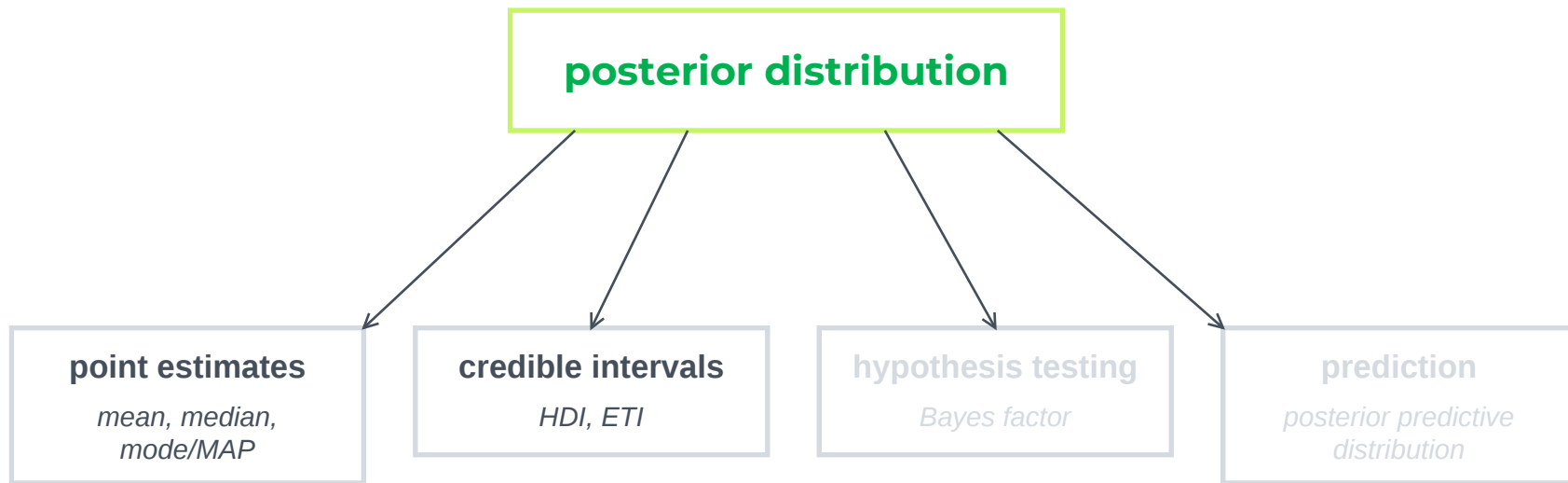
*Bayesian Statistics – CRNL – dec 2024*

1.

# Posterior-based hypothesis testing

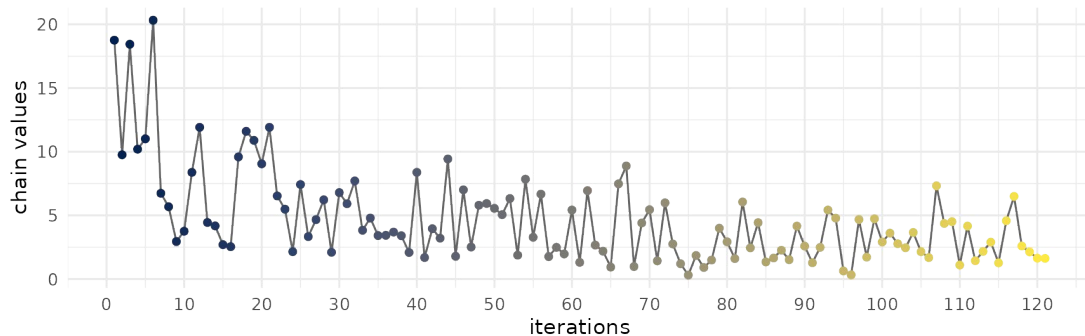
# The central role of the posterior distribution

In Bayesian statistics, all results are derived from the posterior distribution

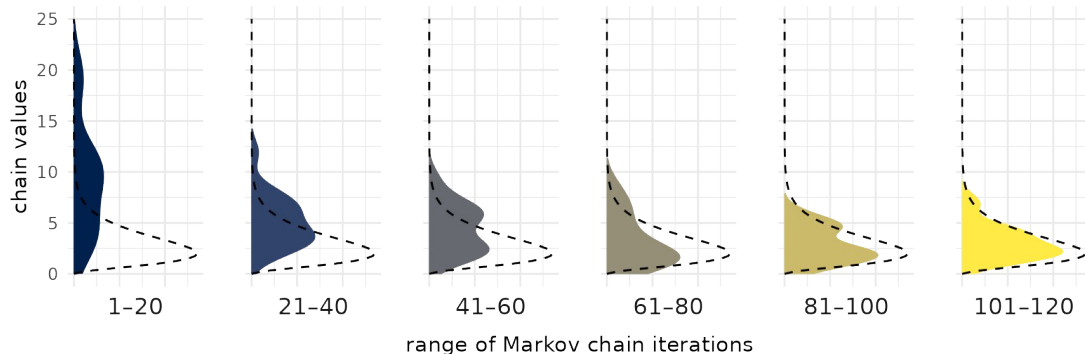


# Numerical simulation of the posterior: MCMC

Markov chain constructed to approximate the posterior distribution



Convergence of the Markov chain distribution towards the posterior distribution  
- - - True posterior distribution

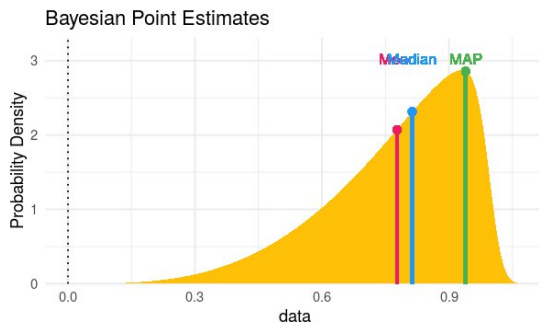


When prior distributions are **not conjugate distributions** of the likelihood, we don't have an explicit expression of the posterior distribution anymore and we need to calculate it **numerically**. We use **Markov chain Monte Carlo** (MCMC) techniques, a family of algorithms sharing the same basic procedure:

1. A **Markov chain** (= random process where each sample depends probabilistically on the previous one) is created such that it, *in the long run*, its distribution converges towards the true posterior distribution.
2. A large number of **samples** (several to tens of thousands) are generated **iteratively** from the Markov chain.
3. Initial samples (typically 1000) are considered as not converged yet and rejected ("**warm up**" phase); the rest of the samples is used as an **approximation of the posterior** distribution.

# Point and interval estimates

**bayestestR** package in the **easystats** ecosystem  
[easystats.github.io/bayestestR/](https://easystats.github.io/bayestestR/)



## point estimates

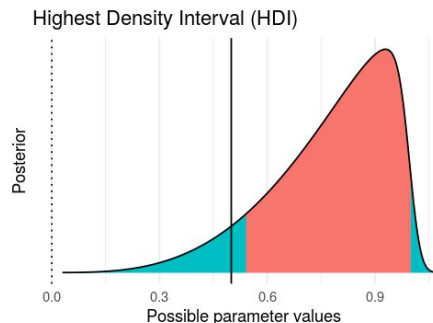
*mean, median,  
mode/MAP*

`point_estimate()`

## credible intervals

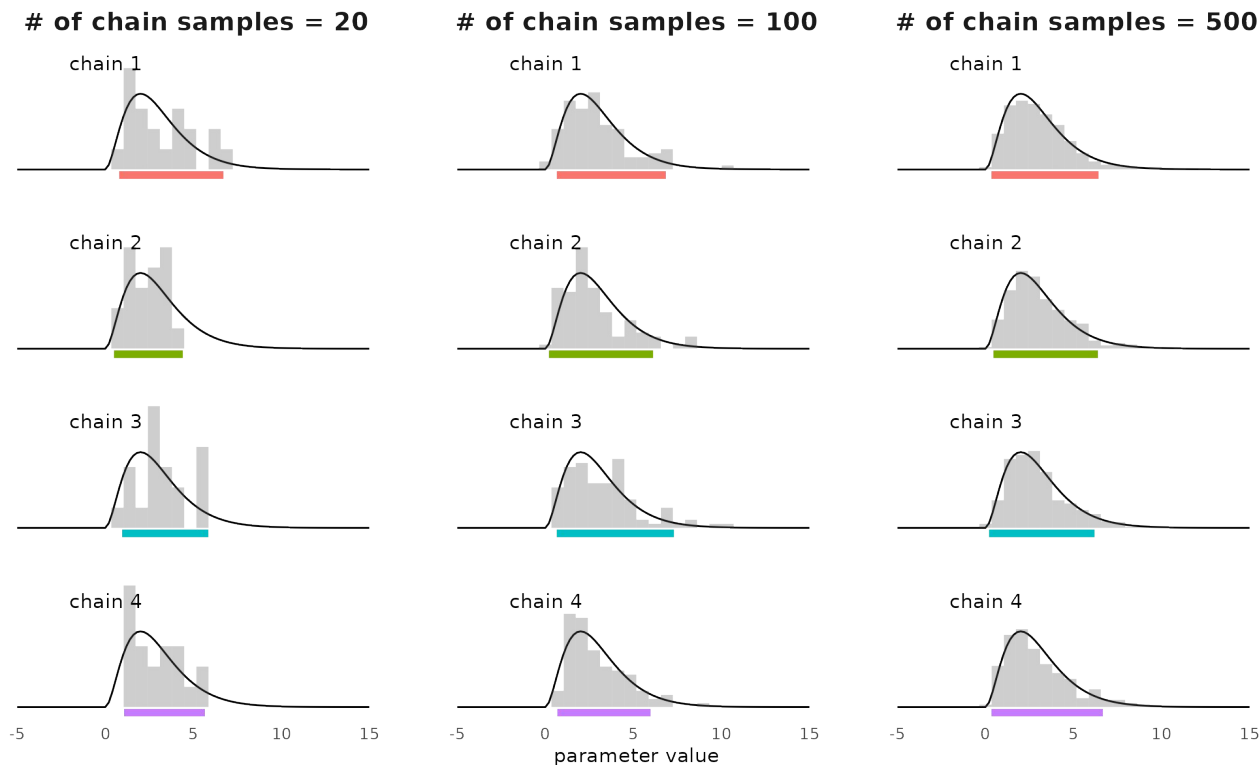
*HDI, (ETI)*

`hdi()`

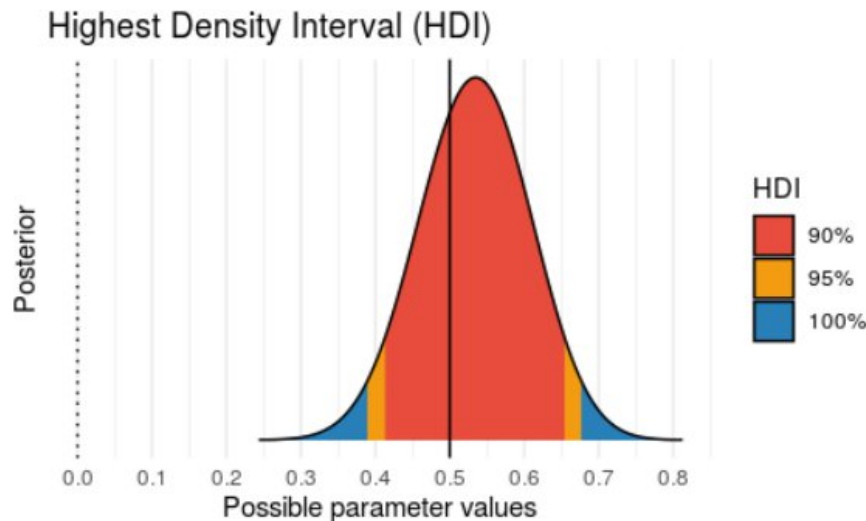


# Credible intervals & numerical simulations

**The more samples**  
in the posterior  
distribution, **the**  
**more stable** the  
credible interval



# Credible interval: 95% or 90%?



Compared to the 95%, the 90% credible interval is...

+ **more stable** to numerical errors  
- **less conservative**

→ Use **95%** if there are more than **10.000 samples** of the posterior distribution

In bayestestR, the default is **89%** (!) to highlight the arbitrariness of the confidence level.

# Frequentist vs. Bayesian statistics

## Frequentist

## Bayesian

### Definition of probability

Long-run frequency of events

Degree of belief / certainty

### View on model parameters

*True value:* unknown  
*Estimate:* fixed

*True value:* unknown  
*Estimate:* probabilistic

### Method of estimation

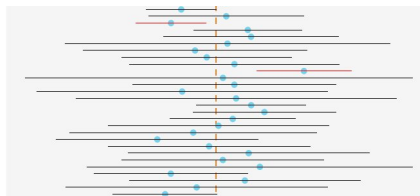
From the data only

From the posterior (data + prior)

### Uncertainty interval

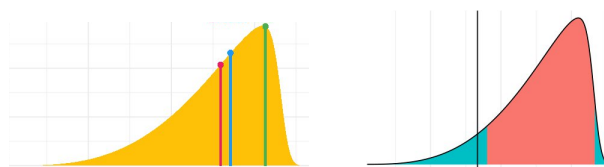
#### "Confidence intervals"

Confidence level (e.g. 95%) is a property of the procedure, not of the interval



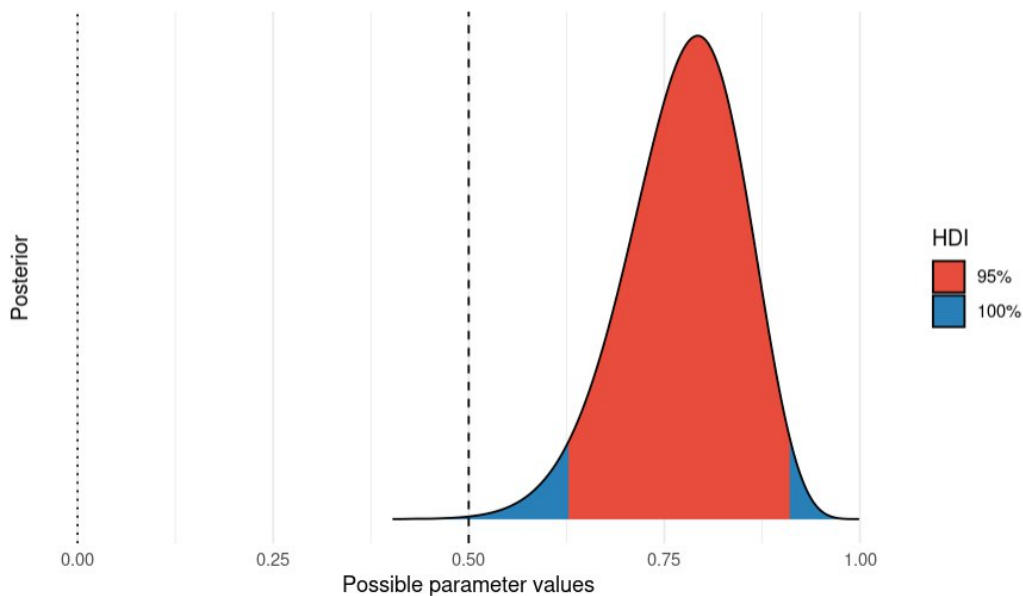
#### "Credibility intervals"

Confidence level (e.g. 95%) is a measure of the uncertainty around the estimate





# Can we test a hypothesis on a parameter from its posterior distribution?



# Hypothesis testing based on the posterior

## *Exact/point/precise hypothesis*

The **credible interval** defines a whole **range of exact hypotheses** that can be **rejected** with high confidence.

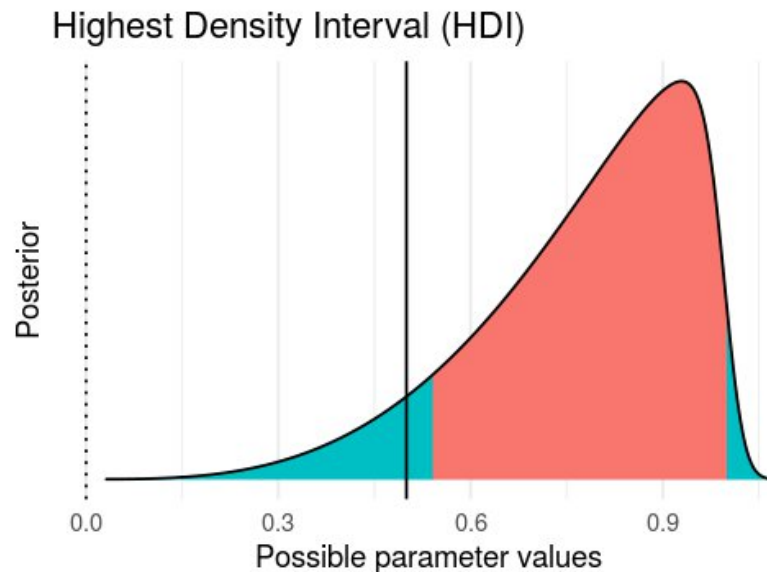
$$p(\theta_{low} \leq \theta \leq \theta_{high}) = .95$$

$$\Leftrightarrow p(\theta \notin [\theta_{low}, \theta_{high}]) = .05$$

$$\Rightarrow p(\theta = \theta_0) < .05$$

**Note:** this is very different from the frequentist  $p$ -value which is  $p(= y_{obs} | \theta = \theta_0)$

However, it does not allow to **accept** an exact hypothesis, only to reject it (at best).



# Hypothesis testing based on the posterior

## *Exact/point/precise hypothesis*

For a continuous parameter, the **absolute probability** of an exact hypothesis,  $p(H_0: \theta = \theta_0)$  is meaningless. That's why probability values are called **densities**.

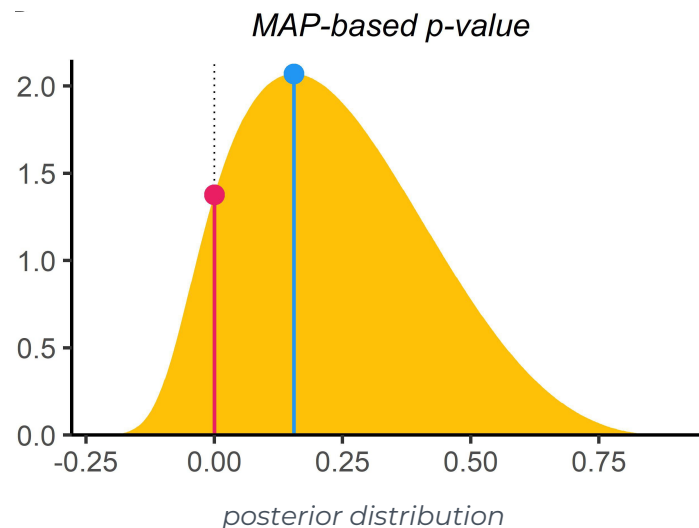
But we could compare the probability of the null hypothesis with the probability of the most likely value (the MAP):  $p(\theta = \theta_0)/p(\theta = \theta_{MAP})$

### Limitations:

- ignores most of the information contained in the posterior distribution
- can not provide evidence *for* the null: at best,  $\theta_0$  is the MAP and  $p = 1$

### Strengths:

- no need for hypothesis-specific priors (unlike BF)
- no Jeffreys-Lindley-Bartlett paradox (unlike BF)



`p_pointnull()` in **bayestestR**



# Hypothesis testing based on the posterior

## Range hypothesis

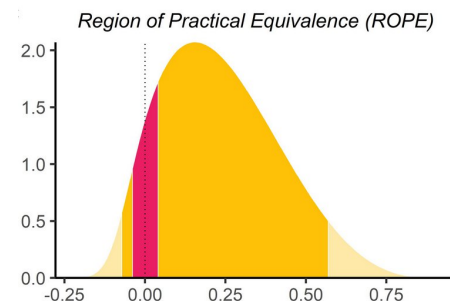
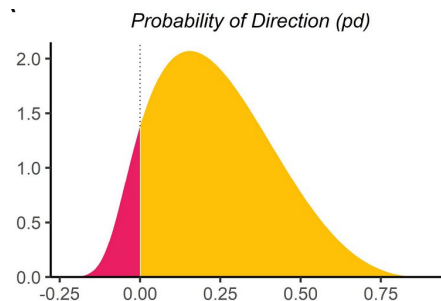
### Arguments against point hypotheses

- **the null hypothesis can always be rejected:** trivial deviations due to measurement bias, sampling error and other uncontrolled factors will come out as statistically significant given sufficiently large sample sizes (Meehl 1978, Cohen 1994)
- rejecting the exact null hypothesis (the frequentist NHST approach) is a **weak form of theory testing**
- **theories rarely make exact hypotheses** (except when they specify a complete and detailed mechanism, e.g. physical equation).

**Alternative:** use a **range hypothesis** as a more powerful statement. Two possibilities:

- test a **direction hypothesis**
- test a null range, also called the **region of practical equivalence (ROPE)**, that encodes negligible effects

Range hypotheses can be rejected, accepted or neither (no conclusion).



`p_direction()`  
in **bayestestR**



`p_rope()`  
in **bayestestR**

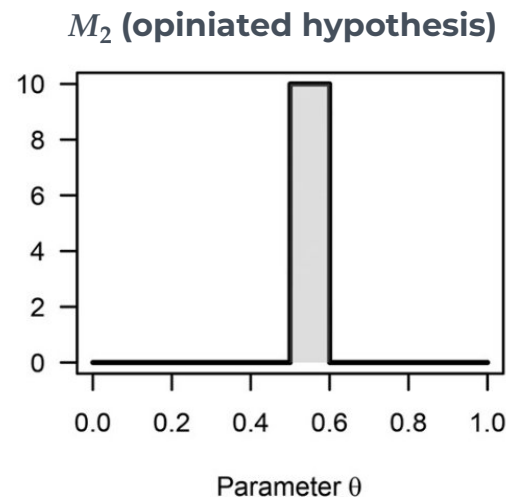
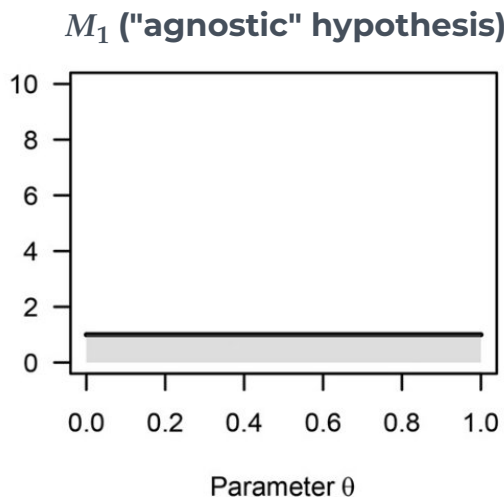
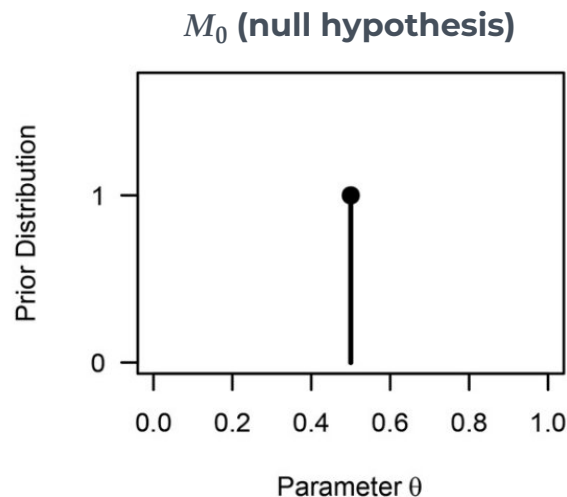
# 2.

## Bayes Factors

# Hypotheses as generative models

Different **hypotheses can be encoded as distinct models making specific predictions** for the data. Here, a model = statistical model + prior distribution for the parameters  $\theta$ .

Examples for a binary outcome experiment (coin toss, medical treatment, test accuracy, etc.):



# Bayes Factor

## As a (marginal) likelihood ratio

**Definition: of isolated hypotheses is of little interest.** We are usually interested in how *competing* hypotheses are *differentially* supported by data.

**Definition:** the **Bayes Factor** is the ratio of the likelihoods of two statistical models, integrated over the prior probabilities of their parameters (“marginal likelihood):

$$BF_{21} = \frac{p(y|M_2)}{p(y|M_1)} = \int \frac{p(y|\theta_2)p(\theta_2)d\theta_2}{\int p(y|\theta_1)p(\theta_1)d\theta_1}$$

**Interpretation:** “Model  $M_2$  supports the data  $BF_{21}$  times more than  $M_1$ ”

If  $M_0$  is a **null hypothesis**, then:

$$BF_{10} = \frac{p(y|M_1)}{p(y|M_0)} = \frac{\int p(y|\theta_1)p(\theta_1)d\theta_1}{p(y|\theta_0)}$$

Compare with the **likelihood ratio test**:

$$LR = \frac{p(y|\theta_{MLE})}{p(y|\theta_0)}$$

# Bayes Factor

## *As relative belief updating*

Applying the Bayes theorem:

$$BF_{10} = \frac{p(y|M_1)}{p(y|M_0)} = \frac{\frac{p(M_1|y)}{p(M_1)}}{\frac{p(M_0|y)}{p(M_0)}}$$

= belief updating of model  $M_1$

= belief updating of model  $M_0$

Which we can rewrite :

$$\frac{p(M_1|y)}{p(M_0|y)} = BF_{10} \times \frac{p(M_1)}{p(M_0)}$$

posterior model odds

prior model odds

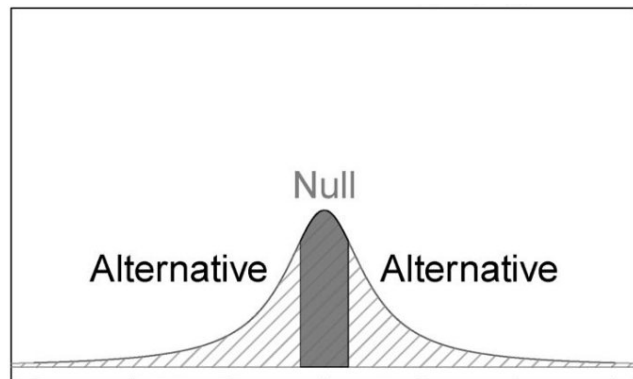
Thus, the Bayes factor is...

- ...a continuous measure of evidence
- ...a predictive updating factor
- ...independent of models' prior probability
- ...equal to the posterior model odds if models are equally probable *a priori*

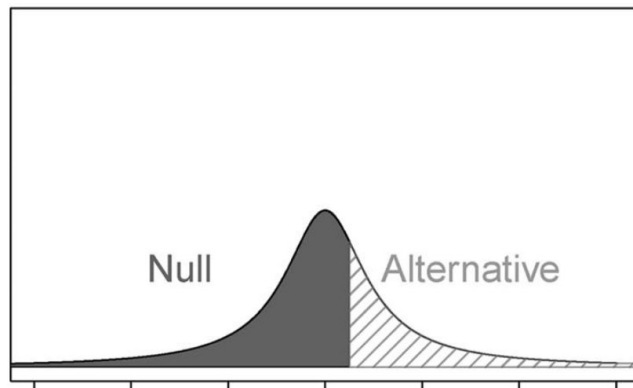


# Bayes Factor for range hypotheses

The calculation of the BF is straightforward when the competing models are **non-overlapping, complementary intervals from the same distribution**



“two-sided”

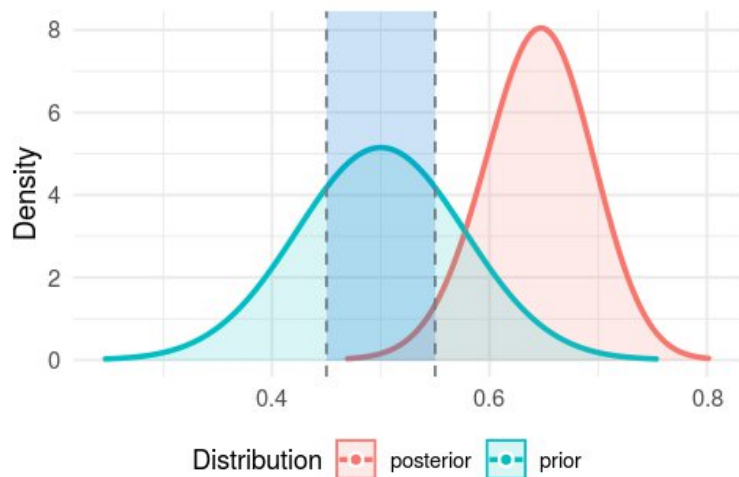


“one-sided”

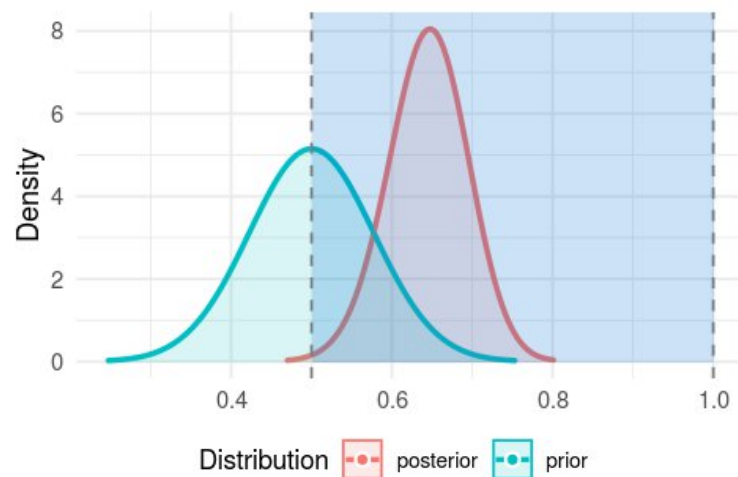
# Bayes Factor for range hypotheses



## Region of practical equivalence (ROPE)



## Probability of direction



`bf_params()` in `bayestestR`

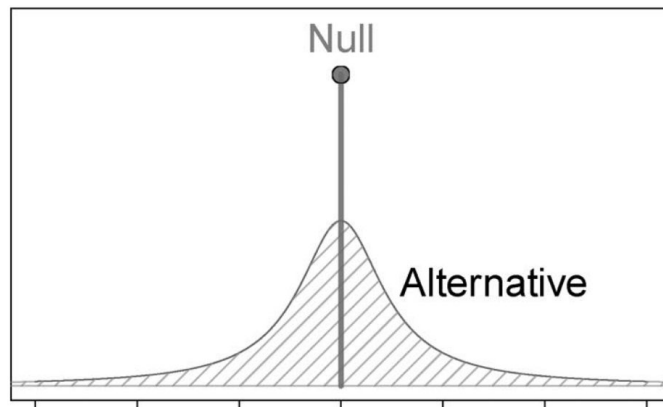
# Bayes Factor for exact hypotheses

## *Savage-Dickey density ratio*

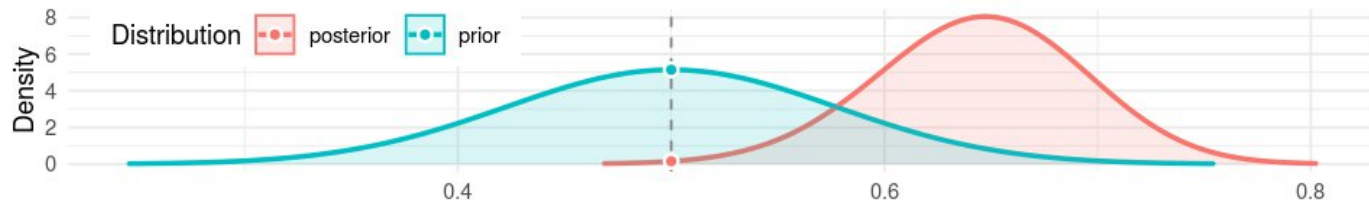


Let  $H_0$  be an exact null hypothesis ( $H_0: \theta = \theta_0$ ) and  $H_1$  the complementary hypothesis ( $H_1: \theta \neq \theta_0$ ). Then:

$$BF_{01} = \frac{1}{p(\theta = \theta_0)}$$



This special case of the Bayes Factor is called the **Savage-Dickey density ratio**.



`bf_params()`  
in `bayestestR`

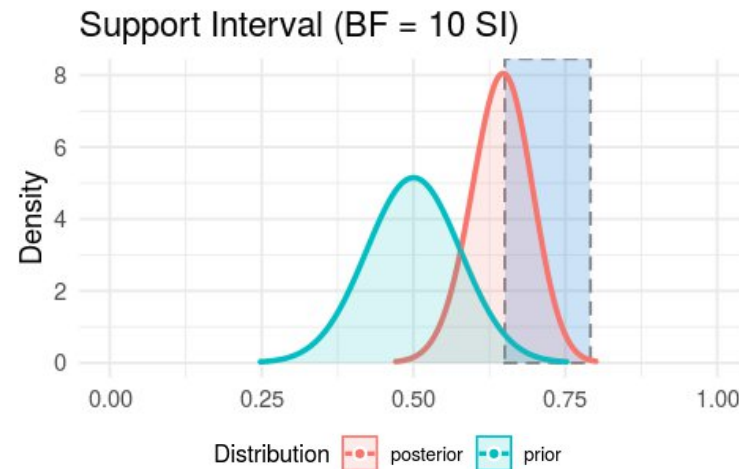
# Bayes Factor for exact hypotheses

## *Support interval*



Which values of the parameter are supported by data?

**Support interval** = all values for which the Savage-Dickey density ratio is above a certain threshold



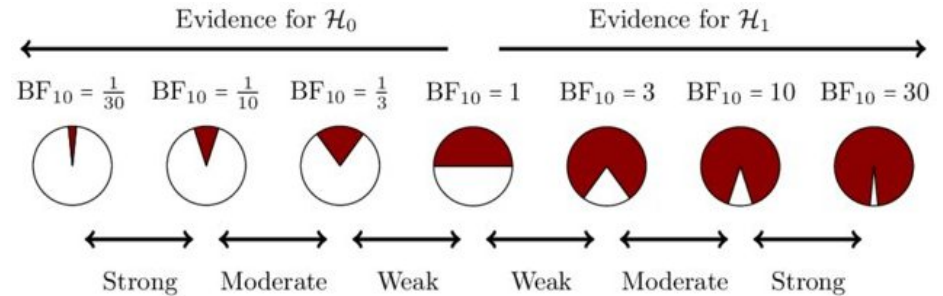
`si()` in `bayestestR`

# Bayes Factor

## Measure of evidence

**Conventional interpretation of Bayes factor values** (Kass & Raftery 1995)

$BF$	$\log_{10}BF$	Strength of evidence
1 to 3	0 to $1/2$	Barely worth mentioning
3 to 10	$1/2$ to 1	Substantial
10 to 100	1 to 2	Strong
$> 100$	$> 2$	Decisive



**Don't replace the p-value dichotomous ritual by a BF multichotomous ritual!!**

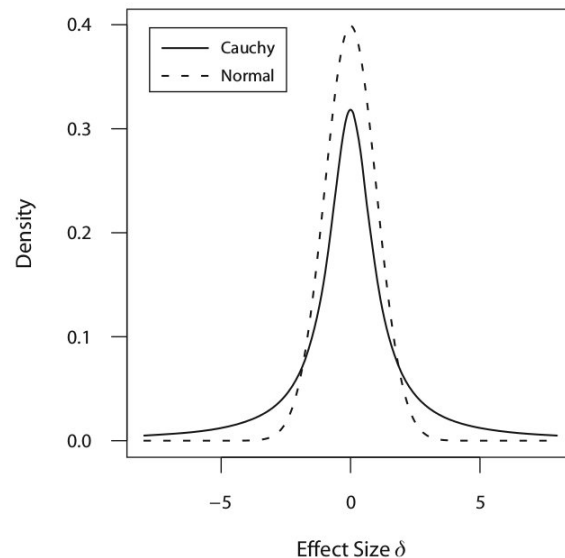
# Bayes Factor

## Application to the two-sample location test

Approach implemented in **BayesFactor** and **JASP**:

- parametrize the model in terms of the standardized effect size ( $\sim$  Cohen's  $d$ ):  $\delta = \mu / \sigma$
- prior = Cauchy distribution with scale  $r$  ( $\sim$  variance)
- $\Rightarrow$  fatter tails than the normal distribution

`ttestBF()` in the  
**BayesFactor** package

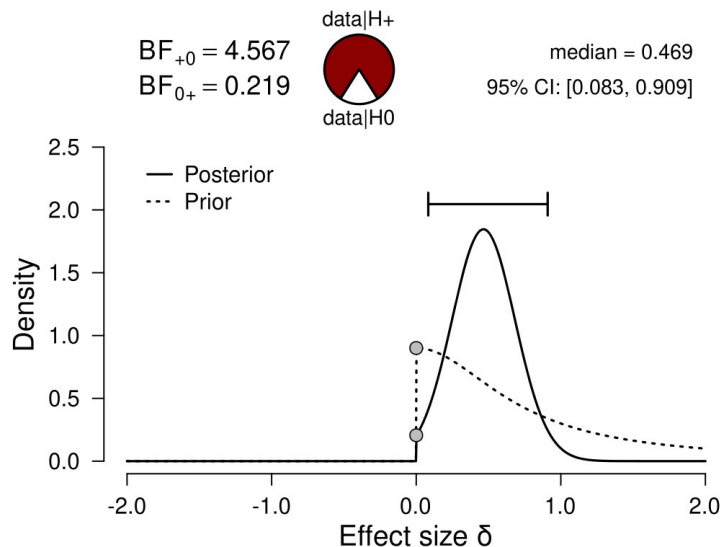


Rouder, Speckman, Sun, Morey & Iverson (2009). *Bayesian  $t$  tests for accepting and rejecting the null hypothesis*.

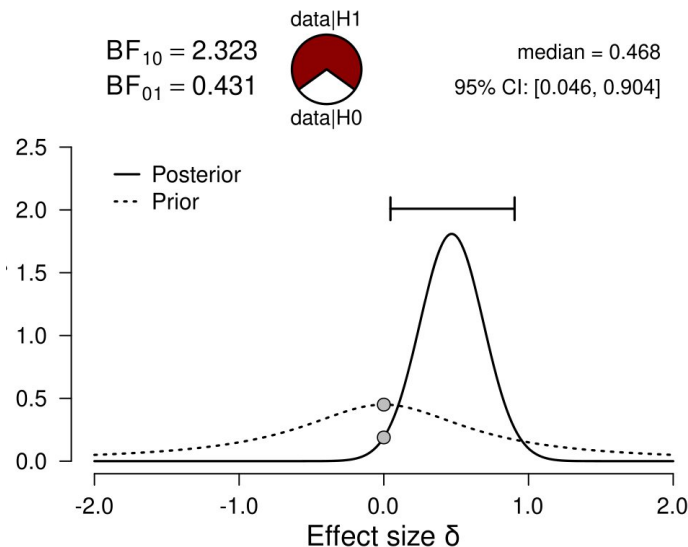
Psychonomic Bulletin & Review doi.org/10.3758/PBR.16.2.225

# Bayes Factor

## Application to the two-sample location test



(a) One-sided analysis for testing:  
 $H_+ : \delta > 0$



(b) Two-sided analysis for estimation:  
 $\mathcal{H}_1 : \delta \sim \text{Cauchy}$

# Bayes Factor

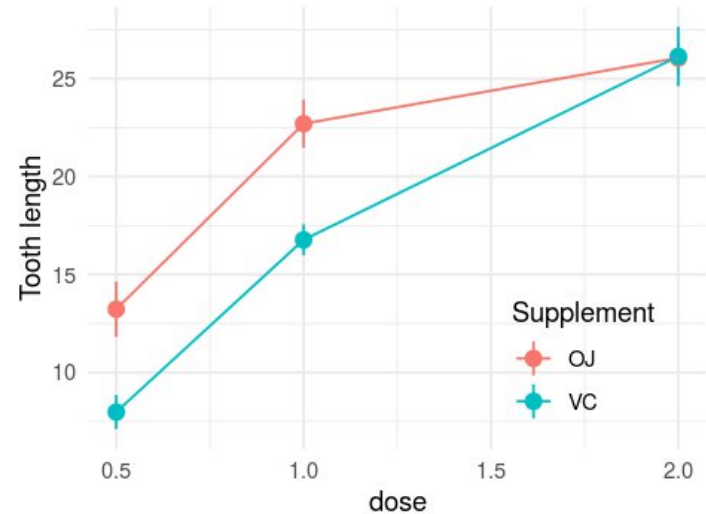
## *Application to the ANOVA*



Guinea pigs are assigned to one of two treatments (vitamin C or orange juice) in one of three doses. The effect on tooth growth is measured.



or





# Bayes Factor

## Application to the ANOVA



### New challenges:

- multiple variables  $\Rightarrow$  multiple parameters
- a single variable (here, *dose*) can be encoded with **2** parameters

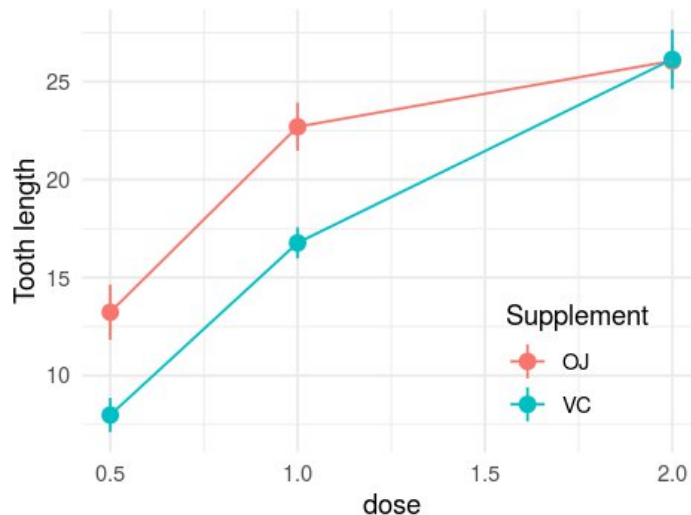
**Solution** implemented in *BayesFactor* and *JASP*

= **comparison between nested models**

e.g. with ( $M_2$ ) and without ( $M_1$ ) the interaction

$$BF_{21} = \frac{p(y|M_2)}{p(y|M_1)}$$

BF apply not only to models with *different priors*,  
but also to models with *different structures*!



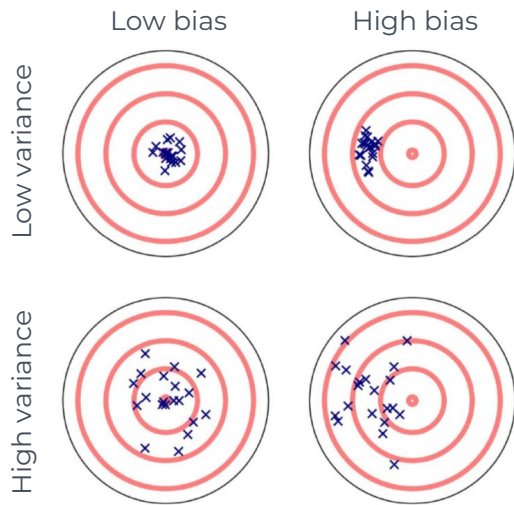
# Bias-variance tradeoff



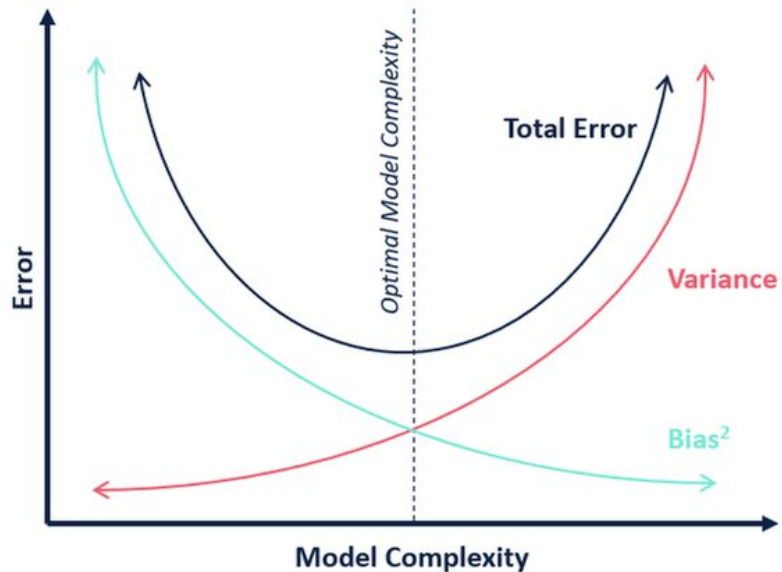
**Bias** = systematic error due to inadequate model (**underfitting**)

**Variance** = sensitivity to small fluctuations in the data (**overfitting**)

→ variability of parameter estimates across replications



Target center = true value  
Crosses = model predictions



# Bayes Factor

## *As an Occam's razor*

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$$BF_{21} = \frac{p(y|M_2)}{p(y|M_1)} = \frac{\int p(y|\theta_2)p(\theta_2)d\theta_2}{\int p(y|\theta_1)p(\theta_1)d\theta_1}$$

**Model complexity** is automatically **penalized** by the Bayes Factor: the more parameters, the more the prior is spread out over “irrelevant” regions, the more “diluted” the predictive power of the model

⇒ diffuse priors follow the same logic

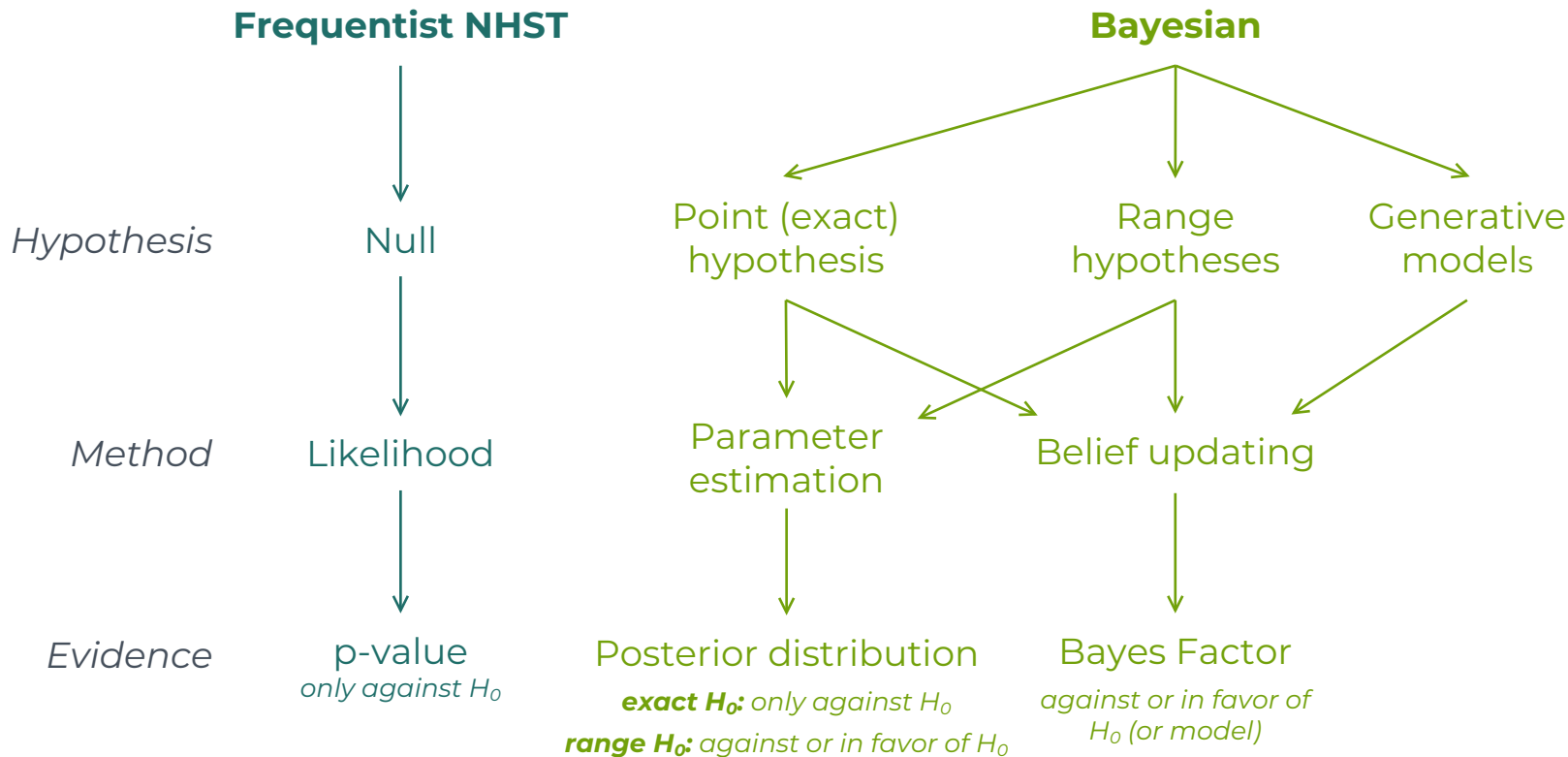
# Bayes Factor

*Application to computational models*

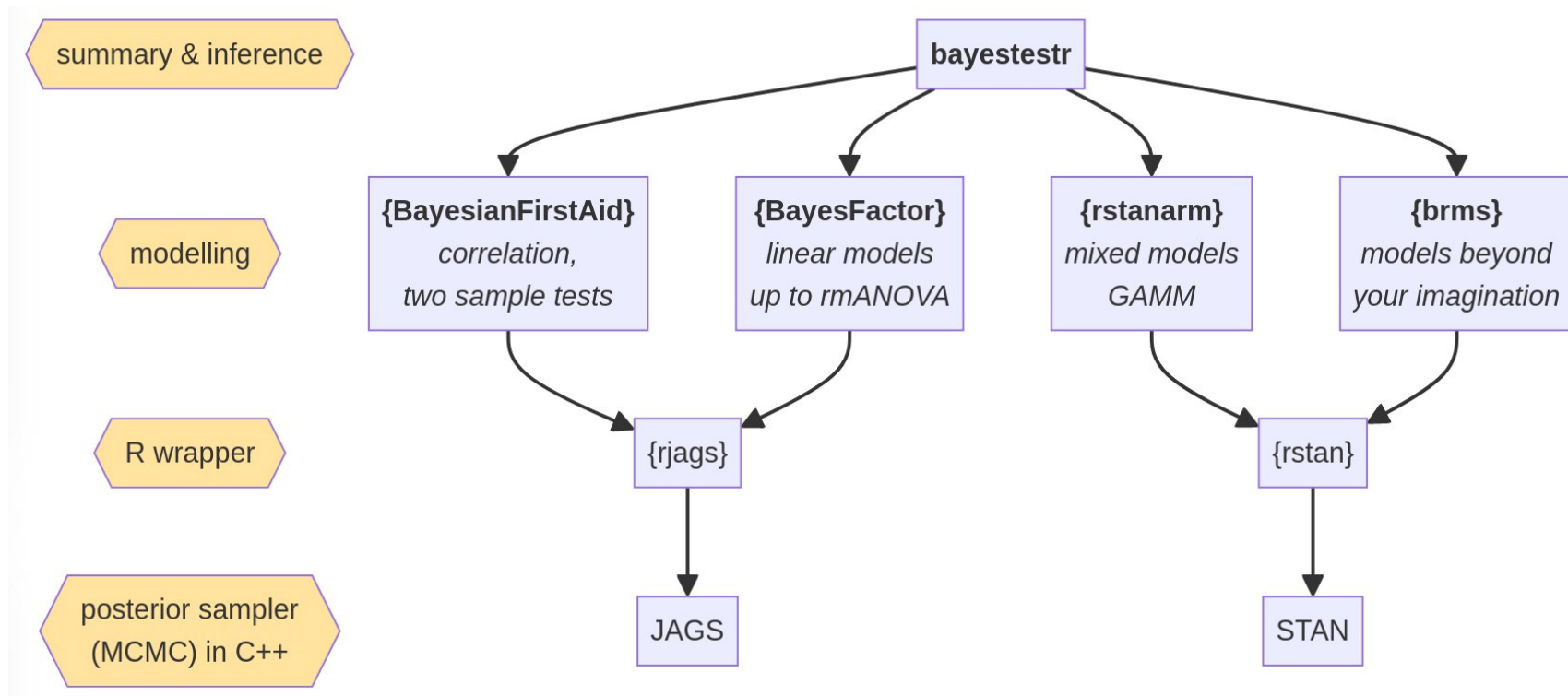
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TO DO

# The many ways of Bayesian hypothesis testing



# Software & package ecosystem



# Frequentist vs. Bayesian statistics

	Frequentist	Bayesian
<b>Definition of probability</b>	Long-run frequency of events	Degree of belief / certainty
<b>View on model parameters</b>	Fixed	Probabilistic
<b>Point estimates</b>	Derived from the sample	Derived from the posterior distribution
<b>Interval estimates</b>	Confidence interval ; confidence level is a property of the procedure, not of the intervals themselves	Credibility intervals ; confidence level is a statement about the uncertainty of the model parameters
<b>Hypothesis testing</b>	Point hypotheses only Can only reject a hypothesis	Point and range hypotheses Can select the best one among multiple
<b>Limitations</b>	<b>Interpretability</b> <b>Usefulness</b>	<b>Time consuming (prior + computation)</b> <b>Lack of standards, rapid evolution</b>