lab3

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0.1 Lab 3: Linear Regression

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```
[]: import numpy as np
import pandas as pd
from matplotlib.pyplot import subplots
import statsmodels.api as sm
```

Here we import some submodules from the statsmodels packages. We rename the named exports to keep the *namespace* clean.

We also use some functions from ISLP

```
[]: from ISLP import load_data from ISLP.models import (ModelSpec as MS, summarize, poly)
```

0.1.1 Objects & Namespaces

We can list the objects in the current namespace using dir() (the following is truncated).

```
[]: dir()[:5]
```

```
[]: ['A', 'Boston', 'Carseats', 'In', 'MS']
```

Each python object has its own namespace. We can also access this namespace of specific objects in the following way. Notice that numpy arrays has a sum property!

```
[]: A = np.arange(0,10,2)
dir(A)[:5]
```

```
[]: ['T', '__abs__', '__add__', '__and__', '__array__']
```

```
[]: A.sum()
```

[]: 20

0.1.2 Simple Linear Regression

First we will create design/model matrices using ModelSpec() transform from ISLP.models.

The task will be to build a model using 13 predictors to predict mev.

```
[]: Boston = load_data("Boston")
Boston.columns
```

```
[]: Index(['crim', 'zn', 'indus', 'chas', 'nox', 'rm', 'age', 'dis', 'rad', 'tax', 'ptratio', 'lstat', 'medv'], dtype='object')
```

We can learn more about the data by typing Boston?. Our first model will just use a single predictor to predict medv.

```
[ ]: Boston?
```

```
[]: intercept lstat
0 1.0 4.98
1 1.0 9.14
2 1.0 4.03
3 1.0 2.94
```

```
[]: # Here we fit the model
y = Boston['medv']
model = sm.OLS(y, X)
results = model.fit()
```

To get more information about the fitted model we can use the **summarize** function. Which gives useful statistics like standard errors, t-statistics, and p-values.

```
[]: summarize(results)
```

```
[]: coef std err t P>|t| intercept 34.5538 0.563 61.415 0.0 lstat -0.9500 0.039 -24.528 0.0
```

Using Transformations: Fit and Transform In practice models usually contain more than one predictor. In addition, transformations of the variables and interactions terms can be added. We can rely on tools from sklearn to create these transforms.

```
[]: design = MS(['lstat'])
  design = design.fit(Boston)
  X = design.transform(Boston)
  X[:4]
```

```
[]: intercept lstat
0 1.0 4.98
1 1.0 9.14
2 1.0 4.03
3 1.0 2.94
```

In our previous data set fit() does not do much, it just checks if the lsat variable exists in the data set.

While this processed was executed in two lines of code, the design object is changed after calling fit().

```
[]: results.summary() results.params
```

[]: intercept 34.553841 lstat -0.950049

dtype: float64

We can also use get_prediction to obtain new labels with confidence intervals from data not in the training set.

```
[ ]: new_df = pd.DataFrame({'lstat':[5, 10, 15]})
newX = design.transform(new_df)
newX
```

```
[]: intercept lstat
0 1.0 5
1 1.0 10
2 1.0 15
```

```
[]: # Here we obtain predictions for the new data.

new_predictions = results.get_prediction(newX)

new_predictions.predicted_mean
```

[]: array([29.80359411, 25.05334734, 20.30310057])

```
[]: # Extract confidence intervals
new_predictions.conf_int(alpha=0.05)
```

```
[]: # Extract prediction intervals new_predictions.conf_int(obs=True, alpha=0.05)
```

As expected the prediction intervals are centered around the same values as the confidence intervals, but are wider. This is because the confidence interval relays information on the *average* of our data, while the prediction interval gives an information on the confidence for a new, single city, which incorporates the irreducible error in addition to the reducible error.

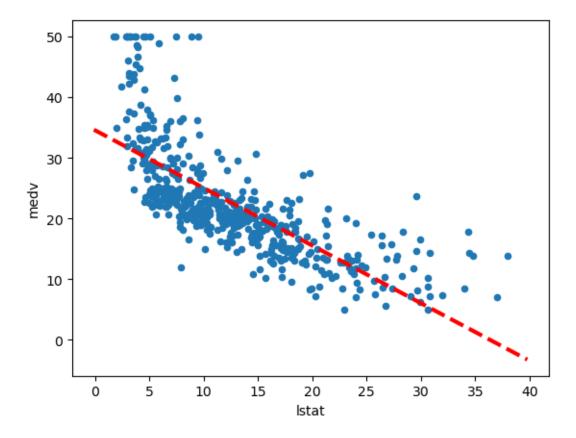
Defining functions Here we define some helper functions to visualize the fits and residuals.

```
[]: def abline(ax, b, m):
    "Add a line with slope m and intercept b to ax"
    xlim = ax.get_xlim()
    ylim = [m * xlim[0] + b, m * xlim[1] + b]
    ax.plot(xlim, ylim)
```

By adding in *args and **kwargs we can any number of non-named arguments and any number of named arguments.

```
[]: def abline(ax, b, m, *args, **kwargs):
    "Add a line with slope m and intercept b to ax"
    xlim = ax.get_xlim()
    ylim = [m * xlim[0] + b, m * xlim[1] + b]
    ax.plot(xlim, ylim, *args, **kwargs)
```

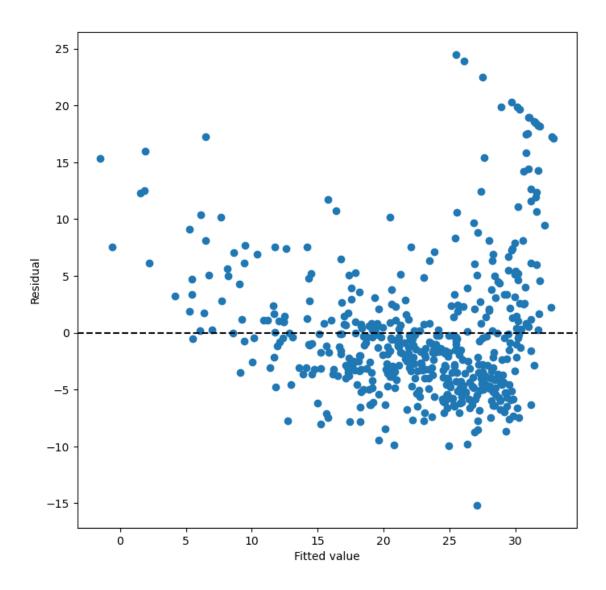
```
[]: ax = Boston.plot.scatter('lstat', 'medv')
abline(ax,
    results.params[0],
    results.params[1],
    'r--', linewidth=3)
```



The figure above shows that there is some evidence of a non-linear relationship between lsat and mev.

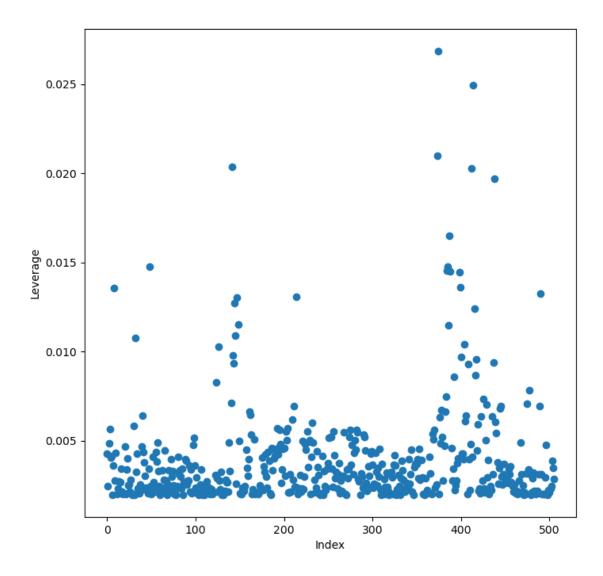
Here we analyze plots that visualize information of the fitted values and residuals. The residual plot also shows evidence of a non-linear relationship. We can also compute leverage statistics by using the hat_matrix_diag attribute from the object returned by get_influence().

```
[]: ax = subplots(figsize=(8,8))[1]
   ax.scatter(results.fittedvalues , results.resid)
   ax.set_xlabel('Fitted value')
   ax.set_ylabel('Residual')
   ax.axhline(0, c='k', ls='--');
```



```
[]: infl = results.get_influence()
ax = subplots(figsize=(8,8))[1]
ax.scatter(np.arange(X.shape[0]), infl.hat_matrix_diag)
ax.set_xlabel('Index')
ax.set_ylabel('Leverage')
```

[]: Text(0, 0.5, 'Leverage')



```
[]: # This identifies the index of the largest element in the array which in turn

→identifies which observation has the largest leverage statistic.

np.argmax(infl.hat_matrix_diag)
```

[]: 374

0.1.3 Multiple Linear Regression

To use multiple linear regression using least squares we can use ModelSpec() transform to construct the required model matrix. In the following we add age to the predictors.

```
[]: X = MS(['lstat', 'age']).fit_transform(Boston)
model1 = sm.OLS(y, X)
results1 = model1.fit()
```

```
[]:
                   coef
                         std err
                                          P>|t|
                           0.731 45.458
                                          0.000
     intercept
                33.2228
     lstat
                -1.0321
                           0.048 -21.416
                                          0.000
                 0.0345
                           0.012
                                   2.826
                                          0.005
     age
    We can easily create a list of all the predictors by dropping a single one instead of typing them all
[]: terms = Boston.columns.drop('medv')
     terms
[]: Index(['crim', 'zn', 'indus', 'chas', 'nox', 'rm', 'age', 'dis', 'rad', 'tax',
            'ptratio', 'lstat'],
           dtype='object')
[]: # Fit the multiple linear regression model.
     X = MS(terms).fit_transform(Boston)
     model = sm.OLS(y, X)
     results = model.fit()
     summarize(results)
[]:
                                       t P>|t|
                        std err
                   coef
     intercept 41.6173
                           4.936
                                   8.431 0.000
                           0.033 -3.678 0.000
     crim
                -0.1214
                 0.0470
                           0.014
                                   3.384 0.001
     zn
     indus
                 0.0135
                           0.062
                                   0.217 0.829
                 2.8400
                           0.870
                                   3.264 0.001
     chas
                           3.851 -4.870 0.000
    nox
               -18.7580
                 3.6581
                           0.420
                                   8.705 0.000
     rm
     age
                 0.0036
                           0.013
                                   0.271 0.787
                           0.202 -7.394 0.000
     dis
                -1.4908
     rad
                 0.2894
                           0.067
                                   4.325 0.000
                           0.004 -3.337 0.001
     tax
                -0.0127
                -0.9375
                           0.132 -7.091 0.000
     ptratio
                -0.5520
                           0.051 -10.897 0.000
     lstat
[]: # Using all the variables as predictors except age
     minus_age = Boston.columns.drop(['medv', 'age'])
     Xma = MS(minus_age).fit_transform(Boston)
     model1 = sm.OLS(y, Xma)
     summarize(model1.fit())
[]:
                                       t P>|t|
                   coef std err
     intercept 41.5251
                           4.920
                                   8.441
                                          0.000
                -0.1214
                           0.033 -3.683 0.000
     crim
                 0.0465
                           0.014
                                   3.379
                                          0.001
     zn
```

summarize(results1)

```
0.217 0.829
indus
            0.0135
                     0.062
            2.8528
                     0.868
                              3.287 0.001
chas
nox
          -18.4851
                     3.714 -4.978 0.000
            3.6811
                     0.411
                              8.951 0.000
rm
           -1.5068
                     0.193 -7.825 0.000
dis
            0.2879
                     0.067
                              4.322 0.000
rad
                     0.004 -3.333 0.001
           -0.0127
tax
ptratio
           -0.9346
                     0.132 -7.099 0.000
lstat
           -0.5474
                     0.048 -11.483 0.000
```

Multivariable goodness of fit. The individual components of results can be accessed by name.

To access this information we can use list comprehension which is a simple and powerful way to form a list in python.

```
[]:
                    vif
     crim
              1.767486
              2.298459
     zn
     indus
              3.987181
     chas
              1.071168
     nox
              4.369093
              1.912532
     rm
              3.088232
     age
     dis
              3.954037
     rad
              7.445301
     tax
              9.002158
              1.797060
     ptratio
     lstat
              2.870777
```

```
[]: # We can use list comprehension to perform repetitive operations
vals = []
for i in range(1, X.values.shape[1]):
    vals.append(VIF(X.values, i))
```

Interaction terms To add an interaction terms we can include a tuple in the model matrix.

```
[]: X = MS(['lstat', 'age', ('lstat', 'age')]).fit_transform(Boston)
model2 = sm.OLS(y, X)
summarize(model2.fit())
```

```
[]:
                          std err
                                        t P>|t|
                    coef
     intercept
                36.0885
                            1.470
                                   24.553
                                           0.000
                -1.3921
                                            0.000
     lstat
                            0.167
                                   -8.313
                -0.0007
                            0.020
                                   -0.036
                                            0.971
     age
                                    2.244
     lstat:age
                 0.0042
                            0.002
                                            0.025
```

Non-linear Transformation of the Predictors We can also include non-linear transformations of the predictors in addition to just the interaction terms and features themeselves.

```
[]: X = MS([poly('lstat', degree=2), 'age']).fit_transform(Boston)
model3 = sm.OLS(y, X)
results3 = model3.fit()
summarize(results3)
```

```
[]:
                                     coef
                                            std err
                                                           t
                                                              P>|t|
     intercept
                                  17.7151
                                              0.781
                                                     22.681
                                                                0.0
     poly(lstat, degree=2)[0] -179.2279
                                              6.733 -26.620
                                                                0.0
     poly(lstat, degree=2)[1]
                                  72.9908
                                              5.482
                                                      13.315
                                                                0.0
                                   0.0703
                                              0.011
     age
                                                       6.471
                                                                0.0
```

We see that the p-value associated with the quadratic term is near zero. This means that the model was improved by adding that term.

```
[]: anova_lm(results1, results3)
```

```
[]:
        df_resid
                                   df_diff
                                                 ss_diff
                                                                     F
                                                                               Pr(>F)
                             ssr
     0
            503.0
                   19168.128609
                                       0.0
                                                     NaN
                                                                   NaN
                                                                                  NaN
     1
            502.0
                   14165.613251
                                       1.0
                                             5002.515357
                                                           177.278785
                                                                        7.468491e-35
```

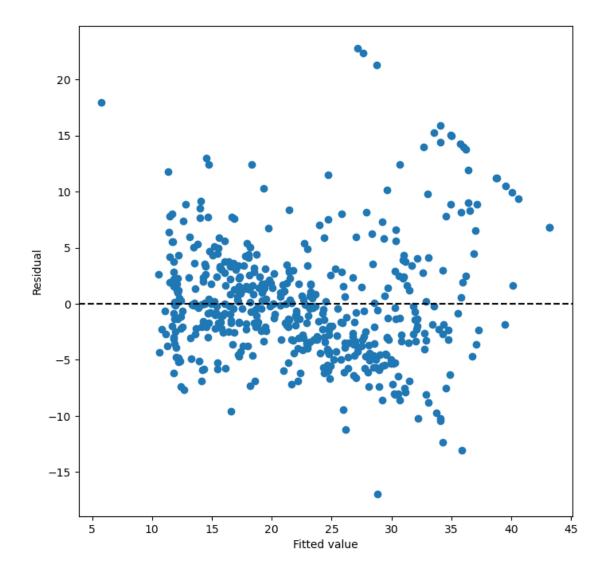
To get a better understanding of how the quadratic fit is superior to the linear fit we can use the anova_lm function, which performs a hypothesis test against the two models.

The F statistics is large and its corresponding p-value is near zero, suggesting ample evidence that the bigger model is better.

In the figure below we see that the residuals do not have a discernible structure or pattern. Therefore, we can be confident that our model can appropriately describe the underlying pattern of the data.

```
[]: ax = subplots(figsize=(8,8))[1]
   ax.scatter(results3.fittedvalues , results3.resid)
   ax.set_xlabel('Fitted value')
   ax.set_ylabel('Residual')
   ax.axhline(0, c='k', ls='--')
```

[]: <matplotlib.lines.Line2D at 0x169535c10>



Qualitative Predictors Instead of the previous data set we will work with the Carseats data which contains qualitative predictors in addition to quantitative predictors. We will work to predict sales in 400 locations.

These data are realized as one-hot-encoded variables.

```
[]: Carseats = load_data('Carseats')
Carseats.columns
```

```
[]: Index(['Sales', 'CompPrice', 'Income', 'Advertising', 'Population', 'Price', 'ShelveLoc', 'Age', 'Education', 'Urban', 'US'], dtype='object')
```

To avoid collinearity we drop the first column.

```
[]: allvars = list(Carseats.columns.drop('Sales'))
    y = Carseats['Sales']
    final = allvars + [('Income', 'Advertising'), ('Price', 'Age')]
    X = MS(final).fit_transform(Carseats)
    model = sm.OLS(y, X)
    summarize(model.fit())
```

[]:		coef	std err	t	P> t
	intercept	6.5756	1.009	6.519	0.000
	CompPrice	0.0929	0.004	22.567	0.000
	Income	0.0109	0.003	4.183	0.000
	Advertising	0.0702	0.023	3.107	0.002
	Population	0.0002	0.000	0.433	0.665
	Price	-0.1008	0.007	-13.549	0.000
	ShelveLoc[Good]	4.8487	0.153	31.724	0.000
	<pre>ShelveLoc[Medium]</pre>	1.9533	0.126	15.531	0.000
	Age	-0.0579	0.016	-3.633	0.000
	Education	-0.0209	0.020	-1.063	0.288
	Urban[Yes]	0.1402	0.112	1.247	0.213
	US[Yes]	-0.1576	0.149	-1.058	0.291
	<pre>Income:Advertising</pre>	0.0008	0.000	2.698	0.007
	Price:Age	0.0001	0.000	0.801	0.424

We have added an interaction term between price and age. To encode the ShelveLoc[Good] dummy variable we insert a 1 to indicate a positive observation and a 0 otherwise.

We see that ShelvLoc[Good] has a positive value, indicating that good shelving location is associated with high sales. It has a higher coefficient than ShelvLoc[Medium] which means it leads to higher sales as well.