## Model Selection

**UofSC Python STEM workshop** 

## AIC / BIC

Bayesian Evidence

# Some questions we should ask when fitting a model to data

Does this model match my data?

Can this model help to **explain** my data?

Can this model **predict** future / different data?

Is this the **best** model for my data?

## Some questions we should ask when fitting a model to data

Does this model match my data?

Can this model help to **explain** my data?

Goodness of fit tests (e.g. chi^2, p-values)

Model validity: apply Physics / Chemistry / Biology / Sociology / ...

Can this model **predict** future / different data?

Cross-validation; New experimentation

Is this the **best** model for my data?

"Best" may mean: good match + least complex

Alternative: use all the models! (Weighted model combination, Bayesian model averaging)

Model selection (e.g. AIC, BIC, Bayesian evidence / Bayes factors)

The Akaike Information Criterion

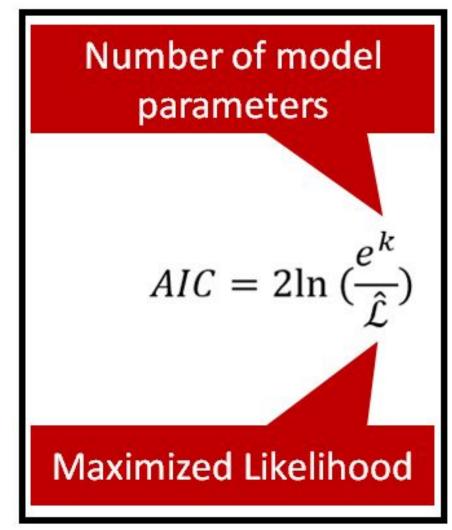
Number of model parameters

$$AIC = 2k - 2\ln(\mathcal{L})$$

 $\mathcal{L} = \mathcal{L}(\widehat{\theta})$  = maximum value of the likelihood function of the model

### Rewriting the AIC

- As model becomes more complex, the numerator grows.
- As max likelihood increases, the denominator grows



## To compare two models, use the difference of the AIC from each.

Reframed as an exponent, this gives the relative likelihood of model 1 to model 2. (can also be phrased as 'model weights' or 'odds ratios')

 $Relative\ likelihood = e^{(AIC_1 - AIC_2)/2}$ 

The
Bayesian
Information
Criterion



Number of data points

$$BIC = k \ln(n) - 2\ln(\mathcal{L})$$

 $\mathcal{L} = \mathcal{L}(\widehat{\theta})$  = maximum value of the likelihood function of the model

Model Selection with Bayesian Evidence

# Bayesian evidence (a.k.a. Bayes factors comparison)

The AIC and BIC use only the maximum likelihood values - not the full likelihood (or posterior) distribution.

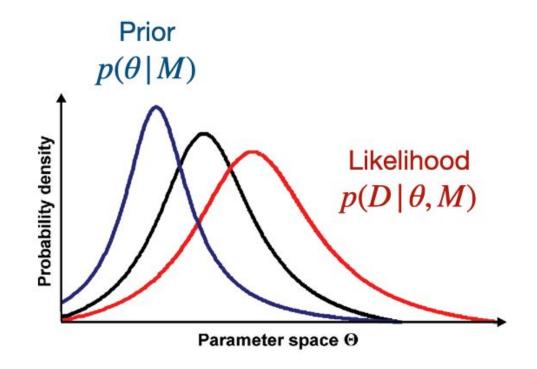
These are therefore useful only when the likelihood is sharply single-peaked.

When the likelihood is more complex, or when you have strong informative priors, you should use a model comparison tool that incorporates that information, such as the Bayesian evidence (Bayes factors).

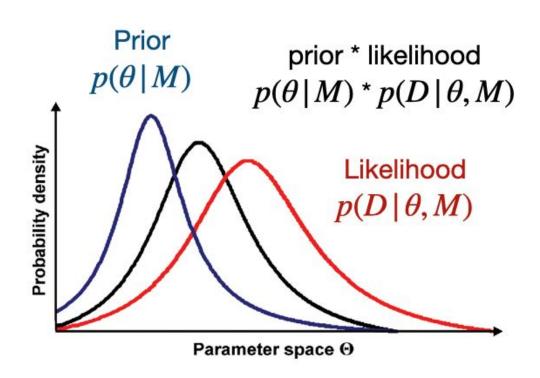
### Bayes theorem:

Posterior
$$P(A|B) = \frac{P(B|A) * P(A)}{P(B)}$$
Evidence

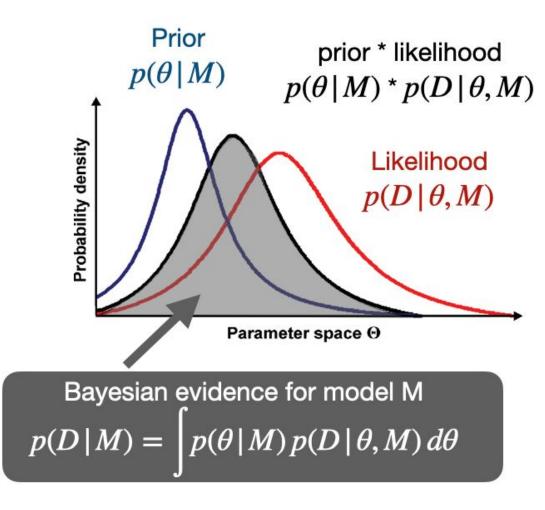
The denominator is the "Bayesian Evidence" found by integrating the likelihood \* prior over all parameter space.



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$$K = \frac{\Pr(D|M_1)}{\Pr(D|M_2)} = \frac{\int \Pr(\theta_1|M_1) \Pr(D|\theta_1, M_1) \, d\theta_1}{\int \Pr(\theta_2|M_2) \Pr(D|\theta_2, M_2) \, d\theta_2} = \frac{\frac{\Pr(M_1|D) \Pr(D)}{\Pr(M_1)}}{\frac{\Pr(M_2|D) \Pr(D)}{\Pr(M_2)}} = \frac{\Pr(M_1|D) \Pr(M_2)}{\Pr(M_2|D) \Pr(M_2)}$$

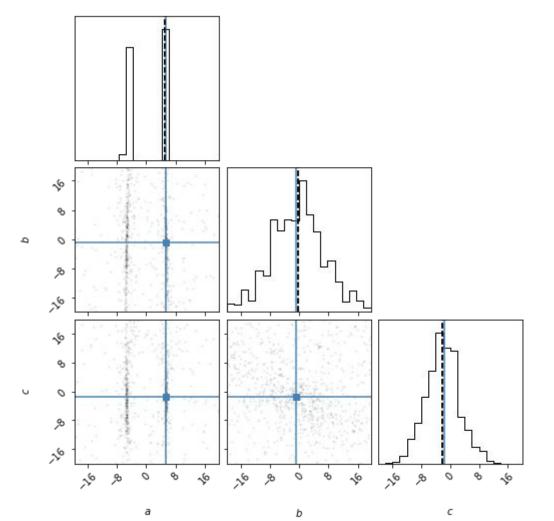
## How to compute the Bayesian evidence

- Define your priors carefully
- Use a tool like MCMC or Nested Sampling to measure the likelihood over all of parameter space
- Approximate the integral from the set of likelihood measurements recorded in your MCMC sampling chain

Some beautiful and instructive MCMC algorithm visualizations:

http://chi-feng.github.io/mcmc-demo

The 'nestle' package used in our nested sampling notebook returns both the array of log(likelihood) values and the integral, the log(evidence).



### Nestle results:

```
# Run nested sampling.
result = nestle.sample(loglike, prior_transform, 2)
result.logz # log evidence
result.logzerr # numerical (sampling) error on logz
result.samples # array of sample parameters
result.weights # array of weights associated with each sample
                                                     $ 8
                                                          0 0
```

### Resources

AIC Article by Sachin Date on the Medium blog "TowardsDataScience"

The <u>emcee</u> and <u>Nestle</u> packages for MCMC and Nested Sampling to measure Bayesian evidence

Book: Sivia, D. and Skilling, J. "Data Analysis: A Bayesian Tutorial"

Some "tutorial" papers:

- On AIC / BIC :
  - Wagenmakers and Farrell 2004
  - Symonds and Moussalli 2010
- Bayesian model averaging
  - Hoeting, Madigan, Raftery & Volinsky 2010