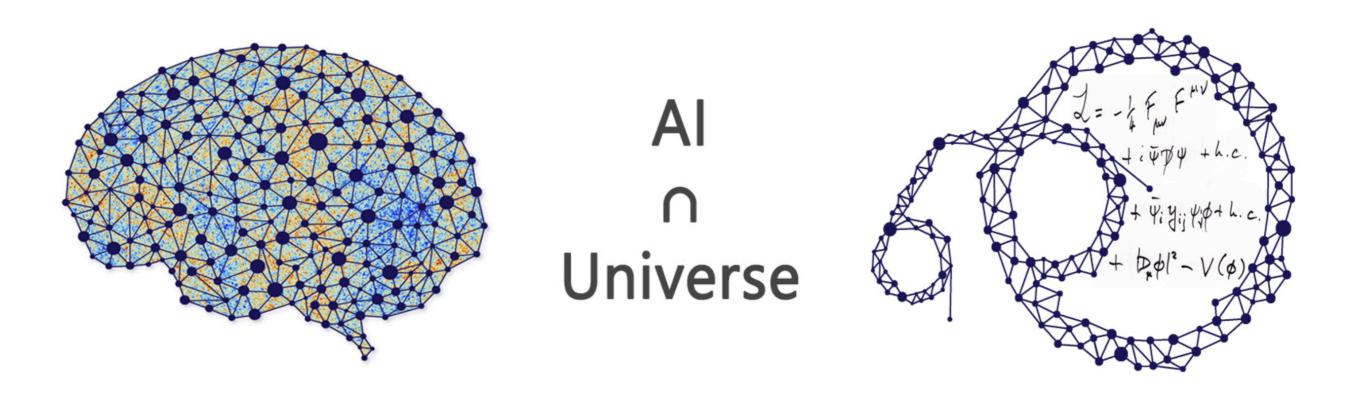
PHY 835: Machine Learning in Physics

Lecture 11: Unsupervised Learning (PCA)

February 27, 2024



Outline for today

- Unsupervised learning
- Challenges of High-dimensional data
- Principal component analysis (PCA)
- Multi-dimensional scaling (MDS)

References: 1803.08823, Deep Learning Book

Unsupervised Learning

- Discovering structure in unlabelled data.
- Two ways: 1) some appropriate numerical measure (e.g. distance in some representation space). 2) with visualizations.
- Need to dimensionally reduce data as it is impractical for datasets involving large number of measured features (e.g. images)
- We call the dimensionally reduced space latent space.
- By dimensional reduction we often loose information. This is not necessarily bad. By loosing only irrelevant information, we can find good representations.

Challenges of High-dimensional Data

- High-dimensional data lives near the edge of sample space.
- Consider data distributed uniformly at random in a D-dimensional hypercube $C = [-e/2,e/2]^D$. Probably of a data point inside a D-dimensional hypersphere S of radius e/2 centered at the origin:

$$p(\|\mathbf{x}\|_2 < e/2) \sim (1/2)^D \rightarrow 0$$
 exponentially as $D \rightarrow \infty$

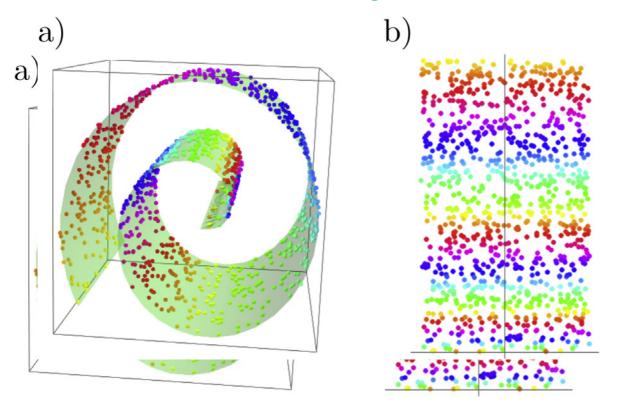
- Most of the data will concentrate outside the hypersphere, in the corners of the hypercube.
- Recall this property underlies properties of statistical systems such as the Maxwell distribution.

Challenges of High-dimensional Data

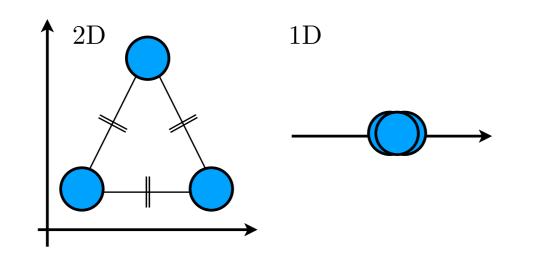
- Real-world data is usually not random or uniformly distributed (data lives in a lower-dim. space compared with original space).
- "Blessing of non-uniformity": Data will typically be locally smooth (local variation will not incur a change in the target variable).
- Examples: thermodynamics variables (temperature, pressure, etc) are not sensitive to variations of the dynamical variables (position and momentum of individual particles); small number of order parameters in statistical systems with large number of dofs.
- Objective: preserve relative pairwise distances between data points when going to latent space.

Challenges of High-dimensional Data

Intrinsic dimensionality and the crowding problem:



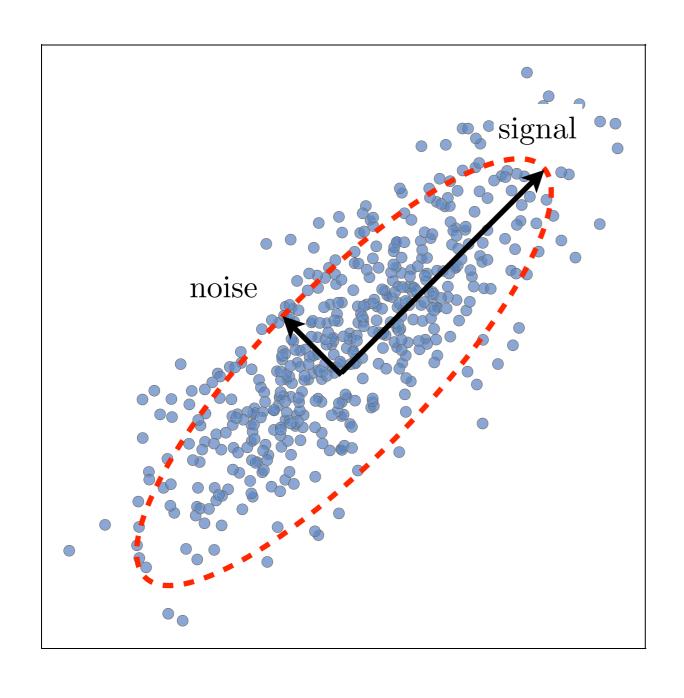
Intrinsic dim = min. # parameters to parametrize the data.

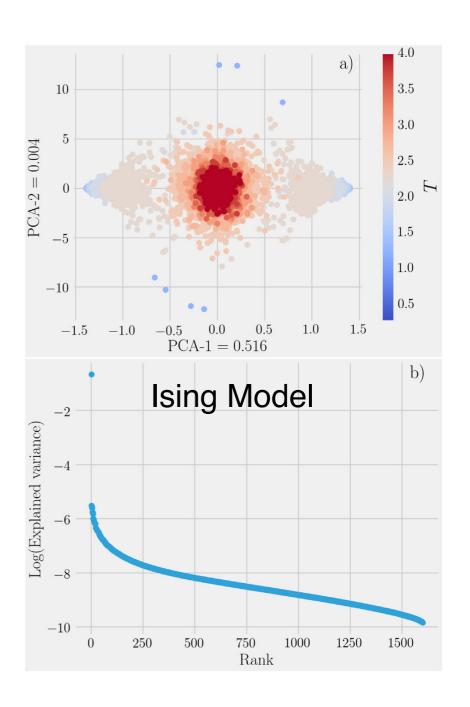


Attempts to represent data in a space with dim < intrinsic dimensionality lead to a "crowding" problem.

Principal Component Ana

 Perform an orthogonal transformation of high variance directions ⇔ minimizing the error in projection.





- Suppose we have a collection of N points $\{\mathbf{x}^{(1)},...,\mathbf{x}^{(N)}\}$ in \mathbb{R}^n .
- Compress them into code vectors $\{\mathbf{c}^{(1)},...,\mathbf{c}^{(N)}\}$ in \mathbb{R}^l with l < n
- Encoding function: $f(\mathbf{x}) = \mathbf{c}$; Decoding function: $\mathbf{x} = g(\mathbf{c})$
- Encoding + decoding: $\tilde{\mathbf{x}} = g(f(\mathbf{x}))$
- A measure of goodness for your compression is how accurate is this encoding+decoding:

$$||\mathbf{x} - \tilde{\mathbf{x}}|| \ll 1$$
 (good commpression)

- Let $g(\mathbf{c}) = \mathbf{D}\mathbf{c}$ where $\mathbf{D} \in \mathbb{R}^{n \times l}$ is a matrix defining the decoding.
- Columns of D are orthogonal to each other and have unit norm.
- Minimizing the loss: $c^* = \arg\min_{c} ||x-g(c)||_2$ or equivalently (and more conveniently): $c^* = \arg\min_{c} ||x-g(c)||_2^2$
- The function to be minimized: $(x g(c))^{\top}(x g(c))$

$$= \boldsymbol{x}^{\top} \boldsymbol{x} - \boldsymbol{x}^{\top} g(\boldsymbol{c}) - g(\boldsymbol{c})^{\top} \boldsymbol{x} + g(\boldsymbol{c})^{\top} g(\boldsymbol{c}) = \boldsymbol{x}^{\top} \boldsymbol{x} - 2 \boldsymbol{x}^{\top} g(\boldsymbol{c}) + g(\boldsymbol{c})^{\top} g(\boldsymbol{c})$$

Omit the first term which does not depend on c:

$$\boldsymbol{c}^* = \arg\min_{\boldsymbol{c}} -2\boldsymbol{x}^{\top} g(\boldsymbol{c}) + g(\boldsymbol{c})^{\top} g(\boldsymbol{c})$$

Using the definition of the decoding function:

$$egin{aligned} oldsymbol{c}^* &= rg \min_{oldsymbol{c}} -2 oldsymbol{x}^ op oldsymbol{D} oldsymbol{c} + oldsymbol{c}^ op oldsymbol{D} oldsymbol{T} oldsymbol{D} oldsymbol{c} + oldsymbol{c}^ op oldsymbol{D} oldsymbol{c} + oldsymbol{c}^ op oldsymbol{I}_l oldsymbol{c} \end{aligned}$$

because the columns of D are orthogonal and have unit norm.

The optimization problem has the solution:

$$abla_{oldsymbol{c}}
abla_{oldsymbol{c}} (-2oldsymbol{x}^{ op} oldsymbol{D} oldsymbol{c} + oldsymbol{c}^{ op} oldsymbol{c}) = oldsymbol{0}$$

$$abla_{oldsymbol{c}} -2oldsymbol{D}^{ op} oldsymbol{x} + 2oldsymbol{c} = oldsymbol{0}$$

- The encoding function: $f(x) = D^{\top}x$
- PCA reconstruction operation: $r(x) = g(f(x)) = DD^{T}x$

• Since we use the same matrix ${f D}$ to decode all the points, we minimize the Frobenius norm of the matrix of errors computed over all dimensions and all points:

$$\boldsymbol{D}^* = \operatorname*{arg\,min}_{\boldsymbol{D}} \sqrt{\sum_{i,j} \left(x_j^{(i)} - r(\boldsymbol{x}^{(i)})_j \right)^2} \text{ subject to } \boldsymbol{D}^\top \boldsymbol{D} = \boldsymbol{I}_l.$$

• Consider l = 1 (exercise: generalization to other l), then $\mathbf{D} = \mathbf{d}$

$$d^* = \underset{\boldsymbol{d}}{\operatorname{arg\,min}} \sum_i ||\boldsymbol{x}^{(i)} - \boldsymbol{d}\boldsymbol{d}^{\top}\boldsymbol{x}^{(i)}||_2^2 \text{ subject to } ||\boldsymbol{d}||_2 = 1.$$

• Some cosmetic changes (noting $\mathbf{d}^T \mathbf{x}^{(i)}$ is a scalar, and so its transpose is equal to itself) give:

$$d^* = \underset{\boldsymbol{d}}{\operatorname{arg\,min}} \sum_i ||\boldsymbol{x}^{(i)} - \boldsymbol{x}^{(i)\top} \boldsymbol{d} \boldsymbol{d}||_2^2 \text{ subject to } ||\boldsymbol{d}||_2 = 1.$$

• Introduce compact notation by defining the matrix \mathbf{X} :

$$oldsymbol{X} \in \mathbb{R}^{m imes n}$$
 $oldsymbol{X}_{i,:} = oldsymbol{x}^{(i)^ op}$

The decoding error is minimized when:

$$d^* = \underset{d}{\operatorname{arg\,min}} || \boldsymbol{X} - \boldsymbol{X} d \boldsymbol{d}^{\top} ||_F^2 \text{ subject to } \boldsymbol{d}^{\top} \boldsymbol{d} = 1.$$

The Frobenius norm part:

$$\begin{split} \arg\min_{\boldsymbol{d}} ||\boldsymbol{X} - \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^\top||_F^2 &= \arg\min_{\boldsymbol{d}} \operatorname{Tr} \left(\left(\boldsymbol{X} - \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^\top \right)^\top \left(\boldsymbol{X} - \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^\top \right) \right) \\ &= \arg\min_{\boldsymbol{d}} \operatorname{Tr} (\boldsymbol{X}^\top \boldsymbol{X} - \boldsymbol{X}^\top \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^\top - \boldsymbol{d} \boldsymbol{d}^\top \boldsymbol{X}^\top \boldsymbol{X} + \boldsymbol{d} \boldsymbol{d}^\top \boldsymbol{X}^\top \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^\top) \\ &= \arg\min_{\boldsymbol{d}} \operatorname{Tr} (\boldsymbol{X}^\top \boldsymbol{X}) - \operatorname{Tr} (\boldsymbol{X}^\top \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^\top) - \operatorname{Tr} (\boldsymbol{d} \boldsymbol{d}^\top \boldsymbol{X}^\top \boldsymbol{X}) + \operatorname{Tr} (\boldsymbol{d} \boldsymbol{d}^\top \boldsymbol{X}^\top \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^\top) \\ &\text{does not depend on } \boldsymbol{d} \end{split}$$

Cycle the order of the matrices inside a trace, the Frobenius norm:

$$= \underset{\boldsymbol{d}}{\operatorname{arg\,min}} - 2\operatorname{Tr}(\boldsymbol{X}^{\top}\boldsymbol{X}\boldsymbol{d}\boldsymbol{d}^{\top}) + \operatorname{Tr}(\boldsymbol{d}\boldsymbol{d}^{\top}\boldsymbol{X}^{\top}\boldsymbol{X}\boldsymbol{d}\boldsymbol{d}^{\top})$$

$$= \underset{\boldsymbol{d}}{\operatorname{arg\,min}} - 2\operatorname{Tr}(\boldsymbol{X}^{\top}\boldsymbol{X}\boldsymbol{d}\boldsymbol{d}^{\top}) + \operatorname{Tr}(\boldsymbol{X}^{\top}\boldsymbol{X}\boldsymbol{d}\boldsymbol{d}^{\top}\boldsymbol{d}\boldsymbol{d}^{\top})$$

• The constraint $\mathbf{d}^T \mathbf{d} = 1$ gives:

$$= \underset{\boldsymbol{d}}{\operatorname{arg\,min}} - 2\operatorname{Tr}(\boldsymbol{X}^{\top}\boldsymbol{X}\boldsymbol{d}\boldsymbol{d}^{\top}) + \operatorname{Tr}(\boldsymbol{X}^{\top}\boldsymbol{X}\boldsymbol{d}\boldsymbol{d}^{\top}) \text{ subject to } \boldsymbol{d}^{\top}\boldsymbol{d} = 1$$

Thus minimizing decoding error is the same as maximizing variance:

$$= \underset{\boldsymbol{d}}{\operatorname{arg\,min}} - \operatorname{Tr}(\boldsymbol{X}^{\top} \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^{\top}) \text{ subject to } \boldsymbol{d}^{\top} \boldsymbol{d} = 1$$

$$= \underset{\boldsymbol{d}}{\operatorname{arg\,max}} \operatorname{Tr}(\boldsymbol{X}^{\top} \boldsymbol{X} \boldsymbol{d} \boldsymbol{d}^{\top}) \text{ subject to } \boldsymbol{d}^{\top} \boldsymbol{d} = 1$$

$$= \underset{\boldsymbol{d}}{\operatorname{arg\,max}} \operatorname{Tr}(\boldsymbol{d}^{\top} \boldsymbol{X}^{\top} \boldsymbol{X} \boldsymbol{d}) \text{ subject to } \boldsymbol{d}^{\top} \boldsymbol{d} = 1.$$

PCA — Maximizing Variance

• The covariance matrix of data matrix X is defined as:

$$\Sigma(\boldsymbol{X}) = \frac{1}{N-1} \boldsymbol{X}^T \boldsymbol{X}$$

- $\Sigma(\mathbf{X})_{jj}$ corresponds to the variance of the j-th feature while $\Sigma(\mathbf{X})_{ij}$ measures the covariance (correlation) between feature i & feature j.
- Find a new basis that emphasizes highly variable directions while reducing redundancy between basis vectors. Perform SVD:

$$\Sigma(X) = \frac{1}{N-1} V S U^T U S V^T$$

$$= V \left(\frac{S^2}{N-1}\right) V^T$$

$$\equiv V \Lambda V^T.$$

• The eigenvalues λ_i of Λ are given by $\lambda_i = s_i^2/(N-1)$.

PCA — Maximizing Variance

• To reduce the dimensionality of data from n to l, construct the $n \times l$ projection matrix \mathbf{V}_l by selecting the singular components with the l largest singular values. The projection is then

$$Y = XV_1$$

- The singular vector with the largest singular value (largest variance)
 is the first principal component; the singular vector with the second
 largest variance is the second principal component, etc.
- Common in data visualization is to project on the first few principal components (as long as a large part of the variance is explained in those components, e.g., Ising Model).
- Low explained variance may imply that the intrinsic dimensionality of the data is high, or it cannot be captured by a linear representation.

Multidimensional Scaling (MDS)

- Non-linear dimensional reduction technique which preserves the pairwise distance (or dissimilarity) d_{ij} between data points.
- Metric MDS: the latent coordinates are obtained by minimizing:

$$\tilde{\mathbf{Y}} = \arg\min_{\mathbf{Y}} \sum_{i < j} w_{ij} |d_{ij}(\mathbf{X}) - d_{ij}(\mathbf{Y})|,$$

• w_{ij} specifies the level of confidence (precision) in the value of $d_{ij}(\mathbf{X})$. If Euclidean metric is used, MDS=PCA; known as classical scaling.

https://stats.stackexchange.com/questions/14002/whats-the-difference-between-principal-component-analysis-and-multidimensional

- MDS (metric or non-metric) is a generalization of PCA.
- Non-metric MDS: d_{ij} can be any distance matrix that preserves the ordination, i.e., if $d_{12}(\mathbf{X}) < d_{13}(\mathbf{X})$ then $d_{12}(\mathbf{Y}) < d_{13}(\mathbf{Y})$.

Multidimensional Scaling (MDS)

- Both MDS and PCA can be implemented using standard Python packages such as Scikit.
- MDS algorithms have a scaling of $\mathcal{O}(N^3)$ where N=# data points.
- Sample-based methods can reduce this scaling to $\mathcal{O}(N \log N)$.
- PCA has a scaling of $\mathcal{O}(Np^2 + p^3)$ for a complete decomposition.

covariance matrix

computation of eigenvalue decomposition where p = # features

• Can be improved to give a $\mathcal{O}(Np^2 + p)$ scaling for PCA if only a few principal components are desired.