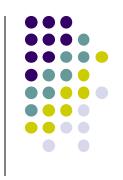


Word2vec and tSNE

https://colab.research.google.com/drive/ 1HJpR8BZh1NK1ugILXgLuWyohQKdx8NZz?usp=sharing

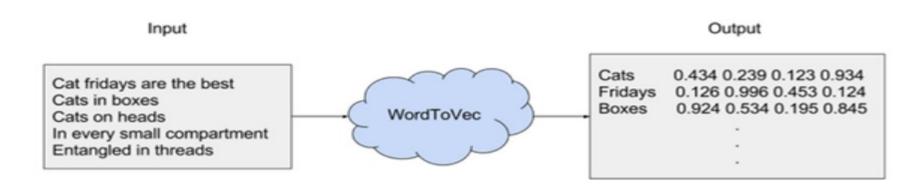




Word2vec can represent words by vectors. These vectors capture contextual sequential information about a word.

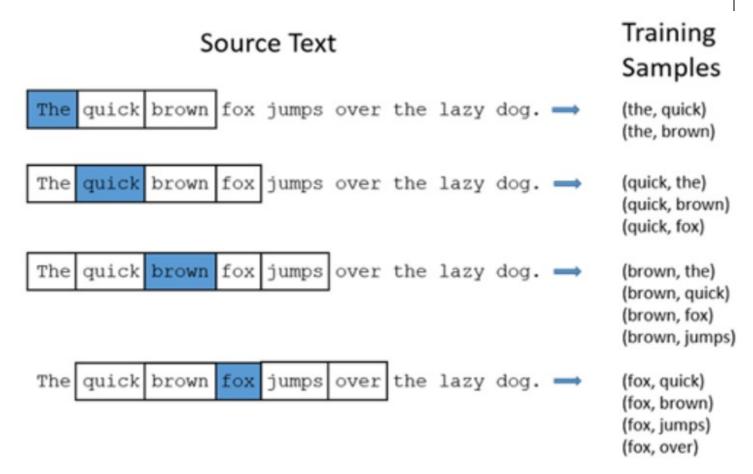
Word2vec tutorial:

https://medium.com/@zafaralibagh6/a-simple-word2vec-tut orial-61e64e38a6a1

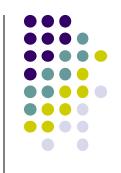


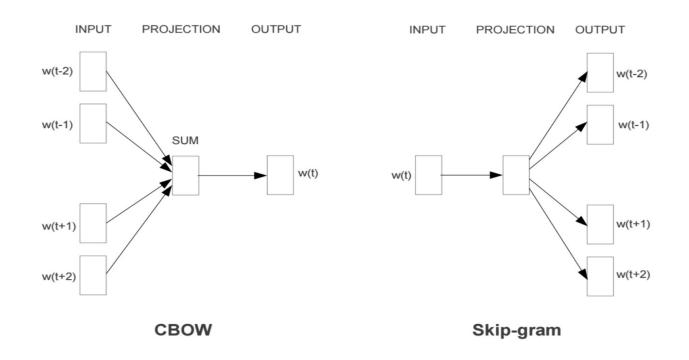


Context and windows





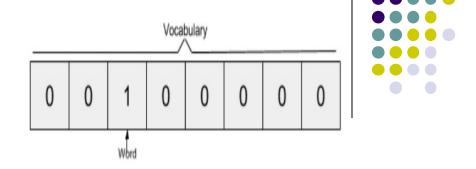




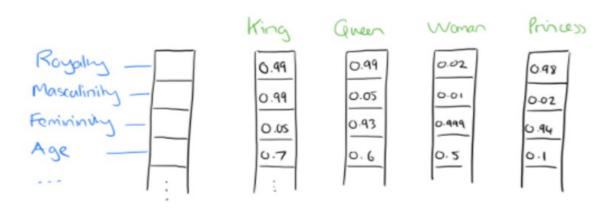
CBOW (Continuous Bag Of Words): using context to predict current word Skip-gram: using word to predict current context

word2vec vector



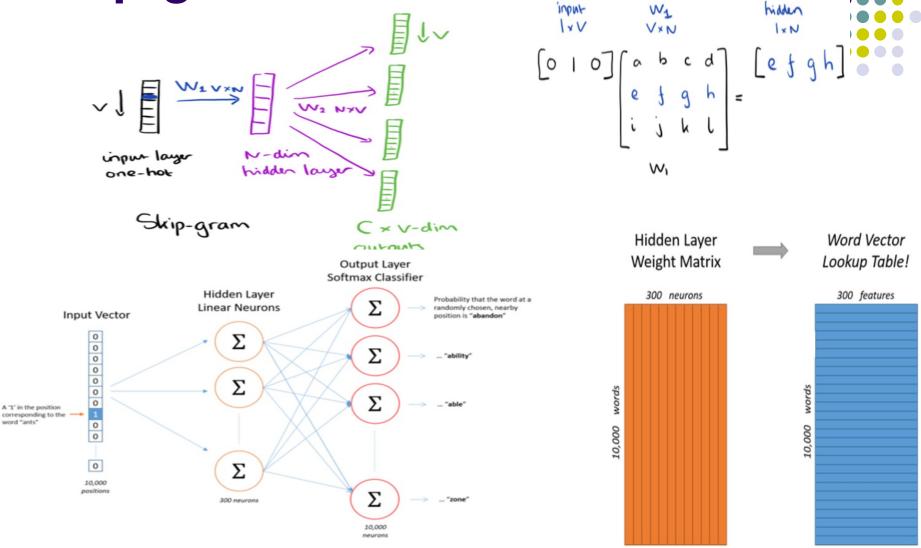


The output of word2vec is a single vector (also with 10,000 components) containing, for every word in our vocabulary, the probability that a randomly selected nearby word is that vocabulary word.



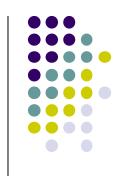
Hypothetical labelone word in blue

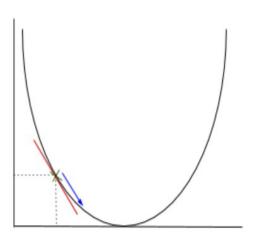
Skip-gram





- we apply stochastic gradient descent to change the values of the weights in order to get a more desirable value for the probability calculated
- In gradient descent we need to calculate the gradient of the function being optimised at the point representing the weight that we are changing. The gradient is then used to choose the direction in which to make a step to move towards the local optimum

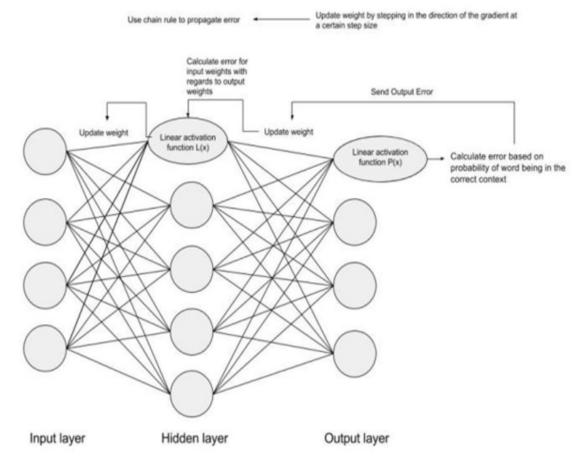








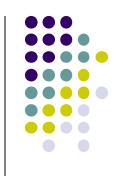
The next step is using Backpropagation, to adjust the weights between multiple layers. The error that is computed at the end of the output layer is passed back from the output layer to the hidden layer by applying the Chain Rule, Gradient descent is used to update the weights between these two layers. The error is then adjusted at each layer and sent back further.





```
!pip install gensim
#!wget -c "http://evexdb.org/pmresources/vec-space-models/PubMed-and-PMC-ri.tar.gz" # this will take some time,
!wget -c "https://s3.amazonaws.com/dl4j-distribution/GoogleNews-vectors-negative300.bin.gz"
import gensim
%matplotlib notebook
import numpy as np
import matplotlib.pyplot as plt
from sklearn.manifold import TSNE
# load pre-trained word2vec embeddings
#model = qensim.models.KeyedVectors.load word2vec format('PubMed-and-PMC-ri.tar.qz', binary=True)
model = gensim.models.KeyedVectors.load word2vec format('GoogleNews-vectors-negative300.bin.gz', binary=True)
Requirement already satisfied: gensim in /usr/local/lib/python3.6/dist-packages (3.6.0)
Requirement already satisfied: scipy>=0.18.1 in /usr/local/lib/python3.6/dist-packages (from gensim) (1.4.1)
Requirement already satisfied: smart-open>=1.2.1 in /usr/local/lib/python3.6/dist-packages (from gensim) (4.0.1)
Requirement already satisfied: six>=1.5.0 in /usr/local/lib/python3.6/dist-packages (from gensim) (1.15.0)
Requirement already satisfied: numpy>=1.11.3 in /usr/local/lib/python3.6/dist-packages (from gensim) (1.19.4)
--2020-12-21 22:34:54-- https://s3.amazonaws.com/dl4j-distribution/GoogleNews-vectors-negative300.bin.gz
Resolving s3.amazonaws.com (s3.amazonaws.com)... 52.217.49.102
Connecting to s3.amazonaws.com (s3.amazonaws.com) | 52.217.49.102 | :443... connected.
HTTP request sent, awaiting response... 416 Requested Range Not Satisfiable
    The file is already fully retrieved; nothing to do.
```





The raw vector for a word

test the loaded word2vec embeddings
print(model['headache'])

```
[ 1.54296875e-01 2.56347656e-02 -1.28906250e-01 3.85742188e-02
-1.74804688e-01
                 1.84570312e-01 -1.37939453e-02 -7.71484375e-02
-7.32421875e-02
                 3.65234375e-01 9.91210938e-02 -3.39843750e-01
 2.89306641e-02 -1.33056641e-02 -1.04980469e-02 4.53125000e-01
 4.98046875e-02 1.75781250e-02 8.15429688e-02 -4.31640625e-01
-7.66601562e-02 1.75781250e-01 2.71484375e-01 7.81250000e-03
-2.11181641e-02 2.22656250e-01 -1.13769531e-01 -2.96875000e-01
-1.46484375e-01 -7.03125000e-02 -5.54199219e-02 -1.59912109e-02
 1.21093750e-01 -1.26953125e-01 -1.09863281e-01 2.64892578e-02
                                 2.69531250e-01
 1.66015625e-01 -1.57226562e-01
                                                 7.17773438e-02
-7.66601562e-02 -3.24218750e-01
                                 2.32421875e-01
                                                 9.46044922e-03
-2.24609375e-01 -7.91015625e-02 6.20117188e-02 1.13769531e-01
```





The words closest to a word

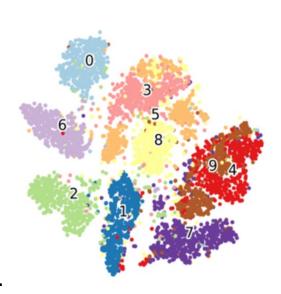
```
# the words closest to a word
model.similar by word('headache')
[('headaches', 0.8292464017868042),
 ('heartburn', 0.574602484703064),
 ('indigestion', 0.5650683641433716),
 ('nightmare', 0.5287497639656067),
 ('nightmares', 0.5257063508033752),
 ('Headaches', 0.5097386837005615),
 ('stomach ache', 0.499220609664917),
 ('problem', 0.4964112341403961),
 ('palpitations', 0.48661914467811584),
 ('earache', 0.4859495460987091)]
```





t-distributed stochastic neighbor embedding

T-distributed Stochastic Neighbor Embedding (t-SNE) is a machine learning algorithm for visualization developed by Laurens van der Maaten and Geoffrey Hinton. It is a nonlinear dimensionality reduction technique well-suited for embedding high-dimensional data for visualization in a low-dimensional space of two or three dimensions. Specifically, it models each high-dimensional object by a two- or three-dimensional point in such a way that similar objects are modeled by nearby points and dissimilar objects are modeled by distant points with high probability.



https://towardsdatascience.com/t-distributed-stochastic-neighbor-embedding-t-sne-bb60ff109561



PCA vs. tSNE

PCA

- •An unsupervised, deterministic algorithm used for feature extraction as well as visualization
- •Applies a linear dimensionality reduction technique where the focus is on keeping the dissimilar points far apart in a lower-dimensional space.
- •Transforms the original data to a new data by preserving the variance in the data using eigenvalues.
- Outliers impact PCA.

t-SNE

- •An unsupervised, randomized algorithm, used only for visualization
- •Applies a non-linear dimensionality reduction technique where the focus is on keeping the very similar data points close together in lower-dimensional space.
- •Preserves the local structure of the data using student t-distribution to compute the similarity between two points in lower-dimensional space.
- •t-SNE uses a heavy-tailed Student-t distribution to compute the similarity between two points in the low-dimensional space rather than a Gaussian distribution, which helps to address the crowding and optimization problems.
- Outliers do not impact t-SNE

Headache in Spacy using word2vec and tSNE



```
import pandas as pd
    pd.options.mode.chained assignment = None
    import numpy as np
    import re
    from gensim.models import word2vec
    from sklearn.manifold import TSNE
    import matplotlib.pyplot as plt
    %matplotlib inline
[2] import spacy
    nlp = spacy.load('en')
[3] # load headache notes
    from google.colab import files
    uploaded = files.upload()
     Browse... notes headache.txt
    notes headache.txt(text/plain) - 9433 bytes, last modified: n/a - 100% done
```

Saving notes headache.txt to notes headache.txt





```
[4] notes = []
    with open ('notes_headache.txt', 'r') as fin:
      lines = fin.readlines()
      for line in lines:
        notes.append(line)
    print(notes)
    print(len(notes))
    ['50 year old female presents after having fallen in t
    11
[5] df=notes
```





```
# Build corpus of all the entities extracted from the notes using spaCy model.
# The corpus is an array of arrays or list of lists where each of the nested lists corresponds to a note.
corpus=[]
for row in range(0, len(df)):
    str_tokens=[]
    tokens= nlp(df[row]).ents
    for i in range(0, len(tokens)):
        str_tokens.append(tokens[i].text)
        corpus.append(list(str_tokens))
```

[['50 year old', 'the bathtub 4 days ago', 'Tylenol', 'the day', 'night', 'morning'], ['23', 'PD', 'RUE', 'F

Get word2vec



- !pip install gensim
- Requirement already satisfied: gensim in /usr/local/lib/python3.6/dist-packages (3.6.0)

 Requirement already satisfied: scipy>=0.18.1 in /usr/local/lib/python3.6/dist-packages (from gensim) (1.4.1)

 Requirement already satisfied: smart-open>=1.2.1 in /usr/local/lib/python3.6/dist-packages (from gensim) (4.0.1)

 Requirement already satisfied: six>=1.5.0 in /usr/local/lib/python3.6/dist-packages (from gensim) (1.15.0)

 Requirement already satisfied: numpy>=1.11.3 in /usr/local/lib/python3.6/dist-packages (from gensim) (1.19.4)

```
[10] import gensim
```

```
[13] model = word2vec.Word2Vec(corpus, min count=1)
```



word2vec

```
model.wv['Tylenol']
```

```
array([ 3.5755248e-03, -7.5043686e-04, -4.0175230e-03, -5.8088248e-04,
      -1.4506313e-03, -1.5477943e-03, 8.4892643e-04, -1.1564353e-03,
       5.8339350e-04, -6.3213066e-04, 4.9648127e-03, 4.2244259e-04,
      -1.5812013e-03, 6.0994085e-04, -3.7646745e-03, 8.8548280e-05,
       2.9612291e-03, 2.5223016e-03, 1.2566439e-03, 3.0257232e-03,
       4.2598490e-03, -3.7888133e-03, 3.4991202e-03, 3.4320420e-03,
       4.6869563e-03, -1.6790380e-03, -4.3789954e-03, 4.1352160e-04,
       2.0200359e-03, 4.0350412e-03, -4.1124565e-03, 2.0835653e-03,
       2.9961050e-03, 2.4687620e-03, 4.6605510e-03, 6.0498028e-04,
       2.3130388e-03, 3.9473213e-03, -3.8562799e-03, 2.5067695e-03,
      -3.9585214e-03, -3.8969840e-03, -1.2115871e-03, -1.3817309e-03,
      -4.6661417e-03, 1.0953289e-03, 4.2884829e-03, 1.8828238e-03,
       2.0013545e-03, -3.4035782e-03, 1.8964872e-03, -2.7806829e-03,
       4.3920926e-03, -2.2318871e-03, -3.1314581e-03, 4.4012973e-03,
       4.0750531e-03, -3.3024163e-03, -1.6775471e-03, -4.1697090e-03,
       4.7896975e-03, 3.7045595e-03, -3.4035568e-06, 1.1239357e-04,
       9.3653216e-05, -4.9044727e-03, -1.8739082e-03, 7.7197549e-04,
      -3.4613409e-03, -1.1089609e-03, -9.2508399e-04, 2.2311143e-03,
      -7.7253534e-04, 5.5203785e-04, -4.1779936e-03, -2.6194218e-03,
```

tSNE



Parameters:

n_components : int, default=2

Dimension of the embedded space.

perplexity: float, default=30.0

The perplexity is related to the number of nearest neighbors that is used in other manifold learning algorithms. Larger datasets usually require a larger perplexity. Consider selecting a value between 5 and 50. Different values can result in significantly different results.

early_exaggeration : float, default=12.0

Controls how tight natural clusters in the original space are in the embedded space and how much space will be between them. For larger values, the space between natural clusters will be larger in the embedded space. Again, the choice of this parameter is not very critical. If the cost function increases during initial optimization, the early exaggeration factor or the learning rate might be too high.

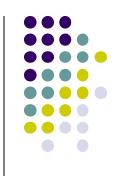
learning_rate : float, default=200.0

The learning rate for t-SNE is usually in the range [10.0, 1000.0]. If the learning rate is too high, the data may look like a 'ball' with any point approximately equidistant from its nearest neighbours. If the learning rate is too low, most points may look compressed in a dense cloud with few outliers. If the cost function gets stuck in a bad local minimum increasing the learning rate may help.

n iter : int. default=1000

https://scikit-learn.org/stable/modules/generated/sklearn.manifold.TSNE.html

tSNE



init: {'random', 'pca'} or ndarray of shape (n_samples, n_components), default='random'

Initialization of embedding. Possible options are 'random', 'pca', and a numpy array of shape (n_samples, n_components). PCA initialization cannot be used with precomputed distances and is usually more globally stable than random initialization.

random_state : int, RandomState instance or None, default=None

Determines the random number generator. Pass an int for reproducible results across multiple function calls. Note that different initializations might result in different local minima of the cost function. See :term:

Glossary <random_state>.

n_iter: int, default=1000

Maximum number of iterations for the optimization. Should be at least 250.



Using tSNE to plot

```
def tsne_plot(model):
    "Creates and TSNE model and plots it"
    labels = []
    tokens = []

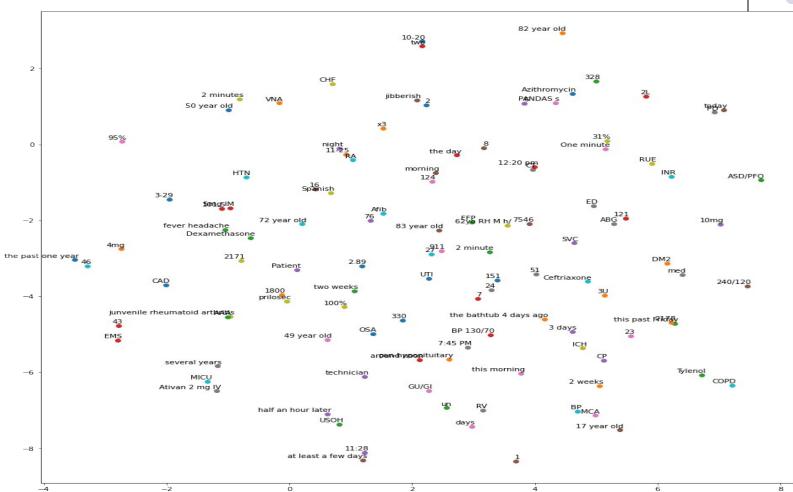
for word in model.wv.vocab:
        tokens.append(model[word])
        labels.append(word)

tsne_model = TSNE(perplexity=30, early_exaggeration=12, n_components=2, init='pca', n_iter=1000, random_state=23)
    new_values = tsne_model.fit_transform(tokens)

x = []
y = []
for value in new_values:
    x.append(value[0])
    y.append(value[1])
```

tsne_plot(model)

tSNE plot



Headache in SciSpacy using word2vec and tSNE



```
!pip install scispacy
                                          51kB 2.7MB/s
 Collecting nmslib>=1.7.3.6
   Downloading https://files.pythonhosted.org/packages/d5/
                                          13.0MB 266kB/s
 Collecting pysbd
   Downloading https://files.pythonhosted.org/packages/26/
                                          71kB 6.6MB/s
 Requirement already satisfied: joblib in /usr/local/lib/p
import scispacy
import spacy
import en ner bc5cdr md
nlp = en ner bc5cdr md.load()
/usr/local/lib/python3.6/dist-packages/spac
 warnings.warn(warn msg)
```

Load notes

```
# load headache notes
from google.colab import files
uploaded = files.upload()
```

Browse... notes_headache.txt
notes_headache.txt(text/plain) - 9433 bytes, last modified: n/a - 100% done
Saving notes headache.txt to notes headache (1).txt

```
notes = []
with open('notes_headache.txt', 'r') as fin:
    lines = fin.readlines()
    for line in lines:
        notes.append(line)
print(notes)
print(len(notes))
['50 year old female presents after having fallen in the bat
```

<

```
] df=notes
```







```
# Build corpus of all the entities extracted from the notes using
# The corpus is an array of arrays or list of lists where each of
corpus=[]
for row in range(0, len(df)):
  str tokens=[]
  tokens= nlp(df[row]).ents
  for i in range(0, len(tokens)):
    str tokens.append(tokens[i].text)
  corpus.append(list(str tokens))
print (corpus)
```

[['headache', 'Tylenol', 'pain', 'dizziness', 'nausea', 'vomiting

Get word2vec

- - !pip install gensim
 - Requirement already satisfied: gensim in /usr/10 Requirement already satisfied: six>=1.5.0 in /us Requirement already satisfied: smart-open>=1.2.1 Requirement already satisfied: scipy>=0.18.1 in Requirement already satisfied: numpy>=1.11.3 in
 - [8] import gensim

```
[10] import pandas as pd
    pd.options.mode.chained assignment = None
     import numpy as np
     import re
     from gensim.models import word2vec
     from sklearn.manifold import TSNE
     import matplotlib.pyplot as plt
     %matplotlib inline
```

[11] model = word2vec.Word2Vec(corpus, min count=1)







```
model.wv['Tylenol']
array([ 0.00421088, 0.00021863, -0.00035665, 0.00322251, 0.00073423,
          -0.00270951, -0.00204165, -0.00214327, 0.0003273, -0.00207853,
          -0.00156233, -0.00453952, 0.00329102, 0.00021155, 0.00089793,
          0.0006254 , 0.0025706 , -0.0007755 , -0.00275109 , -0.0030539 ,
          -0.00065839, 0.00411886, -0.0025017, -0.00214221, -0.0040169,
          0.00014435, -0.00144451, -0.00358639, 0.00494652, 0.00434491,
          -0.0049477 , -0.00241251, 0.00287517, 0.00384575, -0.00263347,
          0.00472971, 0.00027301, 0.002738 , -0.00365123, 0.00489407,
          0.00369412, -0.00314624, -0.00289283, 0.00131375, -0.00289287,
           0.00205069, -0.00407033, 0.00303441, -0.00062189, 0.00429522,
           0.00373649, 0.0030437, -0.00100484, -0.00059962, -0.00244993,
          -0.00459783, 0.00123144, -0.0017654, 0.00143283, -0.0041041,
          -0.00480347, -0.00153869, -0.00160133, 0.00375598, -0.0031069,
          0.0004454 , 0.00474415, -0.00210162, 0.00059001, 0.00245465,
          0.00175198, -0.00054866, 0.00081636, 0.00372542, -0.00212218,
          -0.00335156, 0.00112638, -0.00133766, 0.00219538, 0.00339135,
          0.00437942, 0.00143798, -0.00139074, -0.0019345, -0.00314154,
          -0.00173181, -0.00307571, -0.00368569, -0.00472091, 0.00260101,
```

-0.00292352, 0.00355645, -0.001855 , -0.00047691, 0.00441105, 0.00378569, 0.0006488 , 0.00290273, -0.0008887 , 0.00294835],



tSNE plot

```
def tsne_plot(model):
    "Creates and TSNE model and plots it"
    labels = []
    tokens = []

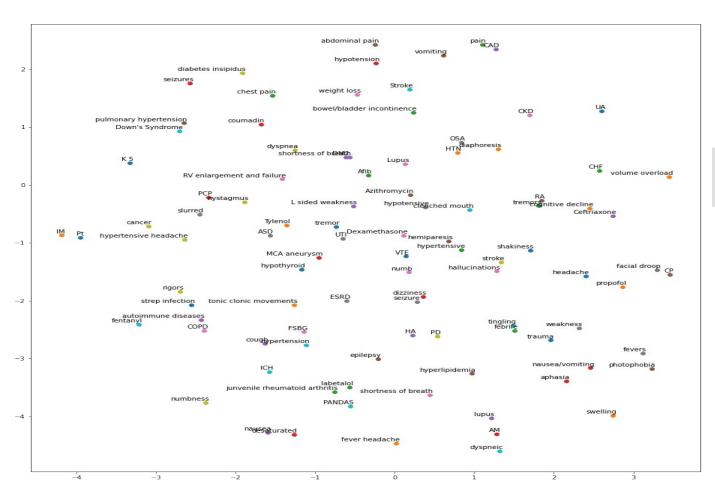
for word in model.wv.vocab:
        tokens.append(model[word])
        labels.append(word)

    tsne_model = TSNE(perplexity=30, early_exaggeration=12, n_components=2, init='pca', n_iter=1000, random_state=23)
    new_values = tsne_model.fit_transform(tokens)

x = []
y = []
for value in new_values:
        x.append(value[0])
        y.append(value[1])
```

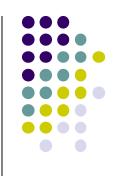






tsne_plot(model)





- Getting headache discharge notes from MIMIC III (around 700 notes)
- Using SciSpacy entity extraction
- Connecting with word2vec
- Plotting them using tSNE



select diagnoses_icd.subject_id, diagnoses_icd.hadm_id, text from noteevents inner join diagnoses_icd on noteevents.hadm_id=diagnoses_icd.hadm_id and noteevents.subject_id=diagnoses_icd.subject_id where diagnoses_icd.ICD9_CODE='430'



https:// querybuilderlcp.mit.edu/

select diagnoses_icd.subject_id, diagnoses_icd.hadm_id, text from noteevents inner join diagnoses_icd on noteevents.hadm_id=diagnoses_icd.hadm_id where diagnoses_icd.ICD9_CODE='430' and noteevents.CATEGORY = 'Discharge summary'

select diagnoses_icd.subject_id, diagnoses_icd.hadm_id, ICD9_CODE from noteevents inner join diagnoses_icd on noteevents.hadm_id=diagnoses_icd.hadm_id where diagnoses_icd.ICD9_CODE='430' and noteevents.CATEGORY = 'Discharge summary'



Thank you!