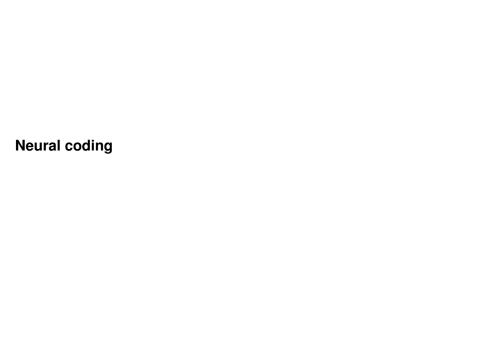
## **Neural Encoding Models**

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February 2019



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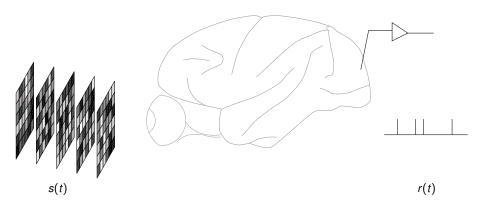
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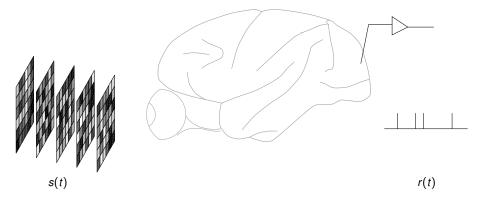
Computation plays a vital part in systematising empirical data.

# Stimulus coding



Decoding:  $\hat{s}(t) = G[r(t)]$  (reconstruction)

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Decoding:  $\hat{s}(t) = G[r(t)]$ 

Encoding:  $\hat{r}(t) = F[s(t)]$ 

(reconstruction)

(systems identification)

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However, on the face of it, mapping *either* the decoding or encoding function does not by itself answer either of our basic questions about coding.

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Goal: Estimate p(spike|s, H) [or *intensity*  $\lambda(t|s[0, t), H(t))$ ] from data.

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- Select stimuli efficiently
- Fit models with smaller numbers of parameters

Most neurons communicate using action potentials — statistically described by a point process:

$$P(\text{spike} \in [t, t + dt)) = \lambda(t|H(t), \text{stimulus}, \text{network activity})dt$$

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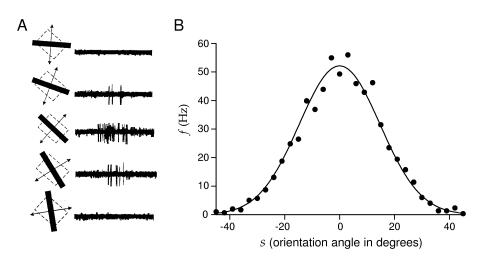
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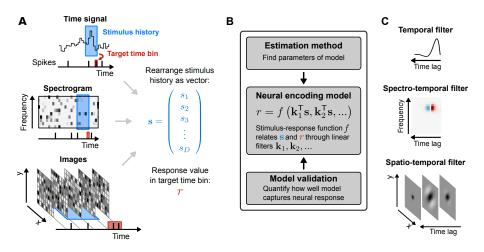
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Attempt to capture history and network effects in simple models.

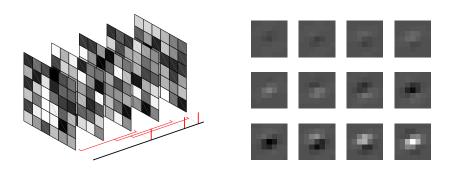
# Tuning – stationary stimuli



## (Nonlinear) filtering – dynamic stimuli

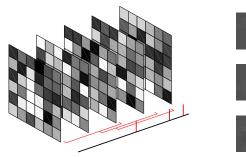


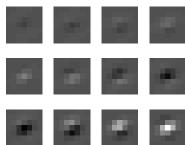
## Spike-triggered average



Decoding: mean of P (s | r = 1)

## Spike-triggered average





Decoding: mean of P (s | r = 1)Encoding: predictive filter

$$s_1$$
  $s_2$   $s_3$   $\ldots$   $s_T$   $s_{T+1}$   $\ldots$ 

$$r(t) = \int_0^\tau s(t-\tau)w(\tau)d\tau$$

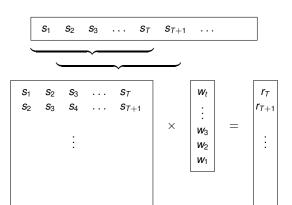
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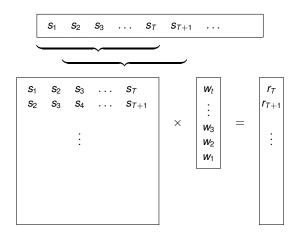
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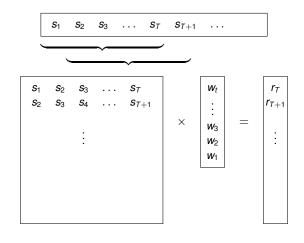
$$r(t) = \int_0^{\tau} s(t-\tau)w(\tau)d\tau$$



$$SW = R$$

# **Linear regression**

$$r(t) = \int_0^T s(t-\tau)w(\tau)d\tau$$



$$SW = R$$

$$W(\omega) = rac{S(\omega)^* R(\omega)}{|S(\omega)|^2}$$

$$W = \underbrace{(S^{\mathsf{T}}S)}_{\Sigma_{SS}}^{-1} \underbrace{(S^{\mathsf{T}}R)}_{\mathsf{STA}}^{\mathsf{T}}$$

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#### Issues:

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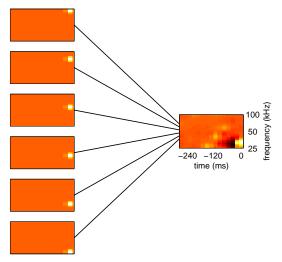
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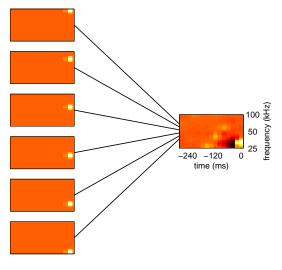
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  - may provide unbiased estimates of cascade filters (see later)

## Likelihood penalties for regularisation

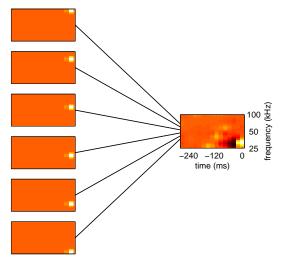
$$\widehat{\mathbf{w}} = \underset{\mathbf{w}}{\operatorname{argmax}} \underbrace{\mathcal{L}(\mathbf{w}; \textit{Data})}_{\text{Likelihood}} \quad - \underbrace{\mathcal{R}(\mathbf{w})}_{\text{Regularise:}}$$

 $\mathcal{R}$  may penalise large values of **w** (e.g.  $\|\mathbf{w}\|^2$  or  $\sum_i |w_i|$ ) or may promote smoothness or other properties.



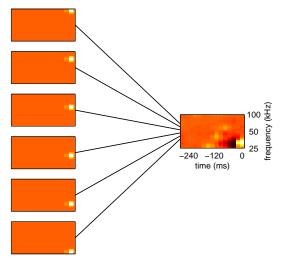


▶ sparsity  $[C_{ii} \text{ zero for many } i]$  ARD



- sparsity
- smoothness

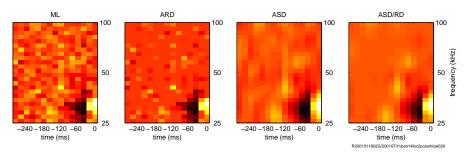
[ $C_{ii}$  zero for many i] [ $C_{ij}$  high for close i and j] ARD ASD

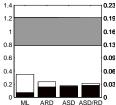


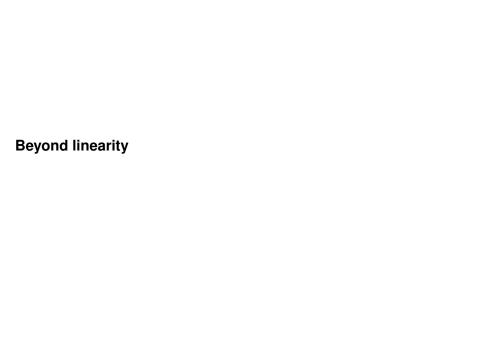
- sparsity
- smoothness
- locality

 $[C_{ii} \text{ zero for many } i]$  $[C_{ij} \text{ high for close } i \text{ and } j]$  $[C_{ii} \text{ high in a single region}]$  ARD ASD ALD

### Smoothness and sparsity (ASD/RD)







### **Beyond linearity**

Linear models often fail to predict well. Alternatives?

- Wiener/Volterra functional expansions
  - M-series
  - Linearised estimation
  - Kernel formulations
- LN (Wiener) cascades
  - Spike-trigger covariance (STC) methods
  - "Maximimally informative" dimensions (MID) 

    ML nonparametric LNP models
  - ML Parametric GLM models
- NL (Hammerstein) cascades
  - Multilinear formulations
- I NI N and more ....

### The Volterra functional expansion

A polynomial-like expansion for functionals (or operators).

Let y(t) = F[x(t)]. Then:

$$y(t) \approx k^{(0)} + \int d\tau \, k^{(1)}(\tau) x(t-\tau) + \iint d\tau_1 \, d\tau_2 \, k^{(2)}(\tau_1, \tau_2) x(t-\tau_1) x(t-\tau_2)$$
$$+ \iiint d\tau_1 \, d\tau_2 \, d\tau_3 \, k^{(3)}(\tau_1, \tau_2, \tau_3) x(t-\tau_1) x(t-\tau_2) x(t-\tau_3) + \dots$$

or (in discretised time)

$$y_{t} = K^{(0)} + \sum_{i} K_{i}^{(1)} x_{t-i} + \sum_{ij} K_{ij}^{(2)} x_{t-i} x_{t-j} + \sum_{ijk} K_{ijk}^{(3)} x_{t-i} x_{t-j} x_{t-k} + \dots$$

For finite expansion, the kernels  $k^{(0)}, k^{(1)}(\cdot), k^{(2)}(\cdot, \cdot), k^{(3)}(\cdot, \cdot, \cdot), \ldots$  are not straightforwardly related to the functional F. Indeed, values of lower-order kernels change as the maximum order of the expansion is increased.

Estimation: model is linear in kernels, so can be estimated just like a linear (first-order) model with expanded "input".

- ► Kernel trick: polynomial kernel  $K(x_1, x_2) = (1 + x_1x_2)^n$ .
- M-series.

### **Wiener Expansion**

The Wiener expansion gives functionals of different orders that are orthogonal for white noise input x(t).

$$G_{0}[x(t); h^{(0)}] = h^{(0)}$$

$$G_{1}[x(t); h^{(1)}] = \int d\tau h^{(1)}(\tau)x(t-\tau)$$

$$G_{2}[x(t); h^{(2)}] = \iint d\tau_{1} d\tau_{2} h^{(2)}(\tau_{1}, \tau_{2})x(t-\tau_{1})x(t-\tau_{2}) - P \int d\tau_{1} h^{(2)}(\tau_{1}, \tau_{1})$$

$$G_{3}[x(t); h^{(3)}] = \iiint d\tau_{1} d\tau_{2} d\tau_{3} h^{(3)}(\tau_{1}, \tau_{2}, \tau_{3})x(t-\tau_{1})x(t-\tau_{2})x(t-\tau_{3})$$

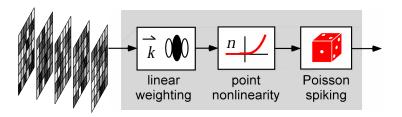
$$-3P \iint d\tau_{1} d\tau_{2} h^{(3)}(\tau_{1}, \tau_{2}, \tau_{2})x(t-\tau_{1})$$

Easy to verify that  $\mathbb{E}[G_i[x(t)]G_j[x(t)]] = 0$  for  $i \neq j$ .

Thus, these kernels can be estimated independently. But, they depend on the stimulus.

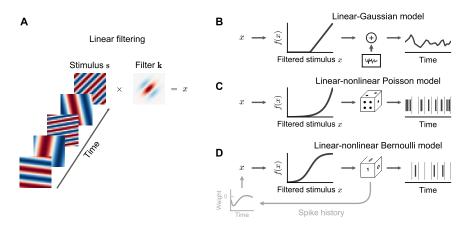
### **Cascade models**

The LNP (Wiener) cascade

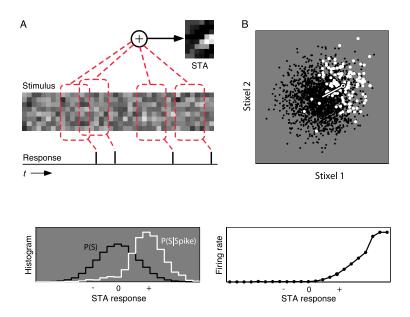


- Rectification addresses negative firing rates.
- Loose biophysical correspondance.

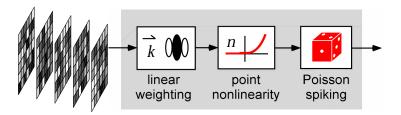
### LNP cascades and noise



# LNP estimation – the Spike-triggered ensemble

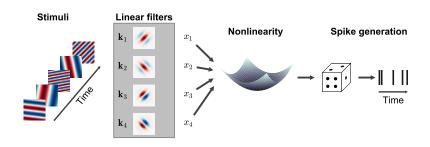


### Single linear filter



- STA is unbiased estimate of filter for spherical input distribution. (Bussgang's theorem)
- lacktriangle Elliptically-distributed data can be whitened  $\Rightarrow$  linear regression weights are unbiased.
- Linear weights are not necessarily maximum-likelihood (or otherwise optimal), even for spherical/elliptical stimulus distributions.
- Linear weights may be biased for general stimuli (binary/uniform or natural).

### **Multiple filters**

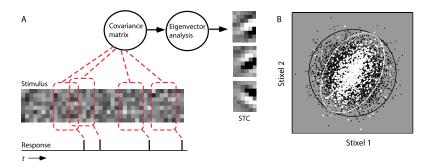


Distribution changes along relevant directions (and, usually, along all linear combinations of relevant directions).

Proxies to measure change in distribution:

- mean: STA (can only reveal a single direction)
- variance: STC
- binned (or kernel) KL divergence: MID "maximally informative directions" (equivalent to ML in LNP model with binned nonlinearity)

### **STC**



Project out STA:

$$\widetilde{S} = S - (S\mathbf{k}_{\mathrm{sta}})\mathbf{k}_{\mathrm{sta}}^{\mathsf{T}}; \quad C_{\mathrm{prior}} = \dfrac{\widetilde{S}^{\mathsf{T}}\widetilde{S}}{N}; C_{\mathrm{spike}} = \dfrac{\widetilde{S}^{\mathsf{T}}\mathrm{diag}(R)\widetilde{S}}{N_{\mathrm{spike}}}$$

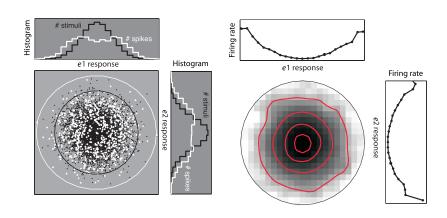
Choose directions with greatest change in variance:

k- argmax 
$$\mathbf{v}^{\mathsf{T}}(C_{\mathsf{prior}} - C_{\mathsf{spike}})\mathbf{v}$$

 $\Rightarrow$  find eigenvectors of  $(C_{prior} - C_{spike})$  with large (absolute) eigvals.

### **STC**

Reconstruct nonlinearity (may assume separability)



### **Biases**

STC (obviously) requires that the nonlinearity alter variance. If so, subspace is unbiased provided distribution is

- radially (elliptically) symmetric
- AND independent

 $\Rightarrow$  Gaussian.

May be possible to correct for non-Gaussian stimulus by transformation, subsampling or weighting (latter two at cost of variance).

### More LNP methods

Non-parametric non-linearities:

"Maximally informative dimensions" (MID)  $\Leftrightarrow$  "non-parametric" maximum likelihood.

 Intuitively, extends the variance difference idea to arbitrary differences between marginal and spike-conditioned stimulus distributions.

$$\mathbf{k}_{\text{MID}} = \operatorname*{argmax} \mathbf{KL}[P(\mathbf{k} \cdot \mathbf{x}) || P(\mathbf{k} \cdot \mathbf{x} | \text{spike})]$$

- Measuring KL requires binning or smoothing—turns out to be equivalent to fitting a non-parametric nonlinearity by binning or smoothing (Williamson, Sahani, Pillow PLoSCB 2015).
- Difficult to use for high-dimensional LNP models (but ML viewpoint suggests separable or "cylindrical" basis functions – see Williamson et al.).
- Parametric non-linearities: the "generalised linear model" (GLM).

### **Generalised linear models**

LN models with specified nonlinearities and exponential-family noise.

In general (for monotonic g):

$$y \sim \text{ExpFamily}[\mu(\mathbf{x})]; \qquad g(\mu) = \beta \mathbf{x}$$

For our purposes easier to write

$$y \sim \text{ExpFamily}[f(\beta \mathbf{x})]$$

(Continuous time) point process likelihood with GLM-like dependence of  $\lambda$  on covariates is approached in limit of bins  $\to$  0 by either Poisson or Bernoulli GLM.

Mark Berman and T. Rolf Turner (1992) Approximating Point Process Likelihoods with GLIM Journal of the Royal Statistical Society. Series C (Applied Statistics), 41(1):31-38.

### **Generalised linear models**

Poisson distribution  $\Rightarrow f = \exp()$  is canonical (natural params  $= \beta \mathbf{x}$ ).

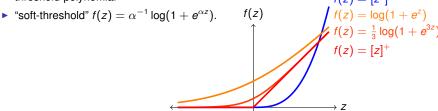
Canonical link functions give concave likelihoods  $\Rightarrow$  unique maxima.

Generalises (for Poisson) to any *f* which is convex and log-concave:

$$\log\text{-likelihood} = c - f(\beta \mathbf{x}) + y \log f(\beta \mathbf{x})$$

#### Includes:

- threshold-linear
- threshold-polynomial



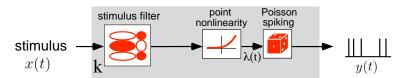
### **Generalised linear models**

ML parameters found by

- gradient ascent
- ► IRLS

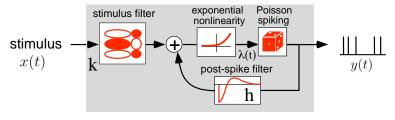
Regularisation by  $L_2$  (quadratic) or  $L_1$  (absolute value – sparse) penalties (MAP with Gaussian/Laplacian priors) preserves concavity.

# Linear-Nonlinear-Poisson (GLM)



# GLM with history-dependence

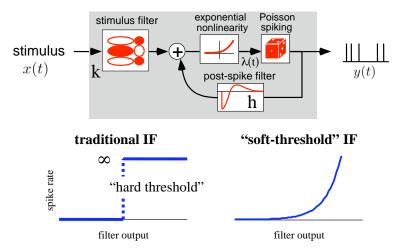
(Truccolo et al 04)



conditional intensity 
$$\lambda(t) = f(k \cdot x(t) + h \cdot y(t))$$
 
$$= e^{k \cdot x(t)} \cdot e^{h \cdot y(t)}$$

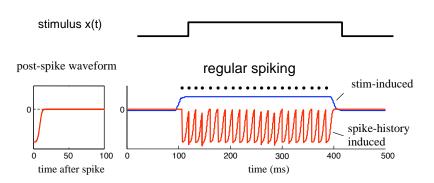
- rate is a product of stim- and spike-history dependent terms
- output no longer a Poisson process
- also known as "soft-threshold" Integrate-and-Fire model

# GLM with history-dependence

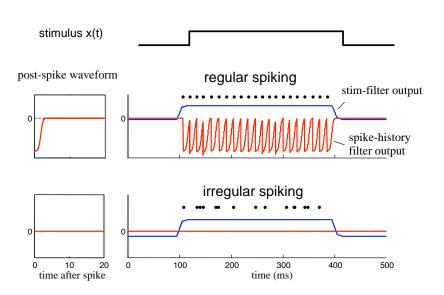


"soft-threshold" approximation to Integrate-and-Fire model

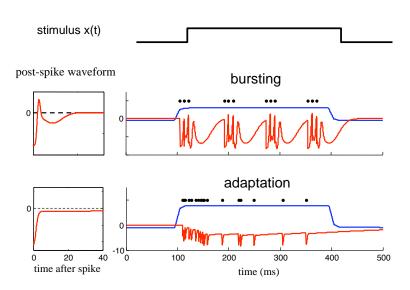
# GLM dynamic behaviors



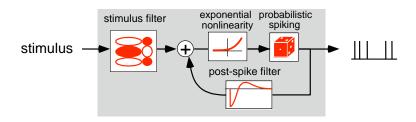
# GLM dynamic behaviors



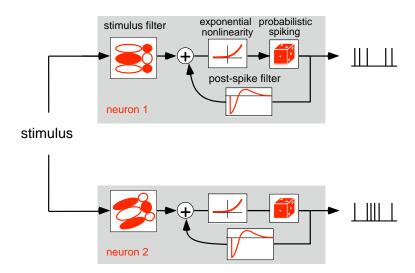
# GLM dynamic behaviors



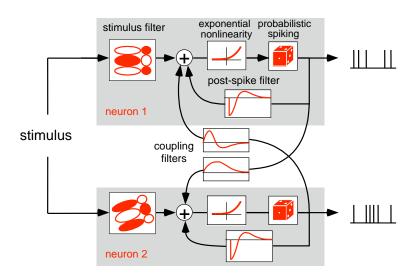
# Generalized Linear Model (GLM)



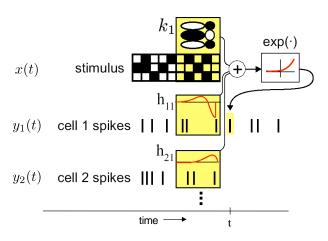
## multi-neuron GLM



## multi-neuron GLM



# GLM equivalent diagram:



conditional intensity (spike rate) 
$$\lambda_i(t) = \exp(k_i \cdot x(t) \ + \ \sum_j h_{ij} \cdot y(t))$$

#### Non-LN models?

The idea of responses depending on one or a few linear stimulus projections has been dominant, but cannot capture all non-linearities.

- ► Contrast sensitivity might require normalisation by  $\|\mathbf{s}\|$ .
- Linear weighting may depend on units of stimulus measurement: amplitude? energy? logarithms? thresholds? (NL models – Hammerstein cascades)
- Neurons, particularly in the auditory system are known to be sensitive to combinations of inputs: forward suppression; spectral patterns (Young); time-frequency interactions (Sadogopan and Wang).
- Experiments with realistic stimuli reveal nonlinear sensivity to parts/whole (Bar-Yosef and Nelken).

Many of these questions can be tackled using a multilinear (cartesian tensor) framework.

The basic linear model (for sounds):

$$\widehat{r}(i) = \sum_{jk} \underbrace{w_{jk}^{\text{tf}}}_{\text{STRF weights}} \underbrace{s(i-j,k)}_{\text{stimulus power}} \; ,$$

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Define: basis functions  $\{g_i\}$  such that  $g(s) = \sum_i w_i^j g_i(s)$  and stimulus array  $M_{ijkl} = g_i(s(i-j,k))$ . Now the model is

$$\hat{r}(i) = \sum_{i \neq l} w_{jk}^{\mathsf{tf}} w_{l}^{\mathsf{I}} M_{ijkl}$$

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Define: basis functions  $\{g_i\}$  such that  $g(s) = \sum_i w_i^j g_i(s)$  and stimulus array  $M_{ijkl} = g_l(s(i-j,k))$ . Now the model is

$$\hat{r}(i) = \sum_{jkl} w_{jk}^{\mathsf{tf}} w_{l}^{\mathsf{l}} M_{ijkl} \quad \mathsf{or} \quad \hat{\mathbf{r}} = (\mathbf{w}^{\mathsf{tf}} \otimes \mathbf{w}^{\mathsf{l}}) \bullet \mathbf{M}.$$

#### **Multilinear models**

Multilinear forms are straightforward to optimise by alternating least squares.

Cost function:

$$\mathcal{E} = \left\| \mathbf{r} - (\mathbf{w}^{\mathsf{tf}} \otimes \mathbf{w}^{\mathsf{I}}) \bullet \mathbf{M} \right\|^{2}$$

Minimise iteratively, defining matrices

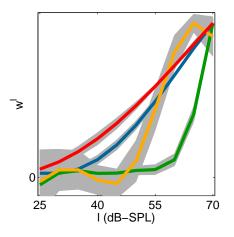
$$\mathbf{B} = \mathbf{w}^{\mathsf{I}} \bullet \mathbf{M}$$
 and  $\mathbf{A} = \mathbf{w}^{\mathsf{tf}} \bullet \mathbf{M}$ 

and updating

$$\mathbf{w}^{tf} = (\mathbf{B}^T \mathbf{B})^{-1} \mathbf{B}^T \mathbf{r}$$
 and  $\mathbf{w}^I = (\mathbf{A}^T \mathbf{A})^{-1} \mathbf{A}^T \mathbf{r}$ .

Each linear regression step can be regularised by evidence optimisation (suboptimal), with uncertainty propagated approximately using *variational* methods.

## Some input non-linearities



Variable (combination-dependent) input gain

Sensitivities to different points in sensory space are not independent.

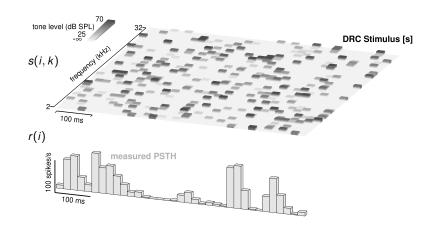
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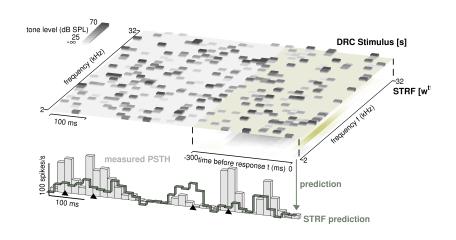
- Sensitivities to different points in sensory space are not independent.
- Rather, the sensitivity at one point depends on other elements of the stimulus that create a local sensory context.
- This context adjusts the input gain of the cell from moment to moment, dynamically refining the shape of the weighted receptive field.

# **Context-sensitive gain**



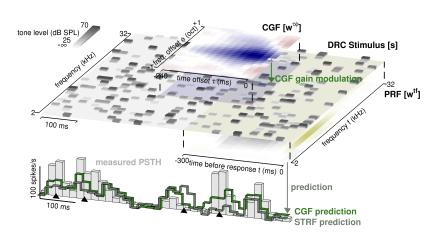
# Context-sensitive gain

$$\hat{r}(i) = c + \sum_{j=0}^{J} \sum_{k=1}^{K} w_{j+1,k}^{tf} s(i-j,k)$$



## Context-sensitive gain

$$\hat{r}(i) = c + \sum_{j=0}^{J} \sum_{k=1}^{K} w_{j+1,k}^{tf} s(i-j,k) \left( 1 + \sum_{m=0}^{M} \sum_{n=-N}^{N} w_{m+1,n+N+1}^{\tau \phi} s(i-j-m,k+n) \right)$$



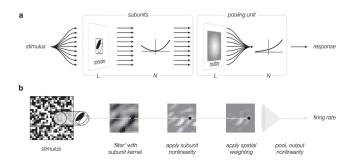
#### **LNLN** cascades

Limited description of 'layered' structure of sensory pathways:

$$\hat{r}(t) = f\left(\sum_{n=1}^{N} w_n g_n(\mathbf{k}_n^{\mathsf{T}} \mathbf{s}(t))\right)$$

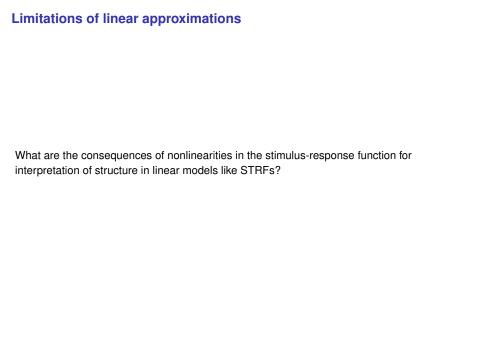
- ▶  $\mathbf{k}_n$  describes the linear filter and  $g_n$  the output nonlinearity of each of N input subunits. The  $g_n$  are usually fixed half-wave rectifiers.
- Called a generalised nonlinear model (GNM; Butts et al. 2007, 2011; Schinkel-Bielefeld et al. 2012)
- Or a nonlinear input model (NIM; McFarland et al. 2013).
- Parameters estimated by maximum-likelihood using inhomogeneous Poisson noise often by alternation (following Ahrens et al. 2008).
- Resembles a (perceptron) "neural network".

#### **Convolutional LNLN**

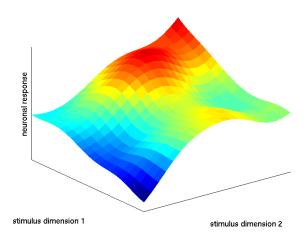


$$\hat{r}(t) = f\left(\sum_{c=1}^{C} \sum_{n=1}^{N} w_{c,n} \sum_{i=1}^{B} b_{c,i} g_i(\mathbf{k}_{c,n}^{\mathsf{T}} \mathbf{s}(t))\right)$$

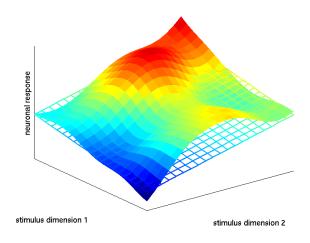
- ightharpoonup C "channels" each uses same kernel  $\mathbf{k}_c$  translated to a different location (convolution).
- ▶ Input nonlinearities learned using basis expansion and alternation (Ahrens et al. 2008).
- Output nonlinearity f fixed.

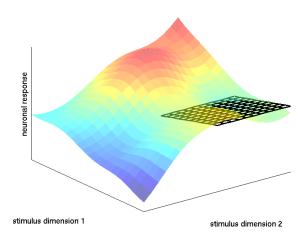


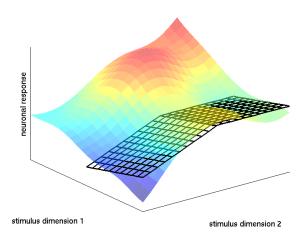
#### Linear fits to non-linear functions

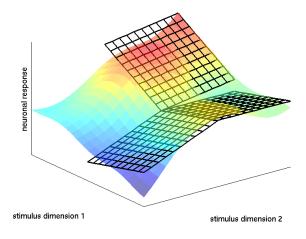


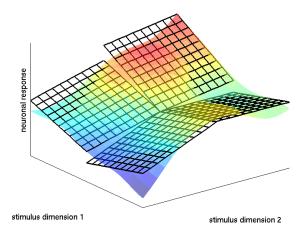
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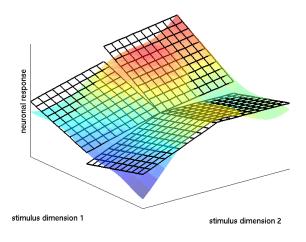




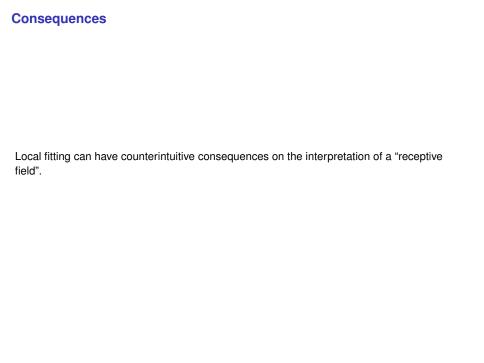






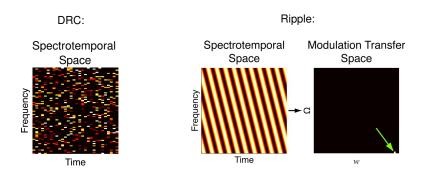


(Stimulus dependence does not always signal response adaptation)



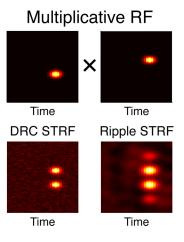
## "Independently distributed" stimuli

Knowing stimulus power at any set of points in analysis space provides no information about stimulus power at any other point.



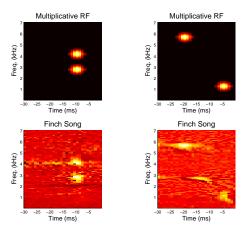
Independence is a property of stimulus and analysis space.

## Nonlinearity & non-independence distort RF estimates



Stimulus may have higher-order correlations in other analysis spaces — and interaction with nonlinearities can produce misleading "receptive fields." (Christianson, Sahani and Linden 2008 J Neurosci)

#### What about natural sounds?



Usually not independent in any space — so STRFs may not be conservative estimates of receptive fields.

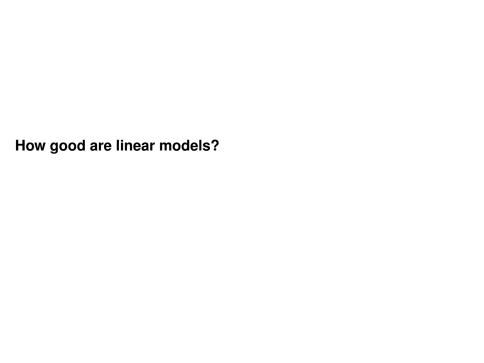
#### **Summary**

How can we use linear models of neuronal stimulus-response functions most effectively to answer biological questions?

Pay a lot of attention to three key issues:

- 1. nature of stimulus
  - ethological/physiological relevance?
  - second-order and/or higher-order autocorrelations?
- 2. choice of stimulus representation
  - appropriate to the biology?
  - appropriate to the question?
- 3. limitations of linear approximation
  - consequences of likely nonlinearities in stimulus-response function?
  - interaction with higher-order autocorrelation in stimulus?

Linear modelling can be a simple and useful tool for answering specific questions about neural coding of stimuli, but results must be interpreted carefully.



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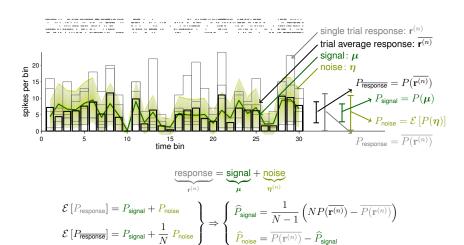
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#### Approaches:

- Compare I(resp; pred) to I(resp; stim).
  - mutual information estimators are biased (and may not be what we really want)
- Compare E(resp pred) to E(resp psth) where psth is gathered over a very large number of trials.
  - may require impractical amounts of data to estimate the psth
- Compare the predictive power to the predictable power (similar to ANOVA).

## **Estimating predictable power**



# Testing a model

For a perfect prediction

$$\left\langle P(\overline{trial}) - P(residual) \right\rangle = P(signal)$$

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$$\left\langle \mathsf{P}(\overline{\mathsf{trial}}) - \mathsf{P}(\mathsf{residual}) \right\rangle = \mathsf{P}(\mathsf{signal})$$

Thus, we can judge the performance of a model by the normalized predictive power

$$\frac{P(\overline{trial}) - P(residual)}{\widehat{P}(signal)}$$

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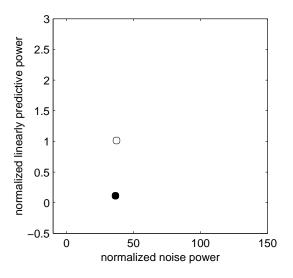
For a perfect prediction

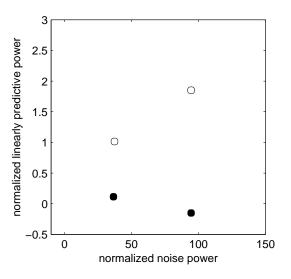
$$\left\langle \mathsf{P}(\overline{\mathsf{trial}}) - \mathsf{P}(\mathsf{residual}) \right\rangle = \mathsf{P}(\mathsf{signal})$$

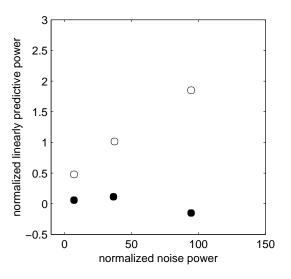
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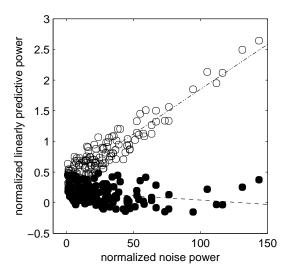
$$\frac{P(\overline{trial}) - P(residual)}{\widehat{P}(signal)}$$

Similar to coefficient of determination  $(r^2)$ , but the denominator is the predictable variance.

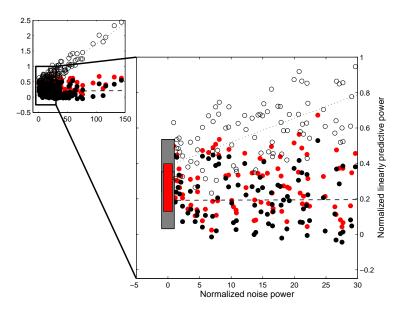






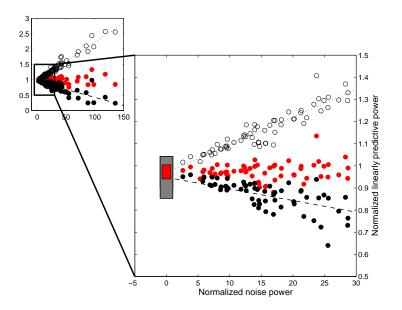


# **Extrapolated linearity**



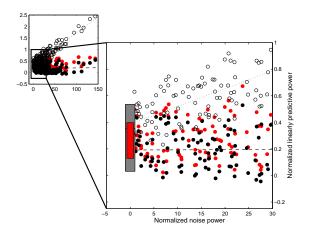
[extrapolated range: (0.19,0.39); mean Jackknife estimate: 0.29]

### Simulated (almost) linear data



[extrapolated range: (0.95,0.97); mean Jackknife estimate: 0.97]

## Linearity and nonlinearity in auditory cortical responses



So, spectrogram-linear models capture approximately 20–40% of the variability in auditory cortical responses to random chord stimuli (Sahani and Linden 2003 NIPS).

For natural sounds, performance is no better (Machens et al. 2004 J Neurosci).



Spectrogram-linear models perform better in the thalamus than in the cortex (more on this later).

Not just because cortex is noisier but because cortical representations are more nonlinear!

Other studies likewise indicate that linearity of stimulus representation generally decreases as we ascend the auditory pathway (e.g., Chechik and Nelken 2012 PNAS; Atencio et al. 2012 J Neurosci; Williamson et al. 2016 Neuron).