

Mathematics 231

Lecture 33

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Announcements

- Today
 - Model Building

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Model Building

- We have seen several examples of ways to select explanatory variables for a multiple regression model.
- We will examine all of the predictors in the birthweight dataset to determine which are important predictors.
- We had mother's age, toxemia, gestational age, infant length, and head circumference as predictors.

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Continuous Predictors

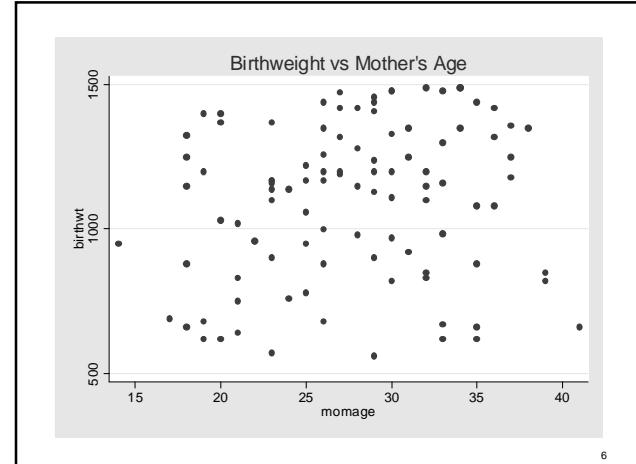
- We should plot the response against each continuous explanatory variable to assess linearity.
- It's often useful to look at all pairwise correlations among the continuous predictors.
- We should then perform all possible simple linear regressions by regressing the outcome on each predictor, separately.
- The variable with the smallest p-value (or largest t-statistic) explains more of the variability of the response and should be included.
- This forward selection process is repeated until no variable adds to the model.
- We can also consider categorical variables in the selection.

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Example

- We have 4 continuous predictors (momage, gestage, length, headcirc).
- We have one categorical (binary) predictor (toxemia).
- We have examined the plots of birthwt vs gestage, birthwt vs length, and birthwt vs headcirc.
- We now need to examine birthwt vs momage.

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Regress Birthwt on Gestage

```
. regress birthwt gestage

      Source |       SS           df          MS
Number of obs =      100
F( 1,    98) =   75.61
Model |  3143019.07     1  3143019.07
Prob > F    = 0.0000
Residual |  4073723.68    98  41568.609
R-squared    = 0.4355
Adj R-squared = 0.4298
Total |  7216742.75    99  72896.3914
Root MSE    = 203.88

      birthwt |      Coef.    Std. Err.      t    P>|t| [95% Conf. Interval]
gestage |  70.30993  8.085854     8.70  0.000    54.26382  86.35604
_cons | -932.4039 234.4884    -3.98  0.000   -1397.738  -467.0693
```

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Regress Birthwt on Headcirc

```
. regress birthwt headcirc

      Source |       SS           df          MS
Number of obs =      100
F( 1,    98) = 172.82
Prob > F    = 0.0000
Residual |  2611443.88    98  26647.3866
R-squared    = 0.6381
Adj R-squared = 0.6344
Total |  7216742.75    99  72896.3914
Root MSE    = 163.24

      birthwt |      Coef.    Std. Err.      t    P>|t| [95% Conf. Interval]
headcirc |  85.17802  6.479268     13.15  0.000    72.32013  98.03592
_cons | -1154.109 172.1523    -6.70  0.000   -1495.739  -812.478
```

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Regress Birthwt on Length

```
. regress birthwt length

Source |      SS       df      MS
Model | 4800034.87     1 4800034.87
Residual | 2416707.88    98 24660.2845
Total | 7216742.75    99 72896.3914

Number of obs = 100
F( 1, 98) = 194.65
Prob > F = 0.0000
R-squared = 0.6651
Adj R-squared = 0.6617
Root MSE = 157.04

birthwt |   Coef.  Std. Err.      t  P>|t|  [95% Conf. Interval]
length |  61.65408  4.419149  13.95  0.000   52.88442  70.42373
_cons | -1171.253 163.4691  -7.16  0.000  -1495.652  -846.854
```

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Regress Birthwt on Momage

```
. regress birthwt momage

Source |      SS       df      MS
Model | 172425.647     1 172425.647
Residual | 7044317.1    98 71880.7868
Total | 7216742.75    99 72896.3914

Number of obs = 100
F( 1, 98) = 2.40
Prob > F = 0.1247
R-squared = 0.0239
Adj R-squared = 0.0139
Root MSE = 268.11

birthwt |   Coef.  Std. Err.      t  P>|t|  [95% Conf. Interval]
momage |  6.975444  4.503782   1.55  0.125  -1.962164  15.91205
_cons |  905.4209 127.7352   7.09  0.000   651.9346  1158.907
```

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Example

- Infant length resulted in the largest t-statistic (13.95) just beating out head circumference.
- We include length in the model and add one additional continuous variable at a time to the model.
- We will also take notice to see if the predictive ability of length decreases with the addition of each variable (collinearity).

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Add Headcirc

```
. regress birthwt length headcirc

Source |      SS       df      MS
Model | 5494338.63     2 2747169.31
Residual | 1722404.12    97 17756.7435
Total | 7216742.75    99 72896.3914

Number of obs = 100
F( 2, 97) = 154.71
Prob > F = 0.0000
R-squared = 0.7613
Adj R-squared = 0.7564
Root MSE = 133.25

birthwt |   Coef.  Std. Err.      t  P>|t|  [95% Conf. Interval]
length |  37.82793  5.346058   7.08  0.000   27.21749  48.43838
headcirc |  47.15045  7.540371   6.25  0.000   32.1849  62.116
_cons | -1541.104 150.7971  -10.22  0.000  -1840.394  -1241.813
```

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Add Gestage

```
. regress birthwt length gestage

      Source |       SS          df          MS
-----+---- Model | 4958359.24        2  2479179.62
Residual | 2258383.51       97  23282.3043
-----+---- Total | 7216742.75       99  72896.3914

      Number of obs =     100
F(  2,    97) = 106.48
Prob > F    = 0.0000
R-squared   = 0.6871
Adj R-squared = 0.6806
Root MSE    = 152.59

      birthwt |      Coef.    Std. Err.      t    P>|t|   [95% Conf. Interval]
-----+---- length |  51.40366  5.821408   8.83  0.000   39.84978  62.95754
gestage |  21.39402  8.204112   2.61  0.011   5.111133  37.67692
_cons | -1411.906 183.6993  -7.69  0.000  -1776.499 -1047.314
```

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Add Momage

```
. regress birthwt length momage

      Source |       SS          df          MS
-----+---- Model | 4804117.61        2  2402058.8
Residual | 2412625.14       97  24872.4242
-----+---- Total | 7216742.75       99  72896.3914

      Number of obs =     100
F(  2,    97) = 96.58
Prob > F    = 0.0000
R-squared   = 0.6657
Adj R-squared = 0.6588
Root MSE    = 157.71

      birthwt |      Coef.    Std. Err.      t    P>|t|   [95% Conf. Interval]
-----+---- length |  62.05671  4.547482  13.65  0.000   53.02022  71.0912
momage | -1.099813  2.714578  -0.41  0.686  -6.487499  4.287872
_cons | -1155.543 168.6876  -6.85  0.000  -1490.342 -820.7452
```

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Example

- Adding head circumference has slightly decreased the usefulness of length (the t-statistic decreased for length).
- Adding gestational age didn't decrease the usefulness of length as much, but it isn't as a good a predictor as head circumference.
- Notice the model with headcirc has a larger R² value than the model with gestage (0.76 to 0.69) and also a larger adjusted R².
- We will keep length and headcirc, and examine the remaining variables.

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Add Gestage

```
. regress birthwt length headcirc gestage

      Source |       SS          df          MS
-----+---- Model | 5505033.78        3  1835011.26
Residual | 1711708.97       96  17830.3017
-----+---- Total | 7216742.75       99  72896.3914

      Number of obs =     100
F(  3,    96) = 102.92
Prob > F    = 0.0000
R-squared   = 0.7628
Adj R-squared = 0.7554
Root MSE    = 133.53

      birthwt |      Coef.    Std. Err.      t    P>|t|   [95% Conf. Interval]
-----+---- length |  38.9962  5.565441    7.01  0.000   27.94889  50.04352
headcirc |  51.30345  9.265343    5.54  0.000   32.91189  69.69501
gestage | -6.818412  8.803777   -0.77  0.441  -24.29377  10.65695
_cons | -1496.982 161.4911   -9.27  0.000  -1817.54 -1176.425
```

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Add Momage

```
. regress birthwt length headcirc momage
      Source |       SS          df          MS
-----+-----+-----+
      Model | 5495606.98        3   1831868.99
      Residual | 1721135.77       96   17928.4976
-----+
      Total | 7216742.75       99   72896.3914
      Number of obs =    100
      F(  3,     96) = 102.18
      Prob > F    = 0.0000
      R-squared    = 0.7615
      Adj R-squared = 0.7541
      Root MSE     = 133.9

      birthwt |       Coef.      Std. Err.      t      P>|t|      [95% Conf. Interval]
-----+-----+-----+-----+-----+-----+
      length | 38.08653  5.459126   6.98  0.000   27.25025  48.92281
      headcirc | 47.08196  7.581126   6.21  0.000   32.03354  62.13037
      momage | -.6133579  2.306035  -0.27  0.791  -5.190801  3.964085
      _cons | -1531.805 155.5052  -9.85  0.000  -1840.481  -1223.13
```

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Add Toxemia

```
. regress birthwt length headcirc toxemia
      Source |       SS          df          MS
-----+-----+-----+
      Model | 5569504.96        3   1856501.65
      Residual | 1647237.79       96   17158.727
-----+
      Total | 7216742.75       99   72896.3914
      Number of obs =    100
      F(  3,     96) = 108.20
      Prob > F    = 0.0000
      R-squared    = 0.7717
      Adj R-squared = 0.7646
      Root MSE     = 130.99

      birthwt |       Coef.      Std. Err.      t      P>|t|      [95% Conf. Interval]
-----+-----+-----+-----+-----+-----+
      length | 38.0629  5.256472   7.24  0.000   27.63989  48.49792
      headcirc | 48.36315 7.434922   6.50  0.000   33.60495  63.12135
      toxemia | -67.92159 32.45179  -2.09  0.039  -132.3379  -3.50529
      _cons | -1567.605 148.7759  -10.54 0.000  -1862.923  -1272.287
```

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Example

- We see that adding toxemia has helped in our prediction of birthweight.
- We will include toxemia in the model, but now want to consider its interactions with head circumference and infant length.
- We have seen collinearity with gestational age and both head circumference and length and will not consider it further.
- Mother's age has given us no indication that it is a useful predictor at all.
- We consider toxemia as a modifying variable.

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Add Length*Toxemia

```
. regress birthwt length headcirc toxemia lentox
      Source |       SS          df          MS
-----+-----+-----+
      Model | 5641405.82        4   1410351.46
      Residual | 1575336.93       95   16582.494
-----+
      Total | 7216742.75       99   72896.3914
      Number of obs =    100
      F(  4,     95) =  85.05
      Prob > F    = 0.0000
      R-squared    = 0.7817
      Adj R-squared = 0.7725
      Root MSE     = 128.77

      birthwt |       Coef.      Std. Err.      t      P>|t|      [95% Conf. Interval]
-----+-----+-----+-----+-----+-----+
      length | 35.89579  5.271313   6.81  0.000   25.43091  46.36067
      headcirc | 46.05232  7.392782   6.23  0.000   31.3758  60.72885
      toxemia | -854.8818 379.2735  -2.25  0.026  -1607.835  -101.9288
      lentox | 21.05083 10.10944   2.08  0.040   .981055  41.12061
      _cons | -1427.483 160.9943  -8.87 0.000  -1747.097  -1107.869
```

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Add Headcirc*Toxemia

```
. regress birthwt length headcirc toxemia headtox
      Source |       SS          df        MS
      Model |  5593029.44         4   1398257.36
      Residual | 1623713.31        95   17091.7191
      Total |  7216742.75        99   72896.3914
      Number of obs =     100
      F(  4,    95) =   81.81
      Prob > F = 0.0000
      R-squared =  0.7750
      Adj R-squared =  0.7655
      Root MSE = 130.74
      birthwt |      Coef.    Std. Err.      t    P>|t| [95% Conf. Interval]
      length |  37.99741   5.246506    7.24  0.000   27.59179   48.41304
      headcirc |  45.28098   7.871737    5.75  0.000   29.65361   60.90836
      toxemia | -475.261   348.715   -1.36  0.176  -1167.548   217.0258
      headtox |  15.12886   12.89553    1.17  0.244  -10.47199   40.72971
      _cons | -1484.175   164.6359   -9.01  0.000  -1811.019  -1157.331
```

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Example

- It looks like headcirc, length, toxemia, and length*toxemia give us the best model.
- We should now at least look at the residual vs predicted value plot (looking at residual vs predictor plots when there is an interaction is difficult).

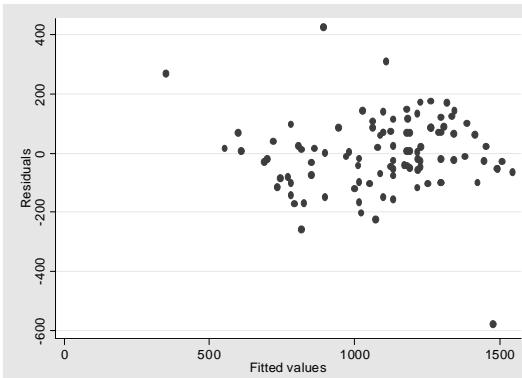
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Regression Diagnostics in Stata

- Click on **Graphics > Regression diagnostic plots**.
- For a residual vs fitted value plot choose **residual-vs-fitted** plot.
- For an added variable plot choose the **added variable** plot and the “all variables” option. Otherwise you can examine predictors individually.
- For a leverage plot choose the **leverage versus fitted value** plot.
- For any of these plots, you may choose an identifying variance to put in the “Marker Label” box under the Plot tab.

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Residuals versus Fitted Values



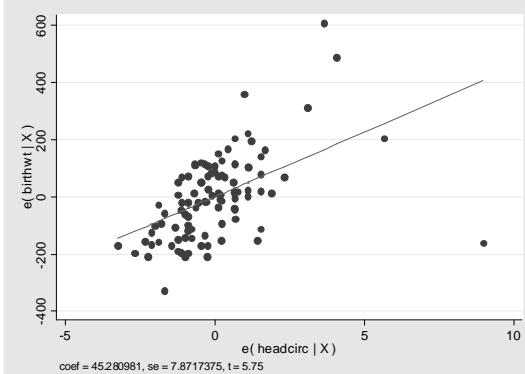
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Added Component Plot

- These plots examine the relationship between the predicted response with each explanatory variable, averaging over the explanatory variables not under immediate consideration.
- These plots should not depart markedly from linearity.
- These aren't generally appropriate to run for a model with an interaction term.
- We will consider this plot for the predictor not involved in the interaction – head circumference.

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Added Component Plot



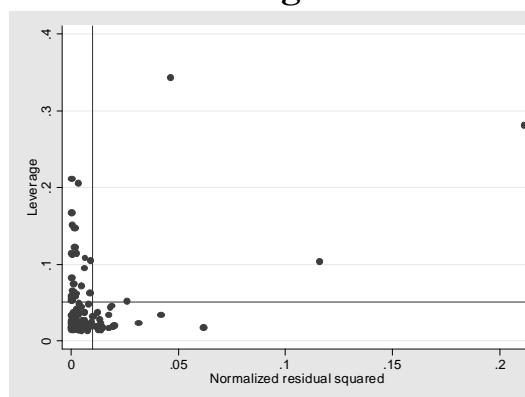
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More Diagnostics

- We still need to consider whether we have points that have had undue influence on our model.
- This can be checked by looking at leverage plots.
- We consider the leverage plot for the final model we've obtained here.

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Leverage Plot



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Leverage Plot

- The horizontal line represents the average “leverage”
- Points far above it have a possibly undue influence on the regression.
- The vertical line represents the average normalized squared residual.
- Points far to the right of this line represent points with large residuals.

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