ML for SS: Ensembling and Clustering

Session 3

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Overview

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Papers

Paper 1: Easton et al. 2020

- Introduces a well motivated use for clustering
- Takes a standard approach to the introduction of a new technique
- The points the paper makes are applicable broadly in any archival/empirical discipline

Paper 2: Qiu, Xie and Jun (2020 working)

• A fairly straightforward paper introducing the concept of ensembling

Technical Discussion: Ensembling

Python

- Rolling your own is pretty doable
- sklearn is the primary tool for constructing them in python

- - SuperLearner
 - EnsembleML

Python is generally a bit stronger for these topics.

There is a fully worked out solution for using python, data and pretrained models are on eLearn.

R

 Rolling your own is pretty doable There are some packages for automating ensemble construction:

Main application: Ensembling

- Idea: Predict instances of intentional misreporting?
- Testing: Predicting 10-K/A irregularities using finance, textual style, and topics

Dependent Variable

Intentional misreporting as stated in 10-K/A filings

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This test mirrors a subset of Brown, Crowley and Elliott (2020 JAR)

We will **combine** the models from the past two weeks

Independent Variables

• 17 Financial measures 20 Style characteristics • 31 10-K discussion topics



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Technical Discussion: Clustering

Python

- sklearn is still good for this
 - k-means and KNN
 - t-SNE
- umap-learn for UMAP

- For UMAP, umap works

Python is generally a bit stronger for these topics.

There is a fully worked out solution for using python, data is on elearn

R

 For standard clustering, caret is a good choice For t-SNE, Rtsne works well

Main application: Clustering

Idea: Industry classification based on the text of annual reports

Dependent Variable

SIC Codes

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Somewhat in the vein of Hoberg and Phillips (2016 JPE), though less precise



Independent Variables

31 10-K discussion topics

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Ensembling

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What are ensembles?

- Ensembles are models made out of models
- Ex.: You train 3 models using different techniques, and each seems to work well in certain cases and poorly in others
 - If you use the models in isolation, then any of them would do an OK (but not great) job
 - If you make a model using all three, you can get better performance if their strengths all shine through
- Ensembles range from simple to complex
 - Simple: a (weighted) average of a few model's predictions



When are ensembles useful?

1. You have multiple models that are all decent, but none are great

• And, ideally, the models' predictions are not highly correlated



ut none are great highly correlated

When are ensembles useful?

2. You have a really good model and a bunch of mediocre models

• And, ideally the mediocre models are not highly correlated



nediocre models ighly correlated



When are ensembles useful?

- 3. You really need to get just a bit more accuracy/less error out of the model, and you have some other models lying around
- 4. You want a more stable model
 - It helps to stabilize predictions by limiting the effect of errors or outliers produced by any one model on your prediction
 - Think: Diversification



A simple ensemble (averaging)

- For continuous predictions, simple averaging is viable
 - Often you may want to weight the best model a bit higher
- For binary or categorical predictions, consider averaging *ranks*
 - i.e., instead of using a probability from a logit, use ranks 1, 2, 3, etc.
 - Ranks average a bit better, as scores on binary models (particularly when evaluated with measures like AUC) can have extremely different variances across models
 - In which case the ensemble is really just the most volatile model's prediction...
 - Not much of an ensemble





A more complex ensemble (voting model)

- If you have a model the is very good at predicting a binary outcome, ensembling can still help
 - This is particularly true when you have other models that capture different aspects of the problem
- Let the other models vote against the best model, and use their prediction if they are above some threshold of agreement



A lot more complex ensemble

- Stacking models (2 layers)
 - 1. Train models on subsets (folds) of the training data
 - 2. Make predictions for each model on the folds it wasn't applied to
 - 3. Train a new model that takes those predictions as inputs (and optionally the data set as well)
- Blending (similar to stacking)
 - Like stacking, but using predictions on a hold out sample instead of folds (and thus all models are using the same data for predictions)



A simple averaging ensemble of our models

test X ens = pd.DataFrame({'XGBoost': models['XGBoost'].predict proba(models['test X ML'])[:,1], 'SVC': logistic(models['SVC'].decision function(models['test X ML'])), models['ElasticNet'].predict proba(models['test X ML'])[:,1], 'ElasticNet': models['LASSO'].predict proba(models['test X ML'])[:,1], 'LASSO': 'logit': models['logit'].predict(models['test pd'][models['vars']])})

```
rank X ens = test X ens.rank()
arank_X_ens = rank_X_ens.XGBoost + rank_X_ens.SVC + rank_X_ens.ElasticNet + rank_X_ens.LASSO + rank_X_ens.logit
auc = metrics.roc auc score(models['test pd'].Restate Int, arank X ens)
fpr, tpr, thresholds = metrics.roc curve(models['test pd'].Restate Int, arank X ens)
display = metrics.RocCurveDisplay(fpr=fpr, tpr=tpr, roc auc=auc)
display.plot()
```

Practicalities

- Methods like stacking or blending are much more complex than a simple averaging or voting based ensemble
 - But in practice they perform slightly better

Recall the tradeoff between complexity and accuracy!

As such, we may not prefer the complex ensemble in practice, unless we only care about accuracy

Example: In 2009, Netflix awarded a \$1M prize to the BellKor's Pragmatic Chaos team for beating Netflix's own user preference algorithm by >10%. The alogorithm was so complex that Netflix never used it. It instead used a simpler algorithm with an 8% improvement.

Where is ensembling useful in academic work

Where multiple reasonable models exist, and pushing performance (accuracy) is important

• It can also be a reasonable approach when you are already calculating other models anyway



[Geoff Hinton's] Dark knowledge

- Complex ensembles work well
- Complex ensembles are exceedingly computationally intensive
 - This is bad for running on small or constrained devices (like phones)

Dark knowledge

We can (almost) always create a simple model that approximates the complex model
Interpret the above literally – we can train a model to fit the model



ally intensive devices (like phones)

Dark knowledge

- Train the simple model not on the actual DV from the training data, but on the best algorithm's (softened) prediction for the training data
- Somewhat surprisingly, this new, simple algorithm can work almost as well as the full thing!



An example of this dark knowledge

- Google's full model for interpreting human speech is >100GB
 - As of October 2019
- In Google's Pixel 4 phone, they have human speech interpretation running locally on the phone
 - Not in the cloud like it works on any other Android phone

How did they do this?

• They can approximate the output of the complex speech model using a 0.5GB model

• 0.5GB isn't small, but it's small enough to run on a phone



Learning more about Ensembling

- Scikit-learn's documentation on ensemble methods it supports
- Geoff Hinton's Dark Knowledge slides
 - For more details on *dark knowledge*, applications, and the softening transform
 - His interesting (though highly technical) Reddit AMA
- A Kaggler's Guide to Model Stacking in Practice
 - A short guide on stacking with nice visualizations
- Kaggle Ensembling Guide
 - A comprehensive list of ensembling methods with some code samples and applications discussed
- Ensemble Learning to Improve Machine Learning Results
 - Nicely covers bagging and boosting (two other techniques)

There are many ways to ensemble, and there is no specific guide as to what is best. It may prove useful in the group project, however.

Addendum: Using R

- There are a couple interesting packages in R for ensembling:
 - The Superlearner package aims to automate building ensembles
 - Think of it like an automated cross-validation for ensemble construction
 - The EnsembleML package allows you to specify an ensemble and train the underlying models together
- You can also roll your own ensemble as we did in the example earlier

bles Instruction Ind train the underlying models together er





What is k-means?

$$\min_{C_k} \sum_{k=1}^K \sum_{x_i \in C_k} \left(x_i - \mu_k
ight)^2$$

- Minimizes the sum of squared distance between points within groups
- Technically this is a machine learning algorithm, despite its simplicity
- You need to specify the number of groups you want
- Pros:
 - Very fast to run
 - Simple interpretation

Cons

Since the algorithm is unsupervised, optimizing k can be tricky

• Simple algorithm • Need to specify *k*, the number of clusters

Projecting to 2D with UMAP

• Like last session, we will use UMAP to get a sense of how well topics line up with SIC industries



Projecting to 2D with UMAP

- It is also interesting to see how well the topics can be clustered
 - The below colors UMAP by a k=9 kmeans algorithm applied to the LDA output



lustered applied to the LDA output



Why are these graphs different?

- Possibly due to...
 - Data: 10-K disclosure content doesn't fully capture industry inclusion
 - Topic modeling: The measure may be noisy
 - SIC code: The measure doesn't cleanly capture industry inclusion
 - Some firms are essentially misclassified
- Recall, SIC covers Agriculture, Forestry and Fishing; Mining; Construction; Manufacturing; Transportation, Communications, Electric, Gas, and Sanitary Services; Wholesale Trade; Retail Trade; Finance, Insurance, and Real Estate; Services; Public Administration



Optimizing K-means clustering

• K-means clustering is very fast to run, but suffers from the same issue as LDA:

You need to specify the number of clusters!

- Often times the solutions to this are similar to what we discussed for LDA
 - Hand tuning
 - In sample performance
- However, there is a statistics-based, researcher-bias-free method

The Gap Statistic

How does the Gap statistic work?

- Let...
 - *k* be the number of clusters,
 - *B* the number of simulated samples
 - W_k be the K-Means inertia score on actual data
 - $W^*_{k,r}$ be the K-Means inertia score for iteration r with synthetic data
 - \overline{l} be the average of the $W^*_{k,r}$ s

$$egin{aligned} & eap(k) = \left(rac{1}{B}
ight)\sum_{r=1}^B \log\left(W_{k,r}^*
ight) - \log\left(W_k
ight) ext{ and } \ & s_k = sd_k\sqrt{1+rac{1}{B}}, ext{ where } sd_k = \sqrt{\left(rac{1}{B}
ight)\sum_{r=1}^B} \end{aligned}$$

• Select the lowest k such that $Gap(k) \geq Gap$

I.e., select the lowest k s.t. the log-scaled error removed by clustering on real data at k is no worse than 1 SD below the log-scaled error removed at k+1

$${iggstyle 1} \left\{ \log \left(W^*_{k,r} - ar{l}
ight)
ight\}^2$$

$$\left(k+1
ight)-s_{k+1}$$

Implementation in python

- The code is too long to put in the slides, but it is in the code file
- Sketch of the code:
 - 1. Iterate through k values starting at 2
 - 2. Determine performance (inertia) at k with real data
 - 3. Determine performance (inertia) at k with simulated (random) data 10 times
 - 4. Calculate the standard deviation of the log of performance on random data
 - 5. See if the 2x2 difference in log inertia between k and k+1 on real and random data is less than the standard deviation
 - If so, k is optimal, stop iterating
 - If not, k = k + 1 and start again

k=35 for the model presented here

Optimal clustering

```
model = cluster.KMeans(n_clusters=30)
kmeans = model.fit(df[topic_names])
df['cluster_opt'] = kmeans.labels_
```

umap_color(df[topic_names], df.cluster_opt.astype("category"))





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Example companies in the optimized clusters

| ar [ar. | cruster_optoj[[industry j].sam | |
|---------|---------------------------------|---------|
| | | |
| ## | industry | ## |
| ## 977 | 0 Services | ## 1377 |
| ## 862 | 2 Wholesale Trade | ## 7898 |
| ## 114 | 01 Services | ## 9036 |
| ## 188 | 2 Retail Trade | ## 1500 |
| ## 696 | Manufacturing | ## 1227 |
| ## 222 | Manufacturing | ## 1238 |
| ## 675 | 2 Manufacturing | ## 5800 |
| ## 102 | 18 Services | ## 1156 |
| ## 189 | Manufacturing | ## 7256 |
| ## 202 | 1 Retail Trade | ## 7181 |

opt==2][['industry']].sample(n=10)

industry ufacturing 2

Clustering: KNN

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Using k-means for filling in data

- One possible approach we could use is to fill based on the category assigned by k-means
- However, as we saw, k-means and SIC code don't line up perfectly...
 - So using this classification will definitely be noisy



category assigned by k-means erfectly...

A better approach with KNN

- KNN, or K-Nearest Neighbors is a *supervised* approach to clustering
- Since we already have industry classifications for most of our data, we can use that structure to inform our assignment of the missing industry codes
- The way the model uses the information is by letting the nearest labeled points "vote" on what the point should be
 - Points are defined by 10-K content in our case
 - Voting can be weighted by distance or done uniformly



Implementing KNN

- Scikit-learn has a KNN implementation in its neighbors module
- The primary parameter in the model is k: how many points get to vote
 - k is n_neighbors in Scikit-learn

```
knn = neighbors.KNeighborsClassifier(n_neighbors=5)
knn.fit(df[topic_names], df['Industry'])
```

The above is sufficient to fit a simple model

ghbors **module** ny points get to vote

Checking performance

- First, we need to get predictions
- Since this is a multiclass problem, we will not output probabilities, but instead the top guess

in_pred = knn.predict(df[topic_names]) out_pred = knn.predict(testing[topic_names])

• We can quickly check multiclass performance using Scikit-learn as well

print('In sample: {},\nOut of sample: {}'.format(metrics.accuracy_score(df['industry'], in_pred), metrics.accuracy score(testing['industry'], out pred)))

In sample: 1.0, Out of sample: 0.922422954303932

A note on dimensionality reduction techniques



Principle Component Analysis

- PCA is a common technique to see in older studies
- It is reasonably efficient at identifying a lower dimensional representation of a relationship
- It is not good at maintaining relationships in the lower dimensional space



Component 1 These two components explain 26.44 % of the point variability.

epresentation of a relationship nsional space

t-distributed Stochastic Neighbor Embedding

- t-SNE is focused on keeping distances relatively similar between the full dimensional input space and the projected output space
 - If 2 points are close to each other in N dimensions, they will be close to each other in 2 or 3 dimensions as well!
- t-SNE *does not* maintain distances over longer distances!
 - Should not be used as input to a regression



industry

Uniform Manifold Approximation and Projection

- UMAP maintains local distances like t-SNE
- UMAP also maintains global distances, mostly
 - As such, it can be used for isolating data components for regression like PCA



- Retail Trad

- Construction
- 0.9333980511751778

Conclusion







Packages used for these slides

Python

- matplotlib
- numpy
- pandas
- scikit-learn
- seaborn
- umap-learn





- caret
- cluster

References

- Easton, Peter D., Martin Kapons, Steven J. Monahan, Harm H. Schütt, and Eric H. Weisbrod. "Forecasting Earnings Using k-Nearest Neighbor Matching." Available at SSRN (2020).
- Qiu, Yue, Tian Xie, and Y. U. Jun. "Forecast combinations in machine learning." (2020).
- Wang, Bingling, Min-Bin Lin, and Wolfgang Karl Hardle, "Non-fungible Tokens & VizTech." (2021).





Custom code

```
From umap.plot source code on Github
def _get_embedding(umap_object):
   if hasattr(umap_object, "embedding_"):
   return umap_object.embedding_
elif hasattr(umap_object, "embedding"):
       return umap_object.embedding
   else:
       raise ValueError("Could not find embedding attribute of umap_object")
# Cut down version of umap.plot.points to remove dependencies on datashader, bokeh, holoviews, scikit-image, and colorcet
# Introduces a dependency on seaborn though
def umap_color(data_map, data_color, cmap='viridis', subset=None, title=None):
   reducer = umap.UMAP()
   umap_object = reducer.fit(data_map)
   embed = _get_embedding(umap_object)
   if subset is not None:
        embed X = embed[subset,0]
        embed Y = embed[subset,1]
        data_color = np.array(data_color[subset])
   else:
        embed_X = embed[:, 0]
       embed_Y = embed[:, 1]
   point_size = 100.0 / np.sqrt(len(embed_X))
    # color by values
   fig, ax = plt.subplots(figsize=(12,8))
   g = sns.scatterplot(ax=ax, x=embed_X, y=embed_Y, hue=data_color, size=point_size)
     = plt.legend(bbox_to_anchor=(1.05, 1), loc=2, borderaxespad=0.)
    return g
```

