MIMIC-III NLP

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AI in Healthcare

What Disease Did I Pick?

I selected disease codes related to 4010 – Malignant Essential Hypertension. Malignant essential hypertension is a severe and life-threatening form of high blood pressure that develops rapidly and can cause damage to multiple organs.

What About the Text Data?

The objective of this analysis is to extract medical entities using Named Entity Recognition (NER) with SpaCy, SciSpaCy, Word2Vec, and t-SNE plots.

Additionally, used bc5cdr, BlueBert, MedSpacy to perform a similar analysis.

GitHub and Google Colab Links:

https://colab.research.google.com/github/AnkitaSavaliya/AIH/blob/main/MIMIC_III_NLP.ipynb

https://github.com/AnkitaSavaliya/AIH/blob/main/MIMIC_III_NLP.ipynb

https://github.com/AnkitaSavaliya/AIH/blob/main/MIMIC-III_NLP.pptx

Data Preparation

```
from google.colab import auth
auth.authenticate user()
print('Authenticated')
!gcloud projects list
from google.cloud import bigquery
# Construct a BigQuery client object.
client = bigquery.Client(project='clinical-entity-extraction')
ICD codes related to Hypertension:
 4010 - Malignant essential hypertension
 4011 - Benign essential hypertension
 4019 - Unspecified essential hypertension
# Fetch notes only for ICD-9 code 4010(Malignant essential hypertension)
query = """
  SELECT SUBJECT ID, TEXT, CATEGORY
    FROM `physionet-data.mimiciii notes.noteevents`
    WHERE SUBJECT_ID IN (
        SELECT d.SUBJECT ID
        FROM `physionet-data.mimiciii_clinical.diagnoses_icd` d
        WHERE d.ICD9 CODE = '4010' -- Hypertension code
        AND d.SEO NUM = 1 -- Assuming 1 indicates primary diagnosis
    AND CATEGORY LIKE 'Discharge summary';
# Run the query
query job = client.query(query)
# Print the results
noteevents df = query job.to dataframe()
len(noteevents_df)
```

- Fetched rows from noteevents only for ICD-9 CODE 4010 and category
 'Discharge Summary' using the BigQuery client.
- The query returned 162 rows.
- Prepared a DataFrame with the required columns.
- Saved the query result to a CSV/XLSX file to reduce queries to the database.

```
patients_dict = {"SUBJECT_ID":[],"CATEGORY":[],"TEXT":[]};
for i in range(0, len(noteevents_df)):
    patients_dict["SUBJECT_ID"].append(noteevents_df.loc[i, 'SUBJECT_ID'])
    patients_dict["CATEGORY"].append(noteevents_df.loc[i, 'CATEGORY'])
    patients_dict["TEXT"].append(noteevents_df.loc[i, 'TEXT'])

patients_df = pd.DataFrame(patients_dict)

patients_df.shape

(162, 3)

#print first few records
patients_df.head(2)

# Download the patients_df dataframe in .csv and excel format
patients_df.to_csv(r'Patient_Summary_4010.csv', index = False)
patients_df.to_excel("Patient_Summary_4010.xlsx")
```

Spacy

Extract and Visualize SpaCy Entities

```
import spacy

# Function to clean and extract tokens
def extract_cleaned_text(text, nlp_model):
    doc = nlp_model(str(text))
    tokens = [token.text for token in doc if not token.is_punct and not token.is_space and not token.is_stop]
    return " ".join(tokens) # Return cleaned text as a string

#Load Patient Discharge summary
patients df scapy = pd.read csv("/content/drive/MyDrive/Colab Notebooks/AIH/Patient Summary 4010.csv")
```

```
#Load Patient Discharge summary
patients_df_scapy = pd.read_csv("/content/drive/MyDrive/Colab Notebooks/AIH/Patient_Summary_4010.csv")

# Load the spacy model
nlp_spacy = spacy.load('en_core_web_sm')

# Apply token extraction
patients_df_scapy["Processed_Text"] = patients_df_scapy["TEXT"].apply(lambda text: extract_cleaned_text(text, nlp_spacy))
```

```
from spacy import displacy

# Visualize named entities using displacy
for i in range(0, len(patients_df_scapy)):
    doc = nlp_spacy( patients_df_scapy['Processed_Text'][i])
    displacy.render(doc, style="ent")
```



Word2Vec and t-SNE Visualization Using SpaCy-Processed Data

```
def build corpus(df, model="en core web sm"):
    Extracts named entities from the specified text column in a DataFrame using a spaCy model,
    builds a corpus.
    Parameters:
    - df (pd.DataFrame): DataFrame containing text data.
    - text column (str): Column name containing processed text.
    - model (str): spaCy model to use (default: "en core web sm").
    Returns:
    - corpus (list of lists): Extracted entities per document.
    nlp = model
    corpus = []
    for , row in df.iterrows():
       tokens = [ent.text for ent in nlp(row["Processed Text"]).ents]
        corpus.append(tokens)
    # Calculate word counts
    word counts = [len(doc) for doc in corpus]
    return corpus
```

 Created common function to build corpus using given model SpaCy/SciSpaCy/other

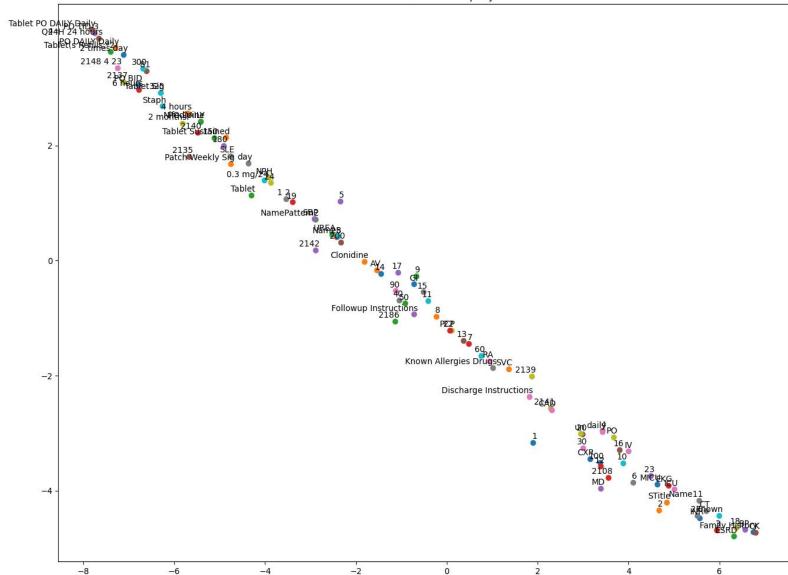
```
from gensim.models import Word2Vec
#Build corpus
corpus spacy = build corpus(patients df scapy, nlp spacy)
model word2vec spacy = Word2Vec(corpus spacy, min count=3, window=2, vector size=100)
model word2vec spacy.wv.similar by key("BP"), model word2vec spacy.wv.similar by key("Clonidine")
([('CT', 0.9996870160102844),
  ('ICU', 0.9996408820152283),
  ('EKG', 0.9996089935302734),
  ('MICU', 0.9996005296707153),
  ('CK', 0.9995817542076111),
  ('Known', 0.9995751976966858),
  ('18', 0.9995639324188232),
  ('27', 0.9995404481887817),
  ('IV', 0.9995347261428833),
  ('INR', 0.999534010887146)],
 [('3', 0.9992740154266357),
  ('100', 0.9992536306381226),
  ('PO', 0.999226987361908),
  ('BP', 0.9992194175720215),
  ('90', 0.9992076754570007),
  ('25', 0.9991985559463501),
  ('EKG', 0.9991586804389954),
  ('30', 0.9991428256034851),
  ('CT', 0.9991336464881897),
  ('18', 0.9991272687911987)])
```

- Defined common function for t-SNE plot.
- Call function using corpus built using Spacy processed text.

```
def tsne plot(model, words, words limit = None, model title="", preTrained=False):
   Creates and displays two t-SNE plots:
   1. Simple scatter plot with labels.
   2. Scatter plot with distance-based coloring.
   Parameters:
   - model: The Word2Vec model or pre-trained model.
   - words: List of words to visualize.
   - words limit : Limit the number of words to visualize.
   - model title: Title of the model.
   - preTrained: Boolean flag to choose between Word2Vec or pre-trained model.
   labels = []
   tokens = []
   # Apply t-SNE for dimensionality reduction
   tsne model = TSNE(perplexity=30, early exaggeration=12, n components=2, init='pca', max iter=1000, random state=23)
   # Prepare tokens and labels
   for word in words[:words limit]:
        if preTrained:
           tokens.append(model[word]) # Pre-trained word vectors
       else:
            tokens.append(model.wv[word]) # Word2Vec model vectors
       labels.append(word)
   tokens = np.array(tokens)
   new values = tsne model.fit transform(tokens)
```

t-SNE Visualization for SpaCy

t-SNE Visualization of Top 100 Words from Word2Vec (SpaCy)



From Word2Vec similarity and above plot we can see that, the entity recognition using SpaCy was limited in extracting hypertension-related terms, likely because it focuses on general English entities rather than clinical ones.

SciSpacy

Extract and Visualize SciSpaCy Entities

patients df SciSpaCy = pd.read csv("/content/drive/MyDrive/Colab Notebooks/AIH/Patient Summary 4010.csv")

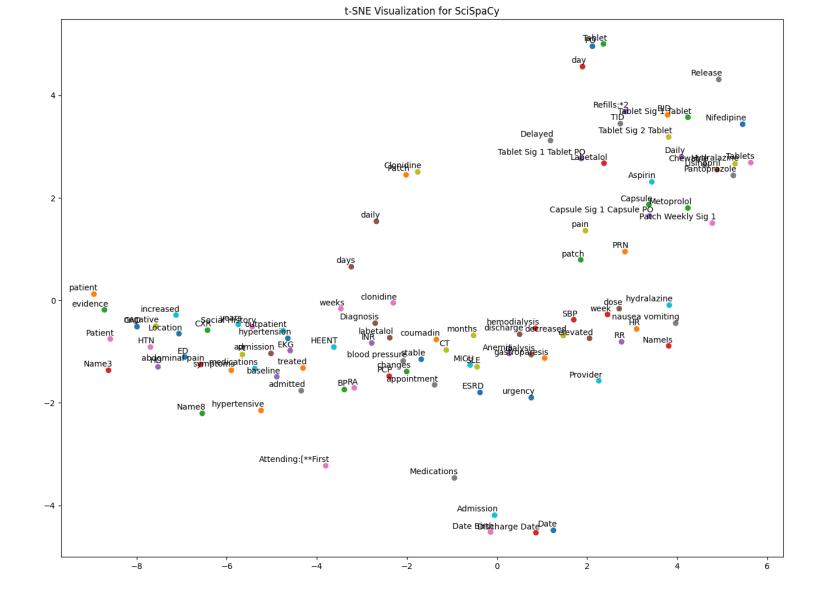
#Load Patient Discharge summary

```
nlp SciSpaCy = spacy.load('en core sci md') # Load the specified NLP model
 # Apply token extraction
 patients df SciSpaCy["Processed Text"] = patients df SciSpaCy["TEXT"].apply(lambda text: extract cleaned text(text, nlp SciSpaCy))
for i in range(0, len(patients df SciSpaCy)):
    doc = nlp SciSpaCy( patients df SciSpaCy['Processed Text'][i])
    displacy.render(doc, style="ent", jupyter=True)
 Admission ENTITY Date 2140 1 19 Discharge Date ENTITY 2140 1 21 Date Birth ENTITY 2117 8 7 Sex F Service ENTITY MEDICINE Allergies Penicillins ENTITY Attending:[**First ENTITY
                                                                                                                                                                              Name3
 ENTITY LF 2297 Chief Complaint headache ENTITY Major Surgical Invasive Procedure Hemodialysis History Present Illness ENTITY Ms. Known lastname ENTITY 22 year old female ENTITY
                                                                                                                                                                            SLE ENTITY
 lupus nephritis ENTITY
                                                                                                      HOCM ENTITY presents HA hypertensive ENTITY
                                       HD ENTITY
                                                     malignant ENTITY
                                                                       HTN ENTITY h/o TTP ENTITY
                                                                                                                                                                       Awoke ENTITY
                                                                                                                                                      urgency ENTITY
         left sided frontal HA sure d/t entity flare uveitis entity started Monday entity d/t HTN entity
                                                                                                      Decided skip ENTITY
                                                                                                                            HD ENTITY come ED ENTITY
                                                                                                                                                           evaluation ENTITY
                                                                                                                                                                               vision
                 numbness weakness ENTITY change gait chest ENTITY pain SOB ENTITY +
                                                                                          Diarrhea ENTITY x 1 day ENTITY
changes ENTITY
                                                                                                                            ED ENTITY
                                                                                                                                         patient ENTITY 217/140 elevated ENTITY 254/152
> received labetolol ENTITY IV 30 mg x 1 MSO4 ENTITY 4 mg pressures ENTITY dropped SBPs ENTITY 208 HA ENTITY improved Repeat labetolol ENTITY 50 mg x 1 repeated dose ENTITY
 morphine ENTITY dropped pressures 193/134 > labetolol ENTITY
                                                             gtt entity started as a given HA entity resolved Head CT entity
                                                                                                                              negative ENTITY
                                                                                                                                                intracranial bleed ENTITY
                                                                                                                                                                         CXR ENTITY
unremarkable ROS cold ENTITY past week fevers chills ENTITY
                                                                                        N/V ENTITY + diarrhea ENTITY
                                                                                                                                        floor patient BP ENTITY 191/126 labetolol ENTITY
                                                            CP ENTITY
                                                                          SOB ENTITY
                                                                                                                        arrival ENTITY
 gtt ENTITY started sxs HA states compliant meds ENTITY
                                                       mother ENTITY
                                                                        cooks salt ENTITY
                                                                                          adherent ENTITY
                                                                                                            diet ENTITY Past Medical History 1 Lupus 2134 ENTITY
                                                                                                                                                                 Diagnosed ENTITY began
swolen fingers rash painful joints 2 ENTITY
                                        ESRD ENTITY
                                                        secodary ENTITY
                                                                         SLE ENTITY 2135 initially cytoxan 1 dose ENTITY 3 months ENTITY 2 years began dialysis ENTITY 3 times week
                                       living donor transplant ENTITY mother 3 HTN ENTITY 2137 Normal BPs ENTITY run 180's/120 1 hypertensive crisis ENTITY
        2137
             T Th Sat Awaiting ENTITY
                                                                                                                                                          precipitated ENTITY
                                                                                                                                                                              seizures
        past 4 Uveitis ENTITY
                               secondary ENTITY
                                                  SLE ENTITY 4 15 5 HOCM ENTITY
                                                                                      Echo entity 2137 6 Vaginal bleeding entity 2139 9 20 7 Mulitple episodes dialysis reactions entity 8
 ENTITY
 Anemia ENTITY 9 Coag neg ENTITY Staph bacteremia ENTITY
                                                               HD line infection ENTITY 6 15 10 H/O UE ENTITY
                                                                                                                            coumadin ENTITY longer Social History ENTITY Lives Location
                                                                                                              clot ENTITY
                                                                  Graduated Name2 NI School ENTITY got sick ENTITY currently working ENTITY attending school Denies ENTITY
 ENTITY 669 mother ENTITY 16 year ENTITY
                                              old brother ENTITY
                                                                                                                                                                           T/E/D. Family
                                   SLE ENTITY -Grandfather HTN ENTITY -Distant history DM ENTITY -No history clotting disorders ENTITY -No history ENTITY
                                                                                                                                                          autoimmune diseases ENTITY
```

Word2Vec and t-SNE Visualization Using SciSpaCy-Processed Data

```
from gensim.models import Word2Vec
corpus scispacy = build corpus(patients df SciSpaCy, nlp SciSpaCy)
model_word2vec_scispacy = Word2Vec(corpus_scispacy, min_count=3, window=2, vector_size=100)
model word2vec scispacy.wv.similar by key("BP"), model word2vec scispacy.wv.similar by key("Clonidine")
([('RA', 0.9994686245918274),
 ('ED', 0.999396562576294),
 ('HR', 0.9993236660957336),
  ('MICU', 0.9991095662117004),
  ('treated', 0.9991006851196289),
  ('patient', 0.9990440011024475),
  ('elevated', 0.998954713344574),
  ('baseline', 0.9989470839500427),
  ('02', 0.9989447593688965),
 ('RR', 0.9989378452301025)],
 [('Patch', 0.9970031380653381),
 ('Prednisone', 0.9962016940116882),
 ('HCl', 0.9951486587524414),
  ('Tablet Sig 1 Tablet PO', 0.9949968457221985),
  ('Labetalol', 0.9949793815612793),
  ('Refills:*0', 0.9945780038833618),
  ('Amlodipine', 0.9940680265426636),
  ('Metoprolol', 0.9939988851547241),
  ('Aspirin', 0.9939936995506287),
  ('Acetaminophen', 0.9935530424118042)])
tsne plot(model word2vec scispacy, np.array(list(model word2vec scispacy.wv.key to index.keys())), 100, 'SciSpaCy')
```

t-SNE Visualization of Top 100 Words from Word2Vec (SciSpaCy)



From Word2Vec similarity and above plot, SciSpaCy primarily recognized medication names and formulations, such as Clonidine and Labetalol, but it did not specifically highlight key hypertension-related entities beyond drug mentions.

BC5CDR (BioCreative V Chemical-Disease Relation)

BC5CDR Entity Visualization Using SciSpaCy-Processed Data

```
nlp__bc5cdr = en_ner_bc5cdr_md.load()

# Visualize named entities using displacy
for i in range(0, len(patients_df_SciSpaCy)):
    doc = nlp__bc5cdr( patients_df_SciSpaCy['Processed_Text'][i])
    displacy.render(doc, style="ent", jupyter=True)
```

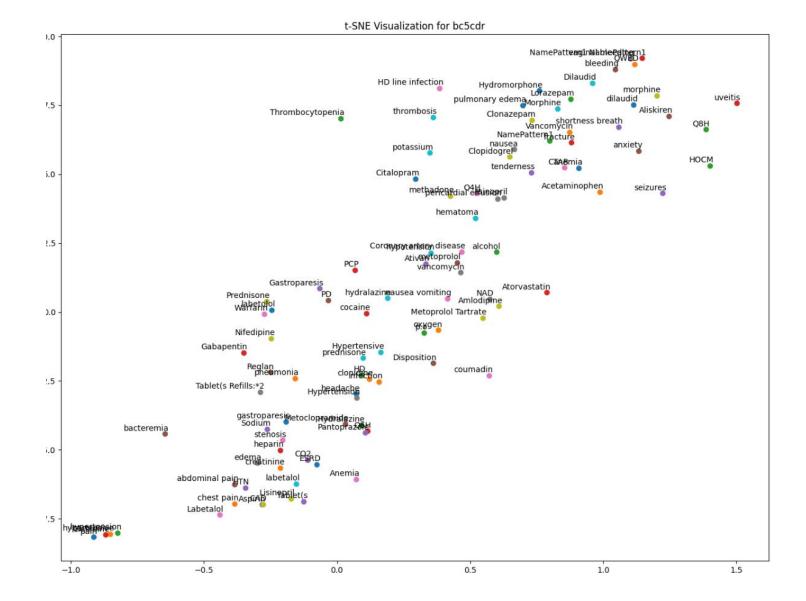
en_ner_bc5cdr_md is a Named Entity Recognition (NER) model from SciSpaCy that specializes in identifying **diseases** and **chemicals** in text



Word2Vec and t-SNE Visualization For BC5CDR

```
from gensim.models import Word2Vec
corpus bc5cdr = build corpus(patients df SciSpaCy, nlp bc5cdr)
model word2vec bc5cdr = Word2Vec(corpus bc5cdr, min count=3, window=2, vector size=100)
model word2vec bc5cdr.wv.similar by word("BP"), model word2vec bc5cdr.wv.similar by word("Clonidine")
([('MSSA bacteremia', 0.8200060129165649),
  ('Metoprolol Succinate', 0.8162233233451843),
  ('dementia', 0.8155735731124878),
  ('fungal infection', 0.8151078820228577),
  ('diabetes brothers diabetes', 0.813332200050354),
  ('lasix', 0.8124308586120605),
  ('Levofloxacin', 0.8123695850372314),
  ('Diabetic ketoacidosis', 0.8123500347137451),
  ('seizure', 0.8114094734191895),
  ('EtOH', 0.810998797416687)],
 [('Labetalol', 0.9987290501594543),
  ('Lisinopril', 0.9986595511436462),
  ('pain', 0.9986243844032288),
  ('hypertensive', 0.9986111521720886),
  ('hypertension', 0.9985666871070862),
  ('HTN', 0.9985529184341431),
  ('Aspirin', 0.9985363483428955),
  ('chest pain', 0.9984893202781677),
  ('Tablet(s', 0.9984655380249023),
  ('Metoclopramide', 0.9984307885169983)])
tsne plot(model word2vec bc5cdr,np.array(list(model word2vec bc5cdr.wv.key to index.keys())), 100, 'bc5cdr')
```

t-SNE Visualization of Top 100 Words from Word2Vec (bc5cdr)



Based on Word2Vec similarity and the above plot, BC5CDR appears to capture disease and medication entities well, with a strong emphasis on hypertension-related terms (e.g., Labetalol, hypertension, headache).

BlueBert

t-SNE Visualization For BlueBert

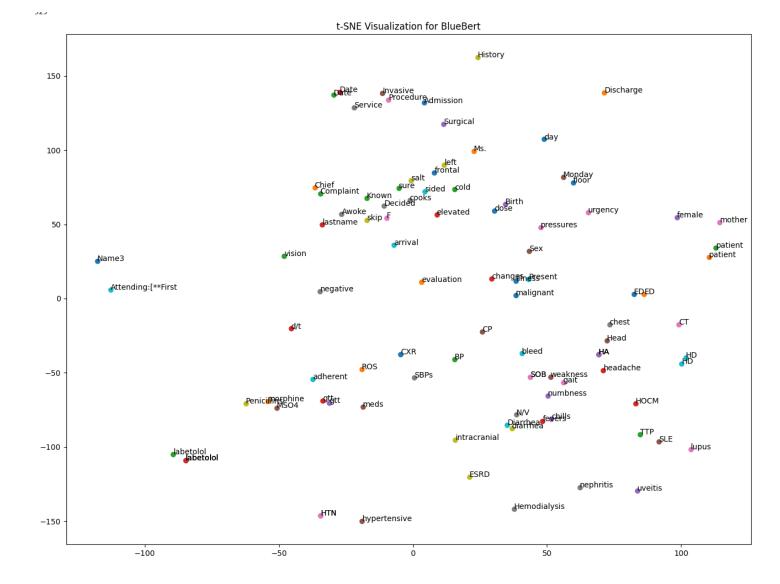
```
# Visualization of notes filtered with SciSpacy using ClinicalBert
import numpy as np
import torch
from sklearn.manifold import TSNE
import string
import matplotlib.pyplot as plt
from transformers import AutoModel, AutoTokenizer, BertModel
# Load the BERT model and tokenizer
model name = "bionlp/bluebert pubmed mimic uncased L-12 H-768 A-12"
tokenizer = AutoTokenizer.from pretrained(model name)
blue bert model = BertModel.from pretrained('bionlp/bluebert pubmed mimic uncased L-12 H-768 A-12')
blue bert model.eval()
# Set first note as text
doc = nlp_SciSpaCy(patients_df_SciSpaCy['Processed_Text'][0])
corpus=
for ent in doc.ents:
    corpus.append(ent.text)
input_text = ' '.join(corpus)
input_tokens = input_text.split()
word embs = []
for token in input tokens:
    # Check if the token is a valid word
    if token not in string.punctuation:
        # Encode the token using the BERT model
       inputs = tokenizer(token, return tensors="pt")
       with torch.no grad():
            outputs = blue bert model(**inputs)
        token emb = outputs.last hidden state.mean(dim=1).squeeze().numpy()
        word embs.append(token emb)
```

- This script utilizes BlueBERT
 (bionlp/bluebert_pubmed_mimic_uncased_L-12_H-768_A-12) to extract word embeddings from clinical notes processed with SciSpaCy.
- Named entities are identified and tokenized, then their embeddings are computed using BlueBERT.
- Only one note was used here because processing all notes with BlueBERT for embedding extraction requires significant time and memory.
- The embeddings are visualized in a 2D space using t-SNE, highlighting relationships among clinical terms.

```
# Perform t-SNE dimensionality reduction
tsne_model = TSNE(n_components=2, perplexity=10, random_state=42)
word_embs_2d = tsne_model.fit_transform(np.array(word_embs))
print(len(word_embs_2d))
# Create a scatter plot of the word embeddings in 2D space
plt.figure(figsize=(16,12))
for i in range(100):
    plt.scatter(word_embs_2d[i, 0], word_embs_2d[i, 1])
    plt.annotate(input_tokens[i], (word_embs_2d[i, 0], word_embs_2d[i, 1]))

plt.title(f"t-SNE Visualization for BlueBert")
plt.show()
```

t-SNE Visualization of Top 100 Words from Word2Vec (BlueBert)



Based on Word2Vec similarity and the above plot, BlueBert appears to capture medical terms(e.g., Labetalol, hypertension, hemodialysis).

MedSpacy

Custom Rule-Based Entity Extraction with MedspaCy NLP Pipeline

```
# Load MedspaCy NLP pipeline
nlp_medspacy = medspacy.load()
# Add rules for target concept extraction
target matcher = nlp medspacy.get pipe("medspacy target matcher")
# Define custom rules for better entity detection
target rules = [
    TargetRule("hyperlipidemia", "DISEASE"),
    TargetRule("02", "CHEMICAL"),
    TargetRule("Fi02", "CHEMICAL"),
    TargetRule("hypertension", "DISEASE"),
    TargetRule("hypertensive urgency", "DISEASE"),
    TargetRule("obesity", "CONDITION"),
    TargetRule("cardiac", "DISEASE"),
    TargetRule("SLE", "DISEASE"),
    TargetRule("lupus nephritis", "DISEASE"),
    TargetRule("ESRD", "DISEASE"),
    TargetRule("dialysis", "TREATMENT"),
    TargetRule("hemodialysis", "TREATMENT"),
    TargetRule("SBP", "MEASUREMENT"),
    TargetRule("HR", "MEASUREMENT"),
    TargetRule("TPN", "TREATMENT"),
    TargetRule("Prednisone", "MEDICATION"),
    TargetRule("Lisinopril", "MEDICATION"),
    TargetRule("Labetalol", "MEDICATION"),
    TargetRule("Clonidine", "MEDICATION"),
    TargetRule("Valsartan", "MEDICATION"),
    TargetRule("Sevelamer", "MEDICATION"),
    TargetRule("Atropine", "MEDICATION"),
    TargetRule("Morphine sulfate", "MEDICATION"),
    TargetRule("Diarrhea", "SYMPTOM"),
    TargetRule("Headache", "SYMPTOM"),
    TargetRule("nausea", "SYMPTOM"),
    TargetRule("vomiting", "SYMPTOM"),
    TargetRule("shortness of breath", "SYMPTOM"),
    TargetRule("fever", "SYMPTOM"),
    TargetRule("chills", "SYMPTOM")
target matcher.add(target rules)
```

- MedSpaCy is a library designed for processing clinical and biomedical text.
- In this code, MedSpaCy is being enhanced by adding custom target rules to better detect specific medical entities such as diseases, treatments, symptoms, and medications in clinical notes.
- Loaded the MedspaCy NLP pipeline. Used the medspacy_target_matcher to add custom rules for extracting medical concepts.
- Defined specific target rules to identify entities like diseases (e.g., hypertension), treatments (e.g., hemodialysis), medications (e.g., Lisinopril), symptoms (e.g., headache), and measurements (e.g., SBP).
- Applied these rules to clinical text for improved entity detection.

MedSpacy Visualization Using SciSpaCy-Processed Data

for i in range(0, len(patients df SciSpaCy)):

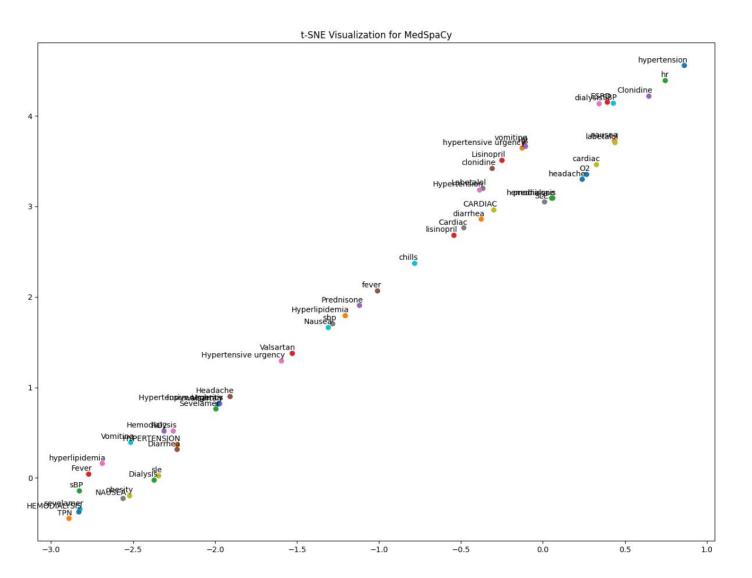
```
# Process the shift note
doc = nlp_medspacy(patients_df_SciSpaCy['Processed_Text'][i])
# visulize
visualize_ent(doc)
```

t-SNE Visualization of Top 100 Words from Word2Vec (MedSpacy)

```
#Build corpus
corpus_medspacy = build_corpus(patients_df_SciSpaCy, nlp_medspacy)

from gensim.models import Word2Vec
model_word2vec_medspacy = Word2Vec(corpus_medspacy, min_count=3, window=2, vector_size=100)

tsne_plot(model_word2vec_medspacy,np.array(list(model_word2vec_medspacy.wv.key_to_index.keys())), 100, 'MedSpaCy')
```



Based on the plot, MedspaCy shows a higher frequency of terms associated with hypertension, indicating that the model is effectively recognizing and extracting a broader range of hypertension-related entities, such as medications, symptoms, and conditions, from the clinical text.

Conclusion

The MIMIC data, especially the free-text notes, contains a lot of shorthand, misspellings, and extra details like dates and measurements that aren't useful for Named Entity Recognition (NER). Pre-trained models like BlueBERT, BC5CDR, and MedSpaCy, tailored for the medical field and charting terminology, tend to extract more relevant and accurate entities in NER than models like SpaCy and SciSpaCy.