MAT 167: Applied Linear Algebra
Lecture 23: Text Mining II

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Outline

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Using Cluster Centroids for Text Mining

- Instead of using the left singular vectors as a basis to approximate a term-document matrix, let's examine the cluster centers (centroids) obtained by k-means algorithm as a basis.
- Let $C_k = [\boldsymbol{c}_1 \dots \boldsymbol{c}_k] \in \mathbb{R}^{m \times k}$ be the k cluster centroids obtained by the k-means algorithm.
- $\boldsymbol{c}_j$'s are non-orthogonal; hence it is more convenient to obtain a set of orthonormal vectors that spans range(C_k).
- To do so, we can use the *reduced QR factorization*: $C_k = \widehat{Q}_k \widehat{R}_k$ where $\widehat{Q}_k \in \mathbb{R}^{m \times k}$, and $\widehat{R}_k \in \mathbb{R}^{k \times k}$.
- Now, let's approximate A using $\widehat{Q}_k$ in the sense of the least squares as:

$$\min_{G_k \in \mathbb{R}^{k \times n}} \|A - \widehat{Q}_k G_k\|_F.$$

- Let $G_k = [\boldsymbol{g}_1 \dots \boldsymbol{g}_n] \in \mathbb{R}^{k \times n}$. Then the above is equivalent to the following set of the LS problems:

$$\min_{\boldsymbol{g}_j \in \mathbb{R}^k} \|\boldsymbol{a}_j - \widehat{Q}_k \boldsymbol{g}_j\|_2, \quad j = 1 : n.$$

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- The inner product between the query vector $\boldsymbol{q}$ and the document vector $\boldsymbol{a}_j$ can be approximated as:

$$\boldsymbol{q}^\mathsf{T} \boldsymbol{a}_j \approx \boldsymbol{q}^\mathsf{T} \widehat{Q}_k \boldsymbol{g}_j = (\widehat{Q}_k^\mathsf{T} \boldsymbol{q})^\mathsf{T} \boldsymbol{g}_j = \boldsymbol{q}_k^\mathsf{T} \boldsymbol{g}_j, \; \boldsymbol{q}_k := \widehat{Q}_k^\mathsf{T} \boldsymbol{q}.$$

- Hence, the cosine similarity can be approximated as:

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An Example Trial with the NIPS Data

- $k = 50$; the same query vector ('entropy', 'minimum', 'maximum').
- The approximation error between $\widehat{Q}_k G_k$ and A was $\|A - \widehat{Q}_k G_k\|_F / \|A\|_F \approx 0.7227$, which was worse than that using the top 100 SVD basis.

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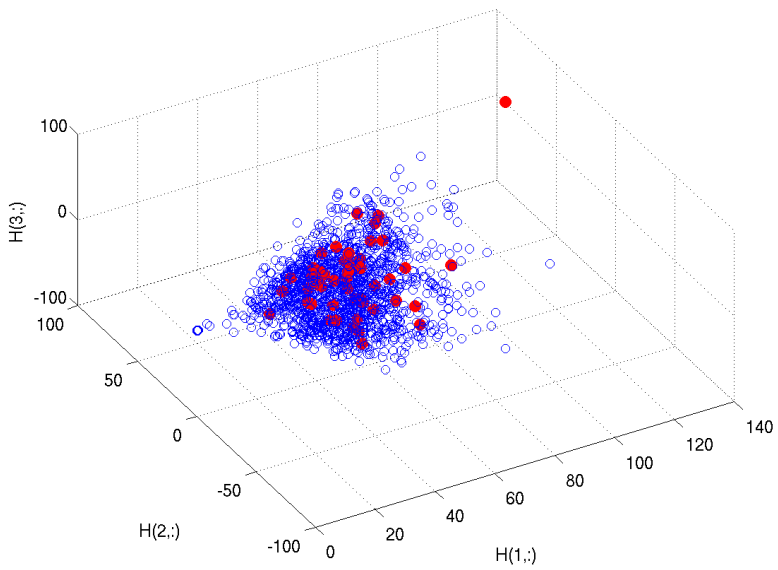
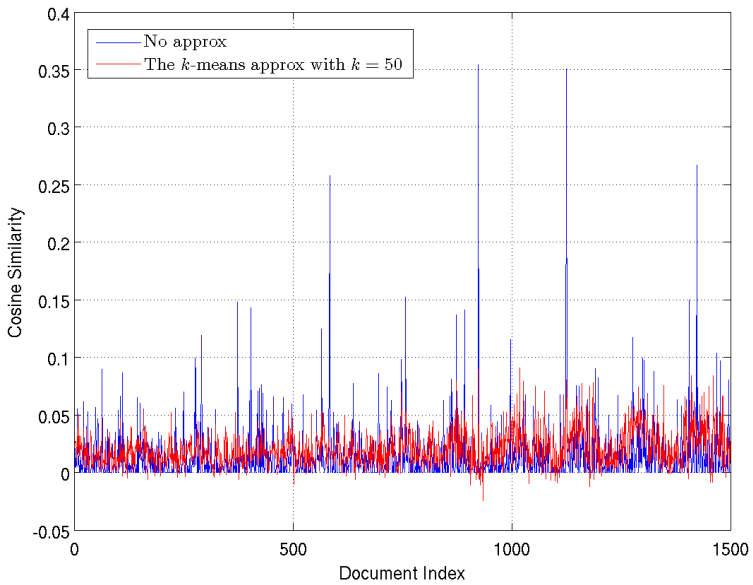


Documents in $U_{100}[:, 1:3]$



Cosine Similarity

With the 50-means based approximation, tol=0.2, 0.1, 0.05 correspond to 0, 0, 81 returned documents; Compare these with the no approximation case: 4, 15, 89; and with the best 100 approximation using SVD: 0, 4, 72.

My Reaction

- Running the k-means algorithm with large m and n is slow in general.
- If your document set really consists of k different topics (or categories), then this k-means-based approach should work well. Example: *The Science News Dataset* consisting of articles in the area of *Anthropology, Astronomy, Behavioral Sciences, Earth Sciences, Life Sciences, Math & CS, Medicine, Physics*. Which value of k should be used is still a question though.
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$$\frac{\widehat{\boldsymbol{q}}^{\mathsf{T}}\boldsymbol{h}_j}{\|\widehat{\boldsymbol{q}}\|_2\|\boldsymbol{h}_j\|_2}, \quad j = 1:n.$$

An Example Trial with the NIPS Data

- $k = 100$ was used.
- $\|A - WH\|_F / \|A\|_F \approx 0.6302$, which was *slightly* worse than that using the top 100 SVD basis (0.6074).
- Each $\boldsymbol{w}_j$ concentrates on one term, and is close to the canonical vector $\boldsymbol{e}_i \in \mathbb{R}^m$ for some i (recall: NNMF applied to the face database in Lecture 20).
- The peaks of $\boldsymbol{w}_j$, $j = 1:10$, correspond to: 'network', 'model', 'learning', 'function', 'unit', 'algorithm', 'input', 'data', 'neuron', 'cell', which are quite similar to the $\boldsymbol{u}_1$ vector or the 10 most frequently used terms.
- On the other hand, because $\boldsymbol{w}_j$'s are localized, the interpretation of the row vectors of H matrix becomes easy.

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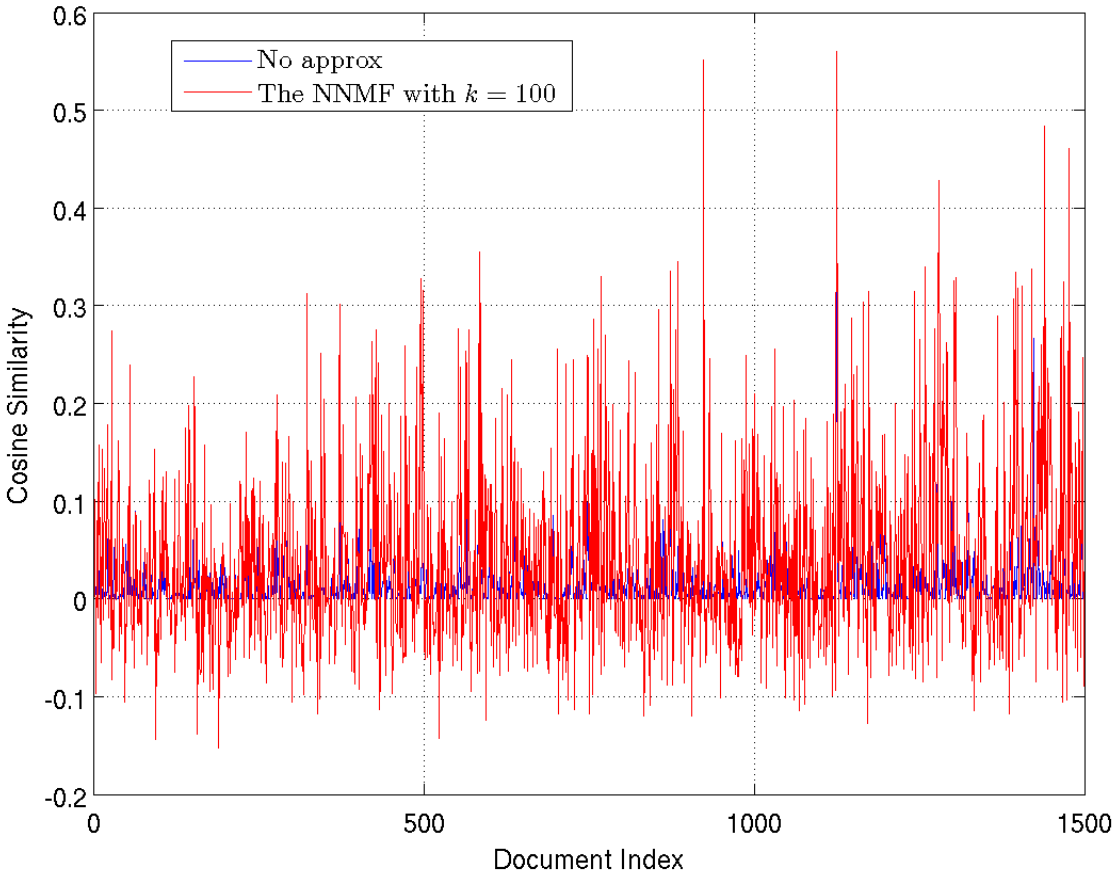
An Example Trial with the NIPS Data ...



With the NNMF-based approach using $k = 100$, tol=0.2, 0.1, 0.05 correspond to 101, 312, 535 returned documents; Compare with the no approximation case: 4, 15, 89. Changing the tol=0.4, 0.3, 0.2 with NNMF returns 5, 26, 101 documents.

My Reaction

- Using the LS solution for the query saves computational cost given the NNMF is already obtained because one can avoid the explicit computation and storage of WH.
- If we can compute and store WH, then we could use the following approximation of the original cosine similarity:

$$\frac{\boldsymbol{q}^\top \boldsymbol{a}_j}{\|\boldsymbol{q}\|_2 \|\boldsymbol{a}_j\|_2} \approx \frac{\boldsymbol{q}^\top W \boldsymbol{h}_j}{\|\boldsymbol{q}\|_2 \|W \boldsymbol{h}_j\|_2}.$$

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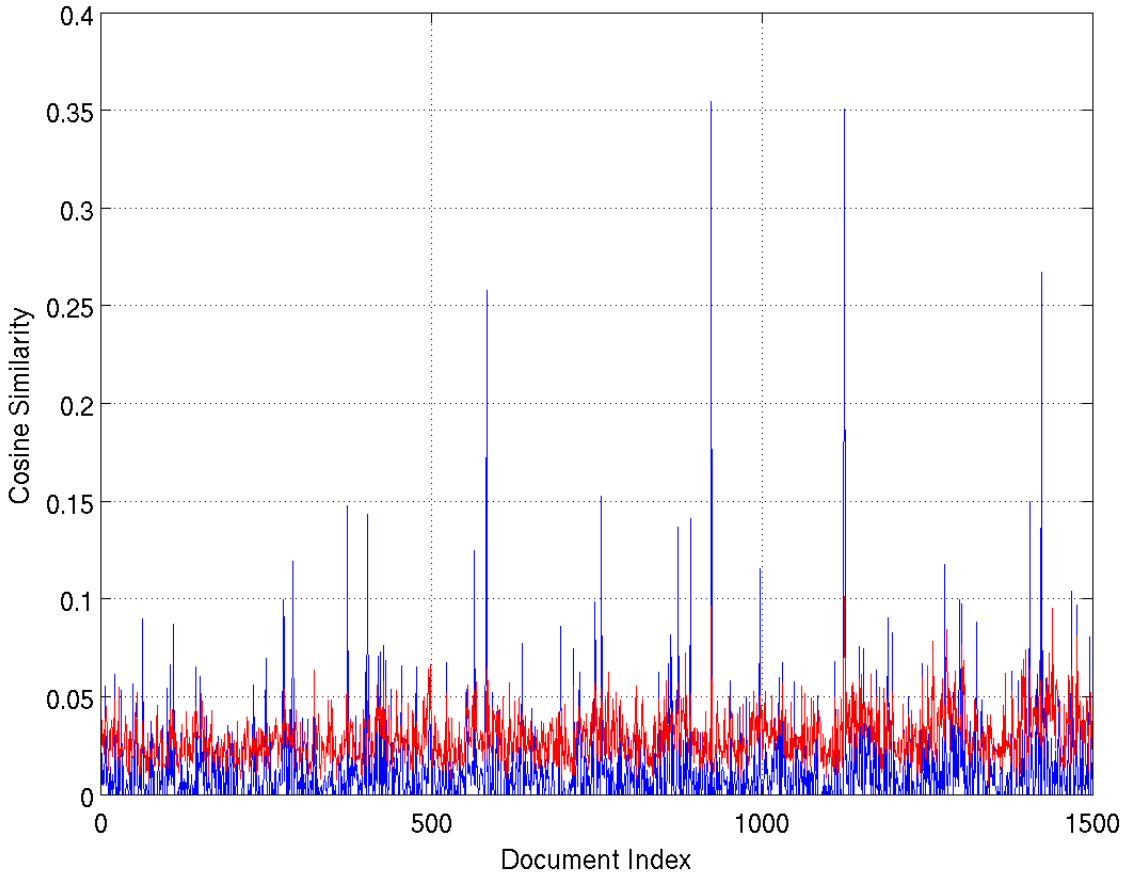

With the NNMF-based approach using $k = 100$ using the above cosine similarity approximation, tol=0.2, 0.1, 0.05 correspond to 0, 1, 97 returned documents; Compare with the no approximation case: 4, 15, 89. Without using the LS query, some of the relevant documents do not stick out clearly.