The neural autoregressive distribution estimator

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CS 598 presentation by Varun Kelkar

Outline

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 - Why is the problem difficult?
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 - Bayesian Networks
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Motivation

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Distribution estimation of high dimensional discrete/binary vectors.

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• Why is this problem important:

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• Why is this problem difficult:

Curse of dimensionality, the PMF is a vector in a d^n -dimensional space, where d is the number of discrete levels, n is the dimensionality of the vector.

Previous approaches

- Mixture of Bernoullis (MoB)
- Restricted Boltzmann Machines (RBMs)
- Bayesian Networks
- Fully visible sigmoid belief network (FVSBN)

Restricted Boltzmann Machines (RBMs)

 $\mathbf{h} = \mathbf{W}\mathbf{v} + \mathbf{b}$

Probabilities evaluated using the energy function:

$$E(\mathbf{v}, \mathbf{h}) = -\mathbf{h}^{\top} \mathbf{W} \mathbf{v} - \mathbf{b}^{\top} \mathbf{v} - \mathbf{c}^{\top} \mathbf{h}$$
(1)

probabilities are assigned to any observation \mathbf{v} as follows:

$$p(\mathbf{v}) = \sum_{\mathbf{h}} \exp(-E(\mathbf{v}, \mathbf{h}))/Z, \qquad (2)$$

Problems:

- Computing partition function *Z* is intractable for all except the small networks. Approximations needed.
- Hence, RBMs cannot be used to model parts of a probabilistic system.
- Difficulty in evaluating the learned distribution

Bayesian networks

Strategy: Decompose the distribution using its conditionals

$$p(\mathbf{v}) = \prod_{i=1}^{D} p(v_i | \mathbf{v}_{\text{parents}(i)}), \qquad (3)$$

Example: Fully visible sigmoid belief networks

$$p(v_i | \mathbf{v}_{\text{parents}(i)}) = \text{sigm}\left(b_i + \sum_{j < i} W_{ij} v_j\right), \quad (4)$$

Converting RBMs into Bayesian networks

Rewrite the RBM PDF in terms of conditionals

$$p(\mathbf{v}) = \prod_{i=1}^{D} p(v_i | \mathbf{v}_{< i})$$
$$= \prod_{i=1}^{D} p(v_i, \mathbf{v}_{< i}) / p(\mathbf{v}_{< i})$$
$$= \prod_{i=1}^{D} \frac{\sum_{\mathbf{v}_{> i}} \sum_{\mathbf{h}} \exp(-E(\mathbf{v}, \mathbf{h}))}{\sum_{\mathbf{v}_{> i}} \sum_{\mathbf{h}} \exp(-E(\mathbf{v}, \mathbf{h}))}, \qquad (5)$$

Use a simplified model for the still intractable conditionals

$$q(v_i, \mathbf{v}_{>i}, \mathbf{h} | \mathbf{v}_{
$$\prod_{j>i} \mu_j(i)^{v_j} (1 - \mu_j(i))^{1 - v_j}$$
$$\prod_k \tau_k(i)^{h_k} (1 - \tau_k(i))^{1 - h_k},$$$$

Converting RBMs into Bayesian networks

Minimize the KL divergence by setting its gradients to 0, which gives the following. Use fixed-point iterations to find the parameters of the distribution:

$$\tau_k(i) = \operatorname{sigm}\left(c_k + \sum_{j \ge i} W_{kj} \mu_j(i) + \sum_{j < i} W_{kj} v_j\right) \quad (7)$$
$$\mu_j(i) = \operatorname{sigm}\left(b_j + \sum_k W_{kj} \tau_k(i)\right) \quad \forall j \ge i \,. \tag{8}$$

Problems:

- Can be slow to converge.
- Needs to be repeated for each component v_i

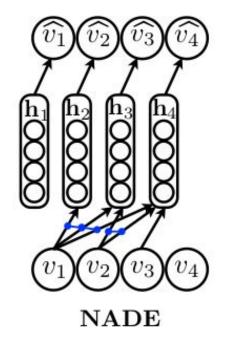
Neural autoregressive distribution estimators (NADE)

Taking inspiration from the first fixed-point iteration above, formulate the network architecture.

$$p(v_i = 1 | \mathbf{v}_{
$$\mathbf{h}_i = \operatorname{sigm} \left(\mathbf{c} + \mathbf{W}_{\cdot,$$$$

Training: Minimize the log-likelihood averaged over a training dataset

$$\frac{1}{T}\sum_{t=1}^{T} -\log p(\mathbf{v}_t) = \frac{1}{T}\sum_{t=1}^{T}\sum_{i=1}^{D} -\log p(v_i|\mathbf{v}_{< i}), \quad (11)$$



Results of experiments

Experiments comparing various baselines using the average test log-likelihood (ALL) relative to the MoB baseline

| Model | ADULT | CONNECT-4 | DNA | MUSHROOMS | NIPS-0-12 | OCR-LETTERS | RCV1 | WEB |
|---------------|------------|------------|------------|------------|------------|-------------|------------|------------|
| MoB | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 | 0.00 |
| | ± 0.10 | ± 0.04 | ± 0.53 | ± 0.10 | ± 1.12 | ± 0.32 | ± 0.11 | ± 0.23 |
| RBM | 4.18 | 0.75 | 1.29 | -0.69 | 12.65 | -2.49 | -1.29 | 0.78 |
| | ± 0.06 | ± 0.02 | ± 0.48 | ± 0.09 | ± 1.07 | ± 0.30 | ± 0.11 | ± 0.20 |
| RBM | 4.15 | -1.72 | 1.45 | -0.69 | 11.25 | 0.99 | -0.04 | 0.02 |
| mult. | ± 0.06 | ± 0.03 | ± 0.40 | ± 0.05 | ± 1.06 | ± 0.29 | ± 0.11 | ± 0.21 |
| RBForest | 4.12 | 0.59 | 1.39 | 0.04 | 12.61 | 3.78 | 0.56 | -0.15 |
| | ± 0.06 | ± 0.02 | ± 0.49 | ± 0.07 | ± 1.07 | ± 0.28 | ± 0.11 | ± 0.21 |
| FVSBN | 7.27 | 11.02 | 14.55 | 4.19 | 13.14 | 1.26 | -2.24 | 0.81 |
| | \pm 0.04 | ± 0.01 | \pm 0.50 | ± 0.05 | ± 0.98 | ± 0.23 | ± 0.11 | ± 0.20 |
| NADE | 7.25 | 11.42 | 13.38 | 4.65 | 16.94 | 13.34 | 0.93 | 1.77 |
| | \pm 0.05 | \pm 0.01 | ± 0.57 | \pm 0.04 | \pm 1.11 | \pm 0.21 | \pm 0.11 | \pm 0.20 |
| Normalization | -20.44 | -23.41 | -98.19 | -14.46 | -290.02 | -40.56 | -47.59 | -30.16 |

Generative performance for binarized images

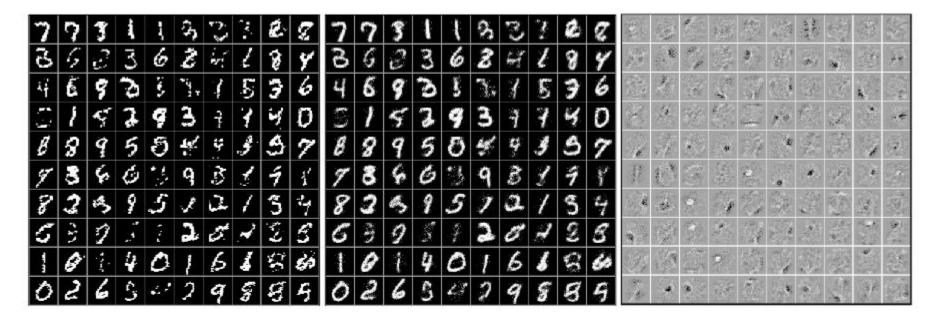


Figure 2: (Left): samples from NADE trained on a binary version of MNIST. (Middle): probabilities from which each pixel was sampled. (Right): visualization of some of the rows of W. This figure is better seen on a computer screen.

Summary

- MoBs not sufficiently expressive to model the complex dependencies in high dimensional distributions
- RBMs can have an intractable computation of the partition function, rendering it difficult to use for downstream applications
- Bayesian networks, such as fully visible sigmoid belief networks may still be less expressive than desired.
- Approaches that convert RBMs to Bayesian networks are slow to converge.
- The proposed approach outperforms other approaches because it is able to utilize the recursiveness in Bayesian network architectures to decompose a complex probability distribution to a tractable form, while still having sufficient generality in the form of the network architecture chosen.

References

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- 2. Fischer, Asja, and Christian Igel. "An introduction to restricted Boltzmann machines." *Iberoamerican congress on pattern recognition*. Springer, Berlin, Heidelberg, 2012.