Needell Coen HW3

February 9, 2020

```
[1]: import numpy as np
import pandas as pd
from matplotlib import pyplot as plt
```

- 0.1 # Conceptual Exercises
- 0.2 ## Training/test error for subset selection

0.2.1 1

0.2.2 2

```
[3]: from sklearn.model_selection import train_test_split

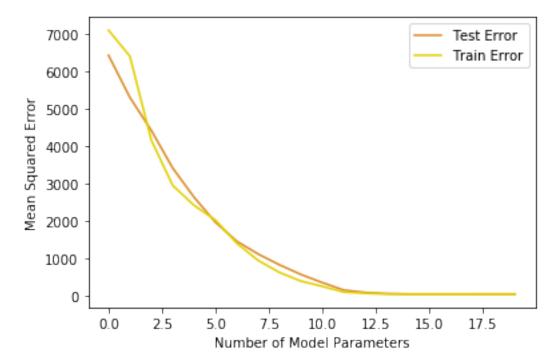
x_train, x_test, y_train, y_test = train_test_split(df1, y, test_size=900)
```

0.2.3 3

```
[4]: from sklearn.linear_model import LinearRegression from sklearn.metrics import mean_squared_error as mse from itertools import combinations from functools import partial import wes
```

```
[5]: def process_subset(feats, xtr, xte, ytr, yte, returnmodel=False):
         train_feats = xtr[feats]
         test_feats = xte[feats]
         model = LinearRegression()
         model.fit(train_feats, ytr)
         if returnmodel:
             return model
         errte = mse(yte, model.predict(test_feats))
         errtr = mse(ytr, model.predict(train_feats))
         return errtr, errte
     # I'm using a forward stepwise algorithm, assuming that the features
     # have some best order which is independent of other features.
     # That assumption is baked into linear models though, so it's safe here.
     # Also we know the data generating function, and we know that they're
     # independant, so we know that both algorithms go to the same result.
     # Also 2 ** 20 = 1,000,000 which is a little out-of-feasibility for python.
     def forward_stepwise(xtr, xte, ytr, yte):
         n = len(xtr.keys())
         model = set() # This is a set to which we will add features.
         params = []
         testerrs = []
         trainerrs = []
         models = \Pi
         # This is a list where we'll record
         #the best process subset outputs
         for k in range(0, n):
             augs = [str(st) for st in xtr.keys() if st not in model]
             results = []
             for x in augs:
                 results.append(process_subset(list(model) + [x],
                               xtr, xte, ytr, yte))
             results = np.array(results)
             best = np.argmin(results[:, 1])
             model.add(augs[best])
             betr, bete = results[best]
             params.append(augs[best])
             testerrs.append(bete)
             trainerrs.append(betr)
         record = pd.DataFrame({"param":params,
                               "test_err":testerrs,
                               "train_err":trainerrs})
         return record
     best_selec = forward_stepwise(x_train, x_test, y_train, y_test)
```

```
[7]: wes.set_palette('FantasticFox1')
   plt.plot(best_selec.test_err, label='Test Error')
   plt.plot(best_selec.train_err, label='Train Error')
   plt.xlabel('Number of Model Parameters')
   plt.ylabel('Mean Squared Error')
   plt.legend()
   plt.show()
   best_selec
```



```
[7]:
        param
                  test_err
                               train_err
               6416.784511
                             7089.241271
     0
           8x
     1
          x16
               5289.490369
                             6390.208384
     2
          x11
               4409.229291
                             4147.456895
     3
          x12
               3402.061119
                             2937.561717
     4
          x19
               2620.542325
                             2405.103896
     5
               1950.391660
                             2010.545592
           x9
     6
           x3
               1443.183115
                             1389.383800
     7
          x10
               1100.965204
                              928.892155
     8
          x15
                811.961551
                              608.203719
     9
           x5
                555.418121
                              378.544639
     10
                337.576521
                              238.096790
          x20
     11
           x2
                138.647058
                               85.752510
     12
          x17
                 74.697425
                               54.693290
     13
                 45.602417
                               30.619249
           x7
     14
           x1
                 34.774484
                               27.710232
```

```
15
             34.737711
                           27.688195
      x6
             34.740764
16
     x13
                           27.671565
17
     x18
             34.782783
                           27.651672
18
     x14
             35.063060
                           27.478137
19
             36.026689
      x4
                           27.269170
```

Unsurprisingly, it starts to even out around 13. If we look at the list of betas it's easy to see why, we know that 5 of the variables are useless, and another two have $|\beta| < 1$. If you look at the parameter that's added in for each step, and compare that to the list of betas, you can see the logic of it. The betas which are set to zero are associated with parameters that are included last.

```
[8]: pd.DataFrame(betas, index=['beta']).transpose()
[8]:
              beta
         -0.717342
     x1
     x2
         -2.844386
     xЗ
         -4.660543
     x4
          0.000000
         -2.933093
     x5
         -0.002561
     x6
          1.038511
     x7
         -9.030795
     8x
     x9
         -5.204751
          3.710081
     x10
     x11
          6.849511
          6.229572
     x12
     x13
          0.00000
     x14
          0.000000
     x15
          3.321539
     x16
          5.709028
     x17
          1.531583
     x18
          0.000000
     x19 -5.046486
     x20 -2.791455
```

The high efficiency models (15 and 14) remove x1 and x6 plus the zeroes. Then the zeroes are added back in in order.

0.2.4 4

See plot for # 3. ### 5

```
[9]: best_test = np.argmin(best_selec.test_err)
m = ','.join(best_selec.param[:best_test].sort_values())
m
```

```
[9]: 'x1,x10,x11,x12,x15,x16,x17,x19,x2,x20,x3,x5,x7,x8,x9'
```

So the best model is the model with 16 parameters. Why exactly it's at this number of parameters and not smaller or larger is somewhat interesting. We're measuring performance on the test set, for one, and chances are good that noise effects detected in the additional variables after the 16th are causing some overfitting when those variables are in the linear model. ### 6 We see that the best model contains all of the features except for those which have coefficients closest to zero. Give or take a few. Because of noise, we see one of the coefficient-zero parameters in the mix, as well as a close to zero parameter treated as if it's coefficient is zero.

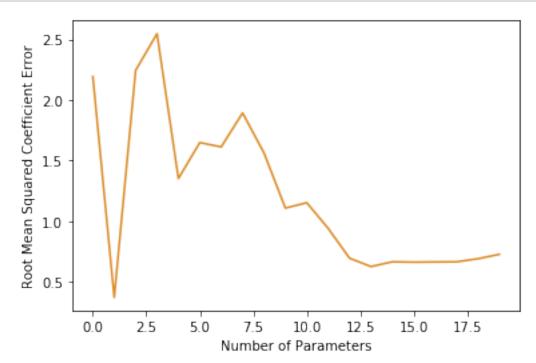
```
Γ10]:
                  coefs
        feats
                            betas
           x8 -9.423881 -9.030795
      0
      1
          x16 5.525469 5.709028
      2
          x11 6.976860 6.849511
      3
          x12 6.230386 6.229572
      4
          x19 -4.830579 -5.046486
           x9 -5.160344 -5.204751
      6
           x3 -4.709259 -4.660543
      7
          x10 3.847959 3.710081
      8
          x15 3.447054 3.321539
      9
           x5 -3.039797 -2.933093
      10
          x20 -2.759123 -2.791455
      11
           x2 -2.932852 -2.844386
      12
          x17 1.387939 1.531583
      13
           x7 0.886101 1.038511
           x1 -0.424171 -0.717342
```

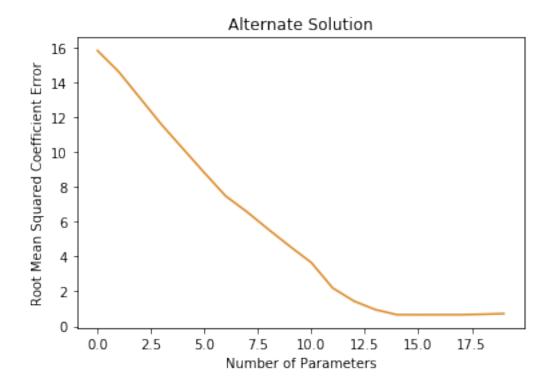
0.2.5 7

```
for r in range(1, 21):
    feats = best_selec.param[:r]
    m = process_subset(feats, x_train, x_test, y_train, y_test,
    returnmodel=True)
    bhat = m.coef_
    b = np.array([betas[f] for f in feats])
    inner = (b - bhat) ** 2
    outer = np.sqrt(np.sum(inner))
    rmcoefaccs.append(outer)

plt.plot(rmcoefaccs)
    plt.ylabel('Root Mean Squared Coefficient Error')
    plt.xlabel('Number of Parameters')
```

```
plt.show()
rmcoefaccs2 = []
for r in range(1, 21):
    feats = best_selec.param[:r]
    m = process_subset(feats, x_train, x_test, y_train, y_test,__
→returnmodel=True)
    coefs = {f: co for f, co in zip(feats, m.coef_)}
    bhat = np.array([coefs[f] if f in coefs else 0 for f in betas])
    b = np.array(list(betas.values()))
    inner = (b - bhat) ** 2
    outer = np.sqrt(np.sum(inner))
    rmcoefaccs2.append(outer)
plt.plot(rmcoefaccs2)
plt.ylabel('Root Mean Squared Coefficient Error')
plt.xlabel('Number of Parameters')
plt.title('Alternate Solution')
plt.show()
```





The first thing we notice is this huge hump between 5 parameters and 10. This is due to the fact that the linear regression algorithm is using the available parameters to spuriously explain other parameters. We can compare this to the MSE plot where the error pretty much declines steadily as we add in more parameters, in this case we see a sort of distribution of error, before we get into the more accurate models about p = 15.

Since there was some widespread confusion about the meaning of this problem, I've included an alternate answer for the case where the Root Mean Squared Coefficient Error should include features that are not in the model. For these cases I have set $\hat{\beta}$ to zero for betas which are not included in the model. In this case it's easier to see that the model is off. But this information seems trivial. If you're judging the model based on things that aren't even included, then you just end up with a value that's remarkably similar to MSE. This measure is artificially inflated for smaller models, whereas the first graph shows how close to 'correct' the model gets for the features that are included.

0.3 # Application Exercises

Please note that sklearn calls the λ parameter α . And the α parameter l_1 . I will use this convention.

```
[15]: from sklearn.linear_model import RidgeCV, LassoCV, ElasticNetCV
    gss_tr = pd.read_csv('data/gss_train.csv')
    gss_te = pd.read_csv('data/gss_test.csv')
    x_tr = gss_tr.drop('egalit_scale', axis=1)
    y_tr = gss_tr.egalit_scale
```

```
x_te = gss_te.drop('egalit_scale', axis=1)
y_te = gss_te.egalit_scale
```

0.3.1 1

```
[16]: def report_mse(model):
    model.fit(x_tr, y_tr)
    err = mse(model.predict(x_te), y_te)
    return err, model
```

```
[17]: print(report_mse(LinearRegression())[0])
```

63.213629623014995

0.3.2 2

```
[18]: print(report_mse(RidgeCV(alphas=np.logspace(-5, 5)))[0])
```

62.202521583380204

A little better...

mse is 62.90446289883912

There are 54 non-zero coefficent estimates

woof

```
[21]: mes, mod = report_mse(ElasticNetCV(l1_ratio=np.linspace(.1,1,11)))
    print(f'mse is {mes}')
    print(f'There are {sum(np.isclose(mod.coef_, 0))} non-zero coefficient
    →estimates')
```

mse is 62.81645879274796

There are 54 non-zero coefficient estimates

Gotta say, we're not exactly killing it on the accuracy front. The feature ranges from 1 to 35, and the best of these methods, RidgeCV, was off by 7.8 points on average. Which I guess isn't so bad, but a more complicated method would, I'm sure, do a lot better. One of the interesting things from this exercise is that all of these MSEs are within 1 point of each other, which tells us that the regression method isn't having too much of an effect on the results. This implies that what we really need is a non-linear model, as we've probably hit the upper bound on what is possible with linear regressions.

I do also find it interesting that the Elastic Net considers more coefficients than the Lasso.