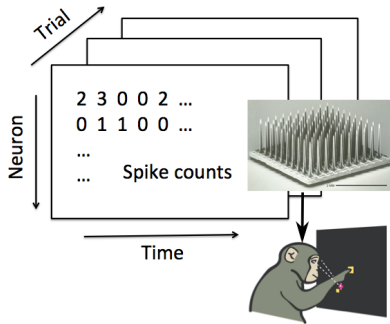


Statistical Machine Learning Methods for High-dimensional Neural Population Data Analysis

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Overview



- Neuroscience + Big data = Opportunities!

Overview

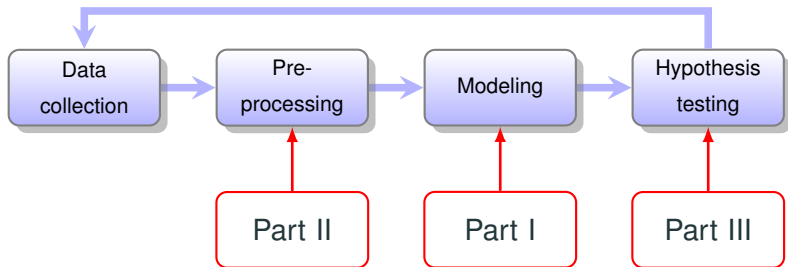


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- Generalized count linear dynamical system
- Linear dynamical neural population models through nonlinear embeddings

II. Region of Interest Detection for Calcium Imaging Data

III. Maximum Entropy Flow Networks

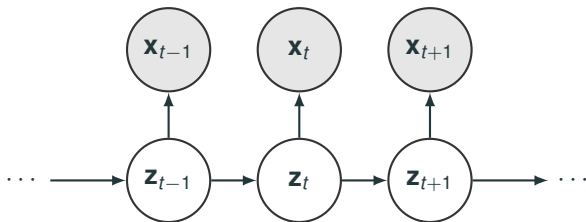
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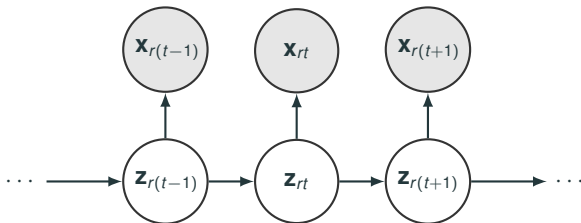
State space models



- $\mathbf{x}_t \in \mathbb{N}^n$: spike counts; $\mathbf{z}_t \in \mathbb{R}^m$: latent variables
- Joint distribution

$$p(\mathbf{x}, \mathbf{z}) = \underbrace{p(\mathbf{z}_1)}_{\text{Initial distribution}} \underbrace{\prod_{t=1}^{T-1} p(\mathbf{z}_{t+1} | \mathbf{z}_t)}_{\text{Transition model}} \underbrace{\prod_{t=1}^T p(\mathbf{x}_t | \mathbf{z}_t)}_{\text{Observation model}}$$

State space models: multiple trials



- $r = 1, \dots, R$: trial number
- $\mathbf{x}_{rt} \in \mathbb{N}^n$: spike counts; $\mathbf{z}_{rt} \in \mathbb{R}^m$: latent variables
- Joint distribution

$$p(\mathbf{x}, \mathbf{z}) = \prod_{r=1}^R \left[\underbrace{p(\mathbf{z}_{r1})}_{\text{Initial distribution}} \underbrace{\prod_{t=1}^{T-1} p(\mathbf{z}_{r(t+1)} | \mathbf{z}_{rt})}_{\text{Transition model}} \underbrace{\prod_{t=1}^T p(\mathbf{x}_{rt} | \mathbf{z}_{rt})}_{\text{Observation model}} \right]$$

Common parameterization and our extensions

- Common assumptions for latent dynamics: linear Gaussian dynamical system (LDS)

$$\mathbf{z}_1 \sim \mathcal{N}(\mu_1, Q_1)$$

$$\mathbf{z}_{t+1} | \mathbf{z}_t \sim \mathcal{N}(A\mathbf{z}_t, Q)$$

- Common observation models:

$$\mathbf{x}_t | \mathbf{z}_t \sim \underbrace{\mathcal{N}(C\mathbf{z}_t + d, \Sigma)}_{\text{model mismatch}} \text{ or } \underbrace{\text{Poisson}(\exp(C\mathbf{z}_t + d))}_{\text{equal dispersion}}$$

stringent assumptions

- Our extensions for observation model:
 - Generalized count distribution (GCLDS) (Gao et al. 2015)
 - Flexible nonlinear observation (fLDS) (Gao et al. 2016)

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Motivation

- Doubly stochastic Poisson model implies **overdispersion**

$$\left. \begin{array}{l} \mathbf{z} \sim p(\mathbf{z}) \\ \mathbf{x} \sim \text{Poisson}(f(\mathbf{z})) \end{array} \right\} \Rightarrow \begin{array}{l} \text{var}(\mathbf{x}|\mathbf{z}) = E(\mathbf{x}|\mathbf{z}) \\ \text{var}(\mathbf{x}) \geq E(\mathbf{x}) \end{array}$$

- Need a more flexible distribution to separate **firing rate variability** with **noise variability**.

$$\text{var}(\mathbf{x}) = \underbrace{\text{var}(E(\mathbf{x}|\mathbf{z}))}_{\text{firing rate variability}} + \underbrace{E(\text{var}(\mathbf{x}|\mathbf{z}))}_{\text{noise variability}}$$

Generalized count distribution family

- Generalized count (GC) distribution family

$$p_{\text{Poisson}}(x; \lambda) \propto \frac{\exp \{ \log \lambda \cdot x \}}{x!}, \quad x \in \mathbb{N}$$

\Downarrow

$$p_{\text{GC}}(x; \theta, g(\cdot)) \propto \frac{\exp(\theta \cdot x + g(x))}{x!}, \quad x \in \mathbb{N}$$

where $\theta \in \mathbb{R}$, $g(\cdot) : \mathbb{N} \rightarrow \mathbb{R}$.

- Parameterizes **all** the count distributions **redundantly**.
- Given $g(\cdot)$, θ controls the expectation.
- $g(\cdot)$ controls the “shape” of the distribution.
Convex/concave/linear $g(\cdot)$ implies
overdispersed/underdispersed/Poisson distribution.

Model formulation

- Linear dynamical systems with generalized count observation

$$\mathbf{z}_{r1} \sim \mathcal{N}(\mu_1, Q_1)$$

$$\mathbf{z}_{r(t+1)} | \mathbf{z}_{rt} \sim \mathcal{N}(\mathbf{A}\mathbf{z}_{rt}, Q)$$

$$x_{rti} \sim \mathcal{GC}(\mathbf{c}_i^T \mathbf{z}_{rt}, g_i(\cdot)), i = 1, \dots, n$$

- Practical considerations
 - Set $g_i(k) = -\infty$ for $k > K$ to facilitate computation;
 - Ridge penalty on the 2nd difference of $g_i(\cdot)$ to avoid overfitting;

Variational Bayes Expectation Maximization (VBEM)

- \mathbf{x} : data, \mathbf{z} : latent variables, θ : model parameters,
- Often hard to compute $p_\theta(\mathbf{x}) = \int p_\theta(\mathbf{x}, \mathbf{z}) d\mathbf{z}$ and $p_\theta(\mathbf{z}|\mathbf{x})$.
- Approximate the posterior by a **tractable** distribution family.

$$p_\theta(\mathbf{z}|\mathbf{x}) \approx q(\mathbf{z}) \in \mathcal{Q}$$

- Optimize a **lower bound of log likelihood**, or ELBO

$$\begin{aligned}\text{ELBO}(\theta, q) &= \int [\log p_\theta(\mathbf{x}, \mathbf{z}) - \log q(\mathbf{z})] q(\mathbf{z}) d\mathbf{z} \\ &= \log p_\theta(\mathbf{x}) - \text{KL}(q(\mathbf{z}) || p_\theta(\mathbf{z}|\mathbf{x})) \leq \log p_\theta(\mathbf{x})\end{aligned}$$

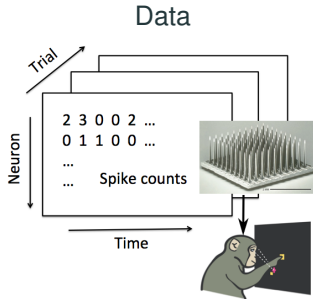
Variational Bayes Expectation Maximization (VBEM)

- VBEM: Optimize $\text{ELBO}(\theta, q) \leq \log p_{\theta}(\mathbf{x})$ iteratively
 - E-step: For a fixed θ , optimize q
 - M-step: For a fixed q , optimize θ
- VBEM for GCLDS
 - We set q to be multivariate Gaussian
 - We derive a looser but tractable ELBO
 - E-step: fast Laplace approximation initialization + dual optimization
 - M-step: convex optimization + analytical solution

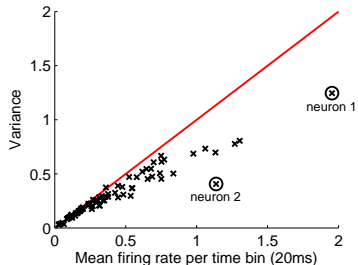
- For both simulated and real dataset, we compare GCLDS with PLDS (Poisson observation model)

	Mean	Variance	Likelihood
PLDS	✓	✗	✗
GCLDS	✓	✓	✓

Real data analysis: data



Variance and mean of spike counts

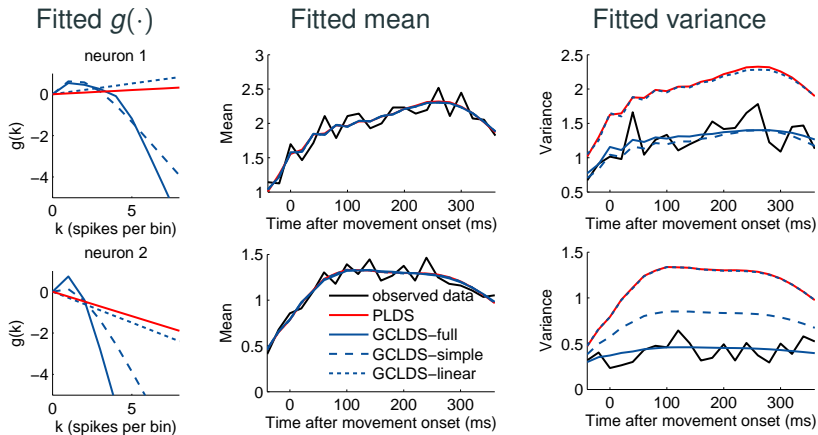


- Center-out reaching experiments
- Multi-electrode array recording
- Strong under-dispersion

Real data analysis: algorithms

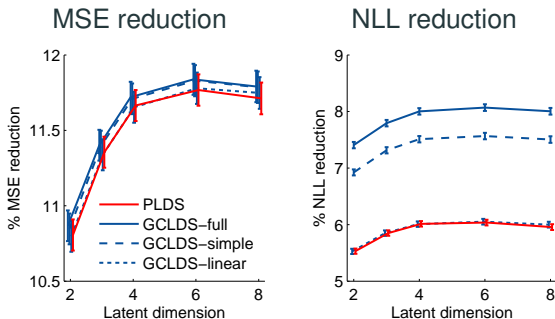
- Main algorithms to be compared
 - **PLDS**: Poisson observation
 - **GCLDS-full**: Generalized count observation, individual $g(\cdot)$ across neurons
- Two control cases for GCLDS
 - **GCLDS-linear**: truncated linear $g(\cdot)$ (truncated Poisson)
 - **GCLDS-simple**: $g(\cdot)$ shared across neurons (up to a linear function)

Real data analysis: single neuron fit



Real data analysis: population fit

- Leave-one-neuron-out prediction



Conclusion and discussion

- Summary
 - Incorporated generalized count family into state space models.
 - Developed VBEM algorithm.
 - Observed superior fitted result on real neural data.
- Extensions
 - $g(\cdot)$ vary across time?
 - Share information of $g(\cdot)$ across neurons? (hierarchical model?)
 - Generative models for under-dispersion?
- Gao Y, Buesing L, Shenoy KV, Cunningham JP (2015) High-dimensional neural spike train analysis with generalized count linear dynamical systems. NIPS 2015.

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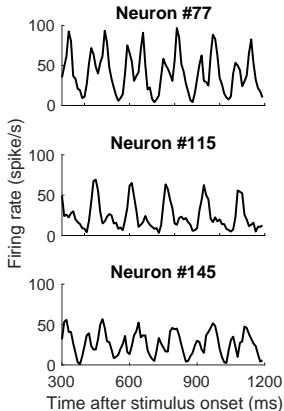
- Generalized count linear dynamical system
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Motivation

- Neural activities lie in a low-dimensional **nonlinear manifold** rather than a **linear subspace**
- Flexible observation model makes the state space model more expressive



Model formulation: fLDS

- Linear dynamical systems with **nonlinear link** and count observation

$$\mathbf{z}_{r1} \sim \mathcal{N}(\mu_1, Q_1)$$

$$\mathbf{z}_{r(t+1)} | \mathbf{z}_{rt} \sim \mathcal{N}(\mathbf{A}\mathbf{z}_{rt}, Q)$$

$$x_{rti} \sim \text{Poisson}(\mathbf{f}_i(\mathbf{z}_{rt})) \text{ (PfLDS)}$$

$$\text{or } \mathcal{GC}(\mathbf{f}_i(\mathbf{z}_{rt}), g_i(\cdot)) \text{ (GCfLDS)}$$

where f_i is a nonlinear function parameterized by a neural network

- Linear latent dynamics: simple, tractable, interpretable
- Nonlinear observation: flexible

Inference algorithm: AEVB (high level idea)

- Auto-encoding Variational Bayes (AEVB)
- Learn a mapping (recognition model) from data to the **approximate posterior distribution of latent variable**.
- Jointly optimize the generative model parameters and recognition model parameters.
- Naturally incorporate stochastic optimization to handle large datasets.
- Tractable for a large class of graphical models

Inference algorithm: AEVB (algorithm)

- Decompose ELBO by trials

$$\text{ELBO}(\theta, q) = \sum_{r=1}^R \int \left[\log \frac{p_{\theta}(\mathbf{x}_r, \mathbf{z}_r)}{q(\mathbf{z}_r)} \right] q(\mathbf{z}_r) d\mathbf{z}_r$$

- Map data \mathbf{x}_r to $q(\mathbf{z}_r)$ by a parameterized function

$$q(\mathbf{z}_r) = q_{\phi}(\mathbf{z}_r | \mathbf{x}_r) = \mathcal{N}(\mu_{\phi}(\mathbf{x}_r), \Sigma_{\phi}(\mathbf{x}_r))$$

- Learn both θ and ϕ by **stochastic** gradient descent on ELBO

$$\begin{aligned} \nabla \text{ELBO}(\theta, q_{\phi}) &\approx R \times \nabla \int \left[\log \frac{p_{\theta}(\mathbf{x}_r, \mathbf{z}_r)}{q_{\phi}(\mathbf{z}_r | \mathbf{x}_r)} \right] q_{\phi}(\mathbf{z}_r | \mathbf{x}_r) d\mathbf{z}_r \\ &\approx R \times \text{an unbiased estimator of gradient} \end{aligned}$$

(Reparameterization trick)

Structure of the recognition model

- Generative model:

$$p_{\theta}(\mathbf{z}_r|\mathbf{x}_r) \propto p_{\theta}(\mathbf{z}_{r1}) \underbrace{\prod_{t=1}^{T-1} p_{\theta}(\mathbf{z}_{t(t+1)}|\mathbf{z}_{rt})}_{\text{Gaussian}} \prod_{t=1}^T \underbrace{p_{\theta}(\mathbf{x}_{rt}|\mathbf{z}_{rt})}_{\text{Complicated}}$$

- Recognition model, product-of-Gaussian form:

$$q_{\phi}(\mathbf{z}_r|\mathbf{x}_r) \propto q_{\phi}(\mathbf{z}_{r1}) \underbrace{\prod_{t=1}^{T-1} q_{\phi}(\mathbf{z}_{r(t+1)} - \tilde{\mathbf{A}}\mathbf{z}_{rt})}_{\text{Gaussian}} \prod_{t=1}^T \underbrace{q_{\phi}(\mathbf{z}_{rt}|\mathbf{x}_{rt})}_{\text{Gaussian}}$$

Approximates a complicated factor with a Gaussian factor dependent on the data in a complicated way.

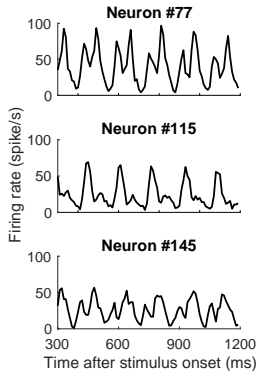
- Jointly Gaussian distribution with block tri-diagonal precision matrix. Maintaining the Markovian structure.

Experiments

	Mean	Variance	Likelihood	Concise representation
PLDS	✓	✗	✗	✗
GCLDS	✓	✓	✓	✗
PfLDS	✓	✗	✗	✓
GCfLDS	✓	✓	✓	✓

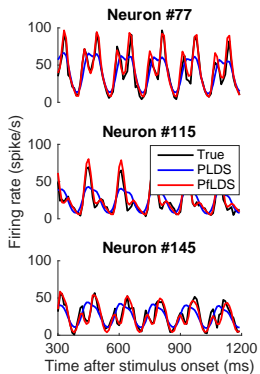
Real data analysis: primate visual cortex

Firing rate

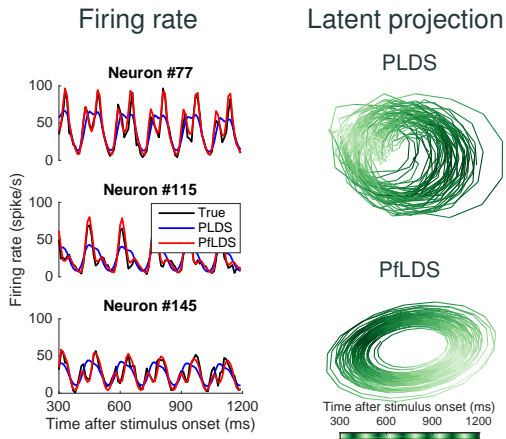


Real data analysis: primate visual cortex

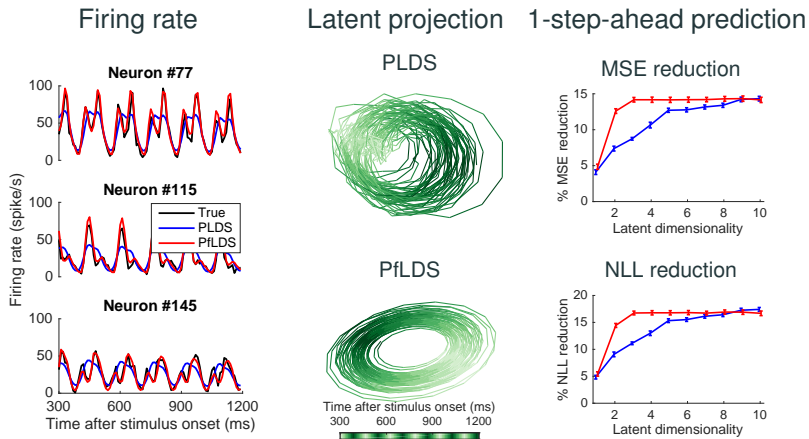
Firing rate



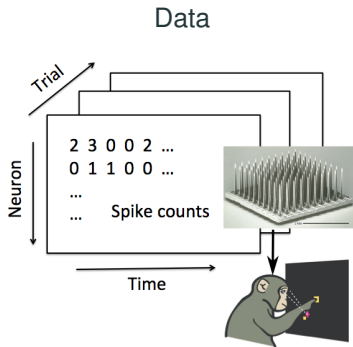
Real data analysis: primate visual cortex



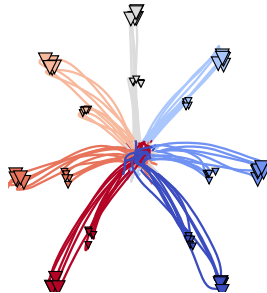
Real data analysis: primate visual cortex



Real data analysis: Primate motor cortex



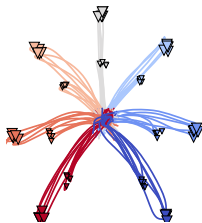
Reaching trajectory



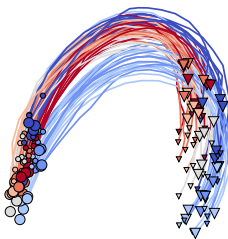
Real data analysis: Primate motor cortex

- Latent projection with 2 latent dimensions

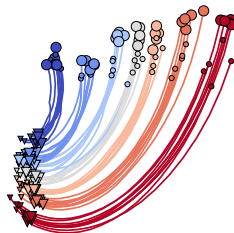
Reaching trajectory



PLDS

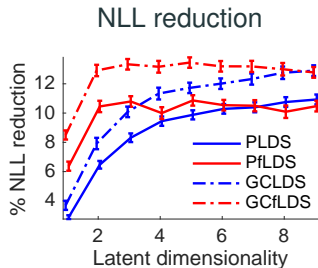
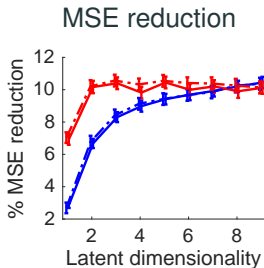


PfLDS



Real data analysis: Primate motor cortex

- One-step-ahead predictive performance



Conclusion and discussion

- Summary
 - Incorporated nonlinear observation into state space models.
 - Developed AEVB algorithm (flexible and scalable).
 - Obtain concise latent representation.
- Future work
 - Better stochastic optimization scheme
 - Interpretable nonlinearity
 - Application on more complex datasets
- Gao Y*, Archer E*, Paninski L, Cunningham JP (2016) Linear dynamical neural population models through nonlinear embeddings. NIPS 2016. (* = equal contribution)

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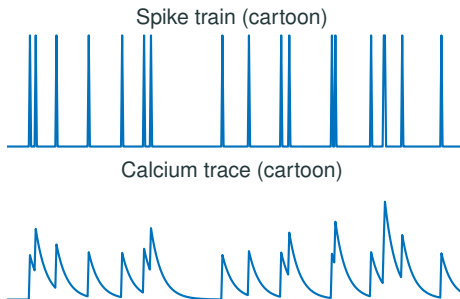
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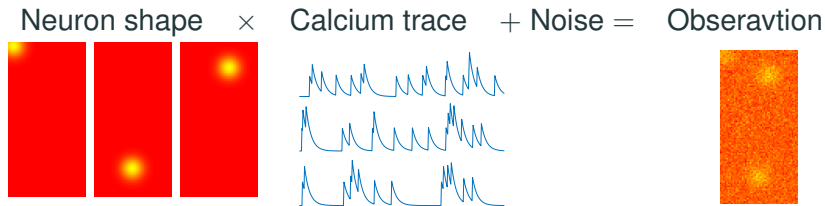
Introduction: calcium imaging data

- Basic principle: the **spiking** activity of a neuron induce a transient increase in **calcium concentration**, which can be indirectly observed by recording the **fluorescent properties** of certain calcium indicators.



Introduction: calcium imaging data

- Calcium imaging enables **simultaneous** recording of **many** neurons.



- Goal:** recover the neuron shape and calcium trace given the observation.

Model formulation

- $X \in \mathbb{R}^{N \times T}$ represents the calcium imaging data, where each column is a (vectorized) frame that contains N pixels
- Decompose X into a product of K **spatial component** and **temporal component**

$$X = D \cdot A^T + \text{noise}$$

- $D = [D_1, \dots, D_K] \in \mathbb{R}^{N \times K}$ represents the neuron shapes
 - $A = [A_1, \dots, A_K] \in \mathbb{R}^{T \times K}$ is the neural activities
- Further exploit structure of the components (localized neuron shapes)

Model formulation: objective

- Structured matrix factorization

$$\begin{aligned} & \underset{D, A}{\text{minimize}} && \|X - DA^T\|_2^2 + f_D(D), \\ & \text{subject to} && D_k \in \mathcal{D}_w^+; k = 1, \dots, K, \\ & && \|A_k\|_2 \leq c_k, \end{aligned}$$

- \mathcal{D}_w^+ : non-negative vectors whose nonzero values is within a $w \times w$ window
- $f_D(D)$ regularizes the neuron shape (will discuss later)
- $\|A_k\|_2 \leq c_k$ avoids degenerate solution

Greedy algorithm

- Identify ROIs one at a time, using the residual un-explained by existing signals.
- At iteration k , given the current residue (unexplained by existing ROIs)
 - **Greedy identification**: Identify the location p_k where the Gaussian kernel explains most of the data (across time)
 - **Shape fine tuning**: Locally optimize the spatial and temporal component
 - **Residue update**: Subtract the newly identified signal

Shape fine tuning

- Given current residue R , an identified center pixel p_k , denote S_k as a $w \times w$ window centered at p_k

$$\underset{D_k, A_k}{\text{minimize}} \quad \|R - D_k A_k^T\|^2 + f(D_k),$$

$$\text{subject to} \quad D_{kp} \geq 0, p \in S_k,$$

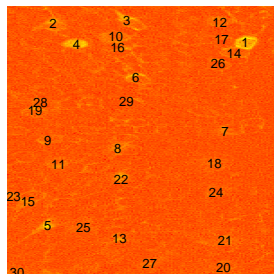
$$D_{kp} = 0, p \notin S_k,$$

$$\|A_k\|_2 \leq c_k,$$

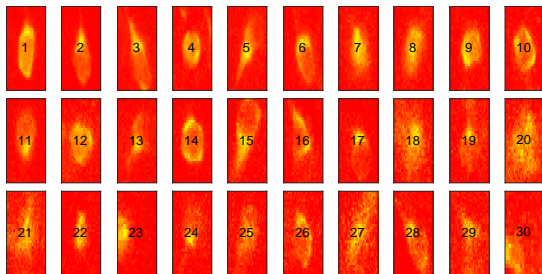
- $f(D_k) = \sum_{i=1}^3 \lambda_i f_i(D_k)$
 - $f_1(D_k) = \sum_p \tau(p, p_k) |D_{kp}|$ encourages sparsity
 - $f_2(D_k) = \sum_p (D_{kp} - G_{p_k})^2$ encourage Gaussian shape
 - $f_3(D_k) = \sum_{p_1 \text{ and } p_2 \text{ are neighbors}} (D_{kp_1} - D_{kp_2})^2$ encourages smoothness
- Optimize by block coordinate descent

Real data analysis: sample patch, no shape regularization

Mean image with fitted ROI locations

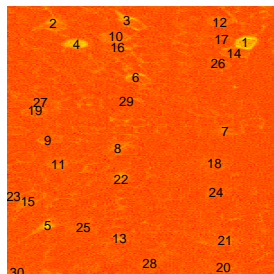


Fitted ROI shape

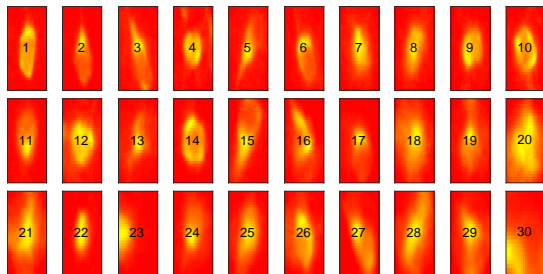


Real data analysis: sample patch, shape regularization

Mean image with fitted ROI locations



Fitted ROI shape



Conclusion and discussion

- Summary
 - Formulating calcium imaging ROI detection as a structured matrix factorization problem
 - Fast greedy algorithm
- Future work
 - More spatial and temporal structure
 - Overlapping neuron
 - Online ROI detection
 - Motion correction, background elimination
- Pnevmatikakis EA, Soudry D, Gao Y, Machado TA, Merel J, Pfau D, Reardon T, Mu Y, Lacefield C, Yang W, Ahrens M, Bruno R, Jessell TM, Peterka DS, Yuste R, Paninski L (2016) Simultaneous denoising, deconvolution, and demixing of calcium imaging data. *Neuron*, 89(2), 285-299.

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Maximum entropy principle

- **Entropy**: for a continuous distribution with density $p(\mathbf{z})$ where $\mathbf{z} \in \mathbb{R}^d$, the entropy is defined as

$$H(p) = - \int p(\mathbf{z}) \log p(\mathbf{z}) d\mathbf{z} = E_{Z \sim p} [-\log p(Z)] .$$

A popular measure of diversity or information content.

- **Maximum entropy (ME) principle**: Subject to some given prior knowledge (moment or support constraints), the distribution that makes **minimal additional assumptions** is that which has the **largest entropy** of any distribution obeying those constraints

Maximum entropy problem

- Maximum entropy (ME) problem

$$\begin{aligned} p^* &= \text{maximize } H(p) \\ &\text{subject to } E_{\mathbf{Z} \sim p}[T(\mathbf{Z})] = 0 \\ &\quad \text{supp}(p) = \mathcal{Z}, \end{aligned}$$

where $T(\mathbf{z}) = (T_1(\mathbf{z}), \dots, T_m(\mathbf{z})) : \mathcal{Z} \rightarrow \mathbb{R}^m$ is the vector of known statistics, and \mathcal{Z} is the given support.

Applications of maximum entropy

- **Texture modeling**: generate an image with a certain texture by specifying expected value of features relevant to texture.
- **Neuroscience**: generate a distribution of neural activity by specifying a set of features (pairwise correlation etc.) for hypothesis testing.
- **Ecology**: Fit species distribution with certain feature constraints (altitude, temperature etc.).
- **Finance**: Fit the risk-neutral distribution of an asset given a list of option prices.
- ...

Gibbs distribution

- Under standard regularity conditions, the maximum entropy problem can be solved by Lagrange multipliers, yielding an exponential family p^* of the form (Gibbs distribution):

$$p^*(\mathbf{z}) \propto e^{\langle \eta, T(\mathbf{z}) \rangle} \mathbb{1}(\mathbf{z} \in \mathcal{Z})$$

- Identifying $\eta \in \mathbb{R}^m$ can be hard in high-dimensional setting.
- Sampling from the distribution can be hard. MCMC methods can take long to mix.
- **Question:** is there a better way to do this?

Idea: normalizing flow

- Considering a family of smooth and invertible transformation (**normalizing flow**)

$$\mathcal{F} = \{f_\phi : \mathbb{R}^d \rightarrow \mathbb{R}^d, \phi \in \mathbb{R}^q\}$$

- Identify a transformation $f_{\phi^*} \in \mathcal{F}$ that transforms a simple distribution p_0 to approximate the maximum entropy distribution.

$$\begin{aligned} \phi^* &= \text{maximize } H(p_\phi) \\ &\text{subject to } E_{\mathbf{Z}_0 \sim p_0}[T(f_\phi(\mathbf{Z}_0))] = 0 \\ &\quad \text{supp}(p_\phi) = \mathcal{Z}. \end{aligned}$$

where $p_\phi(\mathbf{z})$ is the distribution of $f_\phi(Z_0)$ for $Z_0 \sim p_0$.

$$p_\phi(\mathbf{z}) = p_0(f_\phi^{-1}(\mathbf{z})) |\det(J_\phi(\mathbf{z}))|^{-1}$$

Augmented Lagrangian method

- Denote $R(\phi) = E(T(f_\phi(\mathbf{Z}_0))) \in \mathbb{R}^m$, augmented Lagrangian method minimizes the objective

$$L(\phi; \lambda, c) = -H(p_\phi) + \lambda^\top R(\phi) + \frac{c}{2} \|R(\phi)\|^2$$

for a sequence of $\lambda \in \mathbb{R}^m$ and $c \geq 0$.

- Update rule: at iteration k , given λ_k and c_k , suppose ϕ_k optimizes $L(\phi; \lambda_k, c_k)$, update λ and c by

$$\lambda_{k+1} = \lambda_k + c_k R(\phi_k)$$

$$c_{k+1} = \begin{cases} \beta c_k & \|R(\phi_k)\| > \gamma \|R(\phi_{k-1})\| \\ c_k & \text{otherwise} \end{cases}$$

for some $\gamma \in (0, 1)$, $\beta > 1$

Augmented Lagrangian method in stochastic setting

- Denote $R(\phi) = E(T(f_\phi(\mathbf{Z}_0))) \in \mathbb{R}^m$, augmented Lagrangian method minimizes the objective

$$L(\phi; \lambda, c) = -H(p_\phi) + \lambda^\top R(\phi) + \frac{c}{2} \|R(\phi)\|^2$$

for a sequence of $\lambda \in \mathbb{R}^m$ and $c \geq 0$.

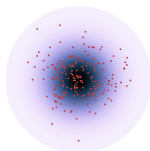
- Here $R(\phi) = E(T(f_\phi(\mathbf{Z}_0))) \in \mathbb{R}^m$ is intractable, but we can approximate with a sampled version.

$$R(\phi) \approx \frac{1}{n} \sum_{i=1}^n T(f_\phi(\mathbf{z}^{(i)})), \mathbf{z}^{(i)} \sim p_0$$

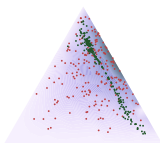
We can then optimize the objective by stochastic gradient descent.

Experiment: Dirichlet

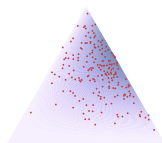
- Dirichlet distribution is the ME distribution on a simplex $\mathcal{S} = \{\mathbf{z} = (z_1, \dots, z_{d-1}) : z_i \geq 0 \text{ and } \sum_{k=1}^{d-1} z_k \leq 1\}$ with expectation on the log of each coordinate $E[\log Z_k] = \kappa_k (k = 1, \dots, d)$, where $Z_d = 1 - \sum_{k=1}^{d-1} Z_k$.
- 10 layers of planar flow (Rezende and Mohamed 2015).



Initial distribution p_0



MEFN result p_{ϕ^*} (Control case: moment matching)



Ground truth p^*

Experiment: Texture modeling

- Sample from the space of images $\mathbf{z} \in [0, 1]^{224 \times 224 \times 3}$ where $T(\mathbf{z}) : [0, 1]^{224 \times 224 \times 3} \rightarrow \mathbb{R}$ is a complicated “texture loss” defined in Ulyanov et al. (2016).
- Ulyanov et al. (2016) proposes texture net, which solves the problem by

$$\min E_{\mathbf{Z} \sim p_0} [T(f_\phi(\mathbf{Z}))]$$

without considering entropy term.

- We compare our formulation with texture net. We use real-nvp (Dinh, Sohl-Dickstein, and Bengio 2016) as the normalizing flow structure

Experiment: Texture modeling

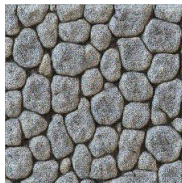
Input



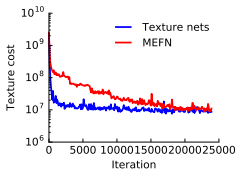
Texture net
(Ulyanov et al. 2016)



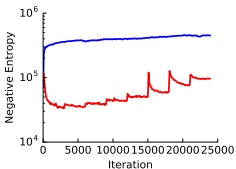
MEFN (ours)



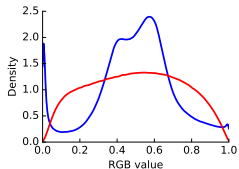
Texture cost



Negative entropy



RGB histogram



Experiment: Texture modeling

- We provide two numerical measure for assessing sample diversity given a set of images $\{\mathbf{z}^{(1)}, \dots, \mathbf{z}^{(n)}\}$.

Method	d_{L^2}	SST	SSW	SSB
Texture net	11534	128680	109577	19103
MEFN	17014	175604	161639	13964

- Mean Euclidean distance: $d_{L^2} = \text{mean}_{i \neq j} \|\mathbf{z}^{(i)} - \mathbf{z}^{(j)}\|_2^2$
- ANOVA: $\text{SST} = \text{SSW} + \text{SSB}$
 - Total: $\text{SST} = \sum_{i,k} (z_k^{(i)} - \bar{z})^2$
 - Within group: $\text{SSW} = \sum_{i,k} (z_k^{(i)} - \bar{z}_k)^2$ (larger \Rightarrow better)
 - Between group: $\text{SSB} = \sum_k n(\bar{z}_k - \bar{z})^2$ (smaller \Rightarrow better)

Conclusion and discussion

- Summary
 - Bridging information theory and deep learning
 - Solve maximum entropy problem by optimizing a normalizing flow
 - Combining augmented Lagrangian optimization with stochastic optimization
 - Promising result on simulation and real data
- Future work
 - Better normalizing flow structure
 - Better constrained stochastic optimization algorithm
- Loaiza G*, Gao Y*, Cunningham JP (2017) Maximum entropy flow networks. ICLR 2017. (*=equal contribution)

Table of Contents

I. Neural Population Data Analysis with Latent Variable Models

- Generalized count linear dynamical system
- Linear dynamical neural population models through nonlinear embeddings

II. Region of Interest Detection for Calcium Imaging Data

III. Maximum Entropy Flow Networks

Backup slides

GCLDS: Supervised case, GCGLM

- Given count data $x_i \in \mathbb{N}$, and associated covariates $z_i \in \mathbb{R}^p$, build Generalized Count Generalized Linear Model (GCGLM).

$$x_i \sim \mathcal{GC}(\theta(z_i), g(\cdot)), \quad \text{where } \theta(z_i) = z_i \beta.$$

- Reminder: generalized count distribution

$$p_{gc}(x; \theta, g(\cdot)) \propto \frac{\exp(\theta \cdot x + g(x))}{x!}, \quad x \in \mathbb{N}$$

GCLDS: special cases of GCGLM

Model Name	Typical Parameterization	GCGLM Parametrization
Logistic regression	$P(x = k) = \frac{\exp(k(\alpha + z\beta))}{1 + \exp(\alpha + x\beta)}$	$g(k) = \alpha k; k = 0, 1$
Poisson regression	$P(x = k) = \frac{\lambda^k}{k!} \exp(-\lambda);$ $\lambda = \exp(\alpha + z\beta)$	$g(k) = \alpha k$
Adjacent category regression	$\frac{P(x = k + 1)}{P(x = k)} = \exp(\alpha_k + z\beta)$	$g(k) = \sum_{i=1}^k (\alpha_{i-1} + \log i);$ $k = 0, 1, \dots, K$
Negative binomial regression	$P(x = k) = \frac{(k + r - 1)!}{k!(r - 1)!} (1 - p)^r p^k$ $p = \exp(\alpha + z\beta)$	$g(k) = \alpha k + \log(k + r - 1)!$
COM-Poisson regression	$P(x = k) = \frac{\lambda^k}{(k!)^\nu} / \sum_{j=1}^{+\infty} \frac{\lambda^j}{(j!)^\nu}$ $\lambda = \exp(\alpha + z\beta)$	$g(k) = \alpha k + (1 - \nu) \log k!$

MEFN: Normalizing flow structures

- Rezende and Mohamed (2015) Proposes two specific families of transformations for variational inference
- Planar flow

$$f_i(\mathbf{z}) = \mathbf{z} + \mathbf{u}_i h(\mathbf{w}_i^T \mathbf{z} + b_i),$$

where $b_i \in \mathbb{R}$, $\mathbf{u}_i, \mathbf{w}_i \in \mathbb{R}^d$ and h is an activation function.

- Circular flow

$$f_i(\mathbf{z}) = \mathbf{z} + \beta_i h(\alpha_i, r_i)(\mathbf{z} - \mathbf{z}'_i),$$

$\beta_i \in \mathbb{R}$, $\alpha_i > 0$, $\mathbf{z}'_i \in \mathbb{R}^d$, $h(\alpha, r) = 1/(\alpha + r)$ and $r_i = \|\mathbf{z} - \mathbf{z}'_i\|$.

MEFN: real-nvp normalizing flow structure

- Used in Dinh, Sohl-Dickstein, and Bengio (2016) for image density estimation.
- Affine coupling layer: split variable $\mathbf{z} \in \mathbb{R}^D$ into $\mathbf{z}_1 \in \mathbb{R}^d$ and $\mathbf{z}_2 \in \mathbb{R}^{D-d}$. linearly transform \mathbf{z}_2 given \mathbf{z}_1 .

$$f \begin{pmatrix} \mathbf{z}_1 \\ \mathbf{z}_2 \end{pmatrix} = \begin{pmatrix} \mathbf{z}_1 \\ \mathbf{z}_2 \odot \exp(s(\mathbf{z}_1)) + t(\mathbf{z}_1) \end{pmatrix}$$

where \odot is element-wise product.

$s(\mathbf{z}_1), t(\mathbf{z}_1) : \mathbb{R}^d \rightarrow \mathbb{R}^{D-d}$.

- The transformation family is flexible because
 - s and t can be complex while maintaining tractability (inversion and Jacobian computation).
 - Partitioning \mathbf{z} into $(\mathbf{z}_1, \mathbf{z}_2)$ can be chosen arbitrarily.