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In [2]:
         from future import division
         import numpy as np
         import matplotlib.pyplot as plt
         from sklearn import datasets, linear model
         import pandas as pd
         from pandas import DataFrame, Series
         import seaborn as sns
         from sklearn.model selection import train test split
         from sklearn.metrics import r2 score
         sns.set(style='ticks', palette='Set2')
         %matplotlib inline
In [3]:
         # Generate data
         np.random.seed(7)
         n_samples, n_features = 100, 200
         X = np.random.randn(n_samples, n_features)
         k = 5
         # beta generated with k nonzeros
         #coef = 10 * np.random.randn(n_features)
         coef = 10 * np.ones(n_features)
         inds = np.arange(n features)
         np.random.shuffle(inds)
         coef[inds[k:]] = 0 # sparsify coef
         y = np.dot(X, coef)
         # add noise
         y += 0.01 * np.random.normal((n samples,))
         # Split data in train set and test set
         n \text{ samples} = X.shape[0]
         X train, y train = X[:25], y[:25]
         X \text{ test, } y \text{ test = } X[25:], y[25:]
In [4]:
         # Problem 2
In [5]:
         #2.1
         #Step forward feature selection starts with the evaluation of each individual fe
         #and selects that which results in the best performing selected algorithm model
         alpha = 0.2
         train samples = 25
         test samples = 75
         ## Lecture 8 explains idea - we need to project y onto the feature then replace
         # use this to find the best error
         def evaluate_feature_error(feature_train, feature_test, y_train=y_train, y_test=
             bias = alpha*np.eye(feature train.shape[1])
             inverse = np.linalg.inv(np.dot(feature train.T, feature train)) + bias
             proj 1 = np.dot(feature train.T, np.reshape(y train, (train samples,1)))
             beta hat = np.dot(inverse, proj 1)
             proj 2 = np.reshape(y test, (test samples,1))
             error_vec = proj_2 - np.dot(feature_test, beta_hat)
```

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return np.linalg.norm(error vec, ord=2)**2
feature selection = []
errors = [None for i in range(0, n features)]
best_scores = [] # best scores will save all the minimum errors from the feature
train_shape = (train_samples, 1)
test shape = (test samples, 1)
for i in range(0, n_features):
   bias_train = np.ones(train_shape)
   bias_test = np.ones(test_shape)
    if len(feature selection) == 0:
       X_feat_train = bias_train
       X_feat_test = bias_test
    else:
        for old feat in feature selection:
            old_feat_train = np.reshape(X_train[:,old_feat], train_shape)
            old_feat_test = np.reshape(X_test[:,old_feat], test_shape)
            X_feat_train = np.hstack((bias_train, old_feat_train))
            X feat test = np.hstack((bias test, old feat test))
    for new feat in range(0, n features):
        if new_feat not in feature_selection:
            new_feat_train = np.reshape(X_train[:,new_feat], train_shape)
            new feat test = np.reshape(X test[:,new feat], test shape)
            feature train = np.hstack((X feat train, new feat train))
            feature test = np.hstack((X feat test, new feat test))
            errors[new_feat] = evaluate_feature_error(feature_train, feature_tes
   min error = min([i for i in errors if i is not None])
    selected feature = errors.index(min error)
   # print(selected feature)
    feature selection.append(selected feature)
   best_scores.append(min_error)
    errors = [None for i in range(0,n features)] # reset the errors
print("The order of the features is:")
print(feature selection)
#print(len(feature selection) == len(set(feature selection)))
```

O2

The order of the features is: [138, 79, 108, 116, 166, 54, 163, 89, 194, 35, 113, 131, 130, 74, 160, 154, 95, 118, 90, 103, 167, 32, 180, 26, 36, 2, 68, 162, 135, 97, 91, 52, 13, 187, 44, 19 2, 179, 117, 188, 144, 119, 37, 47, 149, 29, 127, 101, 7, 28, 139, 141, 43, 94,

146, 189, 92, 198, 129, 31, 159, 60, 99, 122, 51, 12, 110, 62, 45, 140, 64, 121, 61, 88, 115, 169, 14, 148, 3, 65, 145, 5, 120, 58, 55, 105, 171, 137, 0, 151, 18 1, 83, 70, 33, 143, 16, 184, 104, 4, 77, 8, 190, 153, 107, 63, 40, 170, 81, 19, 30, 75, 132, 53, 193, 158, 9, 123, 172, 136, 174, 157, 165, 84, 176, 69, 10, 10 0, 196, 59, 133, 18, 82, 183, 6, 76, 22, 98, 109, 1, 24, 186, 87, 93, 102, 71, 7 3, 199, 48, 85, 39, 86, 38, 17, 66, 57, 126, 155, 164, 168, 80, 177, 25, 152, 2 1, 161, 124, 134, 106, 156, 46, 128, 150, 147, 111, 27, 41, 175, 185, 112, 11, 4 2, 23, 49, 182, 56, 34, 125, 197, 195, 78, 67, 191, 173, 114, 20, 142, 50, 72, 1 78, 15, 96]

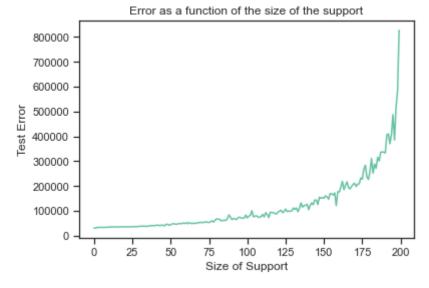
```
In [6]: #2.2
#Plot test error as a function of the size of the support. Can you use this to

txt = plt.plot(best_scores)
txt = plt.xlabel("Size of Support")
txt = plt.ylabel("Test Error")
txt = plt.title("Error as a function of the size of the support")

print("We can use this to recover the true support. We need to find the feature
print("Index " + str(best_scores.index(min(best_scores))) + " error = " + str(m)
```

We can use this to recover the true support. We need to find the feature with the  ${\tt minimal}\ {\tt test-error}$ 

Index 1 error = 30052.233103646933



```
In [7]: # 2.3
    from sklearn.model_selection import GridSearchCV
    from sklearn.metrics import classification_report
    from sklearn.linear_model import Lasso

alphas = np.logspace(-4, -0.5, 30)
    lasso = Lasso(random_state=0, max_iter=10000)

tuned_parameters = [{'alpha': alphas}]
    n_folds = 5

clf = GridSearchCV(lasso, tuned_parameters, cv=n_folds, scoring='r2')

clf.fit(X_train, y_train)

print("Best Alpha:", clf.best_params_)
    means = clf.cv_results_['mean_test_score']
    stds = clf.cv_results_['std_test_score']
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          for mean, std, params in zip(means, stds, clf.cv_results_['params']):
                  print("%0.3f (+/-%0.03f) for %r"
                        % (mean, std * 2, params))
         Best Alpha: {'alpha': 0.0028072162039411755}
         -2.903 (+/-5.040) for {'alpha': 0.0001}
         -2.903 (+/-5.039) for {'alpha': 0.00013203517797162948}
         -2.903 (+/-5.039) for {'alpha': 0.00017433288221999874}
         -2.855 (+/-4.990) for {'alpha': 0.00023018073130224678}
         -2.563 (+/-4.171) for {'alpha': 0.0003039195382313198}
         -2.222 (+/-3.186) for {'alpha': 0.0004012807031942776}
         -1.367 (+/-3.581) for {'alpha': 0.0005298316906283707}
         -0.753 (+/-2.872) for {'alpha': 0.0006995642156712634}
         -0.691 (+/-2.747) for {'alpha': 0.0009236708571873865}
         -0.455 (+/-1.938) for {'alpha': 0.0012195704601594415}
         -0.138 (+/-0.999) for {'alpha': 0.0016102620275609393}
         -0.127 (+/-1.021) for {'alpha': 0.0021261123338996556}
         -0.114 (+/-0.894) for {'alpha': 0.0028072162039411755}
         -0.168 (+/-0.914) for {'alpha': 0.0037065129109221566}
         -0.292 (+/-1.062) for {'alpha': 0.004893900918477494}
         -0.286 (+/-1.009) for {'alpha': 0.006461670787466976}
         -0.407 (+/-0.931) for {'alpha': 0.008531678524172814}
         -0.639 (+/-1.343) for {'alpha': 0.011264816923358867}
         -0.644 (+/-1.348) for {'alpha': 0.014873521072935119}
         -1.106 (+/-2.020) for {'alpha': 0.0196382800192977}
         -1.152 (+/-1.992) for {'alpha': 0.02592943797404667}
         -1.162 (+/-1.986) for {'alpha': 0.03423597957607583}
         -1.211 (+/-2.002) for {'alpha': 0.04520353656360245}
         -1.234 (+/-2.021) for {'alpha': 0.05968456995122311}
         -1.248 (+/-2.078) for {'alpha': 0.07880462815669913}
         -1.260 (+/-2.134) for {'alpha': 0.10404983103657853}
         -1.252 (+/-2.128) for {'alpha': 0.1373823795883264}
         -1.240 (+/-2.121) for {'alpha': 0.1813930693911063}
         -1.230 (+/-2.114) for {'alpha': 0.2395026619987486}
         -1.211 (+/-2.101) for {'alpha': 0.31622776601683794}
In [14]:
          # 2.4
          best alphas = []
          for n fold in range(3,12):
              print("Number of folds: ", n fold)
              clf = GridSearchCV(lasso, tuned parameters, cv=n fold, scoring='r2')
              clf.fit(X train, y train)
              print("Best Alpha:", clf.best_params_)
              best alphas.append(clf.best params ['alpha'])
          print()
          print("As the number of folds increases, the hyper parameter increases till it p
         Number of folds: 3
         Best Alpha: { 'alpha': 0.0021261123338996556}
         Number of folds: 4
         Best Alpha: { 'alpha': 0.0021261123338996556}
         Number of folds: 5
         Best Alpha: { 'alpha': 0.0028072162039411755}
         Number of folds: 6
         Best Alpha: { 'alpha': 0.0021261123338996556}
         Number of folds: 7
         Best Alpha: { 'alpha': 0.008531678524172814}
```

Best Alpha: { 'alpha': 0.31622776601683794}

Number of folds: 8

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Number of folds: 9
Best Alpha: {'alpha': 0.0009236708571873865}
Number of folds: 10
Best Alpha: {'alpha': 0.0009236708571873865}
Number of folds: 11
Best Alpha: {'alpha': 0.0009236708571873865}
```

As the number of folds increases, the hyper parameter increases till it peaked a t 8 folds, then it went started to decrease.

## 2.5

LassoCV will choose a set of alphas and perform comparison searches over then, using cross-validation to generate lasso models and internally score the results. The previous step used GridsearchCV + Lasso model, which is essentially performing the same functions as LassoCV. The results from LassoCV and GridsearchCV + Lasso should agree with each other if the parameters were configured the same way.



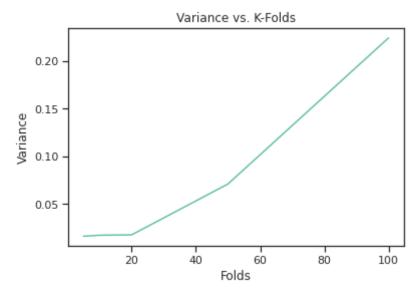
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In [8]:
          import numpy as np
          import matplotlib.pyplot as plt
          from sklearn import datasets, linear_model
          from sklearn.preprocessing import MinMaxScaler
          from sklearn.svm import SVR
          from sklearn.model_selection import KFold
          import pandas as pd
          from pandas import DataFrame, Series
          import seaborn as sns
          sns.set(style='ticks', palette='Set2')
          %matplotlib inline
In [10]:
          #read_file = pd.read_csv ('housing.data.txt', header = None, delim_whitespace= T
          #read file.to csv ('housing data.csv', index=None)
In [11]:
          housing = pd.read_csv('housing_data.csv')
In [12]:
          X = housing.iloc[:, [0, 12]]
          y = housing.iloc[:, 13]
In [13]:
          scaler = MinMaxScaler(feature_range=(0, 1))
          X = scaler.fit transform(X)
In [14]:
          bias = []
          variance total = []
In [15]:
          mean score = []
          variance = []
          best_svr = SVR(kernel='rbf')
          for i in np.arange(1,100):
              scores = []
              cv = KFold(5, shuffle= True)
              for train_index, test_index in cv.split(X):
                  X train, X test, y train, y test = X[train index], X[test index], y[trai
                  best_svr.fit(X_train, y_train)
                  scores.append(best svr.score(X test, y test))
              mean score.append(np.mean(scores))
              variance.append(np.std(scores))
          bias.append(np.mean(mean score))
          variance total.append(np.std(variance))
          print("50 Folds: Mean - " + str(np.mean(mean score)) + " | Variance - " + str(np.mean(mean score))
         50 Folds: Mean - 0.5756046264442204 | Variance - 0.01621214575019245
In [16]:
          mean score = []
          variance = []
          best svr = SVR(kernel='rbf')
          for i in np.arange(1,100):
              scores = []
              cv = KFold(10, shuffle= True)
```

```
for train_index, test_index in cv.split(X):
                  X_train, X_test, y_train, y_test = X[train_index], X[test_index], y[trai
                  best_svr.fit(X_train, y_train)
                  scores.append(best_svr.score(X_test, y_test))
              mean_score.append(np.mean(scores))
              variance.append(np.std(scores))
          bias.append(np.mean(mean_score))
          variance_total.append(np.std(variance))
          print("50 Folds: Mean - " + str(np.mean(mean_score)) + " | Variance - " + str(np.mean(mean_score))
         50 Folds: Mean - 0.5833956249966143 | Variance - 0.017251725110677237
In [17]:
          mean_score = []
          variance = []
          best_svr = SVR(kernel='rbf')
          for i in np.arange(1,100):
              scores = []
              cv = KFold(20, shuffle= True)
              for train_index, test_index in cv.split(X):
                  X_train, X_test, y_train, y_test = X[train_index], X[test_index], y[trai
                  best svr.fit(X train, y train)
                  scores.append(best_svr.score(X_test, y_test))
              mean_score.append(np.mean(scores))
              variance.append(np.std(scores))
          bias.append(np.mean(mean_score))
          variance_total.append(np.std(variance))
          print("50 Folds: Mean - " + str(np.mean(mean_score)) + " | Variance - " + str(np.mean(mean_score))
         50 Folds: Mean - 0.5824509527907801 | Variance - 0.017676315213079788
In [18]:
          mean_score = []
          variance = []
          best_svr = SVR(kernel='rbf')
          for i in np.arange(1,100):
              scores = []
              cv = KFold(50, shuffle= True)
              for train index, test index in cv.split(X):
                  X_train, X_test, y_train, y_test = X[train_index], X[test_index], y[trai
                  best_svr.fit(X_train, y_train)
                  scores.append(best_svr.score(X_test, y_test))
              mean score.append(np.mean(scores))
              variance.append(np.std(scores))
          bias.append(np.mean(mean score))
          variance_total.append(np.std(variance))
          print("50 Folds: Mean - " + str(np.mean(mean_score)) + " | Variance - " + str(np.mean(mean_score))
         50 Folds: Mean - 0.5438518638280582 | Variance - 0.07102601150329856
In [19]:
          mean score = []
          variance = []
          best svr = SVR(kernel='rbf')
          for i in np.arange(1,100):
              scores = []
              cv = KFold(50, shuffle= True)
              for train index, test index in cv.split(X):
                  X_train, X_test, y_train, y_test = X[train_index], X[test_index], y[trai
                  best_svr.fit(X_train, y_train)
                  scores.append(best_svr.score(X_test, y_test))
              mean_score.append(np.mean(scores))
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               variance.append(np.std(scores))
          bias.append(np.mean(mean_score))
          variance_total.append(np.std(variance))
          print("50 Folds: Mean - " + str(np.mean(mean_score)) + " | Variance - " + str(np
          50 Folds: Mean - 0.5426904073913306 | Variance - 0.2241660726495882
In [20]:
          k = [5, 10, 20, 50, 100]
In [21]:
          plt.plot(k, bias)
          plt.title('Bias vs. K-Folds')
          plt.xlabel('Folds')
          plt.ylabel('Bias')
Out[21]: Text(0, 0.5, 'Bias')
                                  Bias vs. K-Folds
            0.585
            0.580
            0.575
            0.570
            0.565
            0.560
            0.555
            0.550
            0.545
                                  40
                         20
                                           60
                                                    80
                                                            100
                                      Folds
In [22]:
          plt.plot(k, variance total)
          plt.title('Variance vs. K-Folds')
          plt.xlabel('Folds')
          plt.ylabel('Variance')
```

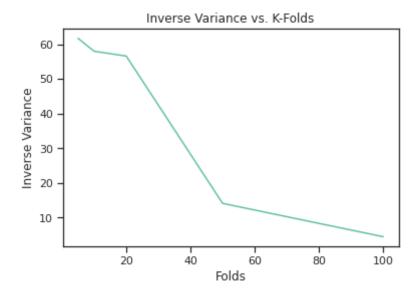
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```
In [26]: v = np.array(variance_total)

In [27]: plt.plot(k, 1/v)
    plt.title('Inverse Variance vs. K-Folds')
    plt.xlabel('Folds')
    plt.ylabel('Inverse Variance')
```

Out[27]: Text(0, 0.5, 'Inverse Variance')



In []: # Looking at this simulation, it seems as though the claim that bias increases a # However, in this graph, at a small K, our bias increases as we get larger rat # This is also sometimes true with our Variance. During some iterations, we can