Econ 491 Assignment 2 Aida Lynck October 8th I. Trung error us. Test cros, in expectation. a.) Give that:  $R_{trum}(B) = \frac{1}{n} \sum_{i=1}^{n} (Y_i - X_i^* B)^2$   $R_{trum}(B) = \frac{1}{n} \sum_{i=1}^{m} (eY_i - eX_i^* B)^2$   $R_{trum}(B) = \frac{1}{n} \sum_{i=1}^{m} (eY_i - eX_i^* B)^2$  $\mathbb{E}[\mathbb{E}_{\text{Hain}}(\hat{\beta})] = \mathbb{E}\left[\frac{1}{n}\sum_{i=1}^{n}(Y_{i} - X_{i}, \hat{\beta})^{2}\right] = \frac{1}{n}\mathbb{E}\left[\sum_{i=1}^{n}(Y_{i} - X_{i}, \hat{\beta})^{2}\right]$ E[R+15+ (B)] = E[= [(eY: -eX: B)] = = E[[:] (eY: -eX: B)] Sma He training error is minimized one the training data, in expectation, E[R+sun (B)] Z E[R+s+ (B)] 6.) The result indicates that the models performance (in terms of error) on the training data, where it has been optimized, is typically better than its Performany on new, unsur data (test data). This reflects a common School in magaine luring: a movel might do really well on the data it's travus on (sory its toilors to it) but myst not perform as well as new data it tousn't sur before This can be day to surfitting, when the model love the rose or spectre quites of the Horang data rather then the sever moberlying pattern

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############################
import pandas as pd
file path = "/Users/aidanlynde/ECON491/Assignment 2/code/data/2021-12.csv"
df = pd.read csv(file path, header=[0, 1])
df = df.dropna(axis=1)
df = df.iloc[:, :105]
df.columns = df.columns.droplevel(1)
transformations = df.iloc[0, 1:].astype(int)
df = df.drop(df.index[0])
df.set index('sasdate', inplace=True)
output and income = ['RPI', 'W875RX1', 'INDPRO', 'IPFPNSS', 'IPFINAL', 'IPCONGD',
labor market = ['HWI', 'HWIURATIO', 'CLF160V', 'CE160V', 'UNRATE', 'UEMPMEAN',
consumption and orders = ['HOUST', 'HOUSTNE', 'HOUSTMW', 'HOUSTS', 'HOUSTW', 'PERMIT',
orders and inventories = ['DPCERA3M086SBEA', 'CMRMTSPLx', 'RETAILx', 'NAPM',
'NAPMNOI',
'NAPMSDI', 'NAPMII', 'ACOGNO', 'AMDMNOx', 'ANDENOx', 'AMDMUOx',
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money_and_credit = ['M1SL', 'M2SL', 'M2REAL', 'AMBSL', 'TOTRESNS', 'NONBORRES',
'BUSLOANS',
'REALLN', 'NONREVSL', 'CONSPI', 'MZMSL', 'DTCOLNVHFNM', 'DTCTHFNM', 'INVEST']
interest rate and exchange rates = ['FEDFUNDS', 'CP3Mx', 'TB3MS', 'TB6MS', 'GS1',
prices = ['PPIFGS', 'PPIFCG', 'PPIITM', 'PPICRM', 'OILPRICEx', 'PPICMM', 'NAPMPRI',
stock market = ['S&P 500', 'S&P: indust', 'S&P div yield', 'S&P PE ratio']
groups = {
'output and income': output and income,
'consumption and orders': consumption and orders,
import numpy as np
for col, transform code in transformations.items():
if transform_code == 1:
pass # No transformation
elif transform code == 2:
df[col] = df[col].diff()
elif transform code == 3:
df[col] = df[col].diff().diff()
elif transform code == 4:
df[col] = df[col].apply(lambda x: np.log(x)).diff()
elif transform code == 5:
df[col] = df[col].apply(lambda x: np.log(x))
elif transform code == 6:
df[col] = df[col].apply(lambda x: np.log(x)).diff()
elif transform code == 7:
df[col] = df[col].apply(lambda x: np.log(x)).diff().diff()
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df.dropna(inplace=True)
df.index = pd.to_datetime(df.index)
df = df.asfreq('MS')
df['CPIAUCSL inflation'] = df['CPIAUCSL'].diff() / df['CPIAUCSL']
df = df.dropna(subset=['CPIAUCSL_inflation'])
from statsmodels.tsa.ar model import AutoReg
best bic = np.inf
best_order = None
endog = df['CPIAUCSL inflation']
for p in range(1, 21):
model = AutoReg(endog, lags=p, trend='c')
results = model.fit()
if results.bic < best bic:</pre>
best bic = results.bic
best_order = p
print(f"Best order by BIC: {best_order}")
from sklearn.preprocessing import StandardScaler
X = df.drop('CPIAUCSL inflation', axis=1)
scaler = StandardScaler()
X_standardized = scaler.fit_transform(X)
from sklearn.decomposition import PCA
pca = PCA()
pca_result = pca.fit_transform(X_standardized)
import matplotlib.pyplot as plt
explained_variance = pca.explained_variance_ratio_.cumsum()
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plt.figure(figsize=(10,5))
plt.plot(range(1, len(explained_variance)+1), explained_variance, marker='o',
linestyle='--')
plt.title('Explained Variance by Components')
plt.xlabel('Number of Components')
plt.ylabel('Cumulative Explained Variance')
plt.show()
num components = np.where(explained variance > 0.95)[0][0] + 1
print(f"Number of components that explain >=95% variance: {num components}")
from sklearn.decomposition import PCA
n components = 17
pca = PCA(n components=n components)
X pca = pca.fit transform(X standardized)
from sklearn.model_selection import train_test_split
y = df['CPIAUCSL inflation']
X train pca, X test pca, y train, y test = train test split(X pca, y, test size=0.2,
random_state=42)
from sklearn.linear model import LinearRegression
model = LinearRegression()
model.fit(X_train_pca, y_train)
from sklearn.metrics import mean squared error
y_pred = model.predict(X_test_pca)
mse = mean_squared_error(y_test, y_pred)
print(f"Mean Squared Error using {n components} principal components: {mse}")
from sklearn.linear model import RidgeCV
X train, X test, y train, y test = train test split(X standardized, y, test size=0.2,
random_state=42)
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alphas = np.logspace(-6, 6, 13)
ridge cv = RidgeCV(alphas=alphas, store cv values=True)
ridge cv.fit(X train, y train)
best_alpha = ridge_cv.alpha_
print(f"Best alpha: {best alpha}")
y pred ridge = ridge cv.predict(X test)
mse_ridge = mean_squared_error(y_test, y_pred_ridge)
print(f"Mean Squared Error using Ridge Regression: {mse ridge}")
from sklearn.linear model import LassoCV
from sklearn.preprocessing import StandardScaler
from sklearn.pipeline import make pipeline
scaler = StandardScaler()
X train scaled = scaler.fit transform(X train)
X test scaled = scaler.transform(X test)
lasso_cv = LassoCV(alphas=alphas, cv=5, max_iter=200000, tol=0.001)
lasso_cv.fit(X_train_scaled, y_train)
lasso_preds = lasso_cv.predict(X_test_scaled)
lasso_mse = mean_squared_error(y_test, lasso_preds)
print(f"Best alpha for LASSO: {lasso cv.alpha }")
print(f"Mean Squared Error using LASSO Regression: {lasso mse}")
print(f"Number of predictors used in the LASSO model: {np.sum(lasso cv.coef != 0)}")
from sklearn.pipeline import Pipeline
from collections import defaultdict
window_size = 492
start pos = 0
end pos = start pos + window size
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ridge_errors = []
lasso_errors = []
pcr errors = []
while end pos < len(df):
train data = df.iloc[start pos:end pos]
test data = df.iloc[end pos:end pos+1]
X train = train data.drop('CPIAUCSL inflation', axis=1)
y train = train data['CPIAUCSL inflation']
X test = test data.drop('CPIAUCSL inflation', axis=1)
y test = test data['CPIAUCSL inflation']
pca = PCA(n_components=n_components)
X_pca_train = pca.fit_transform(scaler.transform(X_train))
model = LinearRegression()
model.fit(X_pca_train, y_train)
ridge_cv = RidgeCV(alphas=alphas, store_cv_values=True)
ridge cv.fit(X train, y train)
X train scaled = scaler.fit transform(X train)
lasso cv = LassoCV(alphas=alphas, cv=5, max iter=200000, tol=0.001)
lasso_cv.fit(X_train_scaled, y_train)
# Append errors
ridge_errors.append((ridge_cv.predict(X_test) - y_test)**2)
lasso_errors.append((lasso_cv.predict(X_test_scaled) - y_test.values)**2)
pcr_errors.append((model.predict(pca.transform(scaler.transform(X test))) -
y test)**2)
start pos += 1
end_pos = start_pos + window_size
ridge importances = []
lasso_importances = []
pcr importances = []
variable importances = {
'ridge': defaultdict(float),
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pcr': defaultdict(float)
for window start in range(0, len(df) - window size):
window end = window start + window size
X_train = df.iloc[window_start:window_end].drop(columns='CPIAUCSL_inflation')
y train = df.iloc[window start:window end]['CPIAUCSL inflation']
pca = PCA(n components=n components)
X pca train = pca.fit transform(scaler.transform(X train))
model = LinearRegression()
model.fit(X pca train, y train)
ridge cv = RidgeCV(alphas=alphas, store cv values=True)
ridge_cv.fit(X_train, y_train)
X train scaled = scaler.fit transform(X train)
lasso cv = LassoCV(alphas=alphas, cv=5, max iter=200000, tol=0.001)
lasso cv.fit(X train scaled, y train)
ridge_importance = ridge_cv.coef_ * X_train.std().values
lasso importance = lasso cv.coef * X train scaled.std(axis=0)
pca_components = pca.components_
pcr importance = np.dot(pca components.T, model.coef )
ridge importances.append(ridge importance)
lasso_importances.append(lasso_importance)
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```
pcr_importances.append(pcr_importance)
for group name, variables in groups.items():
for ridge_imp, lasso_imp, pcr_imp in zip(ridge_importances, lasso_importances,
pcr importances):
for var in variables:
var idx = df.columns.get loc(var)
variable importances['ridge'][group name] += ridge imp[var idx]
variable_importances['lasso'][group_name] += lasso_imp[var_idx]
variable_importances['pcr'][group_name] += pcr_imp[var_idx]
for model name, importances in variable importances.items():
max importance = max(importances.values())
for group_name in importances:
variable_importances[model_name][group_name] /= max_importance / 100
print(variable importances)
```