

GENERALIZED LINEAR MODEL

Linear model

Linear model

Parameters

Measurement noise

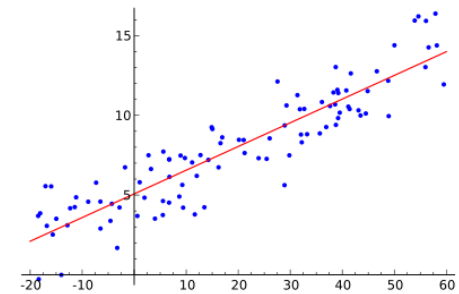
$$Y = X\beta + \varepsilon$$

$\varepsilon \sim N(0, \sigma^2)$

Linear predictor

Dependent variable

Explanatory variables



Probabilistic representation: A normal distribution

$$f(y | \mu, \sigma^2) = \frac{1}{\sqrt{2\pi\sigma^2}} e^{-\frac{(y-\mu)^2}{\sigma^2}},$$

where the mean is modeled as $\mu = X\beta$.

Linear predictor

Generalized linear model

What if the measurement noise follows non-Gaussian distribution?

GLM-analysis extends the regression analysis when the residual is given by an exponential family of distributions.

The exponential family of distributions include many popular distributions: normal, log-normal, exponential, gamma, chi-squared, beta, Dirichlet, Bernoulli, categorical, Poisson, inverse Gaussian, Wishart, Inverse Wishart, von Mises, etc.

Exponential family of distributions

Parameter(s) of the family

(Sufficient statistics)

$$f(y|\theta) = h(y) \exp[\eta(\theta)T(y) - A(\theta)]$$

Natural parameter

log-partition function

By re-parameterizing the distribution by η , an exponential family is simplified to a 'natural form' (canonical form) as

$$f(y|\eta) = h(y) \exp[\eta T(y) - A(\eta)]$$

Relation between a natural parameter and log-partition function:

$$\frac{\partial A(\eta)}{\partial \eta} = E_Y[T(Y)] \quad \frac{\partial^2 A(\eta)}{\partial \eta^2} = E_Y[T(Y) - E_Y T(Y)]^2 = \text{var}[T(Y)]$$

Examples

Bernoulli probability mass function.

$$f(y; \mu) = \mu^y (1 - \mu)^{1-y} = \exp\left[y \log \frac{\mu}{1 - \mu} + \log(1 - \mu)\right] \quad y = \{0, 1\}$$

Poisson probability mass function.

$$f(y; \mu) = \frac{\mu^y}{y!} e^{-\mu} = \frac{1}{y!} \exp[y \log \mu - \mu] \quad y = \{0, 1, 2, \dots\}$$

Generalized linear model

Modeling the mean by a linear predictor.

$$f(y; \theta) = h(y) \exp[\eta(\theta) T(y) - A(\theta)]$$

Mean $\mu \equiv E_Y[Y]$ \longleftrightarrow Linear predictor $X\beta$

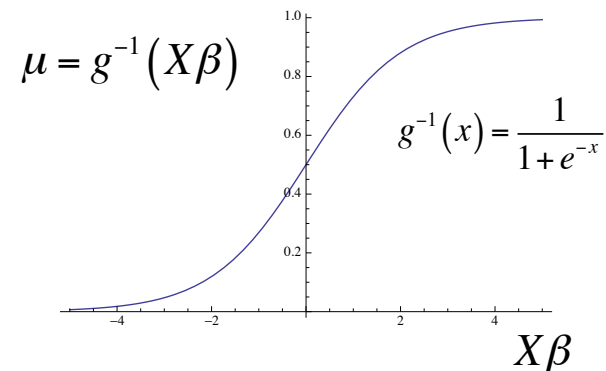
Mean is bounded $0 < \mu < 1$ (Bernoulli)
 $0 < \mu$ (Poisson)

~~$\mu = X\beta$~~

A link function relates the mean and a linear predictor:

$$\mu = g^{-1}(X\beta) \quad X\beta = g(\mu)$$

so that μ is bounded.



A link function can be selected arbitrary. However, there is a function that arises 'naturally', depending on the type of distribution.

Canonical link function: Bernoulli

Poisson probability mass function.

$$f(y; \mu) = \mu^y (1 - \mu)^{1-y} = \exp \left[y \log \frac{\mu}{1 - \mu} + \log(1 - \mu) \right] \quad y = \{0, 1\}$$

Linear predictor: $\eta = X\beta$

Model the mean using a link function: $\mu = g^{-1}(\eta)$

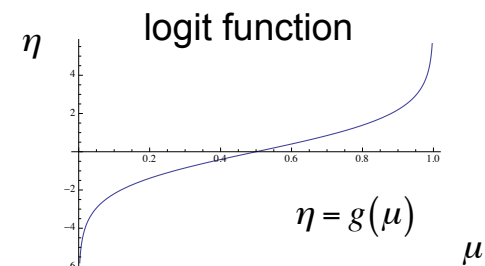
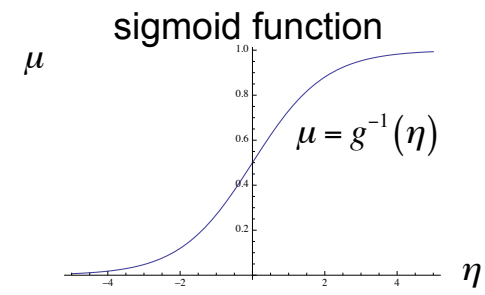
Canonical link function: (logit function)

If we use
$$\mu = g^{-1}(\eta) = \frac{1}{1 + e^{-\eta}},$$

namely
$$\eta = g(\mu) = \log \frac{\mu}{1 - \mu},$$

then the distribution is written in a natural form as

$$f(y; \eta) = \exp[y\eta - A'(\eta)]$$



Canonical link function: Poisson

Bernoulli probability mass function.

$$f(y; \mu) = \frac{\mu^y}{y!} e^{-\mu} = \frac{1}{y!} \exp[y \log \mu - \mu] \quad y = \{0, 1, 2, \dots\}$$

Linear predictor: $\eta = X\beta$

Model the mean using a link function: $\mu = g^{-1}(\eta)$

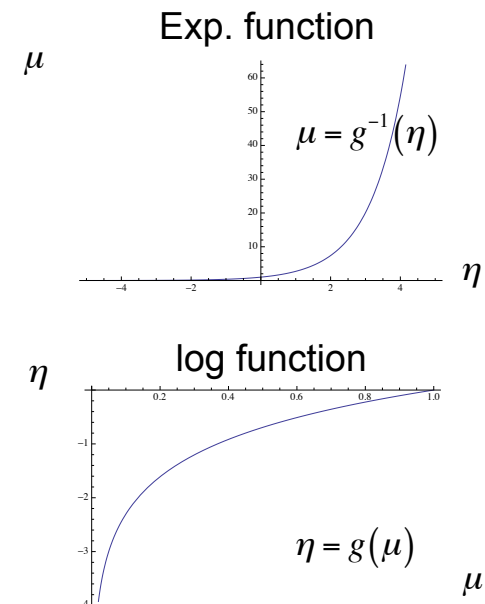
Canonical link function: (log function)

If we use $\mu = g^{-1}(\eta) = e^\eta$,

namely $\eta = g(\mu) = \log \mu$,

then the distribution is written in a natural form as

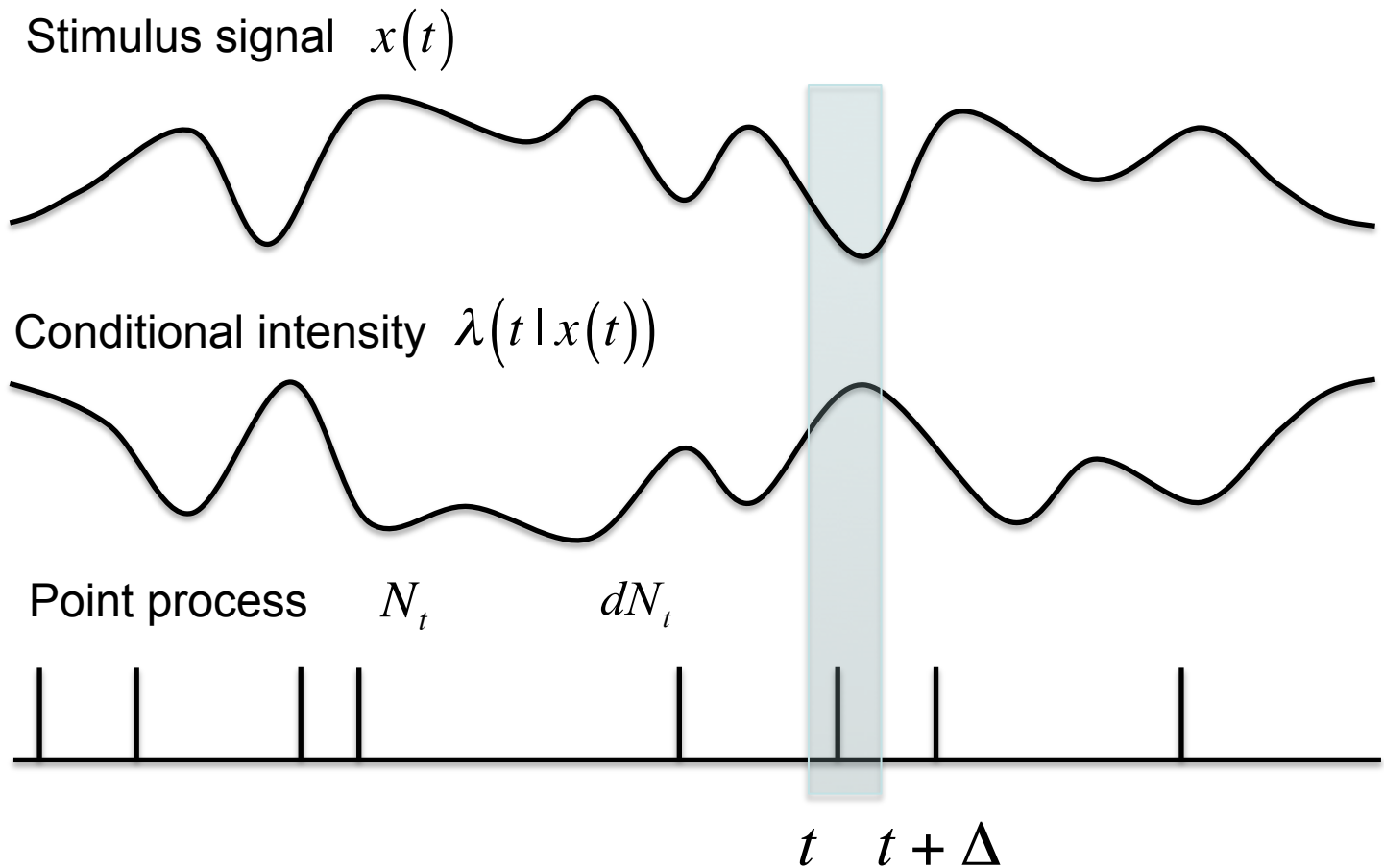
$$f(y; \eta) = \frac{1}{y!} \exp[y\eta - e^\eta]$$



Relating a stimulus input/motor output with neural activity

POINT PROCESS-GLM

Relating stimulus and neural activity

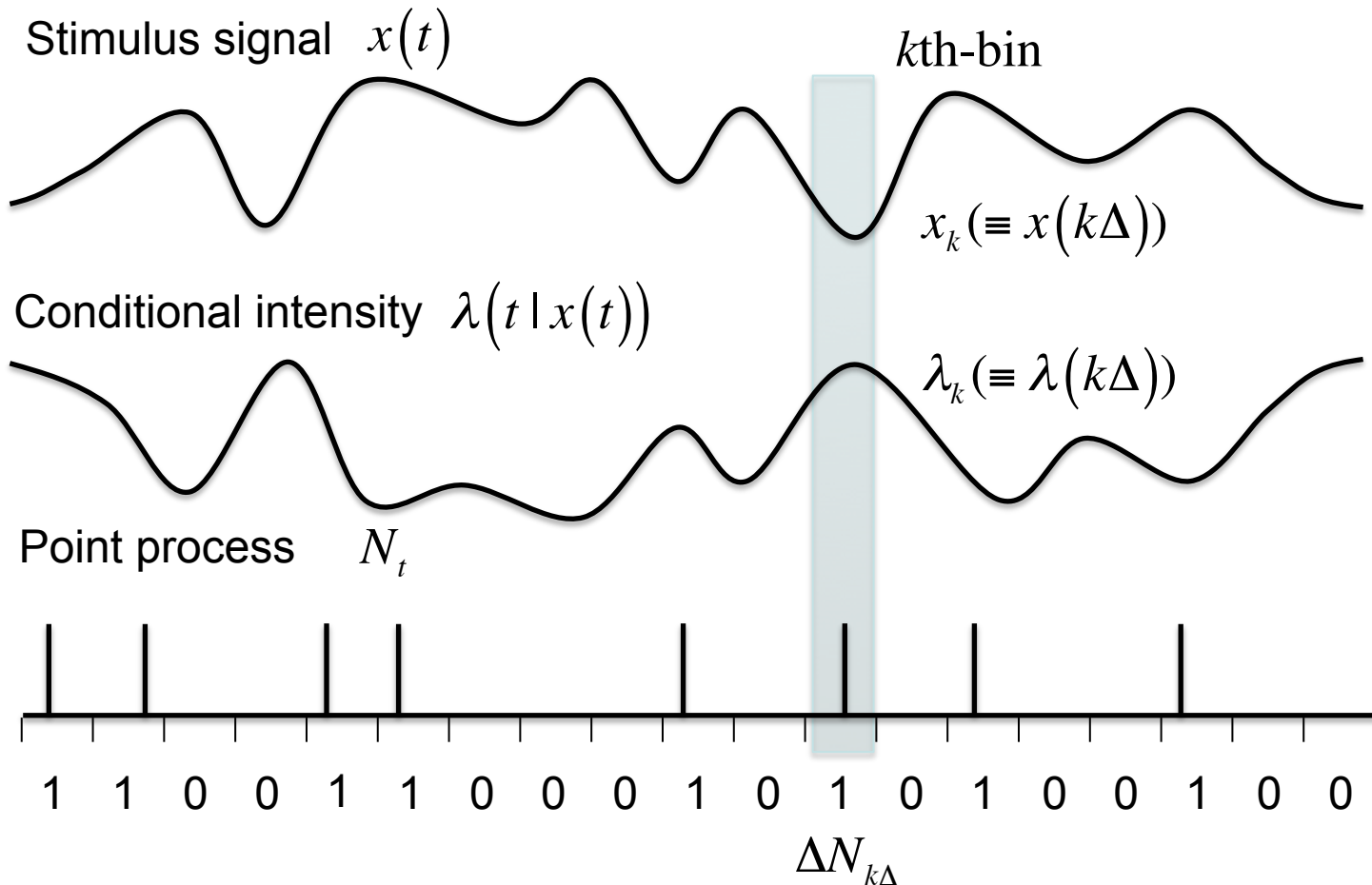


$$P(\text{a spike in } [t, t + \Delta) | x(t)) = \lambda(t | x(t))\Delta + o(\Delta)$$



Stimulus signal (Covariate)

Discretizing time into bins



Likelihood (Joint probability mass function)

$$P(\Delta N_{\Delta}, \Delta N_{2\Delta}, \dots, \Delta N_{N\Delta}) = \prod_{k=1}^N P(\Delta N_{k\Delta} | \Delta N_{\Delta}, \dots, \Delta N_{(k-1)\Delta})$$

Conditional probability mass function

Bernoulli-GLM

Conditional Bernoulli probability:

$$P(\Delta N_{k\Delta} | \Delta N_{\Delta}, \dots, \Delta N_{(k-1)\Delta}) = (\lambda_k \Delta)^{\Delta N_{k\Delta}} (1 - \lambda_k \Delta)^{1 - \Delta N_{k\Delta}} \quad \lambda_k \equiv \lambda(k\Delta | x(k\Delta))$$
$$= \exp \left[\Delta N_{k\Delta} \log \frac{\lambda_k \Delta}{1 - \lambda_k \Delta} + \log \{1 - \lambda_k \Delta\} \right]$$

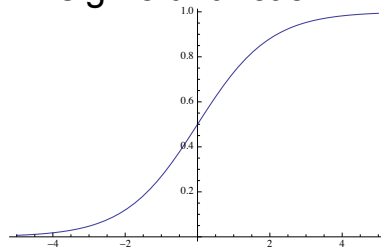
Mean of the Bernoulli probability at k-th bin: $\mu = E(\Delta N_{k\Delta}) = \lambda_k \Delta$

Modeling the mean response via a canonical link function:

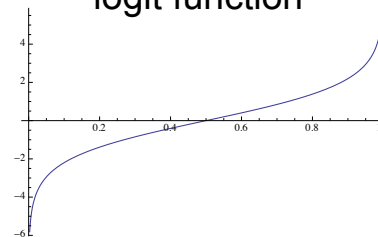
$$\lambda_k \Delta = \frac{1}{1 + e^{-(\beta_0 + \beta x_k)}}$$

$$\log \frac{\lambda_k \Delta}{1 - \lambda_k \Delta} = \beta_0 + \beta x_k$$

sigmoid function



logit function



misprint!

Poisson-GLM

Approximation by Poisson distribution:

$$P(\Delta N_{k\Delta} | \Delta N_{\Delta}, \dots, \Delta N_{(k-1)\Delta}) = \left(\frac{\lambda_k \Delta}{1 - \lambda_k \Delta} \right)^{\Delta N_{k\Delta}} (1 - \lambda_k \Delta) \\ \approx \exp[\Delta N_{k\Delta} \log\{\lambda_k \Delta\} - \lambda_k \Delta]$$

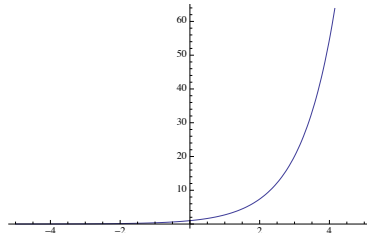
Mean of the Poisson distribution at k-th bin: $\mu = E(\Delta N_{k\Delta}) = \lambda_k \Delta$

Modeling the mean response via a canonical link function:

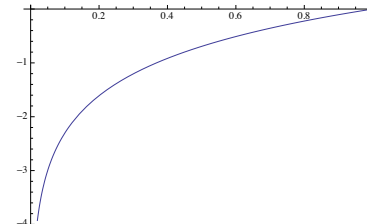
$$\lambda_k \Delta = \exp[\beta_0 + \beta x_k]$$

$$\log[\lambda_k \Delta] = \beta_0 + \beta x_k$$

Exp. function



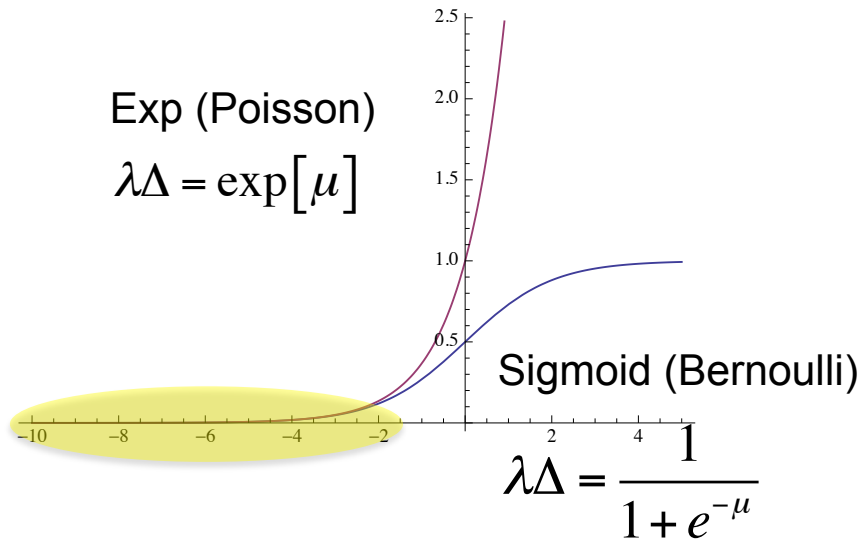
log function



Relation between Bernoulli and Poisson-GLM

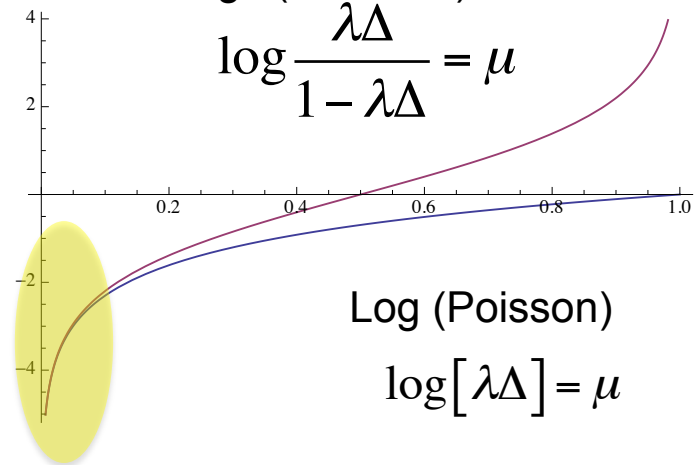
Inverse link function

Exp (Poisson)
 $\lambda\Delta = \exp[\mu]$



Link function

Logit (Bernoulli)
 $\log \frac{\lambda\Delta}{1 - \lambda\Delta} = \mu$



misprint

If the mean $\lambda\Delta$ is small, an exponential and sigmoid functions are nearly identical. Similarly, log and sigmoid functions are nearly identical for small $\lambda\Delta$.

Is the point process a Poisson?

We now recall that the number of spike count in a certain interval follows the Poisson distribution.

The spike count in a small bin is then approximated as

$$P(N_{\Delta} = n) = \frac{(\lambda\Delta)^n}{n!} e^{-\lambda\Delta} = \frac{(\lambda\Delta)^n}{n!} \left[1 - \lambda\Delta + \frac{1}{2}(\lambda\Delta)^2 + \dots \right]$$

In particular,

$$P(N_{\Delta} = 0) = 1 \left[1 - \lambda\Delta + \frac{1}{2}(\lambda\Delta)^2 + \dots \right] = 1 - \lambda\Delta + o(\Delta)$$

$$P(N_{\Delta} = 1) = \lambda\Delta \left[1 - \lambda\Delta + \frac{1}{2}(\lambda\Delta)^2 + \dots \right] = \lambda\Delta + o(\Delta)$$

$$P(N_{\Delta} = 2) = (\lambda\Delta)^2 \left[1 - \lambda\Delta + \frac{1}{2}(\lambda\Delta)^2 + \dots \right] = o(\Delta)$$

The probability of having a spike/no spike is an approximation of the Poisson count distribution for a small time bin.

From discrete to continuous time

Likelihood of discrete-time conditional Poisson distributions:

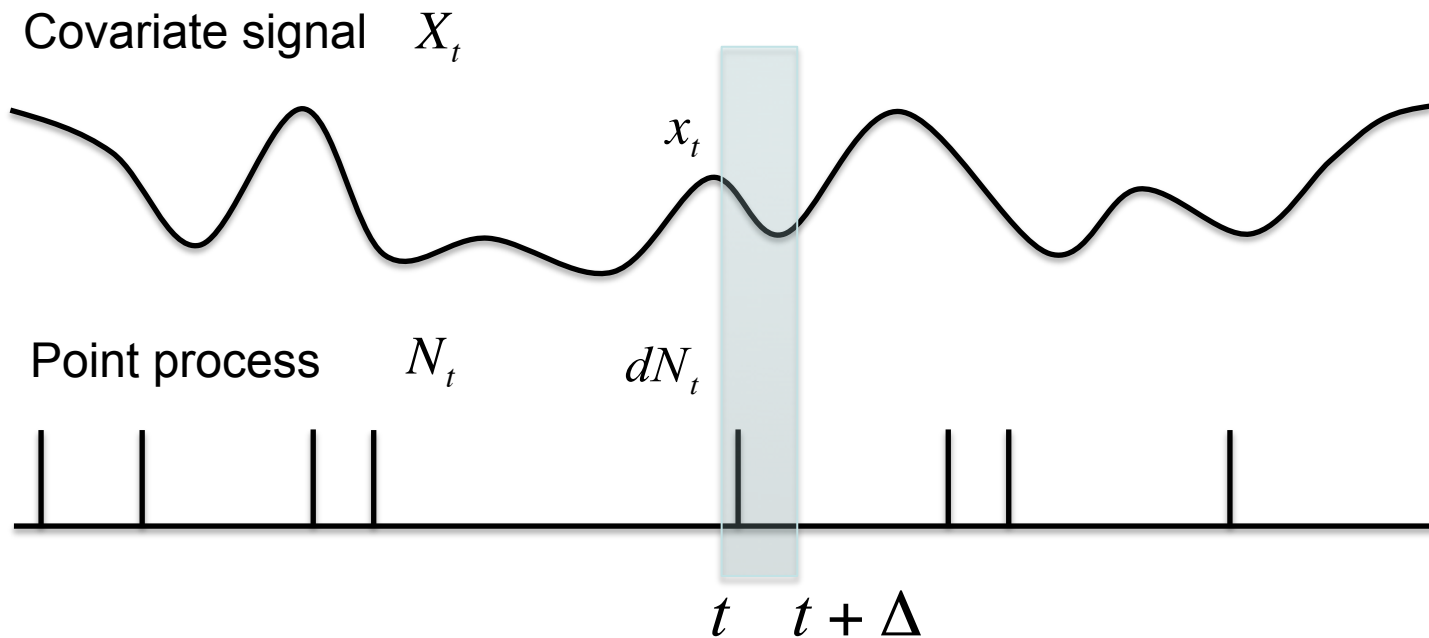
$$\begin{aligned} P(\Delta N_{\Delta}, \Delta N_{2\Delta}, \dots, \Delta N_{N\Delta}) &= \prod_{k=1}^K \exp[\Delta N_{k\Delta} \log\{\lambda_k \Delta\} - \lambda_k \Delta] + o(\Delta^n) \\ &= \Delta^n \exp\left[\sum_{k=1}^K \Delta N_{k\Delta} \log \lambda_k - \lambda_k \Delta\right] + o(\Delta^n) \end{aligned}$$

Likelihood of a point process by continuous-time limit:

$$\begin{aligned} p(t_1, t_2, \dots, t_n \cap N_T = n) &= \lim_{\Delta \rightarrow 0} \frac{P(\Delta N_{\Delta}, \Delta N_{2\Delta}, \dots, \Delta N_{N\Delta})}{\Delta^n} \\ &= \lim_{\Delta \rightarrow 0} \frac{\exp\left[\sum_{k=1}^K \Delta N_{k\Delta} \log\{\lambda_k\} - \sum_{k=1}^K \lambda_k \Delta\right] \Delta^n + o(\Delta^n)}{\Delta^n} \\ &= \exp\left[\int_0^T \log \lambda_u dN_u - \int_0^T \lambda_u du\right] \end{aligned}$$

Truccolo et al. J Neurophysiol (2005)

Point process



$$P(\text{a spike in } [t, t + \Delta) \mid H_t, X_t) = \lambda(t \mid H_t, X_t) \Delta + o(\Delta)$$



History of spiking activity

History of a covariate signal
(Filtration)

$$H_t = \{t_1 < t_2 < \dots < t_n \cap N_t = n\}$$

$$X_t = \{x_u : 0 < u < t\}$$

Point process-GLM

Likelihood function:

$$\begin{aligned} p(t_1, t_2, \dots, t_n \cap N_T = n) &= \prod_{i=1}^n \lambda(t_i | X_{t_i}, H_{t_i}) \exp\left[-\int_0^T \lambda(u | X_u, H_u) du\right] \\ &= \exp\left[\int_0^T \log \lambda(u | X_u, H_u) dN_u - \int_0^T \lambda(u | X_u, H_u) du\right] \end{aligned}$$

Here $dN_t = \sum_{i=1}^n \delta(t - t_i) dt$

Modeling the conditional mean rate via a canonical link function.

$$E\left[\frac{dN_t}{dt}\right] = \lambda(t | H_t, X_t) = \exp[\beta_0 + \beta_1 f(X_t) + \beta_2 g(H_t)]$$



a log-link function

A canonical link function for a Poisson distribution.

Covariates for neural signals

The covariates can be added in a (generalized) linear manner.

$$\begin{aligned} \log \lambda(t | H_t, X_t) = & \beta_0 + \beta_1 [\text{Input stimulus}] \\ \text{Log-link function} & + \beta'_1 [\text{Higher-order feature of stimulus}] \\ & + \beta_2 [\text{Spike history}] \\ & + \beta_3 [\text{Ensemble spike history}] \\ & + \beta_4 [\text{Local field potential}] \end{aligned}$$

Maximum likelihood estimation

Likelihood

$$L(\beta | N_{0:T}) = \exp \left[\int_0^T \log \lambda(u | X_{t_i}, H_{t_i}) dN_u - \int_0^T \lambda(u | X_u, H_u) du \right]$$

where $\lambda(t | H_t, X_t) = \exp[x(t)\beta]$.

Optimize parameters under the principle of maximizing the likelihood:

$$\beta_{\text{MLE}} = \arg \max_{\beta} \log L(\beta | N_{0:T})$$

Newton-Raphson method:

$$l(\beta) = \log L(\beta | N_{0:T})$$

$$\beta^{j+1} = \beta^j - \underbrace{\nabla \nabla l(\beta^j)}_{\text{Hessian}} \cdot \underbrace{\nabla l(\beta^j)}_{\text{Score function}}$$

Matlab:

$$b = \text{glmfit}(X, y, 'poisson')$$

Applications of point process-GLM

Here we review

A Point Process Framework for Relating Neural Spiking Activity to Spiking History, Neural Ensemble, and Extrinsic Covariate Effects

Wilson Truccolo,¹ Uri T. Eden,^{2,3} Matthew R. Fellows,¹ John P. Donoghue,¹ and Emery N. Brown^{2,3}




J Neurophysiol 93: 1074–1089, 2005.

Modeling conditional intensity function

Conditional intensity function:

$$\lambda(t_k | N_{1:k}^{1:C}, \mathbf{x}_{k+\tau}, \theta) = \lambda_I(t_k | N_{1:k}, \theta_I) \lambda_E(t_k | N_{1:k}^{1:C}, \theta_E) \lambda_X(t_k | \mathbf{x}_{k+\tau}, \theta_X)$$



 Auto spike history Ensemble history Extrinsic covariate

Auto spike history Expected to capture refractory effects, recovery periods, and oscillation.

$$\lambda_I(t_k | N_{1:k}, \theta_I) = \exp \left\{ \gamma_0 + \sum_{n=1}^Q \gamma_n \Delta N_{k-n} \right\} \quad \text{Determined by AIC}$$

Ensemble history Expected to capture lagged spike synchrony between target and other cells.

$$\lambda_E(t_k | N_{1:k}^{1:C}, \theta_E) = \exp \left\{ \beta_0 + \sum_c \sum_{r=1}^R \beta_r^c \Delta N_{k-r}^c \right\}$$

Expected to capture ensemble effect on slower time scale

$$\lambda_E(t_k | N_{1:k}^{1:C}, \theta_E) = \exp \left\{ \beta_0 + \sum_c \sum_{r=1}^R \beta_r^c (N_{k-(r-1)W}^c - N_{k-rW}^c) \right\}$$

Extrinsic covariate (hand velocity model)

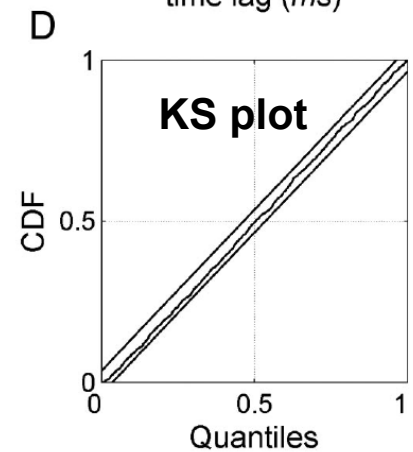
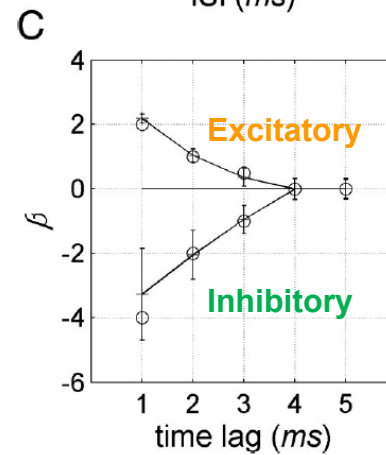
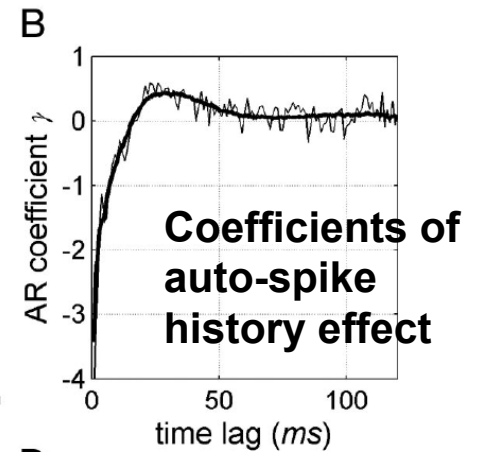
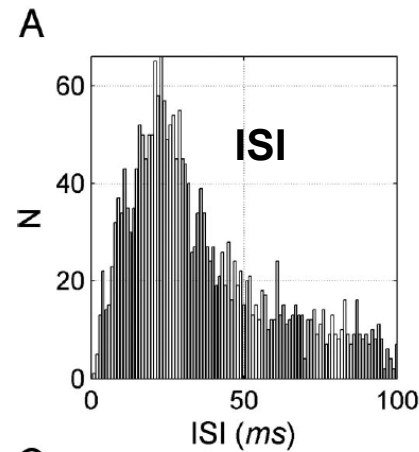
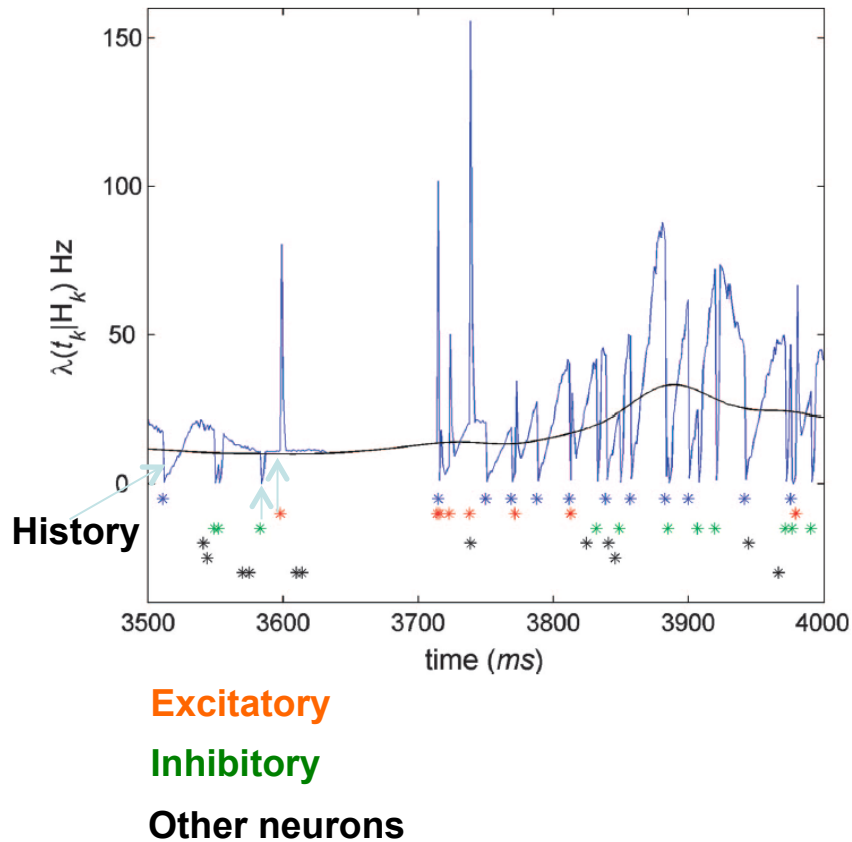
Spike count in a window of size W.

$$\lambda_X(t_k | \mathbf{x}_{k+\tau}, \theta_X) = \exp \left\{ \alpha_0 + |V_{k+\tau}| \left[\alpha_1 \cos(\phi_{k+\tau}) + \alpha_2 \sin(\phi_{k+\tau}) \right] \right\}$$

Angle of hand velocity Magnitude of hand velocity offset $\tau=+150\text{ms}$

Simulation (6 neurons)

Simulated conditional intensity of a **blue** neuron



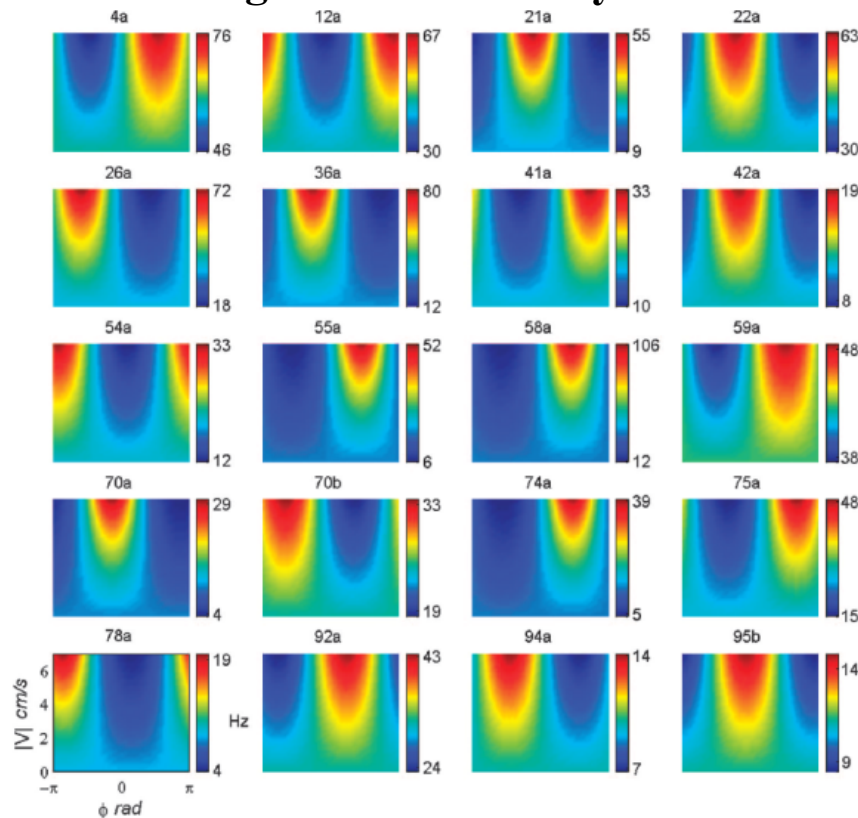
Application to MI spiking data

Two models were compared.

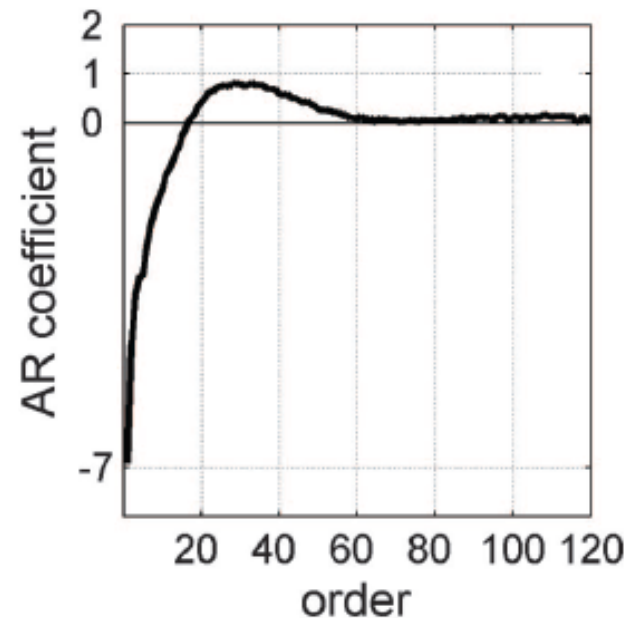
Velocity model

Velocity + Auto spike history model

Tuning curve of velocity model



Auto spike history

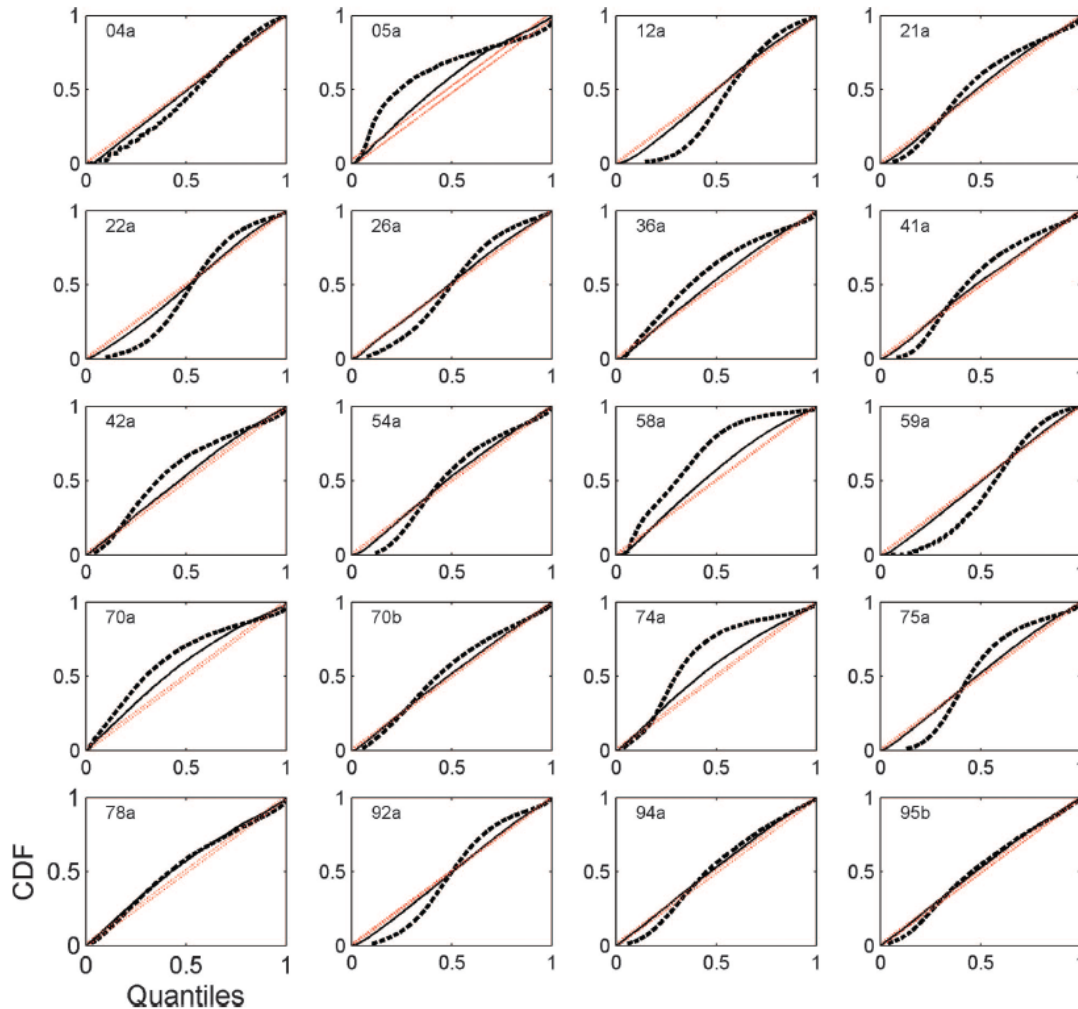


$$\lambda_I(t_k | N_{1:k}, \theta_I) = \exp \left\{ \gamma_0 + \sum_{n=1}^q \gamma_n \Delta N_{k-n} \right\}$$

$$\lambda_X(t_k | \mathbf{x}_{k+\tau}, \theta_X) = \exp \{ \alpha_0 + |V_{k+\tau}| [\alpha_1 \cos(\phi_{k+\tau}) + \alpha_2 \sin(\phi_{k+\tau})] \}$$

Test of goodness-of-fit: KS-plot

KS plot



$$z_j = 1 - \exp \left\{ - \int_{u_j}^{u_{j+1}} \lambda(t | H(t), \hat{\theta}) dt \right\}$$

is expected to be independent uniformly distributed random variables in an unit interval.

Velocity model

Velocity + Auto-spike history model

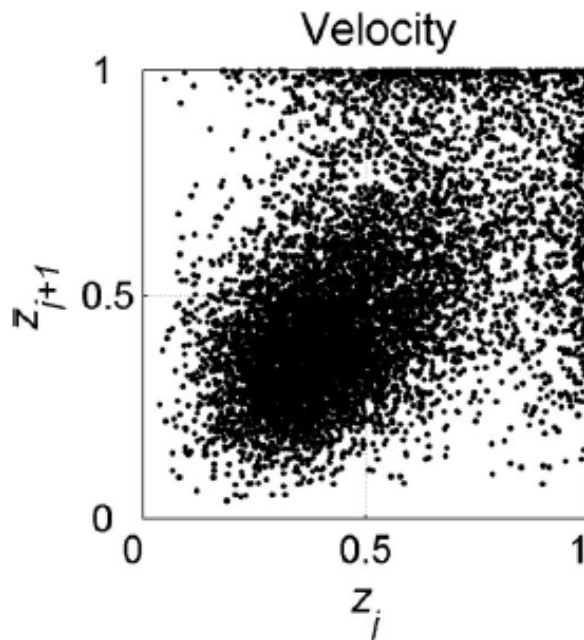
95% confidence bound of KS test

Test of independence for rescaled ISIs

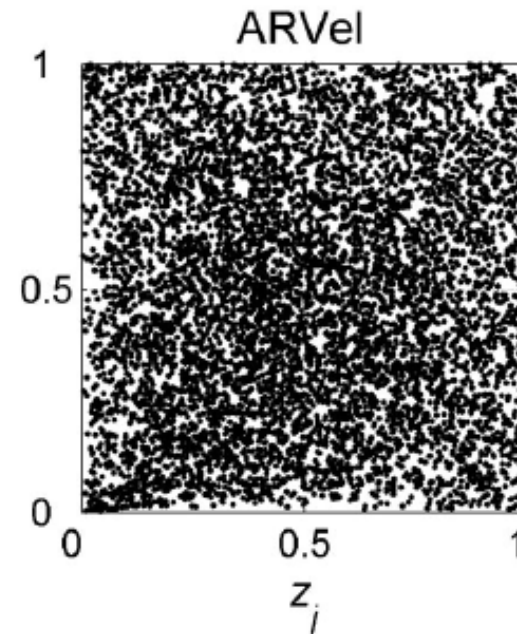
If the conditional intensity model completely explains the data,

$$z_j = 1 - \exp \left\{ - \int_{u_j}^{u_{j+1}} \lambda(t | H(t), \hat{\theta}) dt \right\}$$

is expected to be **independently** and **uniformly** distributed random variables in an unit interval.



There is dependency
between z_j and z_{j+1} .



There is no dependency
between z_j and z_{j+1} .

Model selection by AIC

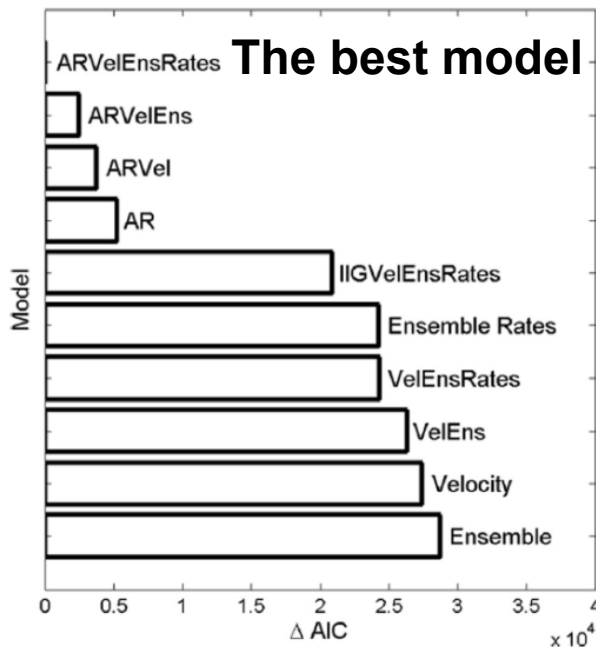
Akaike information criterion

$$\text{AIC} = -2 \log p(\text{DATA} | \beta) + 2 \times \dim(\beta)$$

Parameters of the model

Model dimension
(Number of parameters)

The smaller the AIC, the better the predictive ability of the model is.



AR: Auto-spike history
The models with the auto-spike history significantly decrease AIC.

Point process residual analysis

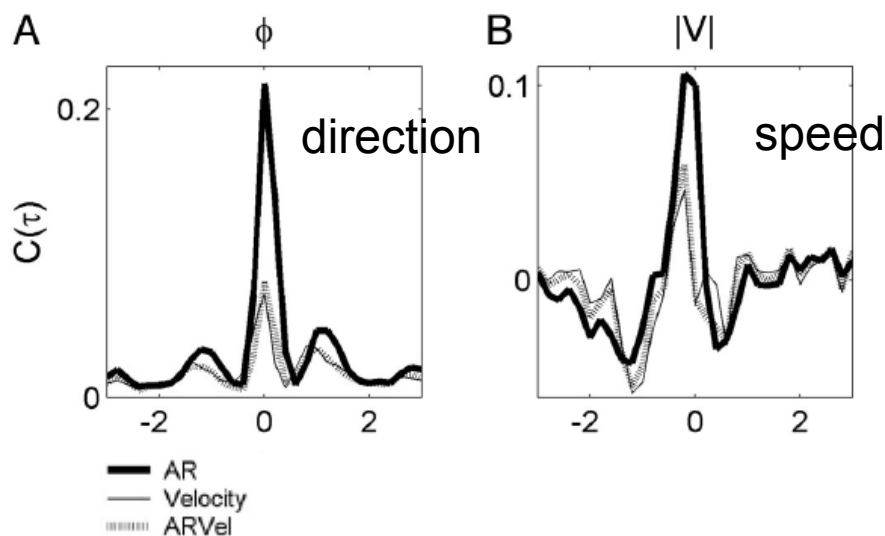
Point process residual

$$M(t_k) = \sum_{i=k-B}^k \Delta N_i - \int_{t_{k-B}}^{t_k} \lambda(t | H(t), \hat{\theta}) dt$$

The difference between the actual spike count and spike count predicted by the model.

If the CIF completely characterizes response characteristics to covariates (stimulus/motor output), the residual should have no dependency with the covariates.

Cross-correlation function between residual and covariates (hand movement).



Inclusion of velocity signal in a CIF greatly reduced the correlation.

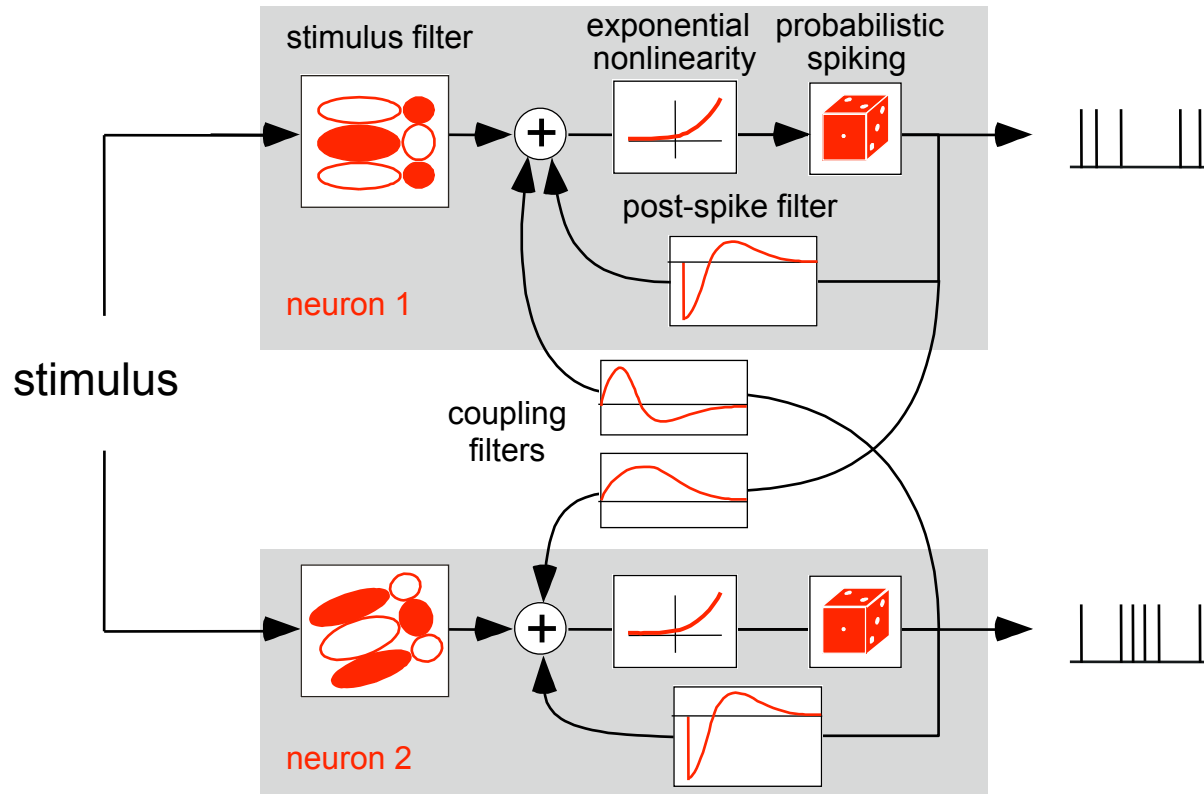
However, the auto-spike history did not further reduce the correlation.

AR component does not carry additional statistical information about hand velocity.

There is a component that was not captured by the point process-GLMs.

We will review

Point process-GLM for the analysis of retinal ganglion cells



Pillow et al (2008). Nature, 454(7207), 995–999.

What we learned

1

- Exponential family of distributions and generalized linear model.

2

- Discrete-time likelihood of conditional Bernoulli and Poisson distributions.

3

- GLM framework for a continuous-time point process.

4

- Review of a paper by Truccolo et al. Focused on methods for model validation (**KS-plot, AIC, residual analysis**).

Course overview

- 1 • Introduction and a **Poisson** point process
- 2 • **Renewal** and **non-Poisson** processes
- 3 • **Point process-GLM**: Stimulus and a point process (Encoding)
- 4 • Inference for a Poisson process: **Spike-rate estimation**
- 5 • **State-space** model and a **point process filter** (Decoding)
- 6 • Paper reviews by SCS 1st year students

From the feedback of the last lecture

