Logistic Regression Example

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This examples demonstrates the binaryReg and other logistic regression support functions in the smwrStats package. The example uses the PugetNitrate dataset from Tesoriero and Voss (1997). The dataset is available from the smwrData package.

- > # Load the smwrStats and smwrData packages
- > library(smwrStats)
- > library(smwrData)
- > # Get the dataset
- > data(PugetNitrate)
- > head(PugetNitrate)

	wellid	110	120	140	surfgeo	date	nitrate	wellmet
1	1000	15.375154	0.000000	57.687577	Coarse	1990-09-06	0.2	60.9600
2	1001	7.839196	77.185930	9.849246	Coarse	1993-06-17	9.4	5.4864
3	1002	7.236181	35.276382	53.969849	Coarse	1991-05-14	0.4	21.9456
4	1003	34.472362	11.155779	53.668342	Coarse	1992-05-11	1.0	113.9952
5	1004	4.623116	13.869347	81.507538	Alluvium	1989-03-17	0.2	30.1752
6	1005	54.974874	0.201005	21.507538	Coarse	1988-09-19	2.8	16.7640

1 Single Variable Model

The hosmerLemeshow.test, leCessie.test, and roc functions performs diagnostic tests on a logistic regression model created by glm. The model can be constructed from either discrete values or counts of successes and failures.

This example follows the assumptions in Tesoriero and Voss (1997). The regression will model the probability that the concentration equals or exceeds 3 mg/L as was done in that report. This example demonstrates the hosmerLemeshow.test and roc functions on one single variable model described by Tesoriero and Voss (1997). The leCessie.test is useful for glm models with fewer than 1000 observations because of the time required to process larger sample sizes.

```
> # Create the logistic regression model
> PSN03.1 <- glm(formula = nitrate >= 3 ~ wellmet, family = binomial,
      data = PugetNitrate, na.action = na.exclude)
> # Print the summary
> print(summary(PSNO3.1))
Call:
glm(formula = nitrate >= 3 ~ wellmet, family = binomial, data = PugetNitrate,
    na.action = na.exclude)
Deviance Residuals:
   Min
             1Q
                 Median
                                3Q
                                        Max
-0.7066 -0.4635 -0.3338 -0.1904
                                    3.0984
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -1.224334
                       0.161778 -7.568 3.79e-14 ***
            -0.029482
                        0.003857 -7.644 2.10e-14 ***
Signif. codes: 0 Ś***Š 0.001 Ś**Š 0.01 Ś*Š 0.05 Ś.Š 0.1 Ś Š 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 1014.85 on 1966 degrees of freedom
Residual deviance: 925.19 on 1965 degrees of freedom
AIC: 929.19
Number of Fisher Scoring iterations: 7
```

The statistics from the printed summary agree reasonably well with table 2 in Tesoriero and Voss (1997). Small differences can be expected among different logistic regression implementations due to differences in convergence criteria for example. The G statistics in table 2 is the difference between the null deviance and the model deviance, 1014.85 - 925.19 = 89.66.

The hosmerLemeshow.test can help diagnose lack of fit and the output can help construct diagnostic plots like figure 2 in Tesoriero and Voss (1997). The code below runs the test and creates a graph to replicate figure 2, very small differences can be noted due to small differences in grouping.

```
> # Run the H-L test
> PSNO3.1.hl <- hosmerLemeshow.test(PSNO3.1)
> print(PSN03.1.hl)
        Hosmer-Lemeshow goodness of fit test
data: nitrate >= 3 ~ wellmet
Chi-square = 22.437, Number of groups = 10, p-value = 0.004167
alternative hypothesis: Some lack of fit
null hypothesis: No lack of fit
sample estimates:
   Size Expected Counts
                        wellmet
   196
          0.751
                    1 172.67231
1
2
   199
          2.965
                     3 101.52597
                   12 82.38760
3
   193
          4.917
                     8 67.11370
   206
          8.104
5
   191
        10.476
                     5 55.11933
6
   188
        12.848
                    14 47.14186
7
   203
                    10 38.15706
         17.736
8
   199
         21.979
                    16 29.28531
   196
9
         26.677
                    28 21.22714
         34.547
                    44 10.89038
  196
> # Added fitted values to dataset for line in figure 2, and order
> PugetNitrate$fits <- fitted(PSN03.1)</pre>
> OrderFits <- order(PugetNitrate$fits)</pre>
> # setSweave is a specialized function that sets up the graphics page for
> # Sweave scripts. For interactive use, it should be removed and the
> # default setting for set.up can be used.
> setSweave("binplot01", 5, 5)
> with(PugetNitrate, xyPlot(wellmet[OrderFits], fits[OrderFits],
      Plot=list(what="lines"),
      xaxis.range=c(0, 200),
      yaxis.range=c(0, .25),
     xtitle="Well Depth, in meters",
      ytitle="Estimated Pobability"))
> # Add the observed frequencies
> with(PSNO3.1.hl$estimate, addXY(wellmet, Counts/Size,
      Plot=list(what="points")))
```

> # Required call to close PDF output graphics
> graphics.off()

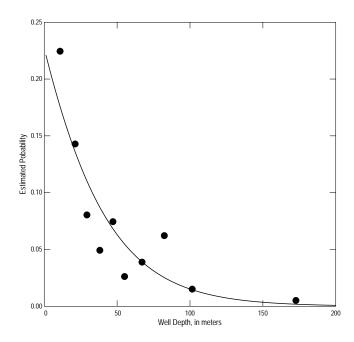


Figure 1. The estimated pobability that nitrate exceeds 3 mg/L as a function of well depth.

The Hosmer-Lemeshow test can be very sensitive to the number of groups. Compare the p-values from the previous test using the default 10 groups with the output below for 12 groups.

```
> # Run the H-L test with 12 groups
> hosmerLemeshow.test(PSNO3.1, 12)
```

Hosmer-Lemeshow goodness of fit test

```
data: nitrate >= 3 ~ wellmet
Chi-square = 15.603, Number of groups = 12, p-value = 0.1116
alternative hypothesis: Some lack of fit
null hypothesis: No lack of fit
sample estimates:
   Size Expected Counts wellmet
1 162 0.466 0 183.632593
```

```
2
    162
            1.942
                        3 109.071363
3
    171
            3.567
                        7
                           89.258274
    160
            4.906
                           75.763755
                        7
5
    166
            7.160
                           63.725234
6
    164
            9.171
                        5
                           54.388215
7
           10.901
                           47.688030
    162
                       10
8
    172
           14.207
                           40.208791
                       12
9
    157
           15.984
                        9
                           32.365101
10
    160
           19.137
                       21
                           26.216610
11
    173
           24.963
                       22
                           18.911695
    158
           28.596
                       38
                            9.761316
12
```

Another quick evaluation of a logisite regression is the area under the receiver-operating-curve (AUROC). It is a measure of the predictive power of the model. The result is a number from varies from 0.5, no predictive power, to 1.0, perfect prediction. Tape, from

http://gim.unmc.edu/dxtests/Default.htm, accessed on 01/27/2009, gives general guidelines for the AUROC: .50-.60, fail; .60-70, poor; .70-80, fair, .80-.90 good, and .90-1.00 excellent. The roc function computes the statistic for any model. The output from the single variable model is shown below. The result indicates fair prediction.

```
> # Compute the area under the ROC
> roc(PSNO3.1)
```

Area under the ROC curve: 0.732

2 Multiple Variable Model

The binaryReg function performs a series of diagnostic tests on a logistic regression model created by glm. The model can be constructed from either discrete values or counts of successes and failures.

This example follows the assumptions in Tesoriero and Voss (1997) for the groundwater vulnerability model for coarse-grained glacial materials. The regression will model the probability that the concentration equals or exceeds 3 mg/L as was done in that report. This example demonstrates the binaryReg function.

```
> # Create the logistic regression model
> PSN03.3 <- glm(formula = nitrate >= 3 ~ wellmet + 120 + 110,
      family = binomial, subset = surfgeo == "Coarse",
      data = PugetNitrate, na.action = na.omit)
> # Create the assessment and print it
> PSN03.3.br <- binaryReg(PSN03.3)
> print(PSN03.3.br)
glm(formula = nitrate >= 3 ~ wellmet + 120 + 110, family = binomial,
    data = PugetNitrate, subset = surfgeo == "Coarse", na.action = na.omit)
Deviance Residuals:
    Min
              1Q
                  Median
                                3Q
                                        Max
-1.5005 -0.4720 -0.3274 -0.1869
                                    3.0998
Coefficients:
            Estimate Std. Error z value Pr(>|z|)
(Intercept) -2.067279
                       0.340674 -6.068 1.29e-09 ***
wellmet
           -0.028260
                       0.005854 -4.827 1.38e-06 ***
120
            0.033697
                       0.006033 5.586 2.33e-08 ***
110
            0.029039
                       0.006281 4.624 3.77e-06 ***
Signif. codes: 0 Ś***Š 0.001 Ś**Š 0.01 Ś*Š 0.05 Ś.Š 0.1 Ś Š 1
(Dispersion parameter for binomial family taken to be 1)
    Null deviance: 518.48 on 718 degrees of freedom
Residual deviance: 409.71 on 715 degrees of freedom
  (23 observations deleted due to missingness)
AIC: 417.71
Number of Fisher Scoring iterations: 6
```

Likelihood ratio test: 108.772 on 3 degrees of freedom, p-value is 0

Response profile:

nitrate >= 3 Response Counts 1 FALSE 0 635 2 TRUE 1 84

Goodness of fit tests

le Cessie-van Houwelingen GOF test

data: nitrate >= 3 ~ wellmet + 120 + 110 Chisq = 22.876, df = 13.509, p-value = 0.0523 alternative hypothesis: Some lack of fit null hypothesis: No lack of fit sample estimates: Q E[Q] se[Q] $58.56150 \ 34.58332 \ 13.30655$

Distance between observations: maximum bandwidth

6.237748 1.471405

Hosmer-Lemeshow goodness of fit test

data: nitrate >= 3 ~ wellmet + 120 + 110
Chi-square = 1.7, Number of groups = 10, p-value = 1
alternative hypothesis: Some lack of fit
null hypothesis: No lack of fit
sample estimates:

Size Expected Counts 0.460 1 72 1.408 2 72 2 3 72 2.329 4 72 3.326 3 5 72 4.335 6 71 5.612 7 72 7.332 7 8 72 9.566 8 9 72 14.846 15

72

10

Predictive power estimates: McFadden R-squared: 0.2098

34.786

37

adjusted R-squared: 0.1982

Classification table.

Percent correct: (1 is sensitivity, 0 is specificity)

1 0 25.0 97.8

Concordance Index, based on 53340 pairs

Discordant Tied Concordant 18.830146 0.001875 81.167979

Area under the ROC curve: 0.812

Influence diagnostic test criteria:

leverage cooksD dfits 0.02086 0.89220 0.34745

Observations exceeding at least one test criterion X...nitrate.X3 yhat resids deviance.res pearson.res

	Xnitrate.X3	yhat	resids	deviance.res	<pre>pearson.res</pre>	leverage	cooksD
2	TRUE	0.6471	0.3529	0.9330	0.7385	0.026464*	4.819e-02
16	TRUE	0.3369	0.6631	1.4752	1.4030	0.009688	3.310e-02
70	FALSE	0.6556	-0.6556	-1.4600	-1.3796	0.025465*	7.237e-02
209	FALSE	0.6308	-0.6308	-1.4117	-1.3071	0.026157*	2.313e-02
324	TRUE	0.5081	0.4919	1.1637	0.9839	0.041930*	6.783e-03
345	TRUE	0.4948	0.5052	1.1862	1.0104	0.016866	3.249e-02
465	FALSE	0.4309	-0.4309	-1.0618	-0.8701	0.024294*	1.002e-02
475	FALSE	0.6252	-0.6252	-1.4010	-1.2916	0.038238*	1.303e-02
503	TRUE	0.6516	0.3484	0.9256	0.7312	0.025533*	6.426e-02
564	TRUE	0.5712	0.4288	1.0584	0.8665	0.021289*	1.799e-03
578	FALSE	0.6716	-0.6716	-1.4923	-1.4300	0.027909*	1.109e-01
584	FALSE	0.5343	-0.5343	-1.2362	-1.0711	0.022086*	2.076e-02
599	FALSE	0.5801	-0.5801	-1.3174	-1.1754	0.022359*	1.255e-04
643	FALSE	0.3427	-0.3427	-0.9161	-0.7220	0.021726*	1.585e-02
687	TRUE	0.6792	0.3208	0.8795	0.6872	0.030449*	1.484e-01
710	FALSE	0.3150	-0.3150	-0.8699	-0.6781	0.009529	3.004e-02
732	FALSE	0.6756	-0.6756	-1.5005	-1.4431	0.024312*	1.232e-01
733	TRUE	0.6718	0.3282	0.8920	0.6990	0.024399*	1.123e-01
734	FALSE	0.6545	-0.6545	-1.4579	-1.3763	0.024823*	6.962e-02
1106	FALSE	0.6027	-0.6027	-1.3587	-1.2317	0.021197*	4.581e-03
1149	TRUE	0.6069	0.3931	0.9994	0.8048	0.023333*	4.907e-03
1298	TRUE	0.5932	0.4068	1.0220	0.8282	0.025341*	2.246e-04
1302	FALSE	0.6519	-0.6519	-1.4527	-1.3683	0.024970*	6.156e-02
1407	FALSE	0.3451	-0.3451	-0.9202	-0.7260	0.011029	4.081e-02
1429	TRUE	0.6115	0.3885	0.9917	0.7970	0.029121*	9.471e-03
1499	FALSE	0.4160	-0.4160	-1.0372	-0.8440	0.032769*	7.507e-03

```
TRUE 0.4799 0.5201
1517
                                         1.2118
                                                      1.0411 0.038195* 1.975e-05
              FALSE 0.4863 -0.4863
                                        -1.1542
                                                     -0.9730 0.038610* 3.318e-04
1518
1524
              FALSE 0.5722 -0.5722
                                        -1.3032
                                                     -1.1566 0.024865* 3.405e-03
1535
               TRUE 0.6894 0.3106
                                         0.8625
                                                      0.6713 0.026537* 1.788e-01
1628
              FALSE 0.5952 -0.5952
                                         -1.3448
                                                     -1.2125 0.025310* 2.732e-04
               TRUE 0.6558 0.3442
                                                      0.7245 0.031254* 7.783e-02
1629
                                         0.9187
              FALSE 0.3710 -0.3710
                                         -0.9630
                                                     -0.7680 0.032507* 7.957e-03
1748
              FALSE 0.1171 -0.1171
                                                     -0.3642 0.022628* 1.879e-02
1775
                                         -0.4991
               TRUE 0.4444 0.5556
                                         1.2736
                                                      1.1181 0.040658* 1.497e-03
1776
              FALSE 0.1137 -0.1137
                                         -0.4913
                                                     -0.3582 0.022933* 1.880e-02
1777
              FALSE 0.1486 -0.1486
1780
                                         -0.5672
                                                     -0.4178 0.025516* 1.562e-02
               TRUE 0.3834 0.6166
                                                      1.2683 0.037391* 6.383e-03
1781
                                         1.3847
              FALSE 0.2802 -0.2802
1782
                                        -0.8109
                                                     -0.6240 0.030746* 3.558e-04
              FALSE 0.4639 -0.4639
1850
                                         -1.1166
                                                     -0.9302 0.038909* 3.632e-04
1904
               TRUE 0.5667 0.4333
                                         1.0658
                                                      0.8745 0.022855* 4.133e-03
                                                      1.2532 0.010635 3.819e-02
1935
               TRUE 0.3890 0.6110
                                         1.3741
```

dfits

- 2 -0.440932*
- 16 0.367118*
- 70 -0.541882*
- 209 -0.304688
- 209 -0.304688
- 324 -0.164669
- 345 0.362151*
- 465 0.200280
- 475 -0.228315
- 503 -0.510145*
- 564 0.084783
- 578 -0.672834*
- 584 0.288699
- 599 0.022392643 0.252151
- 687 -0.780348*
- 710 0.349483*
- 732 -0.711441*
- 733 -0.678452*
- 734 -0.531418*
- 1106 -0.135351
- 1149 -0.140079
- 1298 -0.029952
- 1302 -0.499263*
- 1407 0.407956*
- 1429 -0.194669
- 1499 0.173268
- 1517 -0.008882 1518 -0.036405
- 1524 0.116667

```
1535 -0.861079*
1628 -0.033035
1629 -0.561383*
1748 0.178396
1775 -0.274550
1776 0.077341
1777 -0.274683
1780 -0.250167
1781
     0.159745
1782 0.037698
1850
     0.038091
     0.128556
1904
1935
     0.394507*
```

References

- [1] Tesoriero, A.J., and Voss, F.D., 1997, Predicting the probability of elevated nitrate concentrations in the Puget Sound Basin???Implications for aquifer susceptibility and vulnerability: Groundwater, v. 35, no. 6, p. 1029???1039.
- [2] Helsel, D.R. and Hirsch, R.M., 2002, Statistical methods in water resources: U.S. Geological Survey Techniques of Water-Resources Investigations, book 4, chap. A3, 522 p.