14: Dimensionality Reduction (PCA)

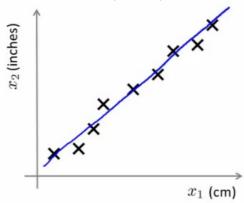
Previous Next Index

Motivation 1: Data compression

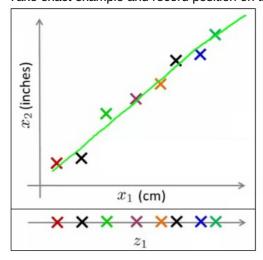
- Start talking about a second type of unsupervised learning problem dimensionality reduction
 - Why should we look at dimensionality reduction?

Compression

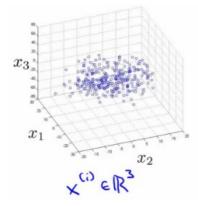
- Speeds up algorithms
- · Reduces space used by data for them
- · What is dimensionality reduction?
 - So you've collected many features maybe more than you need
 - Can you "simply" your data set in a rational and useful way?
 - Example
 - Redundant data set different units for same attribute
 - Reduce data to 1D (2D->1D)



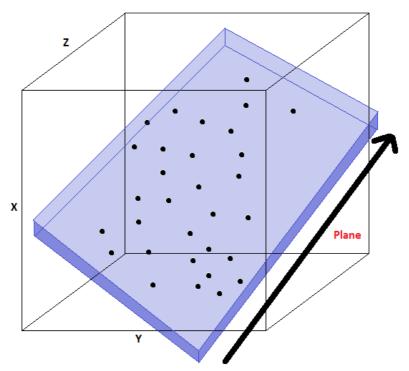
- Example above isn't a perfect straight line because of round-off error
- Data redundancy can happen when different teams are working independently
 - Often generates redundant data (especially if you don't control data collection)
- Another example
 - Helicopter flying do a survey of pilots (x1 = skill, x2 = pilot enjoyment)
 - These features may be highly correlated
 - This correlation can be combined into a single attribute called aptitude (for example)
- What does dimensionality reduction mean?
 - In our example we plot a line
 - Take exact example and record position on that line



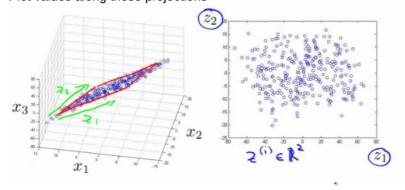
- So before x¹ was a 2D feature vector (X and Y dimensions)
- Now we can represent x¹ as a 1D number (Z dimension)
- So we approximate original examples
 - Allows us to half the amount of storage
 - Gives lossy compression, but an acceptable loss (probably)
 - The loss above comes from the rounding error in the measurement, however
- Another example 3D -> 2D
 - o So here's our data



- Maybe all the data lies in one plane
 - This is sort of hard to explain in 2D graphics, but that plane may be aligned with one of the axis
 - Or or may not...
 - Either way, the plane is a small, a constant 3D space
 - In the diagram below, imagine all our data points are sitting "inside" the blue tray (has a dark blue exterior face and a light blue inside)



- Because they're all in this relative shallow area, we can basically ignore one of the dimension, so we draw two new lines (z1 and z2) along the x and y planes of the box, and plot the locations in that box
- i.e. we loose the data in the z-dimension of our "shallow box" (NB "z-dimensions" here refers to the dimension relative to the box (i.e it's depth) and NOT the z dimension of the axis we've got drawn above) but because the box is shallow it's OK to lose this. Probably....
- Plot values along those projections



- So we've now reduced our 3D vector to a 2D vector
- In reality we'd normally try and do 1000D -> 100D

Motivation 2: Visualization

- It's hard to visualize highly dimensional data
 - o Dimensionality reduction can improve how we display information in a tractable manner for human consumption
 - Why do we care?
 - Often helps to develop algorithms if we can understand our data better
 - Dimensionality reduction helps us do this, see data in a helpful
 - Good for explaining something to someone if you can "show" it in the data
- Example;
 - Collect a large data set about many facts of a country around the world

| | | | | | | Mean | |
|-----------|---------------|--------------|------------|------------|-------------|------------|--|
| | | Per capita | -99 | | Poverty | household | |
| | GDP | GDP | Human | | Index | income | |
| | (trillions of | (thousands | Develop- | Life | (Gini as | (thousands | |
| Country | US\$) | of intl. \$) | ment Index | expectancy | percentage) | of US\$) | |
| Canada | 1.577 | 39.17 | 0.908 | 80.7 | 32.6 | 67.293 | |
| China | 5.878 | 7.54 | 0.687 | 73 | 46.9 | 10.22 | |
| India | 1.632 | 3.41 | 0.547 | 64.7 | 36.8 | 0.735 | |
| Russia | 1.48 | 19.84 | 0.755 | 65.5 | 39.9 | 0.72 | |
| Singapore | 0.223 | `56.69 | 0.866 | 80 | 42.5 | 67.1 | |
| USA | 14.527 | 46.86 | 0.91 | 78.3 | 40.8 | 84.3 | |
| | | | | | | | |

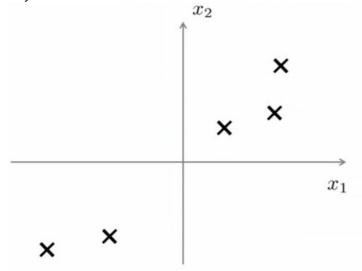
- So
 - x₁ = GDP
 - **.** ...
 - x₆ = mean household
- Say we have 50 features per country
- How can we understand this data better?
 - Very hard to plot 50 dimensional data
- Using dimensionality reduction, instead of each country being represented by a 50-dimensional feature vector
 - Come up with a different feature representation (z values) which summarize these features

| Country | z_1 | z_2 |
|-----------|-------|-------|
| Canada | 1.6 | 1.2 |
| China | 1.7 | 0.3 |
| India | 1.6 | 0.2 |
| Russia | 1.4 | 0.5 |
| Singapore | 0.5 | 1.7 |
| USA | 2 | 1.5 |
| | | |

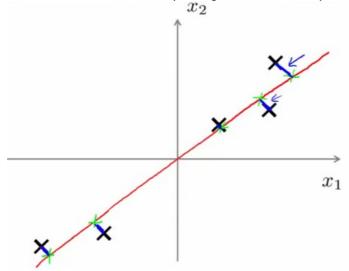
- This gives us a 2-dimensional vector
 - Reduce 50D -> 2D
 - Plot as a 2D plot
- Typically you don't generally ascribe meaning to the new features (so we have to determine what these summary values mean)
 - e.g. may find horizontal axis corresponds to overall country size/economic activity
 - and y axis may be the per-person well being/economic activity
- So despite having 50 features, there may be two "dimensions" of information, with features associated with each
 of those dimensions
 - It's up to you to asses what of the features can be grouped to form summary features, and how best to do that (feature scaling is probably important)
- Helps show the two main dimensions of variation in a way that's easy to understand

Principle Component Analysis (PCA): Problem Formulation

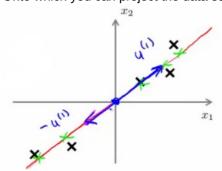
- For the problem of dimensionality reduction the most commonly used algorithm is PCA
 - Here, we'll start talking about how we formulate precisely what we want PCA to do
- So
 - Say we have a 2D data set which we wish to reduce to 1D



- o In other words, find a single line onto which to project this data
 - How do we determine this line?
 - The distance between each point and the projected version should be small (blue lines below are short)
 - PCA tries to find a lower dimensional surface so the sum of squares onto that surface is minimized
 - The blue lines are sometimes called the projection error
 - PCA tries to find the surface (a straight line in this case) which has the minimum projection error

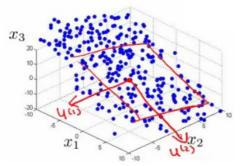


- As an aside, you should normally do mean normalization and feature scaling on your data before PCA
- A more formal description is
 - For 2D-1D, we must find a vector u⁽¹⁾, which is of some dimensionality
 - Onto which you can project the data so as to minimize the projection error



- \circ u⁽¹⁾ can be positive or negative (-u⁽¹⁾) which makes no difference
 - Each of the vectors define the same red line

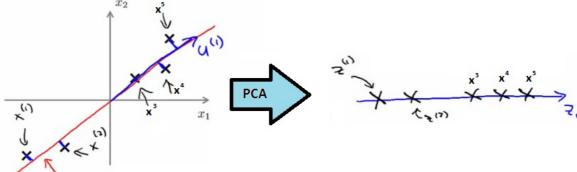
- In the more general case
 - To reduce from nD to kD we
 - Find k vectors $(u^{(1)}, u^{(2)}, \dots u^{(k)})$ onto which to project the data to minimize the projection error
 - So lots of vectors onto which we project the data
 - Find a set of vectors which we project the data onto the linear subspace spanned by that set of vectors
 - We can define a point in a plane with k vectors
 - o e.g. 3D->2D
 - Find pair of vectors which define a 2D plane (surface) onto which you're going to project your data
 - Much like the "shallow box" example in compression, we're trying to create the shallowest box possible (by defining two of it's three dimensions, so the box' depth is minimized)



- · How does PCA relate to linear regression?
 - PCA is **not** linear regression
 - Despite cosmetic similarities, very different
 - For linear regression, fitting a straight line to minimize the straight line between a point and a squared line
 - NB **VERTICAL distance** between point
 - For PCA minimizing the magnitude of the shortest orthogonal distance
 - Gives very different effects
 - More generally
 - With linear regression we're trying to predict "y"
 - With PCA there is no "y" instead we have a list of features and all features are treated equally
 - If we have 3D dimensional data 3D->2D
 - Have 3 features treated symmetrically

PCA Algorithm

- Before applying PCA must do data preprocessing
 - o Given a set of m unlabeled examples we must do
 - Mean normalization
 - Replace each x_i with x_i μ_i
 - In other words, determine the mean of each feature set, and then for each feature subtract the mean from the value, so we re-scale the mean to be 0
 - Feature scaling (depending on data)
 - If features have very different scales then scale so they all have a comparable range of values
 - e.g. x_i¹ is set to (x_i μ_i) / s_i
 - Where s_i is some measure of the range, so could be
 - Biggest smallest
 - Standard deviation (more commonly)
- With preprocessing done, PCA finds the lower dimensional sub-space which minimizes the sum of the square
 - In summary, for 2D->1D we'd be doing something like this;



- Need to compute two things;
 - Compute the u vectors
 - The new planes
 - Need to compute the z vectors
 - z vectors are the new, lower dimensionality feature vectors
- A mathematical derivation for the u vectors is very complicated

• But once you've done it, the procedure to find each u vector is not that hard

Algorithm description

- Reducing data from *n*-dimensional to k-dimensional
 - Compute the covariance matrix

$$\Sigma = \frac{1}{m} \sum_{i=1}^{n} (x^{(i)}) (x^{(i)})^{T}$$

- lacktriangle This is commonly denoted as Σ (greek upper case sigma) NOT summation symbol
- Σ = sigma
 - This is an [n x n] matrix
 - Remember than xⁱ is a [n x 1] matrix
- In MATLAB or octave we can implement this as follows;

$$sigma = (1/m) * (X' * X)$$

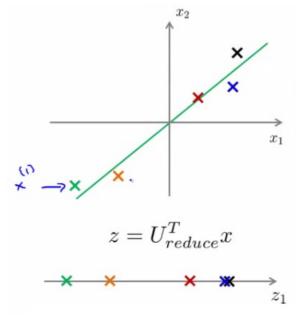
- Compute eigenvectors of matrix Σ
 - [U,S,V] = svd(sigma)
 - svd = singular value decomposition
 - More numerically stable than eig
 - eig = also gives eigenvector
- U,S and V are matrices
 - U matrix is also an [n x n] matrix
 - Turns out the columns of U are the u vectors we want!
 - So to reduce a system from n-dimensions to k-dimensions
 - Just take the first k-vectors from U (first k columns)

$$U = \begin{bmatrix} | & | & | \\ u^{(1)} & u^{(2)} & \dots & u^{(n)} \\ | & | & | \end{bmatrix} \in \mathbb{R}^{n \times n}$$

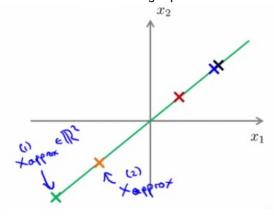
- Next we need to find some way to change x (which is n dimensional) to z (which is k dimensional)
 - (reduce the dimensionality)
 - Take first k columns of the u matrix and stack in columns
 - n x k matrix call this U_{reduce}
 - We calculate z as follows
 - $z = (U_{reduce})^T * x$
 - So [k x n] * [n x 1]
 - Generates a matrix which is
 - k * 1
 - If that's not witchcraft I don't know what is!
- Exactly the same as with supervised learning except we're now doing it with unlabeled data
- So in summary
 - Preprocessing
 - Calculate sigma (covariance matrix)
 - Calculate eigenvectors with svd
 - Take k vectors from U (U_{reduce}= U(:,1:k);)
 - Calculate z (z =U_{reduce}' * x;)
- No mathematical derivation
 - Very complicated
 - But it works

Reconstruction from Compressed Representation

- Earlier spoke about PCA as a compression algorithm
 - If this is the case, is there a way to decompress the data from low dimensionality back to a higher dimensionality format?
- Reconstruction
 - Say we have an example as follows



- \circ We have our examples (x¹, x² etc.)
- Project onto z-surface
- Given a point z¹, how can we go back to the 2D space?
- Considering
 - \circ z (vector) = $(U_{reduce})^T * x$
- To go in the opposite direction we must do
 - x_{approx} = U_{reduce} * z
 - To consider dimensions (and prove this really works)
 - $U_{reduce} = [n \times k]$
 - z[k*1]
 - So
 - x_{approx} = [n x 1]
- · So this creates the following representation



• We lose some of the information (i.e. everything is now perfectly on that line) but it is now projected into 2D space

Choosing the number of Principle Components

- How do we chose *k* ?
 - k = number of principle components
 - Guidelines about how to chose k for PCA
- To chose k think about how PCA works
 - PCA tries to minimize averaged squared projection error

• Total variation in data can be defined as the average over data saying how far are the training examples from the origin

• When we're choosing k typical to use something like this

$$\frac{\frac{1}{m} \sum_{i=1}^{m} \|x^{(i)} - x_{approx}^{(i)}\|^2}{\frac{1}{m} \sum_{i=1}^{m} \|x^{(i)}\|^2} \le 0.01$$
 (1%)

- Ratio between averaged squared projection error with total variation in data
 - Want ratio to be small means we retain 99% of the variance
- o If it's small (0) then this is because the numerator is small
 - The numerator is small when $x^i = x_{approx}^i$
 - i.e. we lose very little information in the dimensionality reduction, so when we decompress we regenerate the same data
- So we chose k in terms of this ratio
- · Often can significantly reduce data dimensionality while retaining the variance
- · How do you do this

Algorithm:

Try PCA with
$$k = 1$$

Compute
$$U_{reduce}, z^{(1)}, z^{(2)},$$

$$\dots, z^{(m)}, x_{approx}^{(1)}, \dots, x_{approx}^{(m)}$$

Check if

$$\frac{\frac{1}{m} \sum_{i=1}^{m} \|x^{(i)} - x_{approx}^{(i)}\|^2}{\frac{1}{m} \sum_{i=1}^{m} \|x^{(i)}\|^2} \le 0.01?$$

Advice for Applying PCA

- Can use PCA to speed up algorithm running time
 - · Explain how
 - · And give general advice

Speeding up supervised learning algorithms

- Say you have a supervised learning problem
 - Input x and v
 - x is a 10 000 dimensional feature vector
 - e.g. 100 x 100 images = 10 000 pixels
 - Such a huge feature vector will make the algorithm slow
 - With PCA we can reduce the dimensionality and make it tractable
 - How
 - 1) Extract xs
 - So we now have an unlabeled training set
 - 2) Apply PCA to x vectors
 - So we now have a reduced dimensional feature vector z
 - 3) This gives you a new training set
 - Each vector can be re-associated with the label
 - 4) Take the reduced dimensionality data set and feed to a learning algorithm
 - Use y as labels and z as feature vector
 - 5) If you have a new example map from higher dimensionality vector to lower dimensionality vector, then feed into learning algorithm
- PCA maps one vector to a lower dimensionality vector
 - ∘ x -> z
 - Defined by PCA only on the training set
 - The mapping computes a set of parameters
 - Feature scaling values
 - U_{reduce}
 - Parameter learned by PCA
 - Should be obtained only by determining PCA on your training set
 - So we use those learned parameters for our
 - Cross validation data
 - Test set
- Typically you can reduce data dimensionality by 5-10x without a major hit to algorithm

Applications of PCA

- Compression
 - Why
 - Reduce memory/disk needed to store data
 - Speed up learning algorithm
 - o How do we chose k?
 - % of variance retained
- Visualization
 - Typically chose k = 2 or k = 3
 - Because we can plot these values!
- · One thing often done wrong regarding PCA
 - · A bad use of PCA: Use it to prevent over-fitting
 - Reasoning
 - If we have xⁱ we have n features, zⁱ has k features which can be lower
 - If we *only* have k features then maybe we're less likely to over fit...
 - This doesn't work
 - BAD APPLICATION
 - Might work OK, but not a good way to address over fitting
 - Better to use regularization
 - PCA throws away some data without knowing what the values it's losing
 - Probably OK if you're keeping most of the data
 - But if you're throwing away some crucial data bad
 - So you have to go to like 95-99% variance retained
 - So here regularization will give you AT LEAST as good a way to solve over fitting
- · A second PCA myth
 - Used for compression or visualization good
 - Sometimes used
 - Design ML system with PCA from the outset
 - But, what if you did the whole thing without PCA
 - See how a system performs without PCA
 - ONLY if you have a reason to believe PCA will help should you then add PCA
 - PCA is easy enough to add on as a processing step
 - Try without first!