

Using data between two subreddits on Reddit, would a Multinomial Naive Bayes model perform better than a Support Vector Machines model to predict which post belonged to the correct subreddit?



### Marvel Studios and the Marvel Cinematic Universe



r/marvelstudios



**Marvel Comics** 



r/Marvel



## PUSH SHIFT



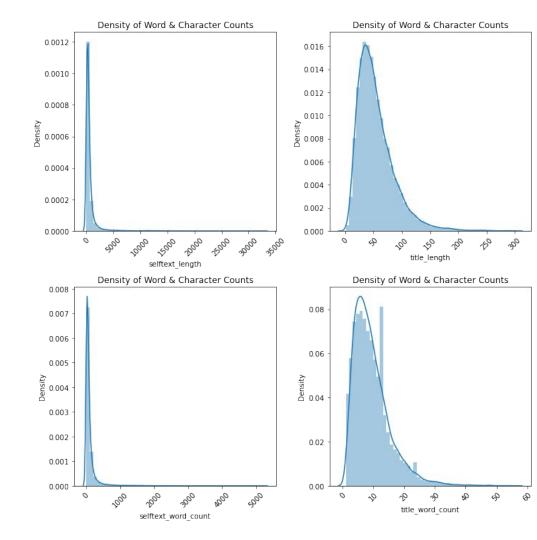
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							46		4685 What are you	•		1523370727		6	1	TRUE	4/10/18
							46	38	4686 Peter Parker			1523375537		4	6	TRUE	4/10/18
							46	39	4687 Post 9/11 he		Marvel	1523375921		3	2	TRUE	4/10/18
							469	90	4688 Let's contact	Hello,	Marvel	1523378686		1	0	TRUE	4/10/18
							469		4689 GOTG 1&am	Hey guys,	Marvel	1523379216	Iml-forwards	6	0	TRUE	4/10/18
							469		4690 All new Guai	Why do the	Marvel	1523380080	varsas	3	1	TRUE	4/10/18
							469	93	4691 I can't descri	I've been wa	Marvel	1523380264	DaveyRocket	7	11	TRUE	4/10/18
							469		4692 Infinity War	[removed]	Marvel	1523381692	BigPapiChee	0	1	TRUF	4/10/18





Dropped Nulls, and all self text labeled "[removed]" because the other data in the rows was not necessary

These graphs show the distribution of words and characters in the text and titles of the posts.



```
Setting up data for modeling
        = df['selftext']
        = df['subreddit']
[16]: # Split data into the training and testing sets.
      X_train1, X_test1, y_train1, y_test1 = train_test_split(X,y,stratify=y)
[17]: #Instantiate a CountVectorizer with a default hyperparamters
      cvec = CountVectorizer(stop_words='english')
[18]: # Fit and transform train
      X_train1 = cvec.fit_transform(X_train1)
      X_test1 = cvec.transform(X_test1)
[19]: # Plotting most used words
      X_train1_df = pd.DataFrame(X train1.todense(),
                               columns = cvec.get_feature_names())
      plt.figure(figsize=(10,10))
      X_train1_df.sum().sort_values(ascending=False).head(40).plot(kind='barh');
```

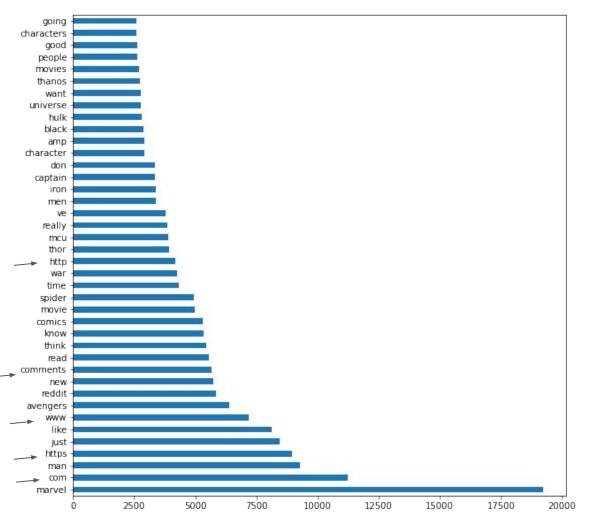


Using very basic parameters, and the basic stop words, didn't get a very good bag of words.

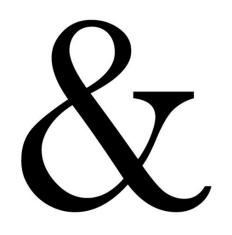


This plot illustrates that the stop
words need to be customized in
order because words like http,
https, www are hyperlinks that give
no distinction between the two
subreddits.

war spider
spider
movie
comics
know
think
read
know
think
read
avengers
sew
subreddits.



### MULTINOMIAL NAIVE BAYES MODEL with CountVectorizer

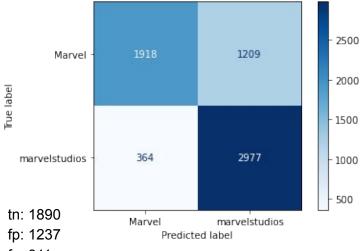




SUPPORT VECTOR MACHINES MODEL with TfidfVectorizer

### **Multinomial Naive Bayes**

The CountVectorizer ran with a Multinomial Naive Bayes model has an accuracy score of 76.3% and 76.1% which is a decently fit model and produces high number of Type I errors. The model performs semi-well with predictions on the test data.



# fn: 311

tp: 3030

### Let's set a pipeline up with two stages:

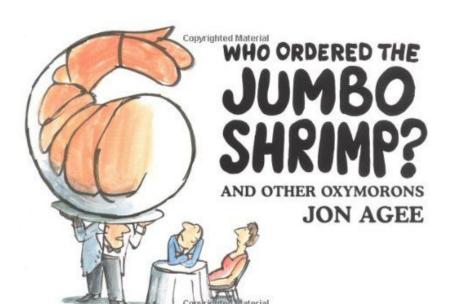
```
[27]: # 1. CountVectorizer (transformer)
      # 2. Multinomial Naive Bayes (estimator)
      pipe = Pipeline([
          ('cvect', CountVectorizer()),
          ('nb', MultinomialNB())
 [*]: pipe_params = {
           'cvect _max features': [2000, 3000, 4000, 5000], # Capping features
           'cvect_min_df': [2, 3], # Word has to show up in more than 2-3 documents
           'cvect__max_df': [.9, .95], # Word can't show up in 90% and 95%
           'cvect ngram range': [(1,1), (1,2)], # 1-gram, 2-gram
           'cvect_stop_words': [None, 'english', stop_words], # Using stop_words
[29]: # Instantiate GridSearchCV.
      gs = GridSearchCV(pipe, # what object are we optimizing?
                        param grid=pipe params, # what parameters values are we searching?
                        cv=5, verbose = 1,
                       n_jobs = -2) # 5-fold cross-validation.
[30]: # Fit GridSearch to training data.
      gs.fit(X_train, y_train)
      Fitting 5 folds for each of 32 candidates, totalling 160 fits
       [Parallel(n jobs=-2)]: Using backend LokyBackend with 15 concurrent workers.
       [Parallel(n jobs=-2)]: Done 20 tasks
                                                              13.05
       [Parallel(n jobs=-2)]: Done 160 out of 160 | elapsed: 1.4min finished
[30]: GridSearchCV(cv=5,
                   estimator=Pipeline(steps=[('cvect', CountVectorizer()),
                                             ('nb', MultinomialNB())]),
                   n jobs=-2,
                   param grid={'cvect max df': [0.9, 0.95],
                                'cvect_max_features': [2000, 3000, 4000, 5000],
                               'cvect__min_df': [2, 3],
                               'cvect_ngram_range': [(1, 1), (1, 2)]},
                   verbose=1)
```

### Pros:

- 1. Very fast modeling
- 2. Excellent Classifier, lesser complication.

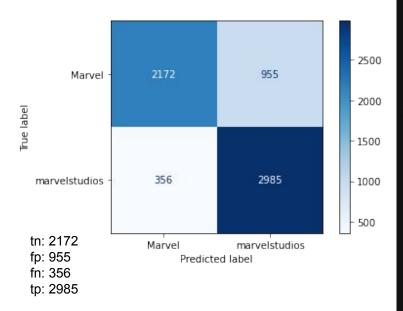
### Cons:

1. Naive Bayes assumes each feature is independent from one another. Text data is never independent. Jumbo Shrimp.



### **Support Vector Machines**

Running a TfidfVectorizer transformer on a Support Vector Machines estimator spat out a decently overfit model. 91.9% accuracy score on training data and 79.7& on testing data with significant Type I errors but better than the Naive Bayes Model.



### Modeling with TfidfVect and Support Vector Machines

```
[61]: # 1. TfidfVectorizer (transformer)
      # 2. Support Vector Machines (estimator)
      pipey = Pipeline([
          ('tvec', TfidfVectorizer()),
          ('svc', SVC())
[65]: pipe_tvec_params = {
          'tvec_max_features': [500, 1_000, 2_000, 3_000, 4_000, 5_000],
          'tvec__stop_words': [None, 'english', stop_words],
          'tvec ngram range': [(1,1), (1,2)],
          'svc_C': np.linspace(0.00001,2,10),
          'svc_kernel': ['poly', 'rbf'],
          'svc gamma': ['scale', 'auto']
[66]: # Instantiate GridSearchCV.
      gsv = GridSearchCV(pipey, # what object are we optimizing?
                        param grid=pipe tvec params, # what parameters values are we searching?
                        cv=5, verbose = 1,
                       n_jobs = -3) # 5-fold cross-validation.
[*]: # Fit and wait
      gsv.fit(X_train,y_train)
      Fitting 5 folds for each of 1440 candidates, totalling 7200 fits
      [Parallel(n jobs=-3)]: Using backend LokyBackend with 14 concurrent workers.
      [Parallel(n jobs=-3)]: Done 22 tasks
                                                   elapsed: 3.7min
      [Parallel(n_jobs=-3)]: Done 172 tasks
                                                   elapsed: 28.9min
      [Parallel(n jobs=-3)]: Done 422 tasks
                                                   elapsed: 70.0min
      [Parallel(n_jobs=-3)]: Done 772 tasks
                                                   elapsed: 129.0min
      [Parallel(n_jobs=-3)]: Done 1222 tasks
                                                    elapsed: 200.0min
```

### Pros:

- The model performs really well.
- 2. Effective in high-dimensional data
- 3. Can work with non-linear boundaries
- Fast to compute with most datasets(kernel trick)

```
Fitting 5 folds for each of 1440 candidates, totalling 7200 fits

[Parallel(n_jobs=-3)]: Using backend LokyBackend with 14 concurrent workers.

[Parallel(n_jobs=-3)]: Done 22 tasks | elapsed: 3.7min

[Parallel(n_jobs=-3)]: Done 172 tasks | elapsed: 28.9min

[Parallel(n_jobs=-3)]: Done 422 tasks | elapsed: 70.0min

[Parallel(n_jobs=-3)]: Done 772 tasks | elapsed: 129.0min

[Parallel(n_jobs=-3)]: Done 1222 tasks | elapsed: 200.0min
```

### Cons:

- Black box model. There isn't an explicable rhyme or reason, it works.
- Very slow to train. Very slow.

(I started my fit before I left for work and it was done in 4 hours. I missed a lot of hyper parameters and ran it again at 1AM. It's almost 6AM.)

If what you were looking for was a quick training model that you can tune based on a mathematical formula within a short deadline, the Naive Bayes model is your model. If you have hours, and I mean hours, to spare then Support Vector is the better performing model of the two.