### **Social Data Science**

Word embeddings I

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### Plan

- 1. Course updates
- 2. The vector-space model review
- 3. Latent semantic analysis
- 4. Language models

### **Course updates**

- Project proposal assignment on Canvas, due Friday at 5pm
  - ► A short description of the planned project
    - Data collection
    - Data cleaning
    - Data analysis
    - Data visualization
    - Team

#### **Vector representations**

- Last week we looked at how we can represent texts as numeric vectors
  - Documents as vectors of words
  - Words as vectors of documents
- ► A document-term matrix (*DTM*) is a matrix where documents are represented as rows and tokens as columns

#### Weighting schemes

- We can use different schemes to weight these vectors
  - $\triangleright$  Binary (Does word  $w_i$  occur in document  $d_i$ ?)
  - $\triangleright$  Counts (How many times does word  $w_i$  occur in document  $d_i$ ?)
  - ▶ TF-IDF (How many times does word  $w_i$  occur in document  $d_j$ , accounting for how often  $w_i$  occurs across all documents  $d \in D$ ?)
    - Recall Zipf's Law: a handful of words account for most words used; such words do little to help us to distinguish between documents

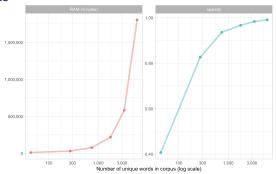
#### **Cosine similarity**

$$cos(\theta) = \frac{\vec{u} \cdot \vec{v}}{\|\vec{u}\| \|\vec{v}\|} = \frac{\sum_{i} \vec{u_i} \vec{v_i}}{\sqrt{\sum_{i} \vec{u_i}_i^2} \sqrt{\sum_{i} \vec{v_i}_i^2}}$$

#### Limitations

- These methods produce high-dimensinal, sparse vector representations
  - Given a vocabulary of unique tokens N the length of each vector |V| = N.
  - Many values will be zero since most documents only contain a small subset of the vocabulary.

#### **Limitations**



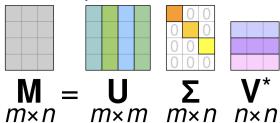
Source: https://smltar.com/embeddings.html

#### **Latent Semantic Analysis**

- One approach to reduce dimensionality and better capture semantics is called Latent Semantic Analysis (LSA)
  - ► We can use a process called *singular value decomposition* to find a *low-rank approximation* of a DTM.
- ▶ This provides *low-dimensional*, *dense* vector representations
  - ightharpoonup Low-dimensional, since  $|V| \ll N$
  - ▶ Dense, since vectors contain real values, with few zeros
- ► In short, we can "squash" a big matrix into a much smaller matrix while retaining important information.

$$DTM = U\Sigma V^T$$

#### **Singular Value Decomposition**



See the Wikipedia page for video of the latent dimensions in a sparse TDM.

### **Example: Shakespeare's writings**

X is a TF-IDF weighted Document-Term Matrix of Shakespeare's writings from Project Gutenberg. There are 11,666 unique tokens (each of which occurs 10 or more times in the corpus) and 66 documents.

```
library(tidyverse)
library(ggplot2)
X <- as.matrix(read.table("data/shakespeare.txt"))
X <- X[, which(colSums(X) != 0)] # Drop zero columns
dim(X)
## [1] 66 11666</pre>
```

#### Creating a lookup dictionary

We can construct a list to allow us to easily find the index of a particular token.

```
lookup.index.from.token <- list()

for (i in 1:length(colnames(X))) {
  lookup.index.from.token[colnames(X)[i]] <- i
}</pre>
```

#### Using the lookup dictionary

This easily allows us to find the vector representation of a particular word. Note how most values are zero since the character Hamlet is only mentioned in a handful of documents.

```
lookup.index.from.token["hamlet"]

## $hamlet

## [1] 8231

round(as.numeric(X[,unlist(lookup.index.from.token["hamlet"])]),3)

## [1] 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.00

## [13] 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.00

## [25] 0.046 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.00

## [37] 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.00

## [49] 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000 0.000

## [61] 0.014 0.000 0.000 0.000 0.000 0.000
```

#### **Calculating similarties**

The following code normalizes each *column* (rather than row normalization seen last lecture) and constructs a word-word cosine-similarity matrix.

```
normalize <- function(X) {
  for (i in 1:dim(X)[2]) {
    X[,i] <- (X[,i]/sqrt(sum(X[,i]^2)))
  }
  return(X)
}

X.n <- normalize(X)

sims <- t(X.n) %*% X.n
dim(sims)

## [1] 11666 11666</pre>
```

#### Most similar function

For a given token, this function allows us to find the n most similar tokens in the similarity matrix, where n defaults to 10.

#### Finding similar words

```
get.top.n("love",sims)
                 fair
                            lie
##
       love
                                    music
                                             sight beauty
                                                                breat
## 1.0000000 0.8583652 0.8533865 0.8425000 0.8213188 0.8098769 0.796472
##
      shine
                 dead
## 0.7616598 0.7616076
get.top.n("hate", sims)
##
       hate
              flatter
                          happy
                                   power
                                         time
                                                      forgot
                                                                  pas
## 1.0000000 0.7925268 0.7785826 0.7709535 0.7473243 0.7378957 0.734621
                kill
##
    friends
## 0.7310338 0.7300051
get.top.n("romeo", sims)
                         tybalt tybalts
                                           capulet benvolio montague
##
      romeo
             mercutio
## 1.0000000 0.9999330 0.9999318 0.9998235 0.9995140 0.9992019 0.998216
##
               juliet
    sampson
## 0.9923716 0.9906526
```

#### Singular value decomposition

The svd function allows us to decompose the DTM. We can then easily reconstruct it using the formula shown above.

```
# Computing the singular value decomposition
lsa <- svd(X)

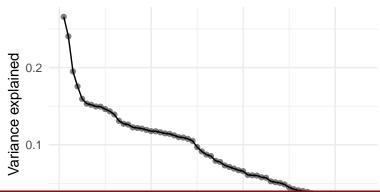
# We can easily recover the original matrix from this representation
X.2 <- lsa$u %*% diag(lsa$d) %*% t(lsa$v) # X = U \Sigma V^T

# Verifying that values are the same, example of first column
sum(round(X-X.2,5))
## [1] 0</pre>
```

### Singular value decomposition

This plot shows the magnitude of the singular values (the diagonal entries of  $\Sigma$ ). The magnitude of the singular value corresponds to the amount of variance explained in the original matrix.

## Scree plot of variance explained



## [1] 15.35817

#### Truncated singular value decomposition

In the example above retained the original matrix dimensions. The point of latent semantic analysis is to compute a truncated SVD such that we have a new matrix in a sub-space of X. In this case we only want to retain the first k dimensions of the matrix.

```
k <- 15 # Dimensions in truncated matrix

# We can take the SVD of X but only retain the first k singular values
lsa.2 <- svd(X, nu=k, nv=k)

# In this case we reconstruct X just using the first k singular values
X.trunc <- lsa.2$u %*% diag(lsa.2$d[1:k]) %*% t(lsa.2$v)

# But the values will be slightly different since it is an approximatio
# Some information is lost due to the compression
sum(round(X-X.trunc,5))</pre>
```

### Recalculating similarties using the LSA matrix

```
words.lsa <- t(lsa.2$v)
colnames(words.lsa) <- colnames(X)

round(as.numeric(words.lsa[,unlist(lookup.index.from.token["hamlet"])])
## [1] -0.00041  0.07355 -0.00271  0.00094 -0.00487  0.02775 -0.03515
## [9]  0.02197 -0.06272  0.00609 -0.01036  0.04939 -0.00635 -0.00269</pre>
```

### Recalculating similarties using the LSA matrix

```
words.lsa.n <- normalize(words.lsa)
sims.lsa <- t(words.lsa.n) %*% words.lsa.n</pre>
```

#### **Comparing similarities**

##

1 love

love

```
get.top.n("love",sims)
                 fair
                            lie
##
       love
                                     music
                                               sight
                                                        beauty
                                                                  breat
## 1.0000000 0.8583652 0.8533865 0.8425000 0.8213188 0.8098769 0.796472
##
       shine
                 dead
## 0.7616598 0.7616076
get.top.n("love",sims.lsa)
##
       love
               counsel
                           loves
                                     tears
                                             modesty
                                                        loving
                                                                    fai
## 1.0000000 0.9514839 0.9406690 0.9385571 0.9377943 0.9318378 0.929927
##
      oaths forsworn
## 0.9141528 0.9133888
bind_cols(names(get.top.n("love",sims)), names(get.top.n("love",sims.ls
## # A tibble: 10 x 2
      ...1 ...2
##
##
     <chr> <chr>
```

#### **Comparing similarities**

```
get.bottom.n <- function(token, sims, n=10) {</pre>
  bottom <- sort(sims[unlist(lookup.index.from.token[token]),],</pre>
                  decreasing=F) [1:n]
  return(bottom)
get.bottom.n("love", sims)
         adde
                  aduerse
##
                                         affayres
                                                       affraid
                                                                      aire
                                    ae
##
                                     0
##
        ascii assortment
                               attyre
##
                                     0
```

#### **Comparing similarities**

1 hate hate

##

```
get.top.n("hate",sims)
##
       hate
              flatter
                         happy power
                                          time
                                                     forgot
                                                                 pas
## 1.0000000 0.7925268 0.7785826 0.7709535 0.7473243 0.7378957 0.734621
##
    friends
                 kill
## 0.7310338 0.7300051
get.top.n("hate",sims.lsa)
##
       hate miserable
                         anger art
                                             bare
                                                     breath
                                                               fault
## 1.0000000 0.9697931 0.9683188 0.9409666 0.9401336 0.9303231 0.929732
##
       aged
                 thee
## 0.9285337 0.9275189
bind_cols(names(get.top.n("hate",sims)), names(get.top.n("hate",sims.ls
## # A tibble: 10 x 2
##
     ...1 ...2
##
     <chr> <chr>
```

#### **Comparing similarities**

```
get.top.n("romeo", sims)

## romeo mercutio tybalt tybalts capulet benvolio montague
## 1.0000000 0.9999330 0.9999318 0.9998235 0.9995140 0.9992019 0.998216
## sampson juliet
## 0.9923716 0.9906526

get.top.n("romeo", sims.lsa)

## romeo montagues tybalts tybalt mercutio mercutios capule
## 1.0000000 0.9999961 0.9999937 0.9999933 0.9999923 0.9999910 0.999988
## capulets romeos
## 0.9999846 0.9999741
```

#### **Comparing similarities**

```
get.top.n("hamlet", sims)
                                  ophelia polonius barnardo laerte
##
     hamlet horatio marcellus
## 1.0000000 0.9829677 0.9824643 0.9607921 0.9600178 0.9591448 0.958729
## voltemand lucianus
## 0.9465517 0.9308989
get.top.n("hamlet", sims.lsa)
##
     hamlet gertrude danish pyrrhus
                                           denmark wittingly
                                                                 polo
## 1.0000000 0.9981001 0.9980634 0.9962985 0.9962098 0.9961391 0.996088
##
       laer
               norwey
## 0.9960141 0.9959639
```

#### Execise

Re-run the code above with a different value of k (try lower or higher). Compare some terms in the original similarity matrix and the new matrix. How does changing k affect the results?

```
get.top.n("", sims)

## [1] NA NA NA NA NA NA NA NA NA NA
get.top.n("", sims.lsa)

## [1] NA NA NA NA NA NA NA NA NA NA
```

#### Inspecting the latent dimensions

We can analyze the meaning of the latent dimensions by looking at the terms with the highest weights in each row. In this case I use the raw LSA matrix without normalizing it. What do you notice about the dimensions?

```
for (i in 1:dim(words.lsa)[1]) {
  top.words <- sort(words.lsa[i,], decreasing=T)[1:5]
  print(paste(c("Dimension: ",i), collapse=" "))
  print(top.words)
## [1] "Dimension: 1"
##
                         bened
                                        bero
                                                      botes
             amv
                                                                     cas
## -1.204978e-06 -1.204978e-06 -1.204978e-06 -1.204978e-06 -1.204978e-0
   [1] "Dimension:
##
     sidenote footnote
                                ham
                                        hamlet
                                                      hane
  0.75168642 0.58659055 0.20541499 0.07354514 0.06784205
   [1] "Dimension:
      sidenote
                  footnote
                                                     antipholus
```

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quarto

syracuse

### **Limitations of Latent Semantic Analysis**

- Bag-of-words assumptions and document-level word associations
  - We still treat words as belonging to documents and lack finer context about their relationships
    - Although we could theoretically treat smaller units like sentences as documents
- ▶ Matrix computations become intractable with large corpora
- ▶ A neat linear algebra trick, but no underlying language model

#### Intuition

- ► A language model is a probabilistic model of language use
- ▶ Given some string of tokens, what is the most likely token?
  - Examples
    - Auto-complete
    - ► Google search completion

### **Bigram models**

- ▶  $P(w_i|w_{i-1})$  = What is the probability of word  $w_i$  given the last word,  $w_{i-1}$ ?
  - ► P(Jersey|New)
  - ► P(Brunswick|New)
  - ► P(York|New)
  - ► P(Sociology | New)

### **Bigram models**

- We use a corpus of text to calculate these probabilities from word co-occurrence.
  - ►  $P(Jersey|New) = \frac{C(New\ Jersey)}{C(New)}$ , e.g. proportion of times "New" is followed by "Jersey", where C() is the count operator.
- More frequently occurring pairs will have a higher probability.
  - We might expect that  $P(York|New) \approx P(Jersey|New) > P(Brunswick|New) >> P(Sociology|New)$

#### **Incorporating more information**

- We can also model the probability of a word, given a sequence of words
- P(x|S) = What is the probability of some word x given a partial sentence S?
- ightharpoonup A = P(Jersey | Rutgers University is in New)
- $ightharpoonup B = P(Brunswick|Rutgers\ University\ is\ in\ New)$
- $ightharpoonup C = P(York|Rutgers\ University\ is\ in\ New)$
- ► In this case we have more information, so "York" is less likely to be the next word. Hence,

 $ightharpoonup A \approx B > C$ .

#### **Estimation**

We can compute the probability of an entire sequence of words by using considering the joint conditional probabilities of each pair of words in the sequence. For a sequence of n words, we want to know the joint probability of  $P(w_1, w_2, w_3, ..., w_n)$ . We can simplify this using the chain rule of probability:

$$P(w_{1:n}) = P(w_1)P(w_2|w_1)P(w_3|w_{1:2})...P(w_n|w_{1:n-1})$$

$$= \prod_{k=1}^{n} P(w_k|w_{1:k-1})$$

#### **Estimation**

The bigram model simplifies this by assuming it is a first-order Markov process, such that the probability  $w_k$  only depends on the previous word,  $w_{k-1}$ .

$$P(w_{1:n}) \approx \prod_{k=1}^{n} P(w_k|w_{k-1})$$

These probabilities can be estimated by using Maximum Likelihood Estimation on a corpus.

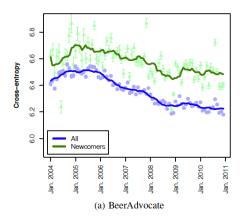
See https://web.stanford.edu/~iurafsky/slp3/3.pdf for an excellent review of language models

### **Empirical applications**

- ▶ Danescu-Niculescu-Mizil et al. 2013 construct a bigram language model for each month on BeerAdvocate and RateBeer to capture the language of the community
  - For any given comment or user, they can then use a measure called *cross-entropy* to calculate how "surprising" the text is, given the assumptions about the language model
- ► The theory is that new users will take time to assimilate into the linguistic norms of the community

https://en.wikipedia.org/wiki/Cross\_entropy

#### **Empirical applications**



Danescu-Niculescu-Mizil, Cristian, Robert West, Dan Jurafsky, Jure Leskovec, and Christopher Potts. 2013. "No Country for Old Members: User Lifecycle and Linguistic Change in Online Communities." In Proceedings of the 22nd International Conference on World Wide Web, 307–18. ACM. https://dl.acm.org/citation.cfm?id=2488416.

### Limitations of N-gram language models

- Language use is much more complex than N-gram language models
- Two limitations
  - Insufficient data
  - Complex models become intractable to compute
- Next lecture we will see how advances in neural network models and the availability of large text corpora have opened up new avenues for language modeling and semantic analysis

# **Summary**

- Limitations of sparse representations of text
  - ► LSA allows us to project sparse matrix into a dense, low-dimensional representation
- Probabilistic language models allow us to directly model language use
- ► Next lecture: How neural language models allow us to create more meaningful semantic representations of texts