Optimization of Designs for fMRI

UCLA Advanced Neuroimaging Summer School August 21, 2007

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Why optimize?

- Scans are expensive.
- Subjects can be difficult to find.
- fMRI data are noisy
- A badly designed experiment is unlikely to yield publishable results.
- Time = Money

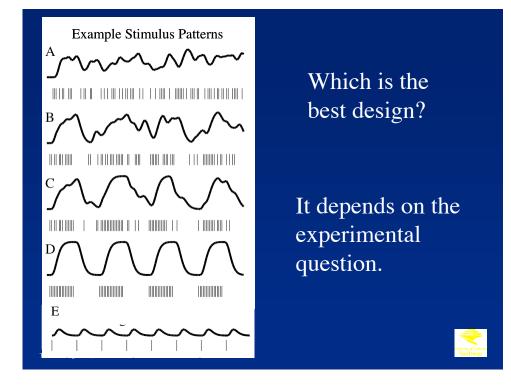
If your result needs a statistician then you should design a better experiment. --*Baron Ernest Rutherford*

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What to optimize?

- Statistical Efficiency: maximize contrast of interest versus noise.
- Psychological factors: is the design too boring? Minimize anticipation, habituation, boredom, etc.





Possible Questions

- Where is the activation?
- What does the hemodynamic response function (HRF) look like?

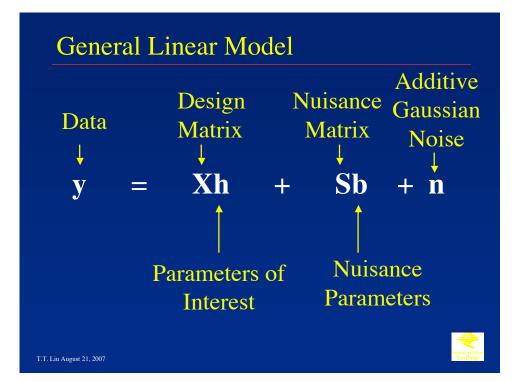


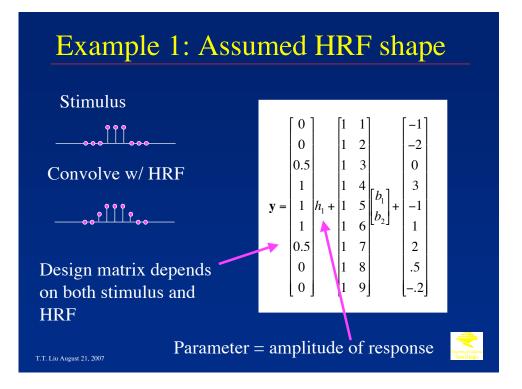
Model Assumptions

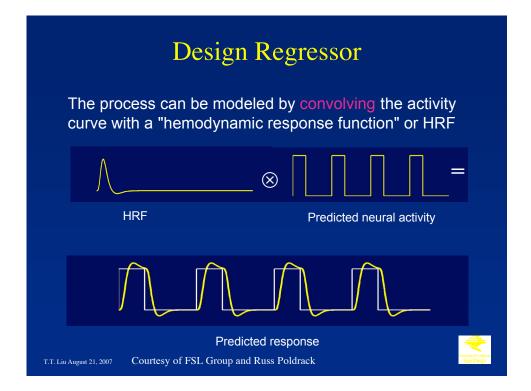
- 1) Assume we know the shape of the HRF but not its amplitude.
- 2) Assume we know nothing about the HRF (neither shape nor amplitude).

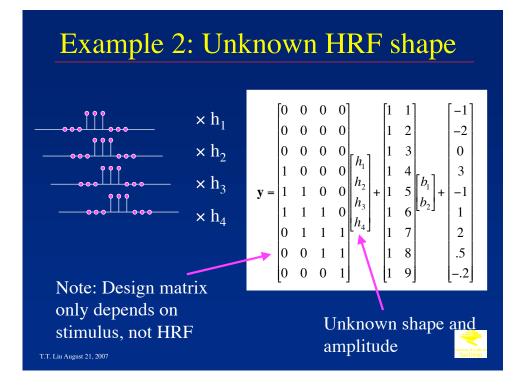
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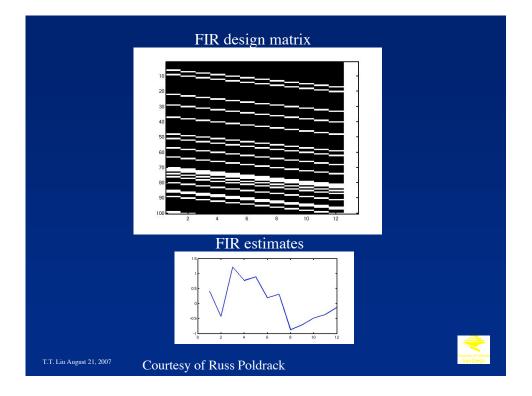
3) Assume we know something about the HRF (e.g. it's smooth).

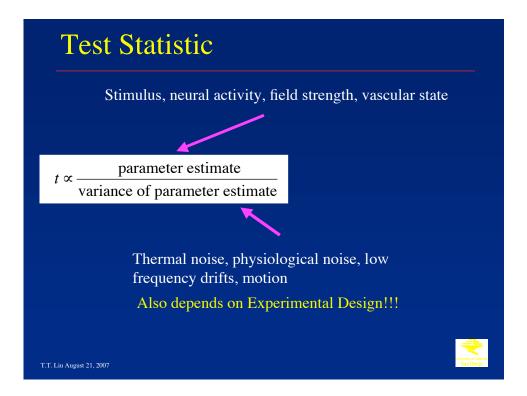


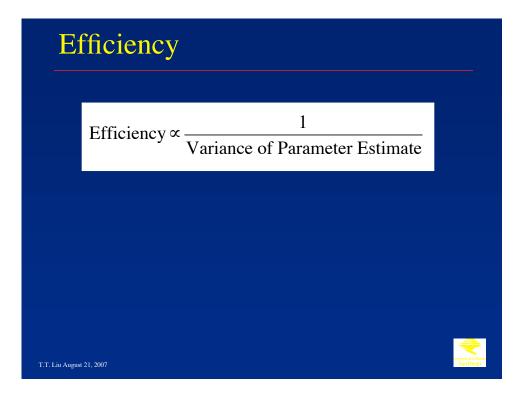


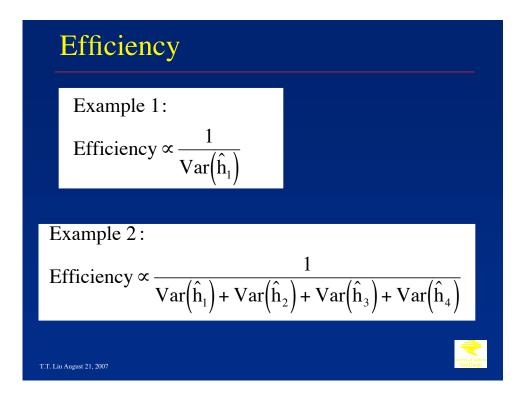


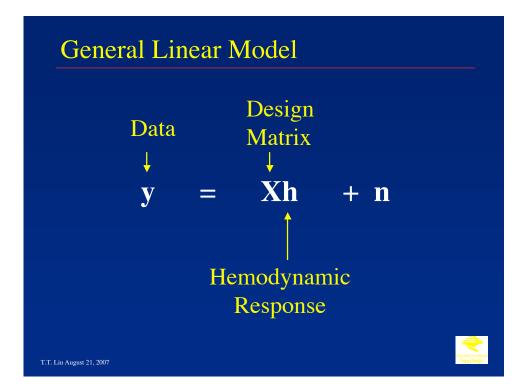












Nuisance Functions

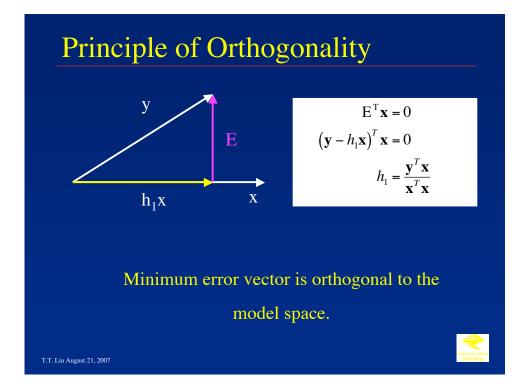
Nuisance terms (constant term, linear drift, etc) are a fact of life in fMRI experiments.

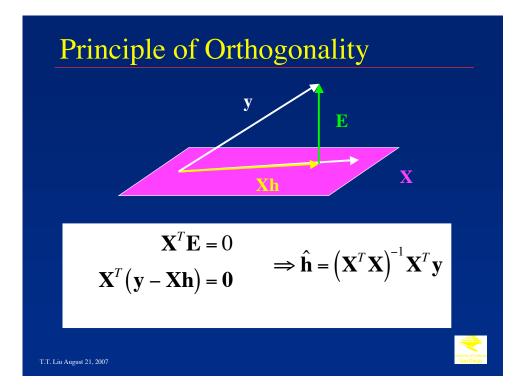
However, to keep things simple, we will ignore the nuisance term Sb in the GLM for this talk.

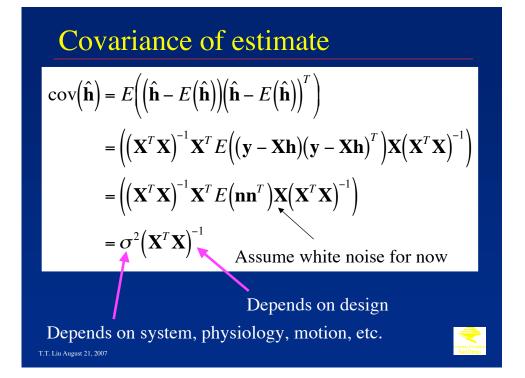
The formulas we derive have the same form when nuisance terms are considered. Just replace X by X_{\perp} , where X_{\perp} is obtained by projecting the nuisance terms out of the columns of X. See Liu et al 2001 and Liu and Frank 2004 for more details.



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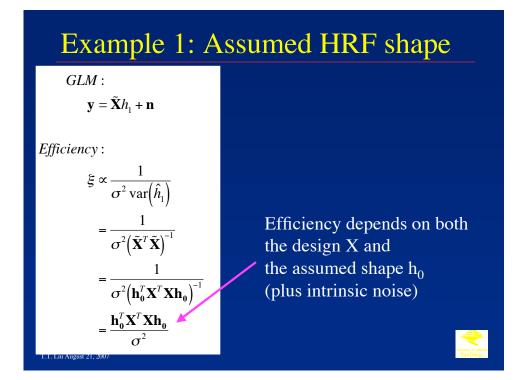


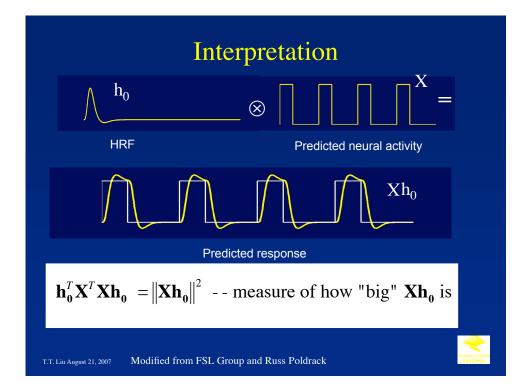


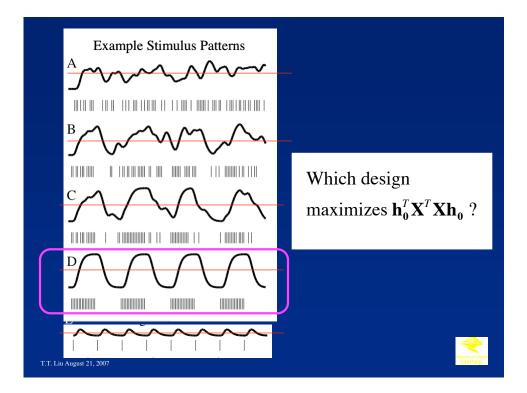
Example 1: Assumed HRF shape

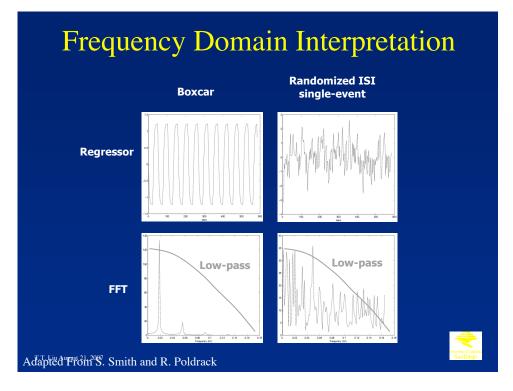
Assume we know the HDR shape \mathbf{h}_0 but not its amplitude h_1 $\mathbf{h} = \mathbf{h}_0 h_1$

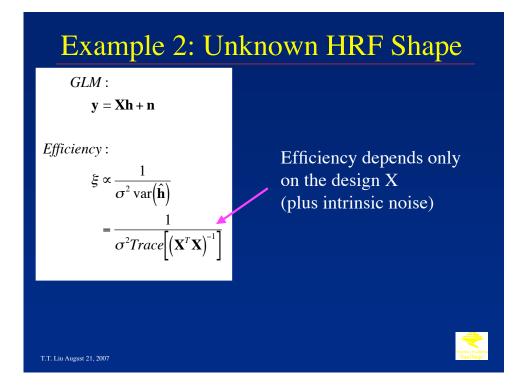
GLM: $\mathbf{y} = \mathbf{X}\mathbf{h} + \mathbf{n}$ $= \mathbf{X}\mathbf{h}_0h_1 + \mathbf{n}$ $= \tilde{\mathbf{X}}h_1 + \mathbf{n} \text{ where } \tilde{\mathbf{X}} = \mathbf{X}\mathbf{h}_0$

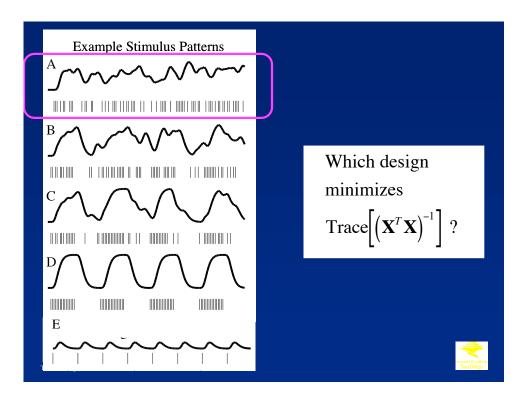


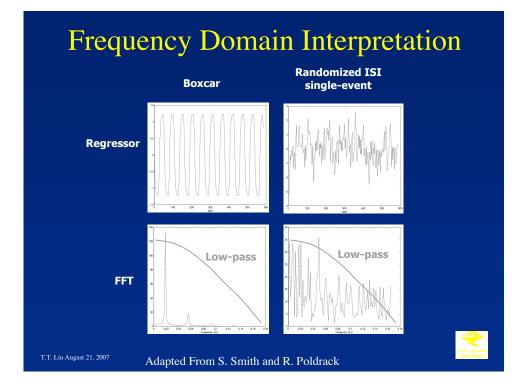


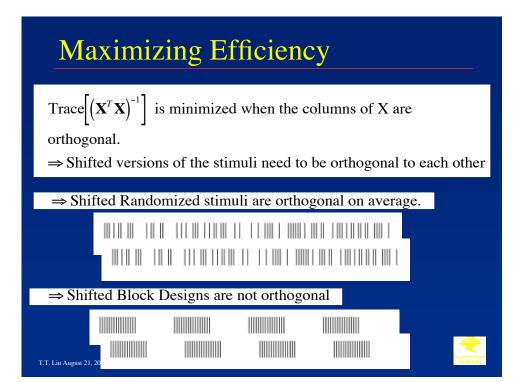


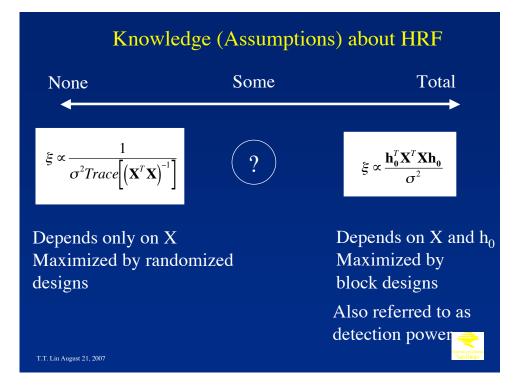




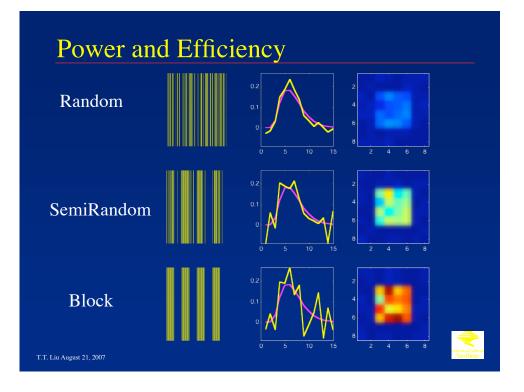


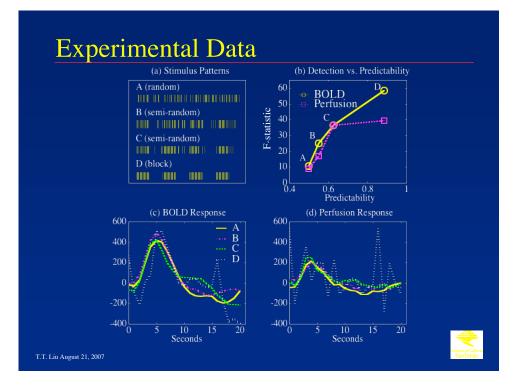


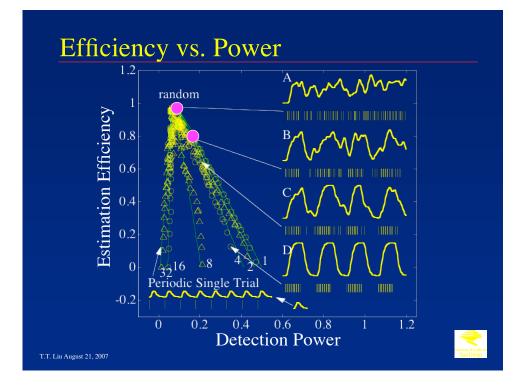


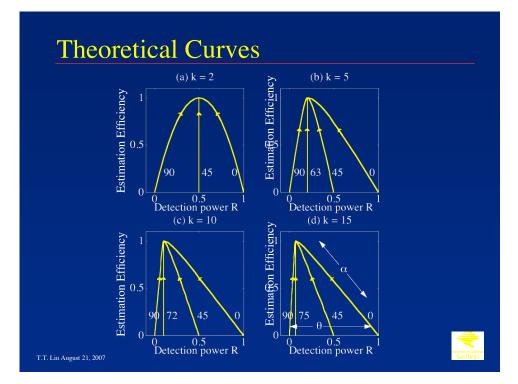


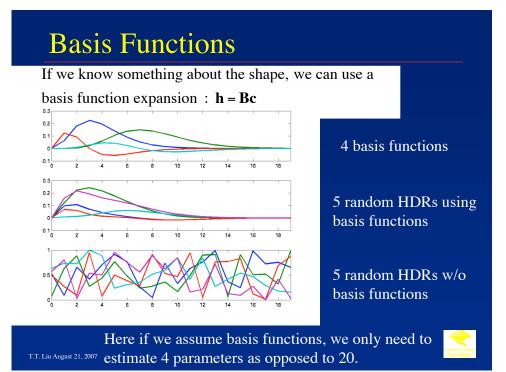












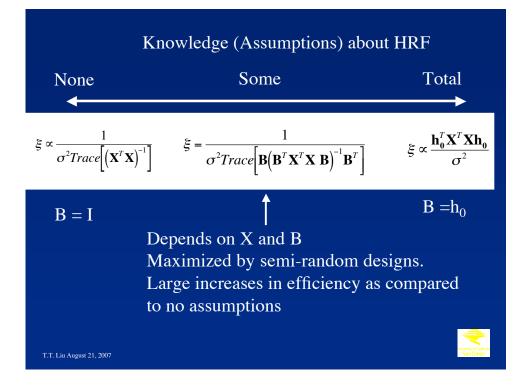
Basis Functions

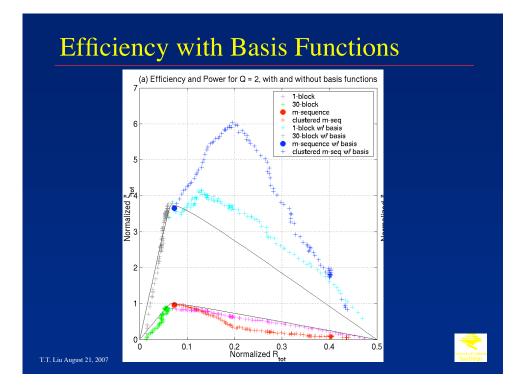
If we know something about the shape, we can use a basis function expansion : $\mathbf{h} = \mathbf{Bc}$

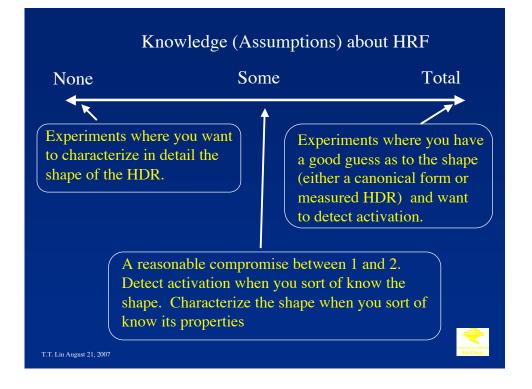
 $GLM : \mathbf{y} = \mathbf{X}\mathbf{B}\mathbf{c} + \mathbf{n} = \tilde{\mathbf{X}}\mathbf{c} + \mathbf{n}$ $Estimate : \hat{\mathbf{c}} = (\mathbf{B}^T\mathbf{X}^T\mathbf{X}\ \mathbf{B})^{-1}\mathbf{B}^T\mathbf{X}^T\mathbf{y}$ $\hat{\mathbf{h}} = \mathbf{B}\hat{\mathbf{c}} = \mathbf{B}(\mathbf{B}^T\mathbf{X}^T\mathbf{X}\ \mathbf{B})^{-1}\mathbf{B}^T\mathbf{X}^T\mathbf{y}$

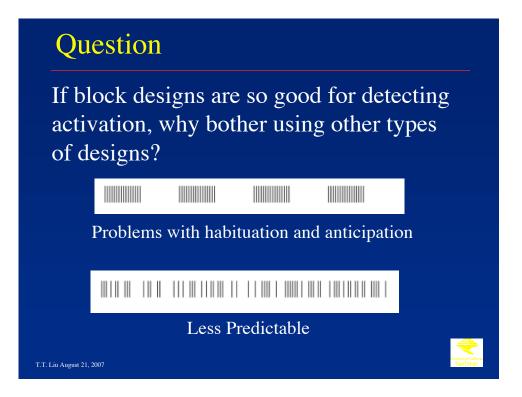
Efficiency:
$$\xi = \frac{1}{\sigma^2 Trace \left[\mathbf{B} \left(\mathbf{B}^T \mathbf{X}^T \mathbf{X} \; \mathbf{B} \right)^{-1} \mathbf{B}^T \right]}$$

Efficiency now depends on both X and B









Entropy

Perceived randomness of an experimental design is an important factor and can be critical for circumventing experimental confounds such as habituation and anticipation.

Conditional entropy is a measure of randomness in units of bits.

Rth order conditional entropy (H_r) is the average number of binary (yes/no) questions required to determine the next trial type given knowledge of the *r* previous trial types.



 2^{H_r} is a measure of the average number of possible outcomes.

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Entropy Example

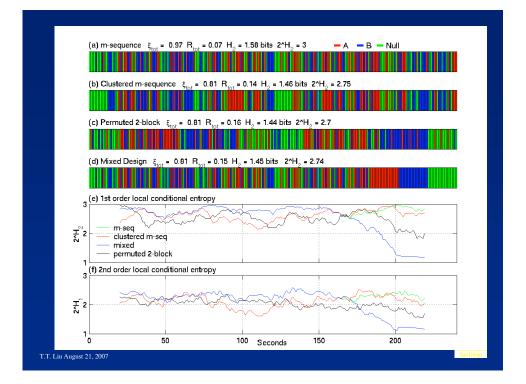
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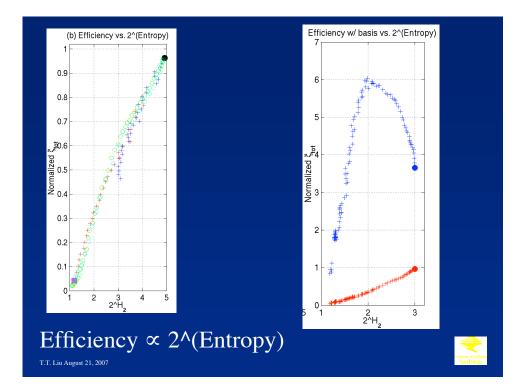
Maximum entropy is 1 bit, since at most one needs to only ask one question to determine what the next trial is (e.g. is the next trial A?). With maximum entropy, $2^1 = 2$ is the number of equally likely outcomes.

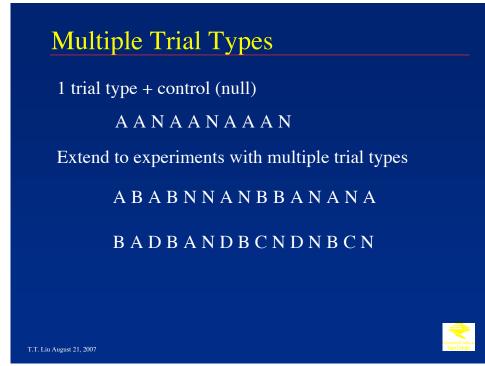
ACBNCBAABCNA

Maximum entropy is 2 bits, since at most one would need to ask 2 questions to determine the next trial type. With maximum entropy, the number of equally likely outcomes to choose from is $4(2^2)$.





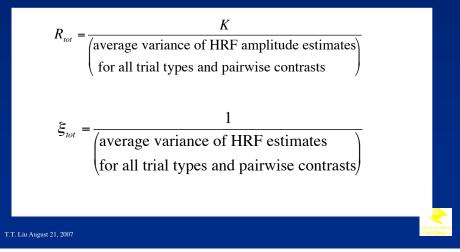


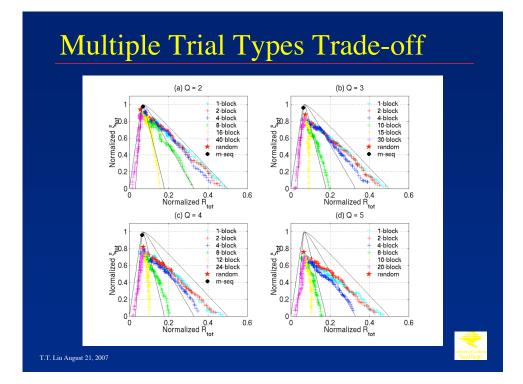


Multiple Trial Types GLM
$$y = Xh + Sb + n$$
 $\mathcal{X} = [X_1 X_2 \dots X_Q]$ $h = [h_1^T h_2^T \dots h_Q^T]^T$

Multiple Trial Types Overview

Efficiency includes individual trials and also contrasts between trials.





Optimal Frequency

Can also weight how much you care about individual trials or contrasts. Or all trials versus events. Optimal frequency of occurrence depends on weighting. Example: With Q = 2 trial types, if only contrasts are of interest p = 0.5. If only trials are of interest, p = 0.2929. If both trials and contrasts are of interest p = 1/3.

$$p = \frac{Q(2k_1 - 1) + Q^2(1 - k_1) + k_1^{1/2} (Q(2k_1 - 1) + Q^2(1 - k_1))^{1/2}}{Q(Q - 1)(k_1Q - Q - k_1)}$$

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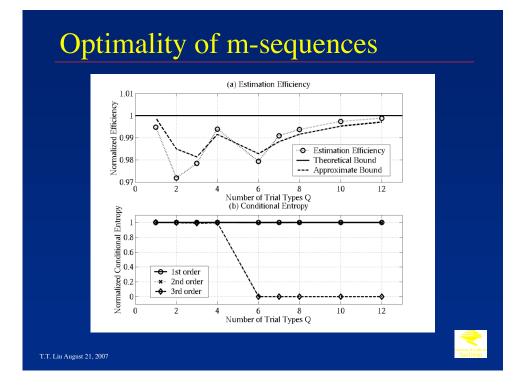
Design

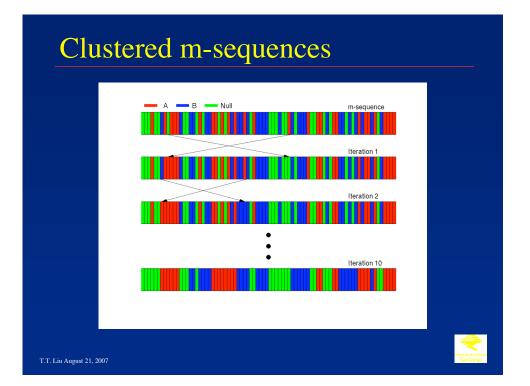
As the number of trial types increases, it becomes more difficult to achieve the theoretical trade-offs. Random search becomes impractical.

For unknown HDR, should use an m-sequence based design when possible.

Designs based on block or m-sequences are useful for obtaining intermediate trade-offs or for optimizing with basis functions or correlated noise.







Topics we haven't covered.

The impact of correlated noise -- this will change the optimal design.

Impact of nonlinearities in the BOLD response.

Other optimization algorithms -- e.g. genetic algorithms.

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Summary

- Efficiency as a metric of design performance.
- Efficiency depends on both experimental design and assumptions about HRF.
- Inherent tradeoff between power (detection of known HRF) and efficiency (estimation of HRF)

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