CS-GY 6923: Lecture 12 Autoencoders, Principal Component Analysis

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State-of-the-art supervised learning models like neural networks learn very good features.

But they require <u>lots and lots</u> of data. Imagenet has 14 million unlabeled images. Mostly of everyday objects.

What if you want to apply deep convolutional networks to a problem where you do not have a lot of **labeled data** in the first place?



Example: Classify images of different Quidditch balls.

Real example: Classify images of insects for use in agricultural applications in new localities.

Zero-Shot Insect Detection via Weak Language Supervision

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ONE-SHOT LEARNING

A human could probably achieve near perfect classification accuracy even given access to a **single labeled example** from each class:



Major question in ML: How? Can we design ML algorithms which can do the same?

Transfer knowledge from one task we already know how to solve to another.



For example, we have learned from past experience that balls used in sports have consistent shapes, colors, and sizes. These <u>features</u> can be used to distinguish balls of different type.

Features

Examples of possible high-level features a human would learn:

Classes

roundness 1 .1 1 .6 1 .4 size relative to human 10 1 7 2 7 5 hand yellowish color .2 .1 1 .1 0 .9

FEATURE LEARNING

If these features are highly informative (i.e. lead to highly separable data) few training examples are needed to learn.



Might suffice to classify ball using nearest training example in feature space, even if just a handful of training examples.

TRANSFER LEARNING

Empirical observation: Features learned when training models like deep neural nets seem to capture exactly these sorts of high-level properties.



Even if we can't put into words what each feature in **z** means...

This is now a common technique in computer vision:

- 1. Download network trained on large image classification dataset (e.g. Imagenet).
- 2. Extract features **z** for any new image **x** by running it through the network up until layer before last.
- 3. Use these features in a simpler machine learning algorithm that requires less data (nearest neighbor, logistic regression, etc.).

This approach has even been used on the quidditch problem: github.com/thatbrguy/Object-Detection-Quidditch **Transfer learning:** Lots of labeled data for one problem makes up for little labeled data for another.

But what if we don't even have labeled data for a sufficiently related problem?

How to extract features in a data-driven way from <u>unlabeled</u> <u>data</u> is one of the central problems in <u>unsupervised learning</u>.

- **Supervised learning:** <u>All</u> input data examples come with targets/labels. What machines have been really good at for the past 8 years.
- **Unsupervised learning:** <u>No</u> input data examples come with targets/labels. Interesting problems to solve include clustering, anomaly detection, semantic embedding, etc.
- Semi-supervised learning: <u>Some</u> (typically very few) input data examples come with targets/labels. What human babies are really good at, and we have recently made machines a lot better at.

AUTOENCODER

Simple but clever idea: If we have inputs $\mathbf{x}_1, \ldots, \mathbf{x}_n \in \mathbb{R}^d$ but few or no targets y_1, \ldots, y_n , just <u>make the inputs the targets</u>.

- Let $f_{\boldsymbol{\theta}} : \mathbb{R}^d \to \mathbb{R}^d$ be our model.
- Let L_{θ} be a loss function. E.g. squared loss: $L_{\theta}(\mathbf{x}) = \|\mathbf{x} - f_{\theta}(\mathbf{x})\|_2^2.$
- Train model: $\theta^* = \min_{\theta} \sum_{i=1}^n L_{\theta}(\mathbf{x}).$

If f_{θ} is a model that incorporates feature learning, then these features can be used for supervised tasks.

 f_{θ} is called an **autoencoder**. It maps input space to input space (e.g. images to images, french to french, PDE solutions to PDE solutions).

Two examples of autoencoder architectures:



Which would lead to better feature learning?

AUTOENCODER

Important property of autoencoders: no matter the architecture, there must always be a **bottleneck** with fewer parameters than the input. The bottleneck ensures information is "distilled" from low-level features to high-level features.



AUTOENCODER

Separately name the mapping from input to bottleneck and from bottleneck to output.

Encoder: $e : \mathbb{R}^d \to \mathbb{R}^k$ Decoder: $d : \mathbb{R}^d \to \mathbb{R}^k$ $f(\mathbf{x}) =$



Often symmetric, but does not have to be.

AUTOENCODER RECONSTRUCTION

Example image reconstructions from autoencoder:



https://www.biorxiv.org/content/10.1101/214247v1.full.pdf

Input parameters: d = 49152. Bottleneck "latent" parameters: k = 1024. The best autoencoders do not work as well as supervised methods for feature extraction, but they require no labeled data.¹

There are a lot of cool applications of autoencoders beyond feature learning!

- Learned data compression.
- Denoising and in-painting.
- Data/image synthesis.

¹Recent progress on **self-supervised** learning achieves the best of both worlds – state-of-the-art feature learning with no labeled data.

Due to their bottleneck design, autoencoders perform dimensionality reduction and thus data compression.



Given input **x**, we can completely recover $f(\mathbf{x})$ from $\mathbf{z} = e(\mathbf{x})$. **z** typically has many fewer dimensions than **x** and for a typical image $f(\mathbf{x})$ will closely approximate **x**.

The best lossy compression algorithms are tailor made for specific types of data:

- JPEG 2000 for images
- MP3 for digital audio.
- MPEG-4 for video.

All of these algorithms take advantage of specific structure in these data sets. E.g. JPEG assumes images are locally "smooth".





AUTOENCODERS FOR IMAGE COMPRESSION

With enough input data, autoencoders can be trained to find this structure on their own.



Proposed method, 5908 bytes (0.167 bit/px), PSNR: luma 23.38 dB/chroma 31.86 dB, MS-SSIM: 0.9219



JPEG 2010, 5908 bytes (0.167 bit/px), PSNR: luma 23.24 dB/chroma 31.04 dB, MS-SSIM: 0.8803



Proposed method, 6021 bytes (0.170 bit/px), PSNR: 24.12 dB, MS-SSIM: 0.9292



JPEG 2000, 6037 bytes (0.171 bit/px), PSNR: 23.47 dB, MS-SSIM: 0.9036

"End-to-end optimized image compression", Ballé, Laparra, Simoncelli

Need to be careful about how you choose loss function, design the network, etc. but can lead to much better image compression than "hand-tuned" algorithms like JPEG.

AUTOENCODERS FOR IMAGE CORRECTION



Image inpainting

Train autoencoder on <u>uncorrupted</u> images (unsupervised). Pass corrupted image \mathbf{x} through autoencoder and return $f(\mathbf{x})$ as repaired result.

AUTOENCODERS LEARN COMPRESSED REPRESENTATIONS

Why does this work?



Consider $128 \times 128 \times 3$ images with pixels values in 0, 1..., 255. How many possible images are there?

If **z** holds *k* values in 0, .1, .2, ..., 1, how many unique images **w** can be output by the autoencoder function *f*?

AUTOENCODERS LEARN COMPRESSED REPRESENTATIONS



For a good (accurate, small bottleneck) autoencoder, S will closely approximate I. Both will be much smaller than A.

AUTOENCODERS LEARN COMPRESSED REPRESENTATIONS



 $f(\mathbf{x}) = d(e(\mathbf{x}))$ projects an image **x** closer to the space of natural images.

Suppose we want to generate a random natural image. How might we do that?

• **Option 1**: Draw each pixel value in **x** uniformly at random. Draws a random image from *A*.



• **Option 2**: Draw **x** randomly from *S*, the space of images representable by the autoencoder.



How do we randomly select an image from \mathcal{S} ?

How do we randomly select an image **x** from *S*?



Randomly select code **z**, then set $\mathbf{x} = d(\mathbf{z})$.²

²Lots of details to think about here. In reality, people use "variational autoencoders" (VAEs), which are a natural modification of AEs.

AUTOENCODERS FOR DATA GENERATION DEMO

Teal created a demo for the "Fashion MNIST" data set:





PRINCIPAL COMPONENT ANALYSIS

Rest of lecture: Deeper dive into understanding a simple, but powerful autoencoder architecture. Specifically we will view **principal component analysis (PCA)** as a type of autoencoder.

PCA is the "linear regression" of unsupervised learning: often the go-to baseline method for feature extraction and dimensionality reduction.

Very important outside machine learning as well.

Consider the simplest possible autoencoder:



- One hidden layer. No non-linearity. No biases.
- Latent space of dimension k.
- Weight matrices are $\mathbf{W}_1 \in \mathbb{R}^{d \times k}$ and $\mathbf{W}_2 \in \mathbb{R}^{k \times d}$.

PRINCIPAL COMPONENT ANALYSIS

Given input $\mathbf{x} \in \mathbb{R}^d$, what is $f(\mathbf{x})$ expressed in linear algebraic terms?



$$f(\mathbf{x})^{\mathsf{T}} = \mathbf{x}^{\mathsf{T}} \mathbf{W}_1 \mathbf{W}_2$$

PRINCIPAL COMPONENT ANALYSIS



Encoder: $e(\mathbf{x}) = \mathbf{x}^T \mathbf{W}_1$. Decoder: $d(\mathbf{z}) = \mathbf{z} \mathbf{W}_2$

Given training data set $x_1,\ldots,x_n,$ let X denote our data matrix. Let $\tilde{X}=XW_1W_2.$



Natural squared autoencoder loss: Minimize $L(X, \tilde{X})$ where:

$$L(\mathbf{X}, \tilde{\mathbf{X}}) = \sum_{i=1}^{n} \|\mathbf{x}_i - f(\mathbf{x}_i)\|_2^2$$
$$= \sum_{i=1}^{n} \sum_{j=1}^{d} (\mathbf{x}_i[j] - f(\mathbf{x}_i)[j])^2$$
$$= \|\mathbf{X} - \tilde{\mathbf{X}}\|_F^2$$

Goal: Find $\mathbf{W}_1, \mathbf{W}_2$ to minimize the Frobenius norm loss $\|\mathbf{X} - \tilde{\mathbf{X}}\|_F^2 = \|\mathbf{X} - \mathbf{X}\mathbf{W}_1\mathbf{W}_2\|_F^2$ (sum of squared entries).

Rank in linear algebra:

- The columns of a matrix with <u>column rank</u> k can all be written as linear combinations of just k columns.
- The rows of a matrix with <u>row rank</u> k can all be written as linear combinations of k rows.
- Column rank = row rank = rank.



 \tilde{X} is a low-rank matrix. It only has rank k for $k \ll d$.
Principal component analysis is the task of finding W_1 , W_2 , which amounts to finding a rank k matrix \tilde{X} which approximates the data matrix X as closely as possible.

Finding the best **W**₁ and **W**₂ is a <u>non-convex</u> problem. We could try running an iterative method like gradient descent anyway. But there is also a direct algorithm!

Any matrix **X** can be written:



Where $\mathbf{U}^T\mathbf{U} = \mathbf{I}$, $\mathbf{V}^T\mathbf{V} = \mathbf{I}$, and $\sigma_1 \ge \sigma_2 \ge \dots \sigma_d \ge 0$. I.e. \mathbf{U} and \mathbf{V} are orthogonal matrices.

This is called the **singular value decomposition**.

Can be computed in $O(nd^2)$ time (faster with approximation algos).

Let $\mathbf{u}_1, \ldots, \mathbf{u}_n \in \mathbb{R}^n$ denote the columns of **U**. I.e. the left singular vectors of **X**.



Can read off optimal low-rank approximations from the SVD:



Eckart-Young-Mirsky Theorem: For any $k \le d$, $X_k = U_k \Sigma_k V_k^T$ is the optimal k rank approximation to X:

$$\mathbf{X}_k = \mathop{\mathrm{arg\,min}}_{ ilde{\mathbf{X}}} \|\mathbf{X} - \hat{\mathbf{X}}\|_F^2.$$

Claim:
$$X_k = U_k \Sigma_k V_k^T = X V_k V_k^T$$
.

So for a model with k hidden variables, we obtain an <u>optimal</u> autoencoder by setting $W_1 = V_k$, $W_2 = V_k^T$. $f(x) = xV_kV_k^T$.

PRINCIPAL COMPONENT ANALYSIS



Usually **x**'s columns (features) are mean centered and normalized to variance 1 before computing principal components.

Computing the SVD.

- Full SVD: U,S,V = scipy.linalg.svd(X). Runs in O(nd²) time.
- Just the top k components: U,S,V = scipy.sparse.linalg.svds(X, k). Runs in roughly O(ndk) time.

Recall that for a matrix $\mathbf{M} \in \mathbb{R}^{p \times p}$, \mathbf{q} is an <u>eigenvector</u> of \mathbf{M} if $\lambda \mathbf{q} = \mathbf{M} \mathbf{q}$ for any scalar λ .

- U's columns (the left singular vectors) are the orthonormal eigenvectors of **XX**^T.
- V's columns (the right singular vectors) are the orthonormal eigenvectors of X^TX.

•
$$\sigma_i^2 = \lambda_i (\mathbf{X}\mathbf{X}^T) = \lambda_i (\mathbf{X}^T\mathbf{X})$$

Exercise: Verify this directly. This means you can use any (symmetric) eigensolver for computing the SVD.

PCA APPLICATIONS

Like any autoencoder, PCA can be used for:

- Feature extraction
- Denoising and rectification
- Data generation
- Compression
- Visualization



denoising



synthetic data generation

The larger we set k, the better approximation we get.



rank 8 approx.

rank 9 approx.

rank 50 approx.

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Error vs. *k* is dictated by X's singular values. The singular values are often called the **spectrum** of X.

$$\|\mathbf{X}-\mathbf{X}_k\|_F^2 = \sum_{i=k}^d \sigma_i^2.$$



COLUMN REDUNDANCY

Colinearity of data features leads to an approximately low-rank data matrix.

	bedrooms	bathrooms	sq.ft.	floors	list price	sale price
home 1	2	2	1800	2	200,000	195,000
home 2	4	2.5	2700	1	300,000	310,000
					•	
	•	•	•	•	•	•
	•	•	•	•	•	•
home n	5	3.5	3600	3	450,000	450,000

sale price $\approx 1.05 \cdot \text{list}$ price. property tax $\approx .01 \cdot \text{list}$ price. Sometimes these relationships are simple, other times more complex. But as long as there exists <u>linear</u> relationships between features, we will have a lower rank matrix.

yard size
$$\approx$$
 lot size $-\frac{1}{2}$ · square footage.

cumulative GPA
$$\approx \frac{1}{4} \cdot$$
 year 1 GPA $+ \frac{1}{4} \cdot$ year 2 GPA
 $+ \frac{1}{4} \cdot$ year 3 GPA $+ \frac{1}{4} \cdot$ year 4 GPA.

LOW-RANK INTUITION

Two other examples of data with good low-rank approximations:

1. Genetic data:

	single nucleotide polymorphisms (SNPs) loc							
	144	312	436	800	943			
individual 1	Α	т	Т	С	G			
individual 2	т	G	G	С	С			
 individual n	С	A	т	A	G			

2. "Term-document" matrix with bag-of-words data:

	0	ar "	lan no	nuse		•••			%g	car
	doc_1	0	0	1	0	0	1	1	0	0
	doc_2	0	0	0	1	0	1	0	0	0
	:	1	1	0	1	0	0	0	1	0
l≡EÞ		0	0	0	0	0	0	0	1	1
	doc_n	1	0	0	0	0	0	0	1	1

SNPs matrices tend to be very low-rank.

	single nucleotide polymorphisms (SNPs) loc						
	144	312	436	800	943		
individual 1	Α	т	Т	С	G		
individual 2	Т	G	G	С	С		
 individual n	С	A	т	A	G		

Most of the information in **x** is explained by just a few **latent variable**.



"Genes Mirror Geography Within Europe" – Nature, 2008.



In data collected from European populations, latent variables capture information about geography.

$$\label{eq:z1} \begin{split} z[1] &\approx \mbox{relative north-south position of birth place} \\ z[2] &\approx \mbox{relative east-west position of birth place} \end{split}$$

Individuals born in similar places tend to have similar genes.

PCA FOR DATA VISUALIZATION

"Genes Mirror Geography Within Europe" – Nature, 2008.



Genetic data can be nicely visualized using PCA! Plot each data example **x** using two loading variables in **z**.

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For more complex data, what do principal components and loading vectors look like?

PRINCIPAL COMPONENTS

MNIST principal components:



Often principal components are difficult to interpret.

What do the loading vectors looks like?

The loading vector z for an example x contains coefficients which recombine the top k principal components v_1, \ldots, v_k to approximately reconstruct x.



Provide a short "finger print" for any image **x** which can be used to reconstruct that image.

For any **x** with loading vector **z**, z_i is the inner product similarity between **x** and the *i*th principal component **v**_{*i*}.



So we approximate $\mathbf{x} \approx \mathbf{\tilde{x}} = \langle \mathbf{x}, \mathbf{v}_1 \rangle \cdot \mathbf{v}_1 + \ldots + \langle \mathbf{x}, \mathbf{v}_k \rangle \cdot \mathbf{v}_k$.



Since $\mathbf{v}_1, \ldots, \mathbf{v}_k$ are orthonormal, this operation is a projection onto first *k* principal components.

I.e. we are projecting **x** onto the *k*-dimensional subspace spanned by $\mathbf{v}_1, \ldots, \mathbf{v}_k$.

For an example \mathbf{x}_i , the loading vector \mathbf{z}_i contains the coordinates in the projection space:



Important takeaway for data visualization and more: Latent feature vectors preserve similarity and distance information in the original data.

Let $\mathbf{x}_1 \dots, \mathbf{x}_n \in \mathbb{R}^d$ be our original data vectors, $\mathbf{z}_1 \dots, \mathbf{z}_n \in \mathbb{R}^k$ be our loading vectors (encoding), and $\tilde{\mathbf{x}}_1 \dots, \tilde{\mathbf{x}}_n \in \mathbb{R}^d$ be our low-rank approximated data.

We have:

$$\begin{split} \|\tilde{\mathbf{x}}_{i}\|_{2}^{2} &= \|\mathbf{z}_{i}\|_{2}^{2} \\ \langle \tilde{\mathbf{x}}_{i}, \tilde{\mathbf{x}}_{j} \rangle &= \langle \mathbf{z}_{i}, \mathbf{z}_{j} \rangle \\ \|\tilde{\mathbf{x}}_{i} - \tilde{\mathbf{x}}_{j}\|_{2}^{2} &= \|\mathbf{z}_{i} - \mathbf{z}_{j}\|_{2}^{2} \end{split}$$

Conclusion: If our data had a good low rank approximation, we expect that:

$$\begin{aligned} \|\mathbf{x}_{i}\|_{2}^{2} &\approx \|\mathbf{z}_{i}\|_{2}^{2} \\ \langle \mathbf{x}_{i}, \mathbf{x}_{j} \rangle &\approx \langle \mathbf{z}_{i}, \mathbf{z}_{j} \rangle \\ \|\mathbf{x}_{i} - \mathbf{x}_{j}\|_{2}^{2} &\approx \|\mathbf{z}_{i} - \mathbf{z}_{j}\|_{2}^{2} \end{aligned}$$

TERM DOCUMENT MATRIX

Word-document matrices tend to be low rank.



Documents tend to fall into a relatively small number of different categories, which use similar sets of words:

- Financial news: markets, analysts, dow, rates, stocks
- US Politics: president, senate, pass, slams, twitter, media
- StackOverflow posts: python, help, convert, javascript

Latent semantic analysis = PCA applied to a word-document matrix (usually from a large corpus). One of the most fundamental techniques in natural language processing (NLP).



Each column of **z** corresponds to a latent "category" or "topic". Corresponding row in **Y** corresponds to the "frequency" with which different words appear in documents on that topic. Similar documents have similar LSA document vectors. I.e. $\langle \mathbf{z}_i, \mathbf{z}_j \rangle$ is large.

- z_i provides a more compact "finger print" for documents than the long bag-of-words vectors. Useful for e.g search engines.
- Comparing document vectors is often <u>more effective</u> than comparing raw BOW features. Two documents can have (z_i, z_j) large even if they have no overlap in words. E.g. because both share a lot of words with words with another document k, or with a bunch of other documents.

Same fingerprinting idea was also important in early facial recognition systems based on "eigenfaces":



Each image above is one of the principal components of a dataset containing images of faces.

WORD EMBEDDINGS



- $\langle \mathbf{y}_i, \mathbf{z}_a \rangle \approx 1$ when doc_a contains $word_i$.
- If word_i and word_j both appear in doc_a , then $\langle \mathbf{y}_i, \mathbf{z}_a \rangle \approx \langle \mathbf{y}_j, \mathbf{z}_a \rangle \approx 1$, so we expect $\langle \mathbf{y}_j, \mathbf{y}_j \rangle$ to be large.



If two words appear in the same document their, word vectors tend to point more in the same direction.

Result: Map words to numerical vectors in a <u>semantically</u> meaningful way. Similar words map to similar vectors. Dissimilar words to dissimilar vectors.



Extremely useful "side-effect" of LSA.

Capture e.g. the fact that "great" and "excellent" are near synonyms. Or that "difficult" and "easy" are antonyms.

Review 1: Very small and handy for traveling or camping. Excellent quality, operation, and appearance.

Review 2: So far this thing is great. Well designed, compact, and easy to use. I'll never use another can opener.

Review 3: Not entirely sure this was worth \$20. Mom couldn't figure out how to use it and it's fairly difficult to turn for someone with arthritis.

Goal is to classify reviews as "positive" or "negative".

Vocabulary: Small, handy, excellent, great, quality, compact, easy, difficult.

Review 1: Very small and handy for traveling or camping. Excellent quality, operation, and appearance.

[, , , , , , , ,]

Review 2: So far this thing is great. Well designed, compact, and easy to use. I'll never use another can opener.

[, , , , , , , ,]

Review 3: Not entirely sure this was worth \$20. Mom couldn't figure out how to use it and it's fairly difficult to turn for someone with arthritis.

[, , , , , , , , ,]

Bag-of-words approach typically only works for large data sets.

The features do not capture the fact that "great" and "excellent" are near synonyms. Or that "difficult" and "easy" are antonyms.



This can be addressed by first mapping words to <u>semantically</u> <u>meaningful vectors</u>. That mapping can be trained using a much large corpus of text than the data set you are working with (e.g. Wikipedia, Twitter, news data sets). How to go from word embeddings to features for a whole sentence or chunk of text?



USING WORD EMBEDDINGS

A few simple options: Feature vector $\mathbf{x} = \frac{1}{q} \sum_{i=1}^{q} \mathbf{y}_{q}$. small average handy useless average heavy Feature vector $\mathbf{x} = [\mathbf{y}_1, \mathbf{y}_2, \dots, \mathbf{y}_q]$. **y**₁**y**₂ ... **y**_q Х
To avoid issues with inconsistent sentence length, word ordering, etc., can concatenate a fixed number of top <u>principal</u> <u>components</u> of the matrix of word vectors:



There are much more complicated approaches that account for word position in a sentence. Lots of pretrained libraries available (e.g. Facebook's **InferSent**).

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Another view on word embeddings from LSA:



We chose Z to equal $XV_k = U_k \Sigma_k$ and $Y = V_k^T$. Could have just as easily set $Z = U_k$ and $Y = \Sigma_k V_k^T$, so Z has orthonormal columns.

WORD EMBEDDINGS

Another view on word embeddings from LSA:



- $\cdot \ {\rm X} \approx {\rm ZY}$
- $\cdot \ \mathbf{X}^{\mathsf{T}}\mathbf{X} \approx \mathbf{Y}^{\mathsf{T}}\mathbf{Z}^{\mathsf{T}}\mathbf{Z}\mathbf{Y} = \mathbf{Y}^{\mathsf{T}}\mathbf{Y}$
- So for word_i and word_j, $\langle \mathbf{y}_i, \mathbf{y}_j \rangle \approx [\mathbf{X}^T \mathbf{X}]_{i,j}$.

What does the i, j entry of $X^T X$ reprent?

WORD EMBEDDINGS



What does the i, j entry of $X^T X$ reprent?

 $\langle \mathbf{y}_i, \mathbf{y}_j \rangle$ is <u>larger</u> if *word*_i and *word*_j appear in more documents together (high value in **word-word co-occurrence matrix**, $\mathbf{X}^T \mathbf{X}$). Similarity of word embeddings mirrors similarity of word context.

General word embedding recipe:

- 1. Choose similarity metric *k*(*word*_{*i*}, *word*_{*j*}) which can be computed for any pair of words.
- 2. Construct similarity matrix $\mathbf{M} \in \mathbb{R}^{n \times n}$ with $\mathbf{M}_{i,j} = k(word_i, word_j)$.
- 3. Find low rank approximation $\mathbf{M} \approx \mathbf{Y}^{\mathsf{T}}\mathbf{Y}$ where $\mathbf{Y} \in \mathbb{R}^{k \times n}$.
- 4. Columns of **Y** are word embedding vectors.

WORD EMBEDDINGS



How do current state-of-the-art methods differ from LSA?

- Similarity based on co-occurrence in smaller chunks of words. E.g. in sentences or in any consecutive sequences of 3, 4, or 10 words.
- Usually transformed in non-linear way. E.g. $k(word_i, word_j) = \frac{p(i,j)}{p(i)p(j)}$ where p(i,j) is the frequency both i, j appeared together, and p(i), p(j) is the frequency either one appeared.

Computing word similarities for "window size" 4:

The girl walks to her dog to the park. It can take a long time to park your car in NYC. The dog park is always crowded on Saturdays.

The girl walks to her dog to the park. It can take a long time to park your car in NYC. The dog park is always crowded on Saturdays.

The girl walks to her dog to the park. It can take a long time to park your car in NYC. The dog park is always crowded on Saturdays.

	dog	park	crowded	the
gop	0	2	0	3
park	2	0	1	2
crowded	0	1	0	0
the	3	2	0	0

Current state of the art models: GloVE, word2vec.

- word2vec was originally presented as a shallow neural network model, but it is equivalent to matrix factorization method (Levy, Goldberg 2014).
- For word2vec, similarity metric is the "point-wise mutual information": $\log \frac{p(i,j)}{p(i)p(j)}$.

CAVEAT ABOUT FACTORIZATION



SVD will not return a symmetric factorization in general. In fact, if **M** is not positive semidefinite³ then the optimal low-rank approximation does not have this form.

³I.e., *k*(*word*_{*i*}, *word*_{*i*}) is not a positive semidefinite kernel.

CAVEAT ABOUT FACTORIZATION



- For each word *i* we get a left and right embedding vector
 w_i and y_i. It's reasonable to just use one or the other.
- If (y_i, y_j) is large and positive, we expect that y_i and y_j have similar similarity scores with other words, so they typically are still related words.
- Another option is to use as your features for a word the concatenation [w_i, y_i]

If you want to use word embeddings for your project, the easiest approach is to use <u>pre-trained</u> word vectors:

- Original gloVe website: https://nlp.stanford.edu/projects/glove/.
- Compilation of many sources: https://github.com/3Top/word2vec-api

Lots of cool demos online for what can be done with these embeddings. E.g. "vector math" to solve analogies.



The same approach used for word embeddings can be used to obtain meaningful numerical features for any other data where there is a natural notion of similarity.



For example, the items could be nodes in a social network graph. Maybe be want to predict an individuals age, level of interest in a particular topic, political leaning, etc.

NODE EMBEDDINGS



Generate random walks (e.g. "sentences" of nodes) and measure similarity by node co-occurence frequency.



Again typically normalized and apply a non-linearity (e.g. log) as in word embeddings.



Popular implementations: **DeepWalk**, **Node2Vec**. Again initially derived as simple neural network models, but are equivalent to matrix-factorization (Qiu et al. 2018).