Coursework 3: document models

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This coursework involves aggregating, summarizing, and joining datasets. This may be done with straight Python, or with MATLAB-style manipulations using numpy, or with pandas dataframes. If you anticipate future work in machine learning and data science then you should learn to use pandas dataframes, and you may find it helpful to follow the walkthrough in Section 3 of IA Scientific Computing. If you prefer not to use dataframes, and you have questions about how they are being used in the code snippets below, ask questions in the LE49 help forum.

```
import numpy as np
import matplotlib
import matplotlib.pyplot as plt
import scipy.io
import pandas
import requests, io
%matplotlib inline
```

Data import

The data is provided as https://teachingfiles.blob.core.windows.net/probml/kos_doc_data.mat. It contains two matrices A and B for training and testing respectively, both matrices with 3 columns: document ID, word ID, and word count. The words themselves are the vector V, where e.g. V[840]='bush'. The following snippet reads in the data, and converts A and B to dataframes.

Question (a): maximum likelihood estimation

We'll model words as independent samples from a multinomial distribution with parameter β , where β_v is the probability of word $v \in V$. Using A as the training set, find the maximum likelihood estimator $\hat{\beta}$, and plot the 20 most-probable words in a histogram. What is the log probability of the test document doc_id=2527, given $\hat{\beta}$? Briefly interpret your answer.

Note: you can plot a histogram with

```
with plt.rc_context({'figure.figsize': (5,8)}):  # set plot size
   plt.barh(np.arange(20), top_20_probs, align='center') # draw bars
plt.yticks(np.arange(20), top_20_words) # label the y axis
plt.xlabel(r'$\hat{\beta}$') # label the x axis
plt.gca().invert_yaxis() # optionally, flip the y-axis
plt.show()
```

Question (b): Bayesian inference

Use Bayesian inference to find the posterior distribution of β given the training set A, using a symmetric Dirichlet distribution with concentration parameter $\alpha = 0.1$ as prior. Let $\tilde{\beta}_v$ be the posterior predictive probability of word $v \in V$, i.e. the posterior probability that a newly chosen word is v. Derive an expression for $\tilde{\beta}_v$, and compare it to $\hat{\beta}_v$. Explain the implications, both for common and for rare words.

```
Hint: \Gamma(z+1) = z \Gamma(z).
```

Question (c)

In information theory, the self-information of a document $w = (w_1, ..., w_n)$ is defined as $i(w) = -\log_2 \mathbb{P}(W = w)$, where W is a random document from some generative probabilistic model. The self-information can be interpreted

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as the number of bits needed to encode or transmit w. The number of bits needed per word is thus i(w)/n. In text modelling, it is more common to use the terms perplexity for $2^{i(w)}$, and per-word perplexity for $2^{i(w)/n}$. Loosely speaking, if the per-word perplexity is g then the uncertainty in the next word is the same as the uncertainty in a g-sided die.

For the trained Bayesian model from part (b), what is the per-word perplexity of the test document doc_id=2000? Plot a histogram showing the distribution of per-word perplexity over all the test documents (using plt.hist). Pick out two documents, one with high per-word perplexity and one with low per-word perplexity, show their contents, and interpret the difference between them.

Question (d): Gibbs sampler for the mixture-of-multinomials model

The Bayesian mixture-of-multinomials model can be described by the following code:

```
def bmm_generate(doc_length, V, α, γ, K):
    θ = np.random.dirichlet(α * np.ones(K))  # prob dist over document classes {1,...,K}
    β = np.random.dirichlet(γ * np.ones(len(V)), size=K) # for each doc class, a prob dist over words
    z = np.random.choice(K, p=θ, size=len(doc_length)) # doc class of each document
    return [np.random.choice(V, p=β[zd], size=nd) for zd,nd in zip(z, doc_length)]

for doc in bmm_generate(doc_length=[5,2,4], V=V, α=10, γ=.1, K=20):
    print(doc)

['critic' 'indymedia' 'susan' 'citizenship' 'cycles']
    ['cool' 'celebrity']
    ['jennings' 'quarter' 'token' 'governance']
```

The following code implements a collapsed Gibbs sampler. Complete the line that defines logp. Use the sampler to find the posterior predictive distribution of document class, and plot how this distribution evolves as a function of the number of Gibbs sweeps. How many iterations does it take to converge?

```
def bmm_gibbs(doc_label, word_id, count, W, \alpha, \gamma, K):
    # doc_labels = distinct values of doc_label
    # doc_index = a list as long as doc_label
                  such that doc_labels[doc_index[j]] = doc_label[j]
    doc_labels, doc_index = np.unique(doc_label, return_inverse=True)
    # z[i] = class of document i, where i enumerates the distinct doc_labels
    # doc_count[k] = number of documents of class k
    z = np.random.choice(K, len(doc_labels))
    doc_count = np.zeros(K, dtype=int)
    for k in z: doc_count[k] += 1
    # occurrences[k,w] = number of occurrences of word_id w in documents of class k
    # word_count[k] = total number of words in documents of class k
    x = pandas.DataFrame({'doc_class': z[doc_index], 'word_id': word_id, 'count': count}) \
        .groupby(['doc_class', 'word_id']) \
        ['count'].apply(sum) \
        .unstack(fill_value=0)
    occurrences = np.zeros((K, len(V)))
    occurrences[x.index.values.reshape((-1,1)), x.columns.values] = x
    word_count = np.sum(occurrences, axis=1)
    while True:
        for i in range(len(doc_labels)):
            # get the words, counts for document i
            # and remove this document from the counts
            w,c = word_id[doc_index==i].values, count[doc_index==i].values
            occurrences[z[i], w] -= c
            word_count[z[i]] -= sum(c)
            doc_count[z[i]] -= 1
```

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Find the log probability that this document belongs to class k, marginalized over θ and β

```
logp = [... for k in range(K)]
    p = np.exp(logp - np.max(logp))
    p = p/sum(p)

# Assign this document to a new class, chosen randomly, and add back the counts
    k = np.random.choice(K, p=p)
    z[i] = k
    occurrences[k, w] += c
    doc_count[k] += 1
    word_count[k] += sum(c)

    yield np.copy(z)

The Gibbs sampler may be run as follows:

g = bmm_gibbs(A['doc_id'], A['word_id'], A['count'], W=len(V), α=10, γ=.1, K=20)

NUM_ITERATIONS = 20

res = np.stack([next(g) for _ in range(NUM_ITERATIONS)])
# this produces a matrix with one row per iteration and a column for each unique doc_id
```

Question (e): interpretation

Let $\alpha=10$, $\gamma=0.1$, K=20. Run the Gibbs sampler, and find the posterior predictive probabilities for topics and words. For the 8 most popular topics, print the probability of the topic, and the 8 most probable words and their probabilities. Display probabilities in $[shannons](https://en.wikipedia.org/wiki/Shannon_(unit))$, i.e. display a probability p as $-\log_2 p$. An increase of 1 shannon corresponds to a 50% decrease in probability.

Rerun with different random seeds. Do you think this method has succeeded in identifying topics?

Optional. There are some words that are very common across all topics. How might we pick out the distinctive words for each topic?

Optional question (f): evaluation

Give a formula for per-word perplexity for the mixture model, in terms of the posterior predictive probabilities for topics and words.

Plot a histogram showing the distribution of per-word perplexity over all the test documents for the model in part (e). Also plot the histogram obtained from K = 8, and the histogram from the plain multinomial model in part (c). Which model do you prefer, and why?

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