# Dense Message Passing for Sparse Principal Component Analysis

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# Outline

#### Introduction

Motivating Application: Gene Regulation

#### 2 Dense Message Passing

- Model Description
- Algorithm Description
- Statistical Mechanics Theory

#### 3 Results

- Simulated Data
- Gene Expression Data
- Marginal Likelihood Estimation

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#### Motivating Application: Gene Regulation

#### Motivation

- Gene regulation inference of explanatory factors.
- Microarray data 'Large *p* small *N*' regime.
- Explanatory factors have truly sparse loadings.
- Zero-norm priors allocate probability mass to truly sparse solutions.
- Easy to encode prior knowledge of sparse structure.
- But, zero-norm priors are problematic for inference.

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#### Model - Probabilistic PCA

• For the  $n^{th}$  data point,  $y_n$ , we assume:

 $\boldsymbol{y}_n = \boldsymbol{w} \boldsymbol{x}_n + \boldsymbol{\epsilon}_n \; ,$ 

where  $x_n \sim \mathcal{N}(0, 1)$ .

- To simplify the description,  $\epsilon_n \sim \mathcal{N}(\mathbf{0}, \mathbf{I})$ .
- Integrate out *x*:

$$P(\boldsymbol{y}_n | \boldsymbol{w}) = \mathcal{N}\left(\boldsymbol{y}_n | \boldsymbol{0}, \boldsymbol{I} + \boldsymbol{w} \boldsymbol{w}^{\mathsf{T}}\right)$$

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#### **Sparsity Mixture Prior**

We use a spike and slab mixture prior:

$$P(\mathbf{w}|C,\lambda) = \prod_{j=1}^{p} \left[ (1-C)\delta(w_j) + C\mathcal{N}(w_j|0,(\lambda)^{-1}) \right]$$



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#### **Sparsity Mixture Prior**

Express in factorised form using binary variables  $z_i \in \{0, 1\}$ :

$$P(\mathbf{v}, \mathbf{z}) = \prod_{j=1}^{p} \{ (1 - C) \mathcal{N}(v_j | 0, 1) \}^{1-z_j} \{ C \mathcal{N}(v_j | 0, \lambda^{-1}) \}^{z_j}$$

where  $w_j = z_j v_j$  and  $v_j \in \mathcal{R}$ 

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Form of the Prior in High Dimensions

- $z_j \sim \text{Bernoulli}(C)$
- $\sum_j z_j \sim \operatorname{Bin}(p, C)$
- For large dimension, *p*, the fraction of non-zero parameters is highly peaked at *C*.

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Form of the Prior in High Dimensions

• 
$$P(w_j|z_j = 1) = \mathcal{N}(w_j|0, \lambda^{-1})$$

• For large dimension, *p*:

$$\|\boldsymbol{w}\|^2 \sim \mathcal{N}(\boldsymbol{0}, \boldsymbol{C}\boldsymbol{p}/\lambda)$$

For large dimension, *p*, this distribution is approximately spherical with radius √Cp/λ.

Conclusion - A constraint-based prior:

$$p(\boldsymbol{w}, \boldsymbol{z} | \boldsymbol{C}, \lambda) \propto \delta \left( \sum_{j=1}^{p} z_j - p \boldsymbol{C} \right) \delta \left( \sum_{j=1}^{p} w_j^2 - \frac{p \boldsymbol{C}}{\lambda} \right)$$

is almost equivalent to the mixture prior in high dimensions.

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 This proves useful for developing the message passing algorithm.

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#### Factor Graph Representation



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#### **Belief Propagation**



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#### Factor to Variable Messages



$$\hat{\mathcal{M}}_{n \to \ell}^{t+1}(\boldsymbol{v}_{\ell}, \boldsymbol{z}_{\ell}) \propto \int \prod_{j \neq \ell} d\boldsymbol{v}_{j} \sum_{\boldsymbol{z} \setminus \boldsymbol{z}_{\ell}} f_{n}(\boldsymbol{y}_{n}, \boldsymbol{z}, \boldsymbol{v}) \prod_{j \neq \ell} \mathcal{M}_{j \to n}^{t}(\boldsymbol{v}_{j}, \boldsymbol{z}_{j})$$

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#### Variable to Factor Messages



 $\mathcal{M}_{\ell \to n}^{t}(\mathbf{v}_{\ell}, z_{\ell}) \propto \mathcal{P}(\mathbf{v}_{\ell}, z_{\ell}) \prod_{m \neq n} \hat{\mathcal{M}}_{m \to \ell}^{t}(\mathbf{v}_{\ell}, z_{\ell})$ 

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#### **Marginal Beliefs**

After *t* iterations, the approximate posterior marginal belief is:

$$p^{t}(z_{\ell}, v_{\ell} \mid \boldsymbol{Y}) = \frac{p(z_{\ell}, v_{\ell}) \prod_{m=1}^{N} \hat{\mathcal{M}}_{m \to \ell}^{t}(v_{\ell}, z_{\ell})}{\int \mathrm{d}v_{\ell} \sum_{z_{\ell}} p(z_{\ell}, v_{\ell}) \prod_{m=1}^{N} \hat{\mathcal{M}}_{m \to \ell}^{t}(v_{\ell}, z_{\ell})}$$

where  $p(z_{\ell}, v_{\ell})$  is the prior.

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#### Two Problems

Unfortunately,

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$$\hat{\mathcal{M}}_{n \to \ell}^{t+1}\left(\mathbf{v}_{\ell}, z_{\ell}\right) \propto \int \prod_{j \neq \ell} d\mathbf{v}_{j} \sum_{\mathbf{z} \setminus z_{\ell}} f_{n}\left(\mathbf{y}_{n}, \mathbf{z}, \mathbf{v}\right) \prod_{j \neq \ell} \mathcal{M}_{j \to n}^{t}\left(\mathbf{v}_{j}, z_{j}\right)$$

is hard to compute.

Belief propagation is not expected to converge for dense graphical models.

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## Solutions

Exploit the high-dimensionality:

Use a Gaussian approximation.

Impose consistency requirements:

Use the constraint-based prior to **enforce sparsity and length constraints** self-consistently at each iteration.

Uda and Kabashima - Statistical Mechanical Development of a Sparse Bayesian Classifier, *J. Phys. Soc. Japan*, 2005

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#### Gaussian Approximation (1)

Notice that likelihood factors may be written as:

$$f_n(\boldsymbol{y}_n, \boldsymbol{z}, \boldsymbol{v}) = \frac{1}{\sqrt{(2\pi)^p \left(1 + \|\boldsymbol{w}\|^2\right)}} \exp\left(-\frac{\boldsymbol{y}_n^{\mathsf{T}} \boldsymbol{y}_n}{2} + \Delta_n^2/2\right) ,$$

with  $\Delta_n$  defined by:  $\Delta_n = \frac{\sum_{j=1}^p y_j^n z_j v_j}{\sqrt{1 + \|\boldsymbol{w}\|^2}}$ 

For large dimension, p, Central Limit Theorem permits a Gaussian approximation.

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#### Gaussian Approximation (2)

For constant  $\|\boldsymbol{w}\|^2$ , we replace  $\Delta_n$  by:

$$\frac{y_{\ell}^{n} z_{\ell} v_{\ell}}{\sqrt{1 + Cp/\lambda}} + \underbrace{\frac{1}{\sqrt{1 + Cp/\lambda}} \sum_{j \neq \ell} y_{j}^{n} m_{j \rightarrow n}^{t}}_{\langle \Delta_{n \setminus \ell} \rangle_{n \setminus \ell}^{t}} u$$

where  $u \sim \mathcal{N}(0, 1)$ .

 $m_{j \to n}^{t}$  is the mean of  $z_{j}v_{j}$  under the cavity distribution with the  $n^{th}$  data point removed.

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#### Gaussian Approximation (3)

• The variance,  $V_{n\setminus\ell}^t$  is given by:

$$\frac{1}{1+C\rho/\lambda}\sum_{j,k\neq\ell}y_j^ny_k^n\left\langle \left(z_jv_j-m_{j\rightarrow n}^t\right)\left(z_kv_k-m_{k\rightarrow n}^t\right)\right\rangle_{n\setminus\ell}^t$$

For large dimension, *p*, fluctuations about the sample mean are O (<sup>1</sup>/<sub>√p</sub>): V<sup>t</sup><sub>n\ℓ</sub> is *self-averaging*.

• 
$$V^t \approx \frac{1}{(1+Cp/\lambda)} \left( Cp/\lambda - \sum_{j=1}^{p} \left( m_j^t \right)^2 \right)$$

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Consistency - Constraint-based Prior

The spike and slab prior can be written:

$$P(\mathbf{v}, \mathbf{z}) \propto \prod_{j=1}^{p} \exp\left(-\frac{1}{2}\left(1 - z_j + Gz_j\right)\mathbf{v}_j^2 + \gamma z_j\right)$$

where 
$$\gamma = \ln \left( \frac{C\sqrt{\lambda}}{1-C} \right)$$
 and  $G = \lambda$ .

Adjust G and  $\gamma$  at each iteration to satisfy the constraint-based prior on average:  $\sum_{j=1}^{p} \langle z_j \rangle^t = Cp \text{ and } \sum_{j=1}^{p} \langle z_j v_j^2 \rangle^t = Cp/\lambda$ 

**③** Note, after convergence,  $G \neq \lambda$  and  $\gamma \neq \ln \left(\frac{C\sqrt{\lambda}}{1-C}\right)$ 

Consistent with replica analysis.

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Onsistent with replica analysis.

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# Replica Analysis (1)

- Compute average of the log marginal likelihood over all possible datasets for  $p \to \infty$
- $\alpha = N/p$  is held constant (where N is the sample size).
- Works well for  $\alpha \ll 1 \text{`large } p \text{ small } N$
- Not mathematically rigorous, but a useful tool.

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# Replica Analysis (2)

- Derive expressions involving the posterior mean (PM) parameter vector, *w*<sup>PM</sup>:
  - squared length,  $||\boldsymbol{w}^{\text{PM}}||^2$
  - overlap with the true parameter vector,  $\boldsymbol{w}^{\text{PM}} \cdot \boldsymbol{w}^t$ .

$$oldsymbol{w}^t \sim \prod_{j=1}^{p} \left[ (1 - \mathcal{C}_t) \delta(oldsymbol{w}_j) + \mathcal{C}_t \mathcal{N}(oldsymbol{w}_j | \mathbf{0}, (\lambda_t)^{-1}) 
ight]$$

- Can show that the algorithm is consistent with this analysis.
- Can compare algorithm performance to theory using

$$\rho^{\mathsf{PM}} = \frac{\boldsymbol{w}^{\mathsf{PM}} \cdot \boldsymbol{w}^{t}}{||\boldsymbol{w}^{\mathsf{PM}}|| \, ||\boldsymbol{w}^{t}||} \, .$$

Simulated Data Gene Expression Data Marginal Likelihood Estimation

#### Simulated Data - DMP vs Theory

DMP

Gibbs



C - fraction of non-zero parameters;

$$N = 200$$
 samples,  $\alpha = N/p$ ;

 $ho^{\rm PM}$  cosine angle between  $\pmb{w}^{\rm PM}$  and  $\pmb{w}^t.$ Results averaged over 50 sample datasets.

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Simulated Data Gene Expression Data Marginal Likelihood Estimation

#### Simulated Data - DMP vs emPCA

DMP

#### emPCA





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0.6

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α=0.25

α=0.2

α=0.15

α=0.1

0.8

Simulated Data Gene Expression Data Marginal Likelihood Estimation

#### Simulated data - DMP vs SPCA

DMP





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Simulated Data Gene Expression Data Marginal Likelihood Estimation

#### Gene Expression Data - DMP vs emPCA and SPCA

#### Armstrong et al.

#### Ramaswamy et al.



p = 12582, N = 72

Simulated Data Gene Expression Data Marginal Likelihood Estimation

Summary

#### Marginal Likelihood Estimation - Simulated Data

DMP





- C fraction of non-zero parameters;
- $\lambda$  assumed signal precision;

True sparsity - 0.1.



N = 200 samples; dimension, p = 2000.

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 $\lambda^t$  - true signal precision.

- Novel message passing algorithm for Sparse Bayesian PCA in high dimensions
- Message updates rendered tractable using a Gaussian approximation
- Convergence achieved by imposing consistency requirements derived from statistical mechanics analysis.
- Inference of posterior marginals exhibits near optimal performance compared to theory.
- Outperforms two other recently published algorithms.
- Approximation to Marginal Likelihood also available.

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#### The Future

#### • Hyperparameter estimation using Marginal likelihood.

#### • Extension to multiple factors:

• Relatively straightforward for orthogonal factors.

(but will require efficient hyperparameter estimation).

 For non-orthogonal factors the best approach is a subject of on-going research.

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#### Explore further

#### Matlab code available from: http://www.cs.man.ac.uk/~sharpk

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