

OPEN DATA SCIENCE CONFERENCE



@ODSC

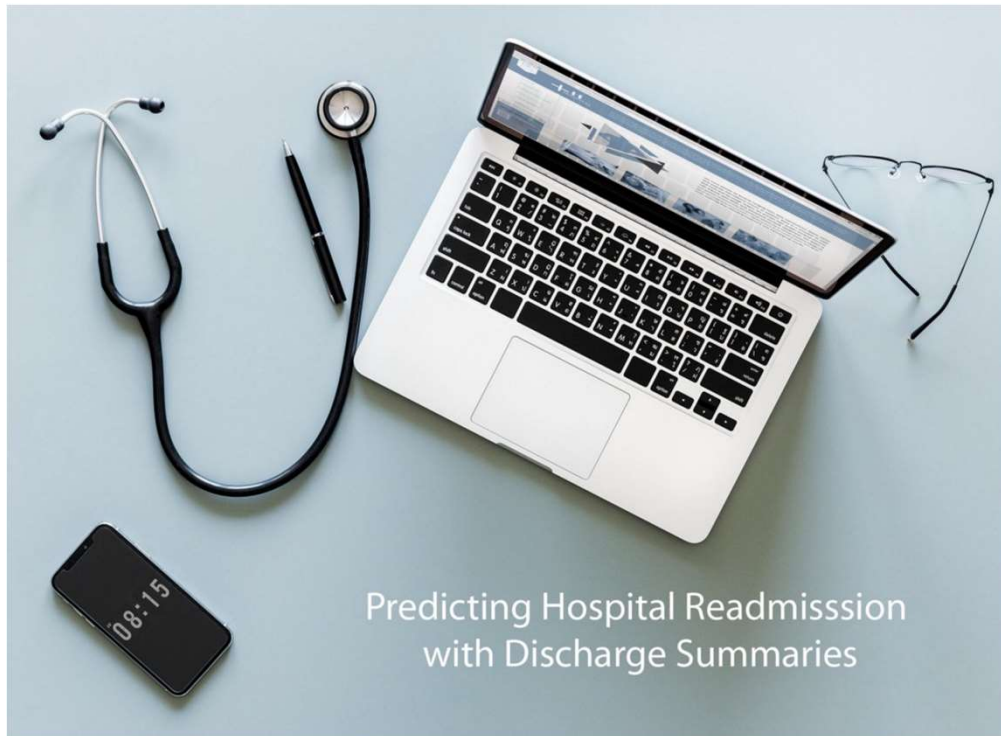
San Francisco | October 31 - Nov. 3 2018



Introduction to Clinical Natural Language Processing

Andrew Long

<https://towardsdatascience.com/introduction-to-clinical-natural-language-processing-predicting-hospital-readmission-with-1736d52bc709>



Predicting Hospital Readmission
with Discharge Summaries

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewlong/odsc_west_2018





FRESENIUS MEDICAL CARE



180,000+
U.S. PATIENTS
SERVED



26M
ANNUAL
HEMODIALYSIS
TREATMENT



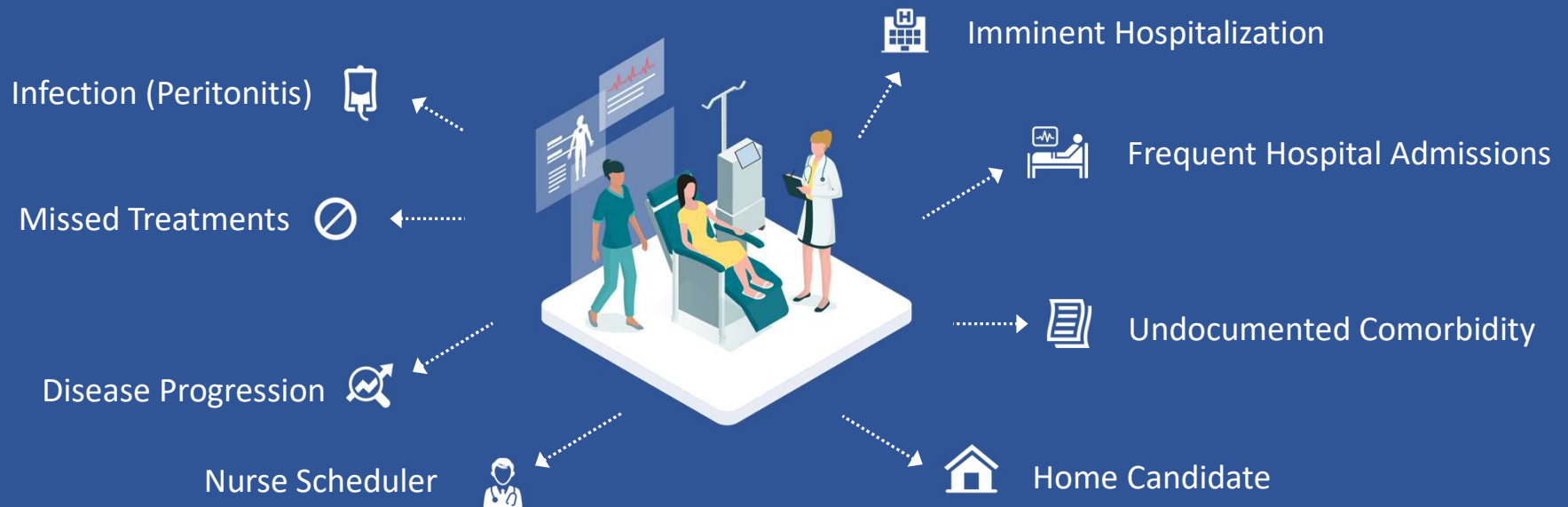
50+
STATES AND
TERRITORIES IN OUR
NETWORK

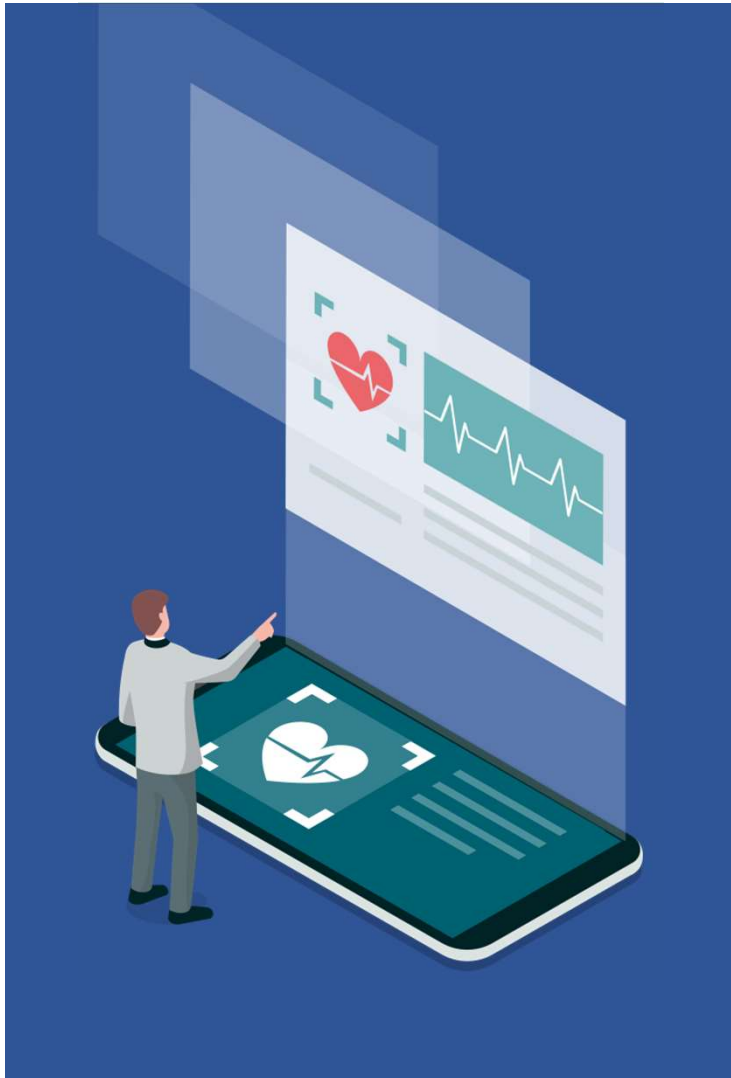


2,200+
U.S. DIALYSIS CLINICS



60,000+
U.S. EMPLOYEES





Clinical Notes

- Chest pain
- Shortness of breathe
- Nausea, vomiting, diarrhea
- Weakness
- Sick
-
-
-

Build predictive models that incorporate free-text clinical notes

Workshop Overview

- Brief overview of clinical dataset (MIMIC III)
 - How to prepare data for a machine learning project
 - How to preprocess the unstructured notes
 - How to build a simple predictive model using a bag-of-words approach
 - How to assess the quality of your model
 - How to decide the next step for improving the model
-
- Note: I created an artificial dataset based on Stanford's IMDB which you can use if you don't have MIMIC access for the workshop

Workshop Project Question

Scalable and accurate deep learning for electronic health records

Rajkomar et al. (paper at <https://arxiv.org/abs/1801.07860>)

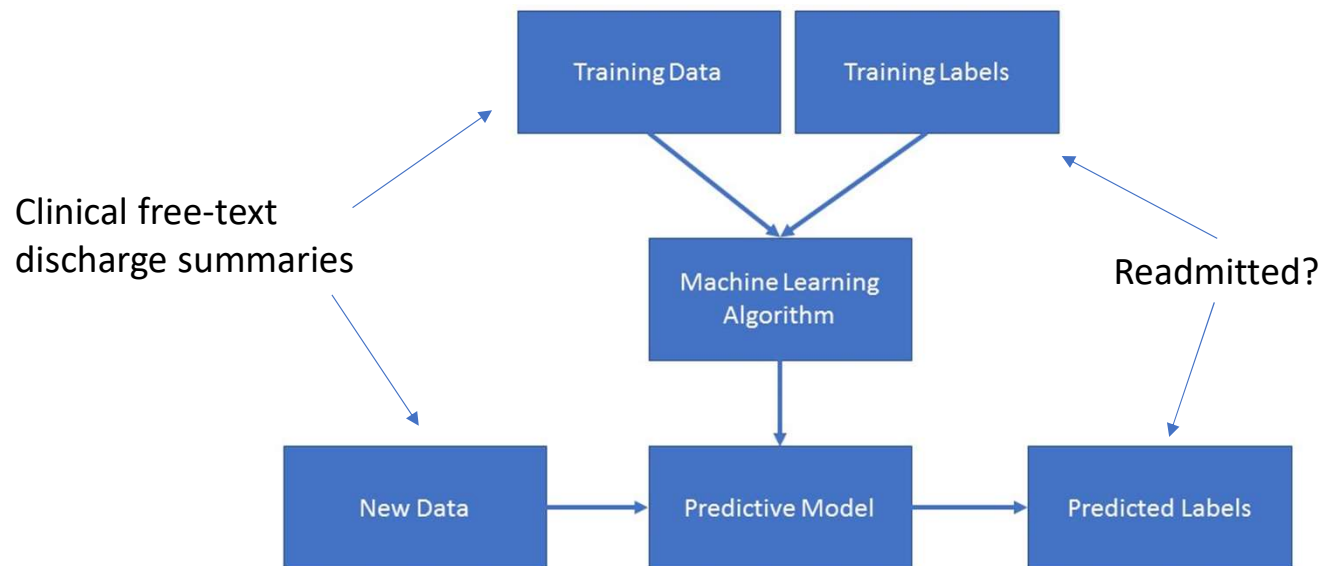
- in-hospital mortality (AUC = 0.93–0.94)
- 30-day unplanned readmission (AUC = 0.75–76)
- prolonged length of stay (AUC = 0.85–0.86)
- discharge diagnoses (AUC = 0.90)

AUC is a data science performance metric where closer to 1 is better

How good of a model can we get if use the discharge free-text summaries with a simple predictive model to predict readmission?

Classification Model Definition

- Predict which patients are at risk for 30-day unplanned readmission utilizing free-text hospital discharge summaries.



Part 0: MIMIC III dataset

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewwlong/odsc_west_2018



MIMIC III dataset

- This database contains de-identified data from over 40,000 patients who were admitted to Beth Israel Deaconess Medical Center in Boston, Massachusetts from 2001 to 2012

Access:

<https://mimic.physionet.org/gettingstarted/access/>

<https://towardsdatascience.com/getting-access-to-mimic-iii-hospital-database-for-data-science-projects-791813feb735>

- Since dataset has restricted access, any single subject data shown in this workshop is artificially created.

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
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MIMIC III datasets for tutorial

<https://physionet.org/works/MIMICIIIClinicalDatabase>

- ADMISSIONS.csv.gz
- NOTEEVENTS.csv.gz
- placed in a 'data' folder in same folder as this workshop's notebook

```
import gzip

for filename in ["data/ADMISSIONS.csv.gz", "data/NOTEVENTS.csv.gz"]:

    with gzip.open(filename, 'rt') as f:
        data = f.read()
    with open(filename[:-3], 'wt') as f:
        f.write(data)
```

1. [checksum_md5_zipped.txt](#) (MD5 checksum for zipped files)
2. [checksum_md5_unzipped.txt](#) (MD5 checksum for unzipped files)
3. [ADMISSIONS.csv.gz](#) (2.5M compressed, 12M decompressed)
4. [CALLOUT.csv.gz](#) (1.2M compressed, 6.1M decompressed)
5. [CAREGIVERS.csv.gz](#) (49K compressed, 199K decompressed)
6. [CHARTEVENTS.csv.gz](#) (4.0G compressed, 33G decompressed)
7. [CPTEVENTS.csv.gz](#) (4.8M compressed, 56M decompressed)
8. [DATETIMEEVENTS.csv.gz](#) (53M compressed, 502M decompressed)
9. [DIAGNOSES_ICD.csv.gz](#) (4.5M compressed, 19M decompressed)
10. [DRGCODES.csv.gz](#) (1.7M compressed, 11M decompressed)
11. [D_CPT.csv.gz](#) (3.9K compressed, 14K decompressed)
12. [D_ICD_DIAGNOSES.csv.gz](#) (279K compressed, 1.4M decompressed)
13. [D_ICD_PROCEDURES.csv.gz](#) (75K compressed, 305K decompressed)
14. [D_ITEMS.csv.gz](#) (184K compressed, 933K decompressed)
15. [D_LABITEMS.csv.gz](#) (12K compressed, 43K decompressed)
16. [ICUSTAYS.csv.gz](#) (1.9M compressed, 6.1M decompressed)
17. [INPUTEVENTS_CV.csv.gz](#) (403M compressed, 2.3G decompressed)
18. [INPUTEVENTS_MV.csv.gz](#) (144M compressed, 931M decompressed)
19. [LABEVENTS.csv.gz](#) (321M compressed, 1.8G decompressed)
20. [MICROBIOLOGYEVENTS.csv.gz](#) (7.3M compressed, 70M decompressed)
21. [NOTEVENTS.csv.gz](#) (1.1G compressed, 3.8G decompressed)
22. [OUTPUTEVENTS.csv.gz](#) (56M compressed, 379M decompressed)
23. [PATIENTS.csv.gz](#) (559K compressed, 2.6M decompressed)
24. [PRESCRIPTIONS.csv.gz](#) (99M compressed, 735M decompressed)
25. [PROCEDUREEVENTS_MV.csv.gz](#) (7.5M compressed, 47M decompressed)
26. [PROCEDURES_ICD.csv.gz](#) (1.8M compressed, 6.5M decompressed)
27. [SERVICES.csv.gz](#) (1.2M compressed, 3.4M decompressed)
28. [TRANSFERS.csv.gz](#) (5.3M compressed, 24M decompressed)

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewwlong/odsc_west_2018



Part 1: How to prepare data for a machine learning project

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewlong/odsc_west_2018



Load, clean, merge dataset

```
In [2]: import pandas as pd
import numpy as np
import matplotlib.pyplot as plt

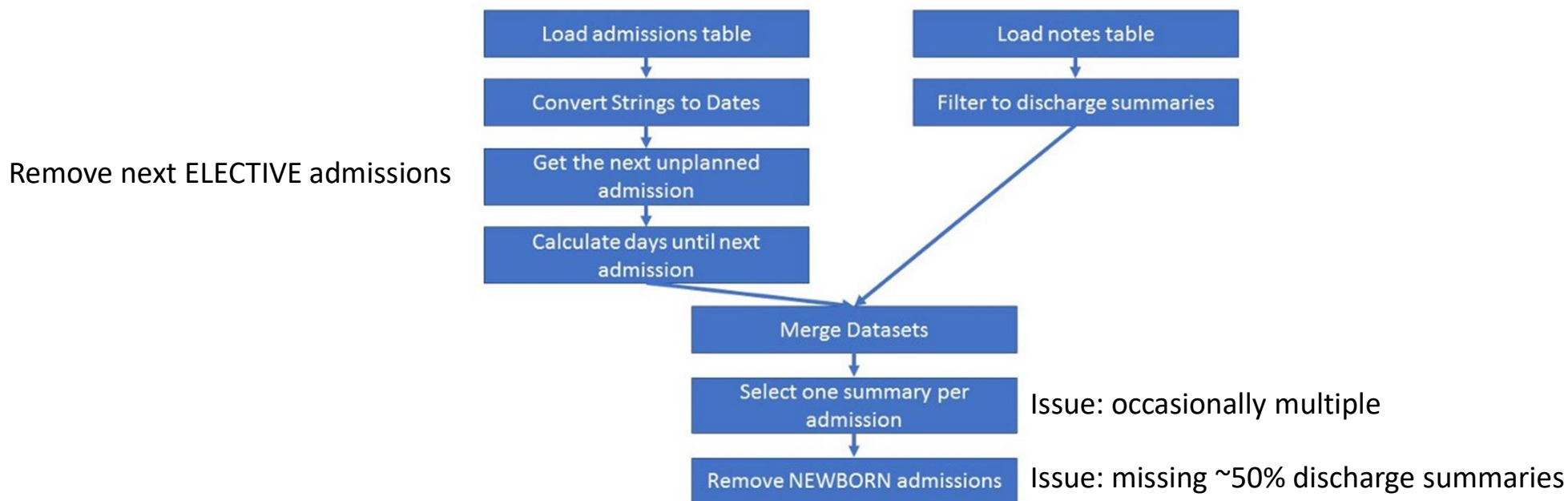
import odsc2018_utils
|
df_adm_notes_clean = odsc2018_utils.load_clean_merge_dataset('data/ADMISSIONS.csv', 'data/NOTEEVENTS.csv')

C:\Users\3236283\AppData\Local\Continuum\Anaconda3\lib\site-packages\IPython\core\interactiveshell.py:2910:
ing: Columns (4,5) have mixed types. Specify dtype option on import or set low_memory=False.
exec(code_obj, self.user_global_ns, self.user_ns)
```

We skip this process to save some time In workshop.

See additional Jupyter Notebook (odsc_2018_mimic_pre) for tutorial on these steps.

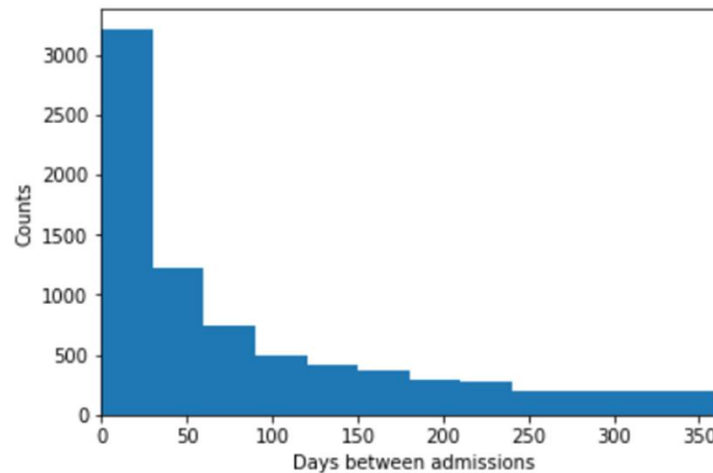
Prepare data for ML Project



df_adm_notes_clean

- SUBJECT_ID – unique patient identifier
- HADM_ID- unique admission identifier
- ADMITTIME – admission date
- DISCHTIME – discharge date
- DEATHTIME – death date
- DAYS_NEXT_ADMIT – days until next admission if it exists
- TEXT – discharge summary for this admission

Add OUTPUT_LABEL



```
In [3]: df_adm_notes_clean['OUTPUT_LABEL'] = (df_adm_notes_clean.DAYS_NEXT_ADMIT < 30).astype('int')
```

```
In [4]: print('Number of positive samples:', (df_adm_notes_clean.OUTPUT_LABEL == 1).sum())
print('Number of negative samples:', (df_adm_notes_clean.OUTPUT_LABEL == 0).sum())
print('Total samples:', len(df_adm_notes_clean))
```

```
Number of positive samples: 3004
Number of negative samples: 48109
Total samples: 51113
```

Make training/validation/test sets



- Training samples: these samples are used to train the model
- Validation samples: these samples are held out from the training data and are used to make decisions on how to improve the model
- Test samples: these samples are held out from all decisions and are used to measure the generalized performance of the model

Make training/validation/test sets

70% Training

15% Validation

15% Test

70/15/15 is a design choice

```
In [5]: # shuffle the samples
df_adm_notes_clean = df_adm_notes_clean.sample(n = len(df_adm_notes_clean), random_state = 42)
df_adm_notes_clean = df_adm_notes_clean.reset_index(drop = True)

# Save 30% of the data as validation and test data
df_valid_test=df_adm_notes_clean.sample(frac=0.30,random_state=42)

df_test = df_valid_test.sample(frac = 0.5, random_state = 42)
df_valid = df_valid_test.drop(df_test.index)

# use the rest of the data as training data
df_train_all=df_adm_notes_clean.drop(df_valid_test.index)
```

Make training/validation/test sets

70% Training

15% Validation

15% Test

70/15/15 is a design choice

Verify that positive prevalence is approximately the same in the 3 groups

```
In [6]: print('Test prevalence(n = %d):%.3f'%(len(df_test),df_test.OUTPUT_LABEL.sum()/ len(df_test)))
print('Valid prevalence(n = %d):%.3f'%(len(df_valid),df_valid.OUTPUT_LABEL.sum()/ len(df_valid)))
print('Train all prevalence(n = %d):%.3f'%(len(df_train_all), df_train_all.OUTPUT_LABEL.sum()/ len(df_train_all)))
print('all samples (n = %d)'%len(df_adm_notes_clean))
assert len(df_adm_notes_clean) == (len(df_test)+len(df_valid)+len(df_train_all)), 'math didnt work'
```

```
Test prevalence(n = 7667):0.062
Valid prevalence(n = 7667):0.057
Train all prevalence(n = 35779):0.058
all samples (n = 51113)
```

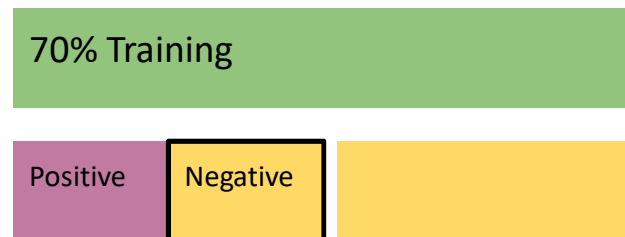
```
In [7]: df_train_all.to_csv('data/df_train_all.csv',index=False)
df_valid.to_csv('data/df_valid.csv',index=False)
df_test.to_csv('data/df_test.csv',index=False)
```

Artificial Dataset

- For those without MIMIC data, I created an artificial dataset based on IMDB dataset.
 - data_artificial/df_train_all_imdb.csv
 - data_artificial/df_valid_imdb.csv
 - data_artificial/df_test_imdb.csv
-
- validation and test sets were created to have approximately same number and prevalence as the MIMIC sets

Imbalanced Classification

- Model that always guesses 'Not readmitted' → 94% accuracy, but never catches any readmissions (0% recall)
- To prevent this from happening, we need to balance the training set
 - **sub-sample the more dominant class: use a random subset of the negatives**
 - over-sample the imbalanced class: use the same positive samples multiple times
 - create synthetic positive data



Subsample Training Dataset

```
In [9]: # split the training data into positive and negative
rows_pos = df_train_all.OUTPUT_LABEL == 1
df_train_pos = df_train_all.loc[rows_pos]
df_train_neg = df_train_all.loc[~rows_pos]

n = np.min([len(df_train_pos), len(df_train_neg)])

# merge the balanced data
df_train = pd.concat([df_train_pos.sample(n = n, random_state = 42), \
                      df_train_neg.sample(n = n, random_state = 42)], axis = 0)

# shuffle the order of training samples
df_train = df_train.sample(n = len(df_train), random_state = 42).reset_index(drop = True)

print('Train prevalence (n = %d):' % len(df_train), df_train.OUTPUT_LABEL.sum() / len(df_train))

Train prevalence (n = 4184): 0.5
```

70% Training

Positive

Negative

Part 2: How to preprocess the unstructured notes

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewlong/odsc_west_2018



Pre-process Text Data

- Occasionally, need to modify the text to make useable (for example drop newlines, carriage returns, numbers, etc)
- Two Methods:
 - Modify the original dataframe TEXT column
 - Pre-process as part of the pipeline

Modify Original Text

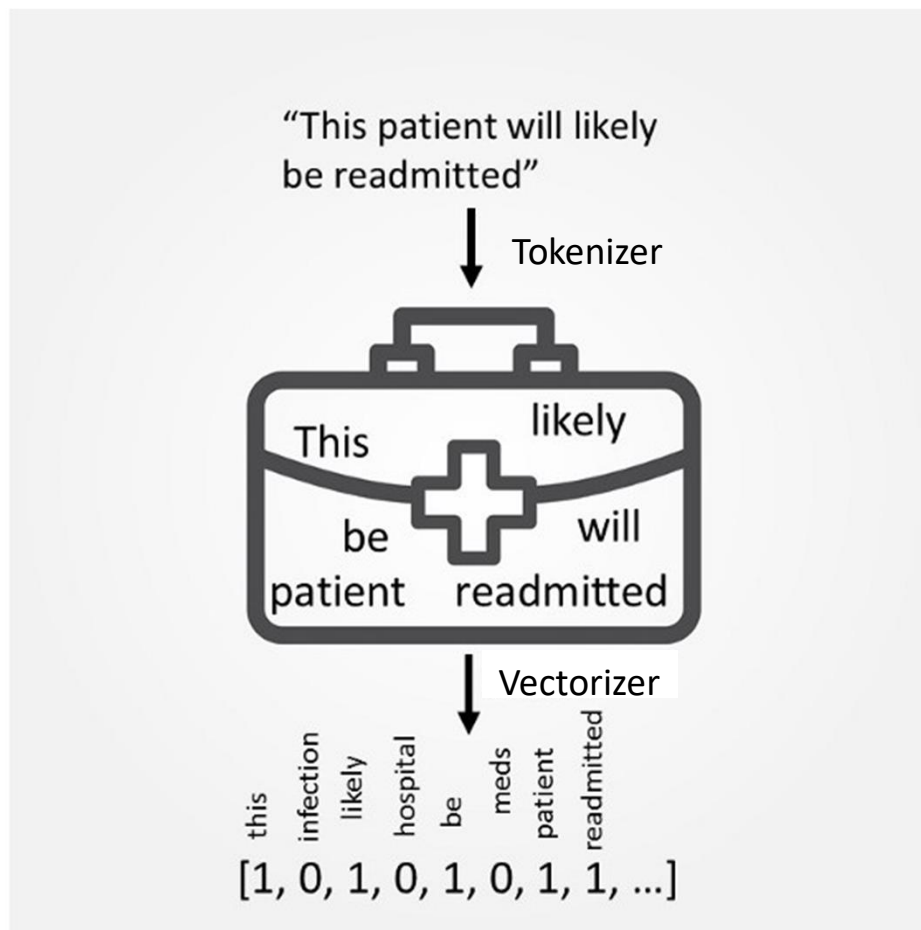
Drop newline, carriage returns
Replace missing notes with a space

```
In [12]: def preprocess_text(df):  
        # This function preprocesses the text by filling  
        df.TEXT = df.TEXT.fillna(' ')  
        df.TEXT = df.TEXT.str.replace('\n', ' ')  
        df.TEXT = df.TEXT.str.replace('\r', ' ')  
        return df
```

```
In [13]: # preprocess the text to deal with known issues  
df_train = preprocess_text(df_train)  
df_valid = preprocess_text(df_valid)  
df_test = preprocess_text(df_test)
```


Bag-of-words

- Split a note into tokens (i.e. words) then 'count' the number of each token
- Use these 'counts' as feature columns
- Note: different techniques for 'counts'



Bag-of-words design choices

- How to preprocess the words into tokens
- How to count the tokens
- Which tokens to use

Build a tokenizer

```
In [14]: import nltk
          from nltk import word_tokenize
          word_tokenize('This should be tokenized. 11/01/2018 sentence has stars**')

Out[14]: ['This',
          'should',
          'be',
          'tokenized',
          '.',
          '11/01/2018',
          'sentence',
          'has',
          'stars**']
```

- Sentence is tokenized by spaces and some punctuation but not all.
- Numbers are also still included
- 'This' would be considered a different token than 'this'
- 'stars**' would be different than 'stars'

Build a custom tokenizer

- Replace punctuation with spaces
- Replace numbers with spaces
- Lowercase all words

```
In [15]: import string
print(string.punctuation)

!"#$%&'()*+,-./:;<=>?@[\\]^_`{|}~
```

```
In [16]: def tokenizer_better(text):
# tokenize the text by replacing punctuation and numbers with spaces and lowercase all words

punc_list = string.punctuation+'0123456789'
t = str.maketrans(dict.fromkeys(punc_list, " "))
text = text.lower().translate(t)
tokens = word_tokenize(text)
return tokens
```

Fast way to replace characters with spaces

```
In [17]: tokenizer_better('This should be tokenized. 11/01/2018 sentence has stars**')
```

```
Out[17]: ['this', 'should', 'be', 'tokenized', 'sentence', 'has', 'stars']
```

Build a simple vectorizer

```
In [18]: sample_text = ['Open Data Science Conference is about learning',  
                        'Data data DATA',  
                        'Learning is part of data science']
```

```
In [19]: from sklearn.feature_extraction.text import CountVectorizer  
vect = CountVectorizer(tokenizer = tokenizer_better)  
vect.fit(sample_text)  
  
# matrix is stored as a sparse matrix (since you have a lot of zeros)  
X = vect.transform(sample_text)
```

Specify custom tokenizer



- CountVectorizer is the simplest method for bag-of-words
- Counts the number of occurrences of each word
- Other common method is TfidfVectorizer which takes into account frequency of word usage across notes

Build a simple vectorizer

```
In [18]: sample_text = ['Open Data Science Conference is about learning',  
                        'Data data DATA',  
                        'Learning is part of data science']
```

```
In [19]: from sklearn.feature_extraction.text import CountVectorizer  
vect = CountVectorizer(tokenizer = tokenizer_better)  
vect.fit(sample_text)  
  
# matrix is stored as a sparse matrix (since you have a lot of zeros)  
X = vect.transform(sample_text)
```

```
In [20]: X
```

```
Out[20]: <3x9 sparse matrix of type '<class 'numpy.int64'>'  
         with 14 stored elements in Compressed Sparse Row format>
```

```
In [21]: # we can visualize this small example if we convert it to an array  
X.toarray()
```

```
Out[21]: array([[1, 1, 1, 1, 1, 0, 1, 0, 1],  
                [0, 0, 3, 0, 0, 0, 0, 0, 0],  
                [0, 0, 1, 1, 1, 1, 0, 1, 1]], dtype=int64)
```

```
: # get the column names  
vect.get_feature_names()  
  
: ['about',  
  'conference',  
  'data',  
  'is',  
  'learning',  
  'of',  
  'open',  
  'part',  
  'science']
```

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewwlong/odsc_west_2018



Train clinical vectorizer

```
In [23]: from sklearn.feature_extraction.text import CountVectorizer
vect = CountVectorizer(max_features = 3000, tokenizer = tokenizer_better)

# this could take a while
vect.fit(df_train.TEXT.values)
```

```
Out[23]: CountVectorizer(analyzer='word', binary=False, decode_error='strict',
dtype=<class 'numpy.int64'>, encoding='utf-8', input='content',
lowercase=True, max_df=1.0, max_features=3000, min_df=1,
ngram_range=(1, 1), preprocessor=None, stop_words=None,
strip_accents=None, token_pattern='(?u)\\b\\w\\w+\\b',
tokenizer=<function tokenizer_better at 0x000001CA1DA7A400>,
vocabulary=None)
```

Good practice to specify the max_features (otherwise it could take a long time with big data set)
Size of max_features is then a hyperparameter to tune

Stop words

“the”, “is”, “are”, “and”

- Stop word – commonly used words with little value to ML model
- Frequency of word use depends on domain (clinical, twitter, Wikipedia)

Stop words

<https://www.linkedin.com/pulse/another-twitter-sentiment-analysis-python-part-2-ricky-kim/>

```
In [24]: neg_doc_matrix = vect.transform(df_train[df_train.OUTPUT_LABEL == 0].TEXT)
pos_doc_matrix = vect.transform(df_train[df_train.OUTPUT_LABEL == 1].TEXT)
neg_tf = np.sum(neg_doc_matrix,axis=0)
pos_tf = np.sum(pos_doc_matrix,axis=0)
neg = np.squeeze(np.asarray(neg_tf))
pos = np.squeeze(np.asarray(pos_tf))

term_freq_df = pd.DataFrame([neg,pos],columns=vect.get_feature_names()).transpose()
term_freq_df.columns = ['negative', 'positive']
term_freq_df['total'] = term_freq_df['negative'] + term_freq_df['positive']
term_freq_df.sort_values(by='total', ascending=False).iloc[:10]
```

Out[24]:

	negative	positive	total
the	71054	76756	147810
and	62455	71658	134113
to	53226	62085	115311
of	51303	60491	111794
was	48074	53521	101595
with	38036	44583	82619
a	35428	41629	77057
on	32290	39765	72055
mg	27718	39045	66763
in	29567	34755	64322

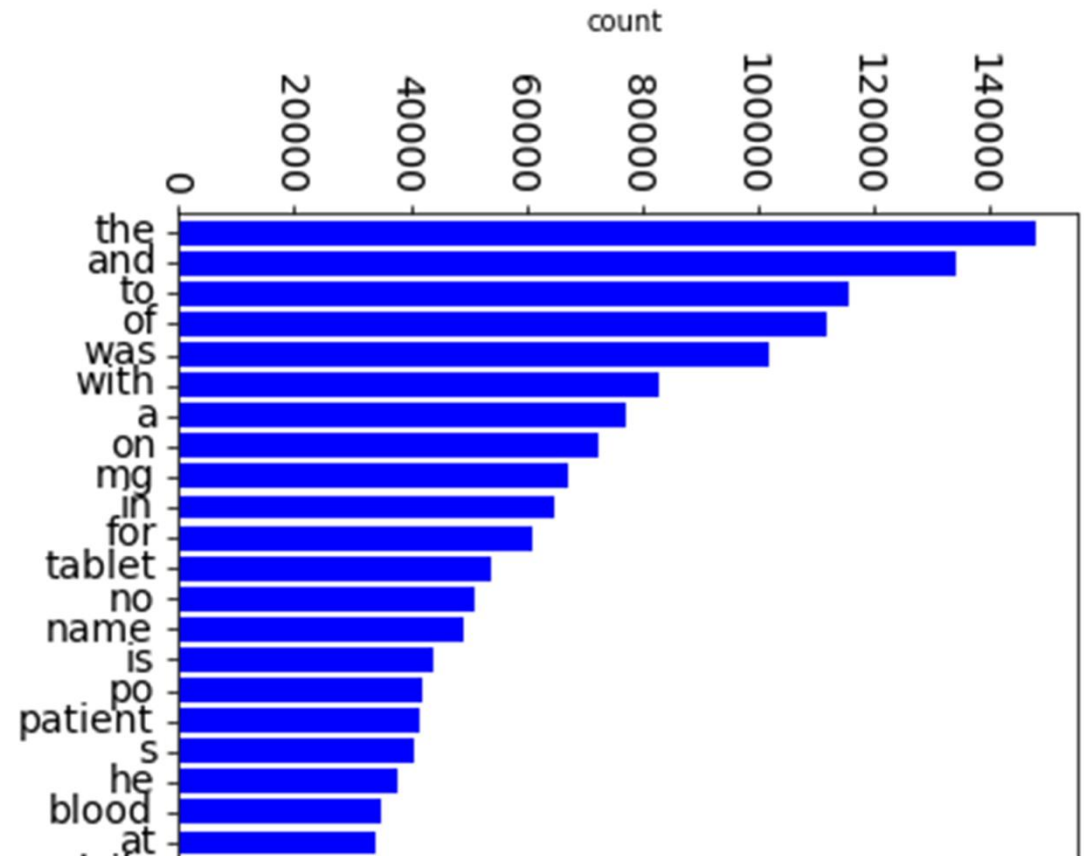
More medications predictive of readmission?

Fast technique for
finding term frequency

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewwlong/odsc_west_2018



Stop words



```
In [26]: my_stop_words = ['the', 'and', 'to', 'of', 'was', 'with', 'a', 'on', 'in', 'for', 'name',  
                          'is', 'patient', 's', 'he', 'at', 'as', 'or', 'one', 'she', 'his', 'her', 'am',  
                          'were', 'you', 'pt', 'pm', 'by', 'be', 'had', 'your', 'this', 'date',  
                          'from', 'there', 'an', 'that', 'p', 'are', 'have', 'has', 'h', 'but', 'o',  
                          'namepattern', 'which', 'every', 'also']
```

Build a vectorizer removing stop words

```
In [27]: from sklearn.feature_extraction.text import CountVectorizer
vect = CountVectorizer(max_features = 3000,
                      tokenizer=tokenizer_better,
                      stop_words = my_stop_words)
# this could take a while
vect.fit(df_train.TEXT.values)
```

```
Out[27]: CountVectorizer(analyzer='word', binary=False, decode_error='strict',
                        dtype=<class 'numpy.int64'>, encoding='utf-8', input='content',
                        lowercase=True, max_df=1.0, max_features=3000, min_df=1,
                        ngram_range=(1, 1), preprocessor=None,
                        stop_words=['the', 'and', 'to', 'of', 'was', 'with', 'a', 'on', 'in', 'for', 'name', 'is', 'patient', 's', 'h
e', 'at', 'as', 'or', 'one', 'she', 'his', 'her', 'am', 'were', 'you', 'pt', 'pm', 'by', 'be', 'had', 'your', 'this',
'date', 'from', 'there', 'an', 'that', 'p', 'are', 'have', 'has', 'h', 'but', 'o', 'namepattern', 'which', 'every', '
also'],
                        strip_accents=None, token_pattern='(?u)\\b\\w\\w+\\b',
                        tokenizer=<function tokenizer_better at 0x000001CA1DA7A400>,
                        vocabulary=None)
```

Create X, y

```
In [28]: X_train_tf = vect.transform(df_train.TEXT.values)
X_valid_tf = vect.transform(df_valid.TEXT.values)
```

Get labels

```
In [29]: y_train = df_train.OUTPUT_LABEL
y_valid = df_valid.OUTPUT_LABEL
```

Part 3: How to build a simple predictive model using a bag-of-words approach

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewlong/odsc_west_2018



Logistic Regression

- Traditional Machine Learning algorithm
- Works well with sparse matrices
- Fast to train
- Interpretable

Logistic Regression

```
In [30]: # logistic regression
from sklearn.linear_model import LogisticRegression
clf=LogisticRegression(C = 0.0001, penalty = 'l2', random_state = 42)
clf.fit(X_train_tf, y_train)

C:\Users\Andy\AppData\Local\conda\conda\envs\odsc_west_2018\lib\site-packag
utureWarning: Default solver will be changed to 'lbfgs' in 0.22. Specify a
FutureWarning)

Out[30]: LogisticRegression(C=0.0001, class_weight=None, dual=False,
    fit_intercept=True, intercept_scaling=1, max_iter=100,
    multi_class='warn', n_jobs=None, penalty='l2', random_state=42,
    solver='warn', tol=0.0001, verbose=0, warm_start=False)
```

Hyperparameter C helps control the effect of regularization

We will discuss how to optimize C

Tip: the same C usually doesn't work as well for both CountVectorizer and TfidfVectorizer

Predictions

```
In [31]: model = clf  
y_train_preds = model.predict_proba(X_train_tf)[: ,1]  
y_valid_preds = model.predict_proba(X_valid_tf)[: ,1]
```

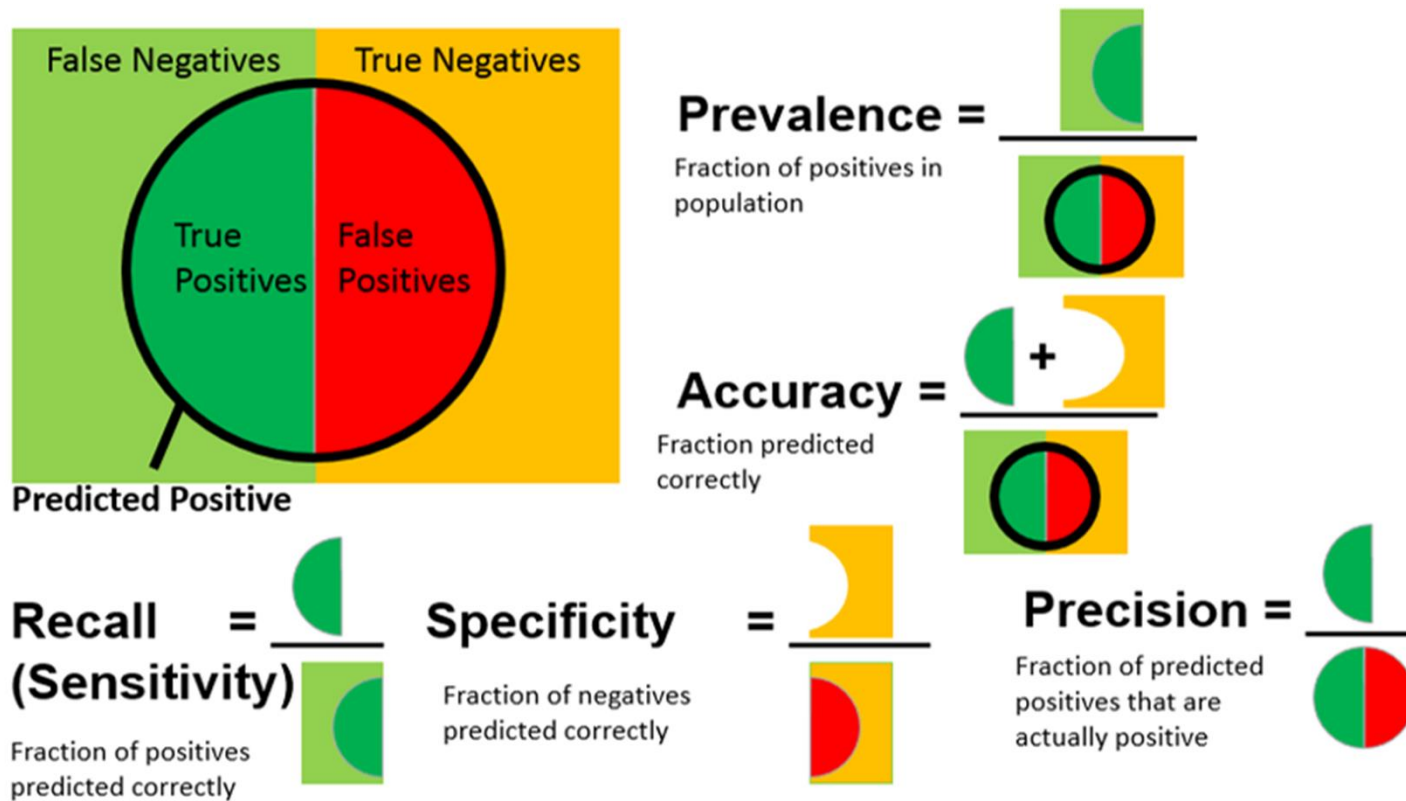
```
In [32]: print(y_train[:10].values)  
print(y_train_preds[:10])  
  
[1 1 0 1 1 1 0 0 1 1]  
[0.76307111 0.63114288 0.29772094 0.77926068 0.59694889 0.55643044  
 0.36292154 0.90984735 0.47806099 0.67622763]
```


Part 4: How to assess the quality of your model

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewlong/odsc_west_2018



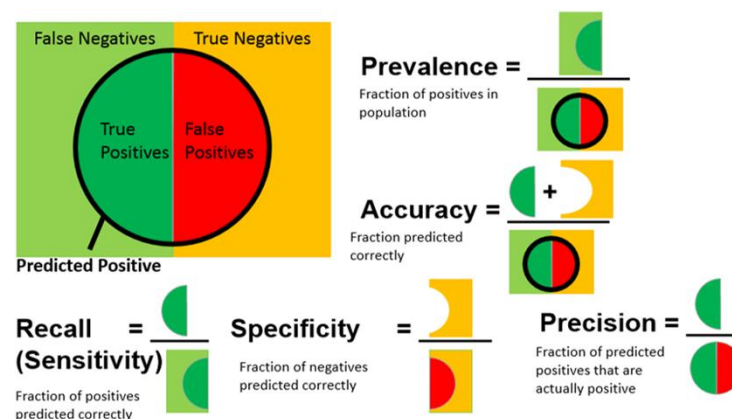
Performance Metrics



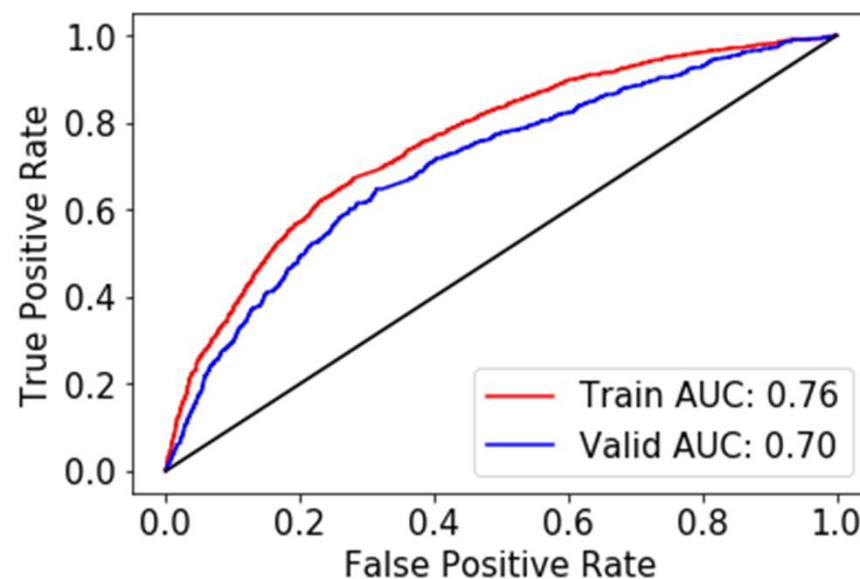
Example: Positive = Hospitalized, Negative = Not Hospitalized

Performance

Metric	Training	Validation
Prevalence	50%	5.7 %
Accuracy	69.5%	68.2 %
Recall	66.6 %	64.8 %
Precision	70.6 %	11.0 %
Specificity	72.3 %	68.4 %
Area Under ROC Curve (AUC)	0.757	0.704



Example: Positive = Hospitalized, Negative = Not Hospitalized



Part 5: How to decide the next step for improving the model

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
https://github.com/andrewlong/odsc_west_2018

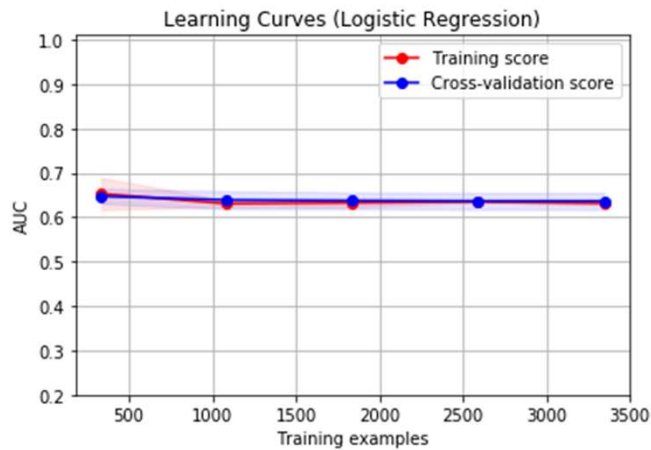


Design decisions

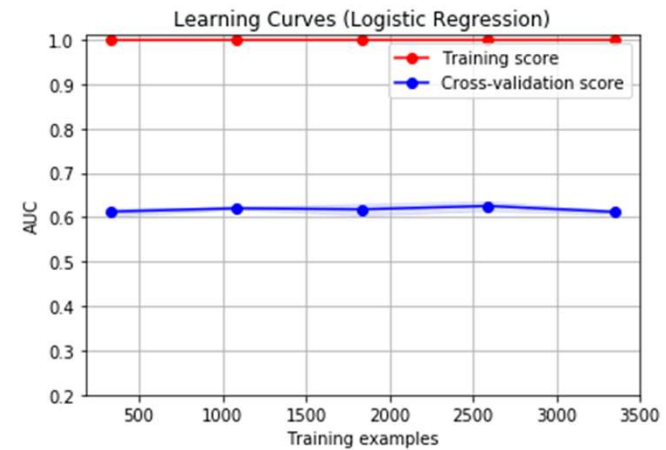
- Which and how much data to use? Should we spend time collecting more data?
- How to tokenize?
 - Should we use stemming? (“stemming” → “stem”)
- How to vectorizer?
 - Change number of words?
 - Switch to tfidfvectorizer?
- How to select hyperparameters in Logistic regression?
- Should we switch to a different ML model?



Learning Curve (diagnose bias/variance)

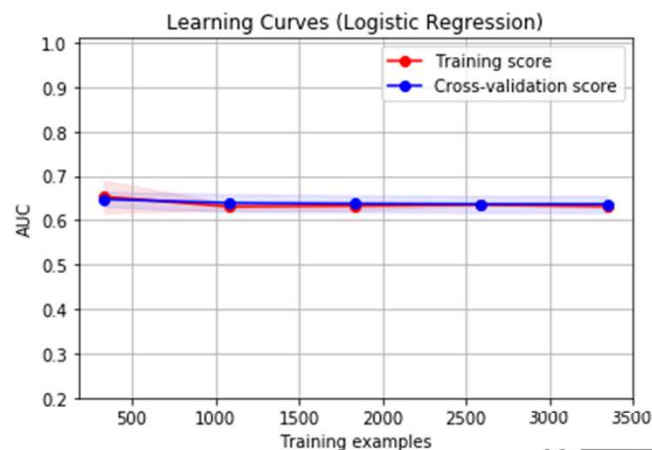


High Bias

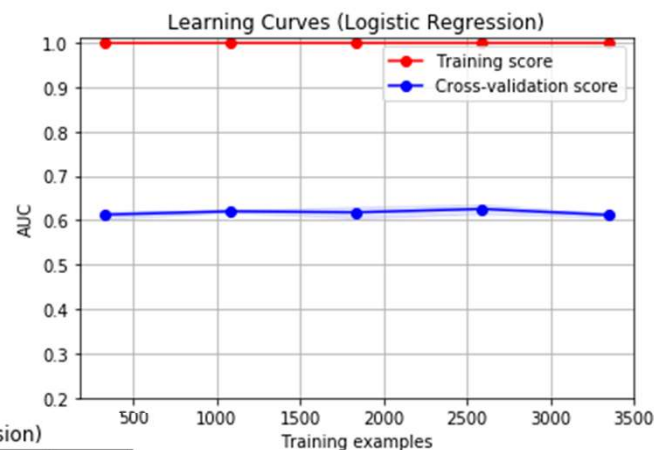


High Variance

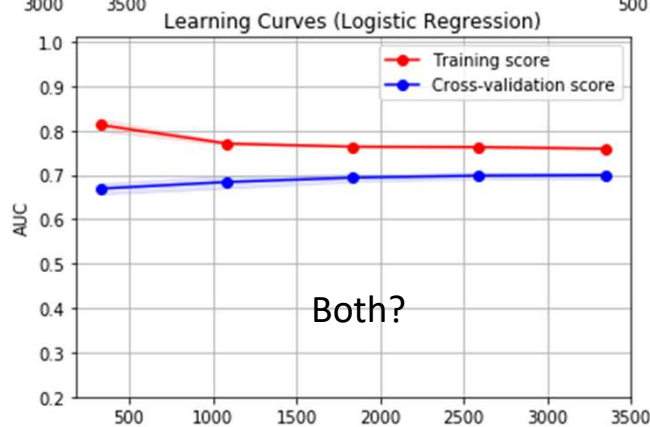
Learning Curve (diagnose bias/variance)



High Bias



High Variance



Both?

Adding more samples
Probably won't help majorly
at this point

Helpful techniques

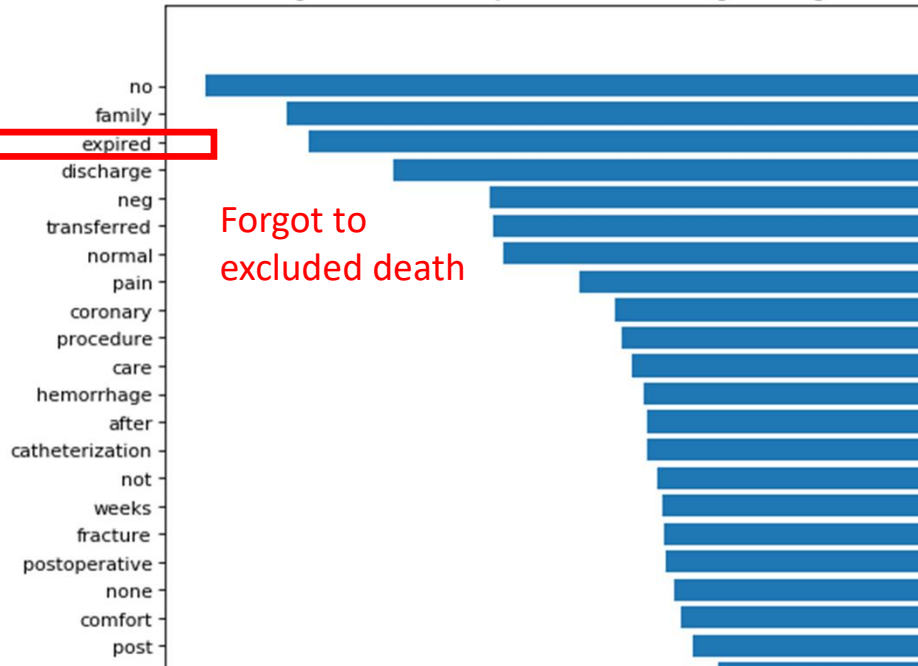
- Techniques for reducing bias (underfitting)
 - Add new features
 - Increase model complexity
 - Reduce regularization
 - Change model architecture
- Techniques for reducing variance (overfitting)
 - Add more samples
 - Add regularization
 - Reduce number of features
 - Decrease model complexity
 - Add better features
 - Change model architecture

Helpful techniques

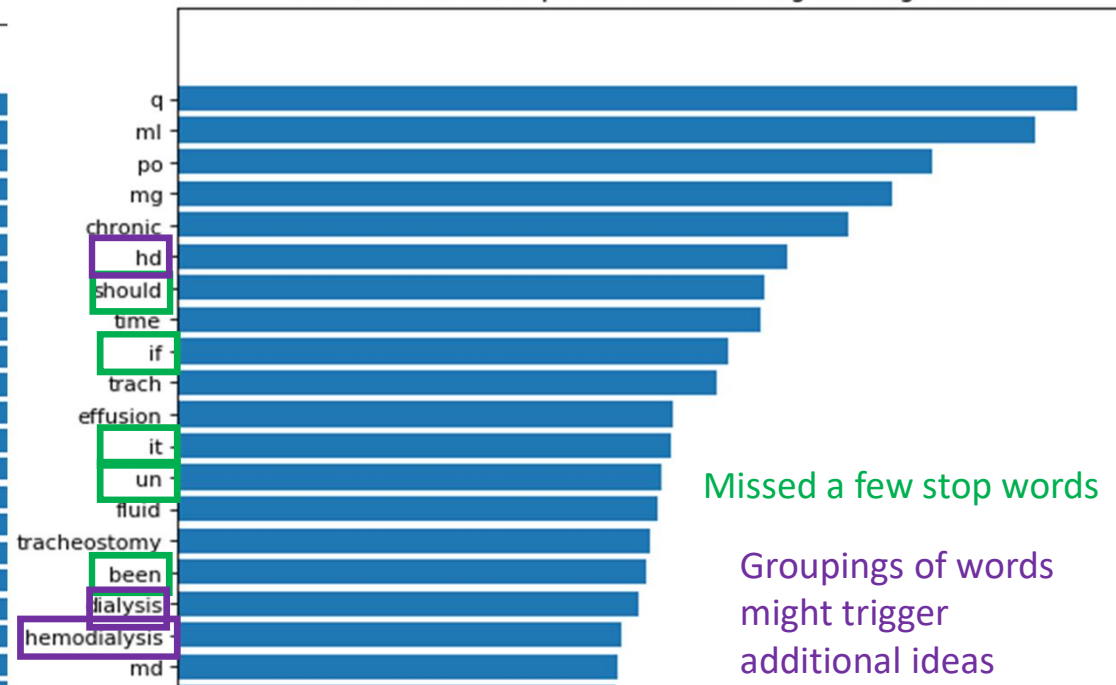
- Techniques for reducing bias (underfitting)
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 - Decrease model complexity
 - Add better features
 - Change model architecture

Feature Importance

Negative Feature Importance Score - Logistic Regression



Positive Feature Importance Score - Logistic Regression



Missed a few stop words

Groupings of words might trigger additional ideas

More Negative

Logistic regression coefficient

More positive

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)

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Helpful techniques

- Techniques for reducing bias (underfitting)
 - Add new features
 - Increase model complexity
 - Reduce regularization
 - Change model architecture
- Techniques for reducing variance (overfitting)
 - Add more samples
 - Add regularization
 - Reduce number of features
 - Decrease model complexity
 - Add better features
 - Change model architecture

Hyper parameter tuning

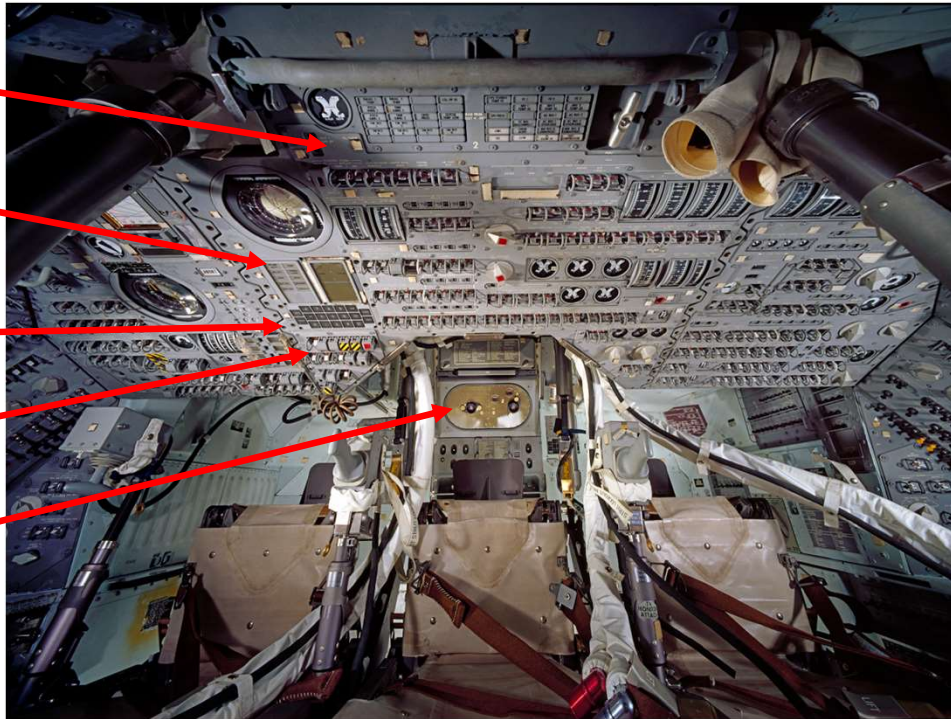
Regularization C

Stemming
(hospitalizations → hospital)

CountVectorizer or
Tfidfvectorizer

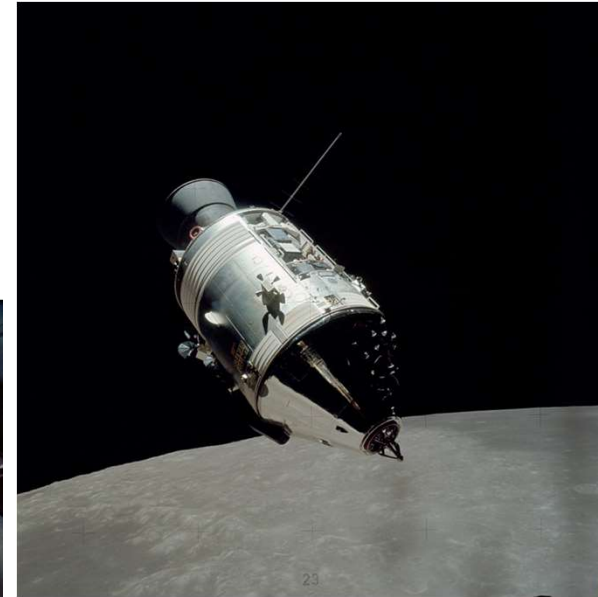
max_features

N grams



Source: <https://airandspace.si.edu/multimedia-gallery/5128hjpg>

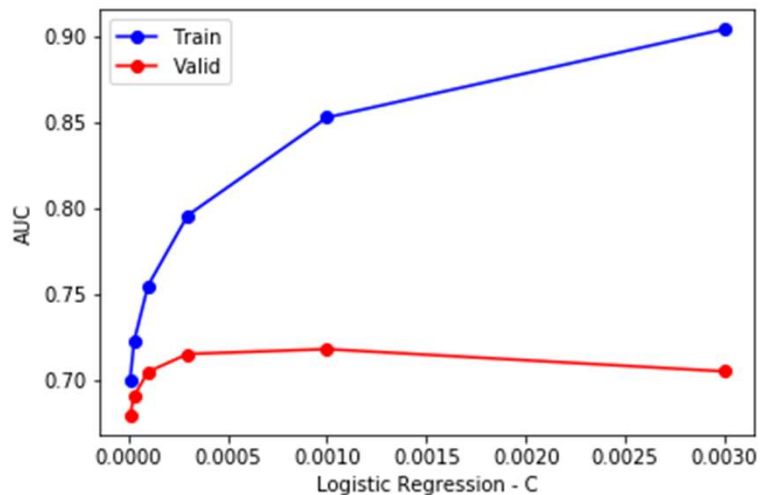
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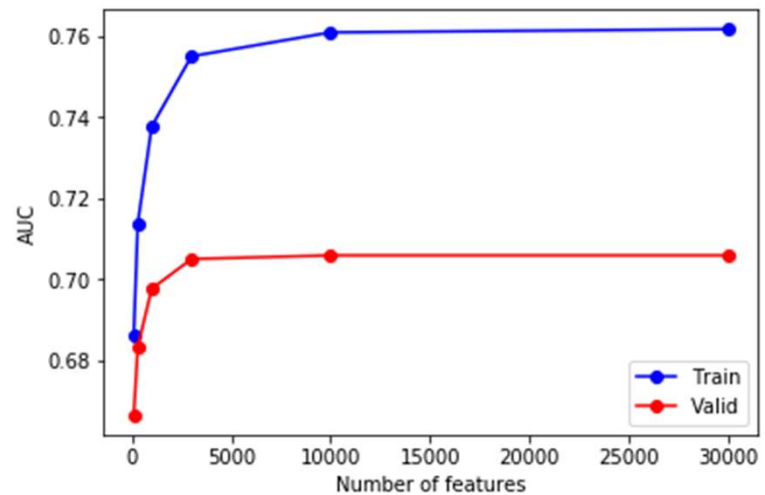
Source: [https://en.wikipedia.org/wiki/Apollo_\(spacecraft\)](https://en.wikipedia.org/wiki/Apollo_(spacecraft))

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Hyper parameter tuning



Higher C = more overfitting



Higher max_features = more overfitting

Model Architecture

- Naïve Bayes
- Neural Networks (CNN, RNN)
with Word2Vec

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
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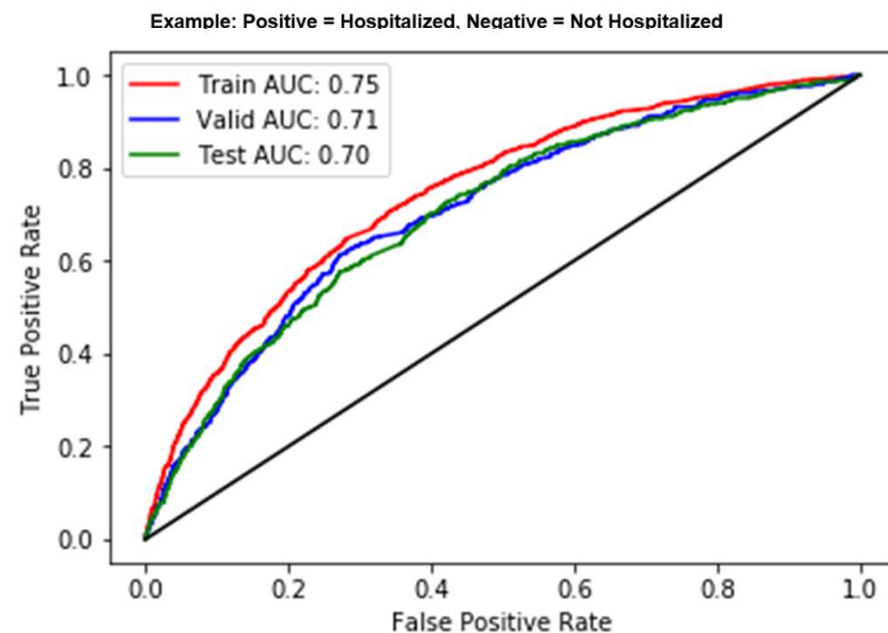
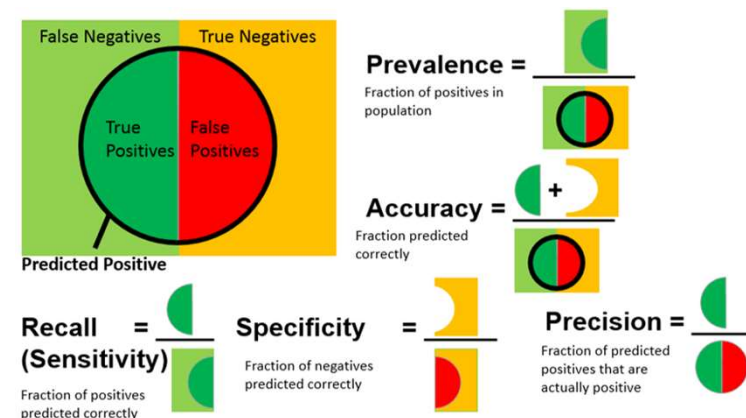


Final Model (excluding deaths)

Metric	Training	Validation	Test
Prevalence	50%	6.9%	6.6%
Accuracy	0.683	0.672	0.681
Recall	0.644	0.651	0.607
Precision	0.698	0.129	0.120
Specificity	0.722	0.674	0.686
Area Under ROC Curve (AUC)	0.745	0.709	0.704

30-day unplanned readmission (AUC = 0.75–76)
(Rajkomar et al 2017)

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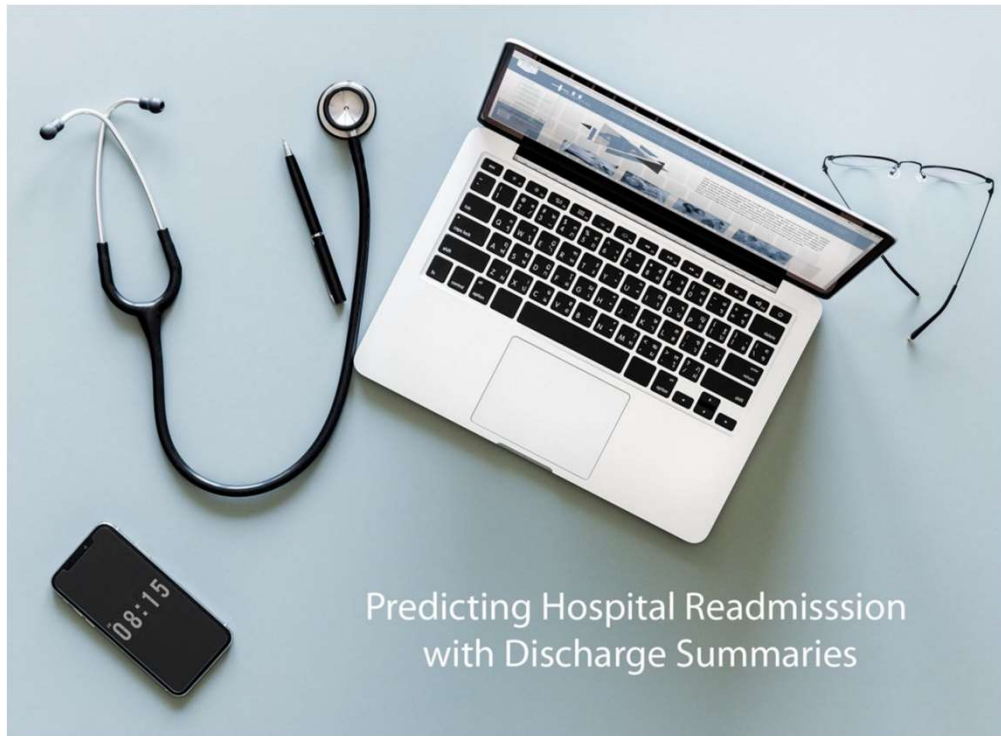
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Introduction to Clinical Natural Language Processing

Andrew Long

<https://towardsdatascience.com/introduction-to-clinical-natural-language-processing-predicting-hospital-readmission-with-1736d52bc709>



Predicting Hospital Readmission
with Discharge Summaries

Andrew Long • awlong20@gmail.com • [linkedin.com/in/awlong/](https://www.linkedin.com/in/awlong/)
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