## **67658 Natural Language Processing**

## Exercise 1

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```
initialize notebook:
import spacy
import textacy
import pickle
import numpy as np
from collections import Counter
from datasets import load dataset
# uncomment to download the module:
# !python -m spacy download en core web sm
# load the data:
nlp = spacy.load("en core web sm")
dataset = load dataset('wikitext', 'wikitext-2-raw-v1', split='train')
corpus = dataset['text']
Found cached dataset wikitext
(/Users/asafshul/.cache/huggingface/datasets/wikitext/wikitext-2-raw-
v1/1.0.0/
a241db52902eaf2c6aa732210bead40c090019a499ceb13bcbfa3f8ab646a126)
functions:
def clean text(text):
    return 'START ' + ' '.join([c.lemma for c in nlp(text) if
c.is alpha])
def train language model(corpus, level=2):
    narams = []
    print('counting words...')
    # extract ngrams:
    for text in corpus:
ngrams.extend(list(textacy.extract.ngrams(nlp(clean text(text)),
level, filter stops=False)))
    print('calculating fregs...')
    if level==1:
        return get unigram(ngrams)
```

```
if level==2:
        return get bigram(ngrams)
def get unigram(ngrams):
    word counts = dict(Counter([str(w) for w in ngrams if str(w) !=
'START'1)
    N = sum(word counts.values())
    for key in word counts:
        word counts[key] = np.log(word counts[key] / N)
    return word counts
def get bigram(ngrams):
    base dict = dict(Counter([(str(w[0])), str(w[1])) for w in
ngrams]))
    freq dict = {}
    # format to bigram dict:
    for key, count in base dict.items():
        base word, next word = key
        if base word not in freq dict:
            freq dict[base word] = {next word : count}
        elif next word not in freq dict[base word]:
            freq dict[base word][next word] = count
        else:
            freq dict[base word][next word] += count
    # change count to relative probability:
    for word counts in freq dict.values():
         N = sum(word counts.values())
        for key in word counts:
            word counts[key] = np.log(word counts[key] / N)
    # return the model:
    return freq_dict
1.
Train maximum-likelihood unigram and bigram language models based on the above
training data.
%%time
# unigram:
uni model = train language model(corpus, 1)
counting words...
calculating freqs...
```

```
CPU times: user 10min 49s, sys: 3.99 s, total: 10min 53s
Wall time: 10min 53s
%%time
# bigram:
bi model = train language model(corpus, 2)
counting words...
calculating freqs...
CPU times: user 10min 53s, sys: 4.22 s, total: 10min 57s
Wall time: 10min 58s
# save models to pickle file:
with open('unigram.pickle', 'wb') as file:
    pickle.dump(uni model, file, protocol=pickle.HIGHEST PROTOCOL)
with open('bigram.pickle', 'wb') as file:
    pickle.dump(bi model, file, protocol=pickle.HIGHEST PROTOCOL)
# load models:
# with open('unigram.pickle', 'rb') as fp:
      uni model = pickle.load(fp)
# with open('bigram.pickle', 'rb') as fp:
      bi model = pickle.load(fp)
     Using the bigram model, continue the following sentence with the most probable
     word predicted by the model: "I have a house in ...".
def pred bigram(model, sentance):
    last word = sentance.split(' ')[-1]
    probs = bi model[last word]
    idx = np.argmax(list(probs.values()))
    return list(probs.keys())[idx]
sentance = 'I have a house in'
pred bigram(bi model, sentance)
'the'
  1. Using the bigram model:
```

- (a) compute the probability of the following two sentences (for each sentence separately).
- (b) compute the perplexity of both the following two sentences (treating them as a single test set with 2 sentences).
  - Brad Pitt was born in Oklahoma
  - The actor was born in USA

```
def calc sentance log prob bigram(sentance, bi model):
    sentance arr = clean text(sentance).split(' ')
    prob = 0
    for word, next word in zip(sentance arr[:-1], sentance arr[1:]):
        if word in bi model and next word in bi model[word]:
            prob += bi model[word][next word]
        else:
            return -np.inf
    return round(prob, 3)
sentance1 = 'Brad Pitt was born in Oklahoma'
sentance2 = 'The actor was born in USA'
prob1 = calc_sentance_log_prob_bigram(sentance1, bi_model)
prob2 = calc sentance log prob bigram(sentance2, bi model)
print(f'- prob for "{sentance1}" is: {prob1}')
print(f'- prob for "{sentance2}" is: {prob2}')
- prob for "Brad Pitt was born in Oklahoma" is: -inf
- prob for "The actor was born in USA" is: -29.687
print(f'prob for both "{sentance1}" and "{sentance2}" is: {prob1 +
prob2}')
prob for both "Brad Pitt was born in Oklahoma" and "The actor was born
in USA" is: -inf
```

- 1. Now we use linear interpolation smoothing between the bigram model and unigram model with:
- $\lambda$ bigram = 2/3
- $\lambda_{\text{unigram}} = 1/3$

using the same training data.

Given this new model, compute the probability and the perplexity of the same sentences such as in the previous question.

```
elif word in bi model and next word in bi model[word]:
            prob *= (lam bi * np.exp(bi model[word][next word]))
        elif next word in uni model:
            prob *= (lam_uni * np.exp(uni_model[next word]))
        else:
            return -np.inf
    return np.log(prob)
lam bi = 2/3
lam uni = 1/3
# probability:
sents1_prob = calc_sentance_log_prob_interpulation(sentance1,
uni model, bi model, lam bi, lam uni)
sents2_prob = calc_sentance_log_prob_interpulation(sentance2,
uni_model, bi_model, lam_bi, lam_uni)
print(f'- prob for "{sentancel}" is: {sentsl_prob}')
print(f'- prob for "{sentance2}" is: {sents2 prob}')
- prob for "Brad Pitt was born in Oklahoma" is: -36.17630614092038
- prob for "The actor was born in USA" is: -30.99632986192533
# perplexity:
M = len(clean text(sentance1).split(' ') +
clean text(sentance2).split(' ')) - 2
print(f'perplexity in test set is is: {np.exp(-((sents1 prob +
sents2_prob) / M))}')
perplexity in test set is is: 269.81044770279806
```