

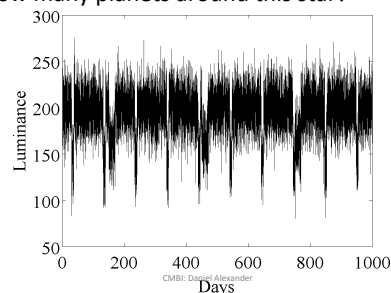
Model Selection

Daniel Alexander

CMBI: Daniel Alexander

What is the problem?

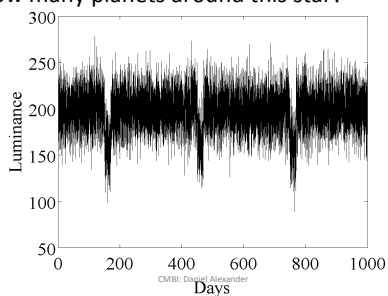
- How many planets around this star?



CMBI: Daniel Alexander

What is the problem?

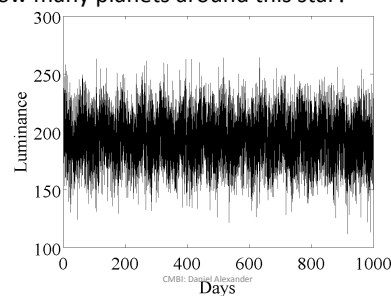
- How many planets around this star?



CMBI: Daniel Alexander

What is the problem?

- How many planets around this star?



CMBI: Daniel Alexander

Occam's Razor

- **Lex parsimoniae** – the law of parsimony
- The best model is the simplest that explains the data.



CMBI: Daniel Alexander

Guiding Principles

- A good model should...
- Predict unseen data closely
- Provide accurate and precise parameter estimates
 - Balance goodness of fit and model complexity
- Reflect what's going on in the world
 - Scientific intuition

CMBI: Daniel Alexander

Classical F-test

- Tests the null hypothesis that nested models are equivalent.
- Simple model M_1 with N_1 parameters x_1, \dots, x_{N_1} .
- Complex model M_2 with $N_2 (> N_1)$ parameters $x_1, \dots, x_{N_1}, \dots, x_{N_2}$.

CMBI: Daniel Alexander

Classical F-test

- The statistic

$$F = \frac{(K - N_2 - 1)(\text{Var}(M_2) - \text{Var}(M_1))}{(N_2 - N_1)E(M_2)}$$

- has F-distribution with degrees of freedom $N_2 - N_1$ and $K - N_2 - 1$ under the null hypothesis.

$$\text{Var}(M) = \frac{1}{K-1} \sum_{i=1}^K (M(\tilde{\mathbf{x}}; \mathbf{y}_i) - \bar{M})^2; \text{ and } \bar{M} = \frac{1}{K} \sum_{i=1}^K M(\tilde{\mathbf{x}}; \mathbf{y}_i)$$

$$E(M) = \frac{1}{K} \sum_{i=1}^K (M(\tilde{\mathbf{x}}; \mathbf{y}_i) - A_i)^2$$

Armitage P, Berry G. Statistical methods in medical research. Oxford, UK: Blackwell Scientific Publications; 1971.

Classical F-test

- Assumes nested models
- Assumes Gaussian noise model
- Rejects or does not reject the null hypothesis that the models are equivalent.
 - No relative likelihood obtained

CMBI: Daniel Alexander

Akaike's information criterion

- The criterion is $AIC = 2N - 2 \log L$
- where N is the number of parameters in the model and L is the likelihood $p(\mathbf{A} | \mathbf{x})$.
- Loss of Kullback-Liebler information using model instead of truth
 - See Burnham and Anderson

CMBI: Daniel Alexander

Akaike IEEE Trans Automatic Control 1974

AIC – Gaussian noise

- Classic formula for Gaussian noise with unknown variance

$$AIC = 2N + K \log \left(K^{-1} \sum_{k=1}^K (A_k - S_k)^2 \right)$$

- K must include the standard deviation, which is also estimated.

CMBI: Daniel Alexander

AIC

- With i models to choose from, compute each AIC_i .
- Smallest is best.
- Or, model i is $\exp((AIC_{\min} - AIC_i)/2)$ times as likely to be correct as the best model.
- These provide *Akaike weights*, which represent model probabilities.
 - See Burnham and Anderson book

CMBI: Daniel Alexander

AIC

- Any noise model
- Models need not be nested

CMBI: Daniel Alexander

AIC Corrected (AICc)

- Basic AIC valid only as the number of data points, K , tends to infinity.
- For finite K ,
$$AICc = AIC + \frac{2N(N+1)}{K-N-1}$$
- Favour over classic AIC for $K/N < 40$.

CMBI: Daniel Alexander

Bayesian information criterion

- Works in a similar way.
- The criterion is
$$BIC = N \log K - 2 \log L$$
- Derived via a Bayesian argument
 - See Burnham and Anderson

Schwarz Annals of Statistics 1978
CMBI: Daniel Alexander

Application in Diffusion MRI

NeuroImage 59 (2012) 2241–2254

Contents lists available at ScienceDirect

NeuroImage

journal homepage: www.elsevier.com/locate/ynimg

Compartment models of the diffusion MR signal in brain white matter: A taxonomy and comparison

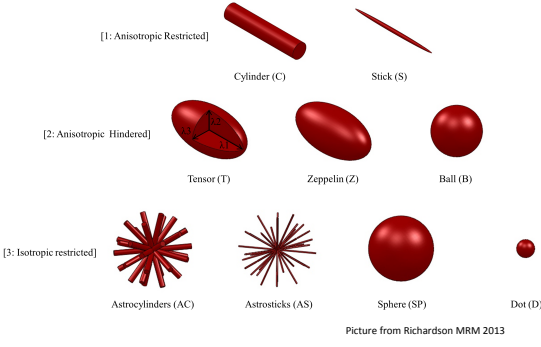
Eleftheria Panagiotaki^{a,*}, Torben Schneider^b, Bernard Siow^{a,c}, Matt G. Hall^a, Mark F. Lythgoe^c, Daniel C. Alexander^a

^a Centre for Medical Image Computing, Department of Computer Science, University College London, Gower Street, London WC1E 6BT, UK
^b NMR Research Unit, Department of Neuroinflammation, UCL Institute of Neurology, University College London, WC1N 3BG, UK
^c Centre for Advanced Biomedical Imaging, University College London, Gower Street, London WC1E 6BT, UK

CMBI: Daniel Alexander

Panagiotaki Models

Panagiotaki et al Neuroimage 2012.



Picture from Richardson MRM 2013

Model ranking

Panagiotaki et al Neuroimage 2012.

Table 4
Table of the models ranked by the Bayesian information criterion (BIC) for dataset A, the ranking of the models according to BIC for dataset B and the number of parameters (counting S_0) for each model.

Models	BIC		No. parameters	
	Dataset A	Rank A	Dataset B	Rank B
TensorCylindersSphere	-2.587	1	-2.636	1
TensorSticksSphere	-2.558	2	-2.586	2
ZeppelinCylindersSphere	-2.495	3	-2.542	5
ZeppelinGDRKylindersSphere	-2.489	4	-2.507	7
ZeppelinStickSphere	-2.482	5	-2.488	9
TensorCylinderDot	-2.481	6	-2.561	3
TensorGDRKylindersDot	-2.446	7	-2.557	4
TensorGDRKylindersSphere	-2.431	8	-2.518	6
ZeppelinCylinderDot	-2.429	9	-2.499	8
ZeppelinGDRKylindersDot	-2.412	10	-2.484	10
BallGDRKylindersDot	-2.382	11	-2.466	11
BallCylindersSphere	-2.377	12	-2.448	13
BallCylinderDot	-2.375	13	-2.454	12
BallGDRKylindersSphere	-2.366	14	-2.420	14
ZeppelinStickDot	-2.329	15	-2.396	15
TensorCylinderAstrocyllinders	-2.290	16	-2.331	17
TensorCylinderAstrosticks	-2.289	17	-2.331	18
TensorGDRKylindersAstrosticks	-2.286	18	-2.329	19
TensorStickAstrocyllinders	-2.282	19	-2.324	20
TensorStickAstrosticks	-2.279	20	-2.324	21
ZeppelinGDRKylindersAstrocyllinders	-2.278	21	-2.293	36
BallStickSphere	-2.272	22	-2.332	16

Other criteria

- Deviance information criterion
 - Designed to work with MCMC
 - Large sample size
 - Posterior distribution approximately normal.
- Minimum description length
 - Model selection as a coding problem
- Minimum message length
 - Bayesian version of MDL.

CMBI: Daniel Alexander

Bayesian Model Selection

- Suppose we have two models M_1 and M_2 and some measured data \mathbf{A} .

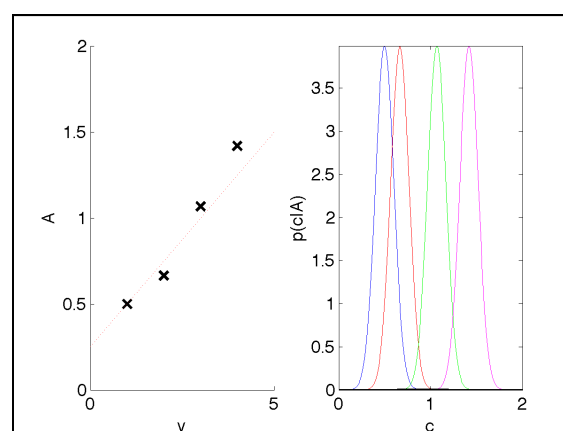
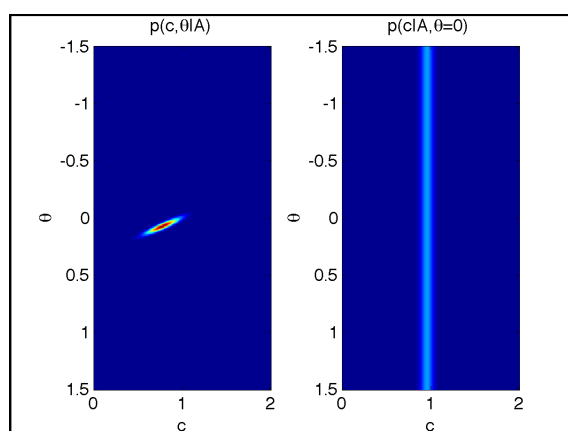
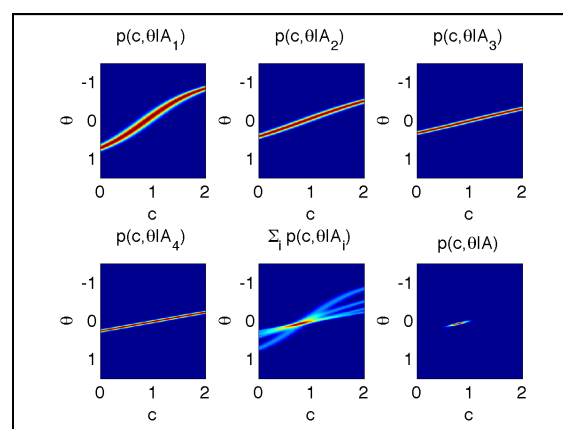
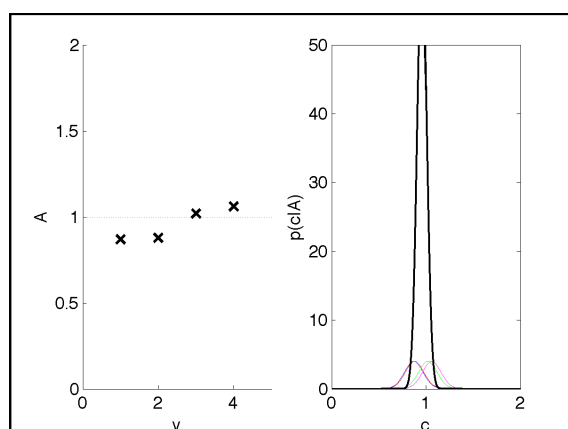
$$p(M_i | \mathbf{A}) = \frac{p(\mathbf{A} | M_i) p(M_i)}{p(\mathbf{A})} = \frac{p(\mathbf{A} | M_i) p(M_i)}{\sum_j p(\mathbf{A} | M_j) p(M_j)}$$

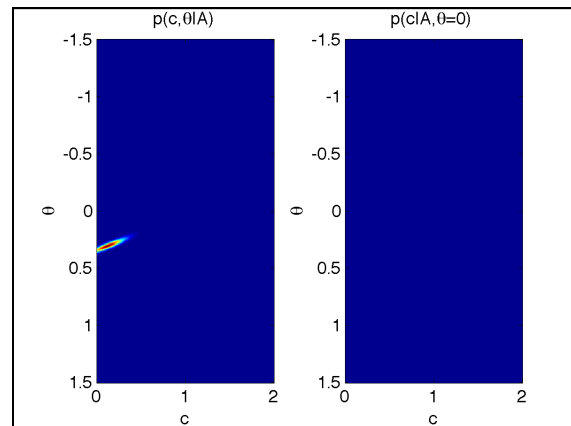
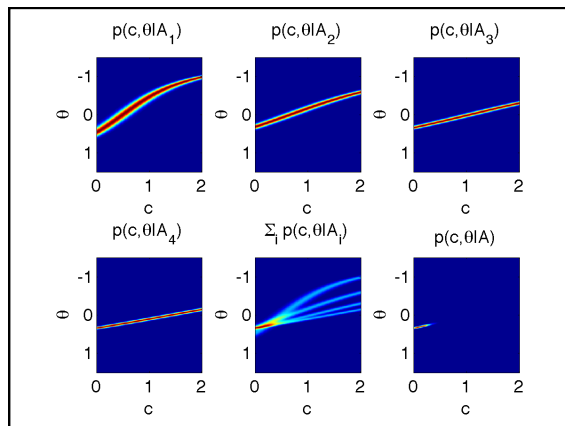
- $p(M_i)$ is the prior belief in M_i .
- $p(\mathbf{A} | M_i)$ is the likelihood of M_i .
- If M_i has parameters \mathbf{x} , then

$$p(\mathbf{A} | M_i) = \int p(\mathbf{A} | M_i, \mathbf{x}) p(\mathbf{x}) d\mathbf{x}$$

- where $p(\mathbf{x})$ is the term for the prior on \mathbf{x} .

CMBI: Daniel Alexander





The Bayes Factor

- The Bayes factor is the likelihood ratio

$$K = \frac{p(\mathbf{A} | M_1)}{p(\mathbf{A} | M_2)} = \frac{\int p(\mathbf{A} | M_1, \mathbf{x}_1) p(\mathbf{x}_1) d\mathbf{x}_1}{\int p(\mathbf{A} | M_2, \mathbf{x}_2) p(\mathbf{x}_2) d\mathbf{x}_2}$$

- Rule of thumb: if $K > 10$, accept M_1 over M_2 .

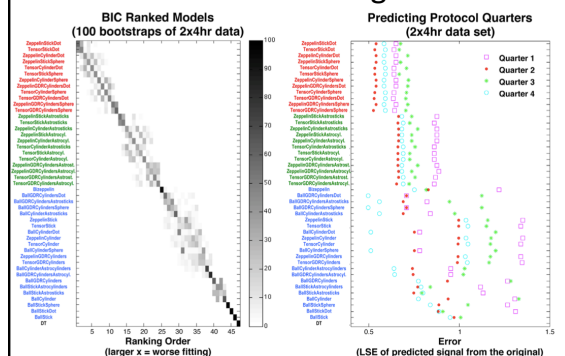
CMBI: Daniel Alexander

Cross validation

- Ultimate empirical model selection:
 - Estimate parameters on training set
 - Evaluate fit on unseen test set
- Cross validation divides the available data into multiple pairs of training and test sets:
 - k-fold cross-validation**: randomly divide into k equal-sized subsets. Use $k-1$ sets to fit; compute error on remainder; average error over all k subsets.
 - Repeated random subsampling**: as above, but draw a random sample each time.
 - Leave-one-out validation**: $k = K-1$.

CMBI: Daniel Alexander

Ferizi Ranking

Ferizi et al
MRM 2014.

Guiding Principles Revisited

- A good model should...
- Predict unseen data closely
- Provide accurate and precise parameter estimates
- Reflect what's going on in the world

CMBI: Daniel Alexander