# Logistic Regression Example

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### Abstract

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.

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## 1 Introduction

These examples use data from the smwrData package. The data are retrieved in the following code.

- > # 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

### 2 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
> PSNO3.1 <- glm(formula = nitrate >= 3 ~ wellmet, family = binomial,
     data = PugetNitrate, na.action = na.exclude)
> # Print the summary
> print(summary(PSN03.1))
Call:
glm(formula = nitrate >= 3 ~ wellmet, family = binomial, data = PugetNitrate,
   na.action = na.exclude)
Deviance Residuals:
   Min
            10
                 Median
                             30
                                    Max
-0.7066 -0.4635 -0.3338 -0.1904
                                  3.0984
Coefficients:
           Estimate Std. Error z value Pr(>|z|)
wellmet
          -0.029482
                     0.003857 -7.644 2.10e-14 ***
Signif. codes: 0 Ś***Š 0.001 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
```

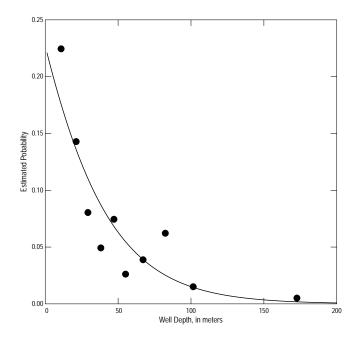
The statistics from the printed summary agree reasonably well with table 2 in Tesoriero and Voss (1997). Small differences can be expected among different

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
> PSN03.1.hl <- hosmerLemeshow.test(PSN03.1)
> 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
4.917 12 82.38760
          2.965
                     3 101.52597
3
   193
                     8 67.11370
4
   206
          8.104
5
   191
        10.476
                     5 55.11933
6
   188 12.848
                    14 47.14186
        17.7361038.1570621.9791629.2853126.6772821.22714
7
   203
8
   199
9
    196
10
   196
         34.547
                    44 10.89038
> # 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(PSN03.1.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)
```

 ${\tt Hosmer-Lemeshow} \ {\tt goodness} \ {\tt of} \ {\tt fit} \ {\tt test}$ 

```
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
4	160	4.906	7	75.763755
5	166	7.160	7	63.725234
6	164	9.171	5	54.388215
7	162	10.901	10	47.688030
8	172	14.207	12	40.208791
9	157	15.984	9	32.365101
10	160	19.137	21	26.216610
11	173	24.963	22	18.911695
12	158	28.596	38	9.761316

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

### 3 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
> PSNO3.3 <- glm(formula = nitrate >= 3 ~ wellmet + 120 + 110,
     family = binomial, subset = surfgeo == "Coarse",
     data = PugetNitrate, na.action = na.omit)
> # Create the assessment and print it
> PSNO3.3.br <- binaryReg(PSNO3.3)</pre>
> print(PSN03.3.br)
Call:
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
           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 0.05 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
1	72	0.460	1
2	72	1.408	2
3	72	2.329	2
4	72	3.326	3
5	72	4.335	4
6	71	5.612	5
7	72	7.332	7
8	72	9.566	8
9	72	14.846	15
10	72	34.786	37

#### Predictive power estimates:

McFadden R-squared: 0.2098 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

			•	least one tes				
	Xnitrate.X3	yhat	resids	deviance.res	${\tt pearson.res}$	leverage	cooksD	dfits
2	TRUE	0.6471	0.3529	0.9330	0.7385	0.026464*	4.819e-02	-0.440932*
16	TRUE	0.3369	0.6631	1.4752	1.4030	0.009688	3.310e-02	0.367118*
70	FALSE	0.6556	-0.6556	-1.4600	-1.3796	0.025465*	7.237e-02	-0.541882*
209	FALSE	0.6308	-0.6308	-1.4117	-1.3071	0.026157*	2.313e-02	-0.304688
324	TRUE	0.5081	0.4919	1.1637	0.9839	0.041930*	6.783e-03	-0.164669
345	TRUE	0.4948	0.5052	1.1862	1.0104	0.016866	3.249e-02	0.362151*
465	FALSE	0.4309	-0.4309	-1.0618	-0.8701	0.024294*	1.002e-02	0.200280
475	FALSE	0.6252	-0.6252	-1.4010	-1.2916	0.038238*	1.303e-02	-0.228315
503	TRUE	0.6516	0.3484	0.9256	0.7312	0.025533*	6.426e-02	-0.510145*
564	TRUE	0.5712	0.4288	1.0584	0.8665	0.021289*	1.799e-03	0.084783
578	FALSE	0.6716	-0.6716	-1.4923	-1.4300	0.027909*	1.109e-01	-0.672834*
584	FALSE	0.5343	-0.5343	-1.2362	-1.0711	0.022086*	2.076e-02	0.288699
599	FALSE	0.5801	-0.5801	-1.3174	-1.1754	0.022359*	1.255e-04	0.022392
643	FALSE	0.3427	-0.3427	-0.9161	-0.7220	0.021726*	1.585e-02	0.252151
687	TRUE	0.6792	0.3208	0.8795	0.6872	0.030449*	1.484e-01	-0.780348*
710	FALSE	0.3150	-0.3150	-0.8699	-0.6781	0.009529	3.004e-02	0.349483*
732	FALSE	0.6756	-0.6756	-1.5005	-1.4431	0.024312*	1.232e-01	-0.711441*
733	TRUE	0.6718	0.3282	0.8920	0.6990	0.024399*	1.123e-01	-0.678452*
734	FALSE	0.6545	-0.6545	-1.4579	-1.3763	0.024823*	6.962e-02	-0.531418*
1106	FALSE	0.6027	-0.6027	-1.3587	-1.2317	0.021197*	4.581e-03	-0.135351
1149	TRUE	0.6069	0.3931	0.9994	0.8048	0.023333*	4.907e-03	-0.140079
1298	TRUE	0.5932	0.4068	1.0220	0.8282	0.025341*	2.246e-04	-0.029952
1302	FALSE	0.6519	-0.6519	-1.4527	-1.3683	0.024970*	6.156e-02	-0.499263*
1407	FALSE	0.3451	-0.3451	-0.9202	-0.7260	0.011029	4.081e-02	0.407956*
1429	TRUE	0.6115	0.3885	0.9917	0.7970	0.029121*	9.471e-03	-0.194669

1499	FALSE 0.4160 -0.4160	-1.0372	-0.8440 0.032769* 7.507e-03	0.173268
1517	TRUE 0.4799 0.5201	1.2118	1.0411 0.038195* 1.975e-05	-0.008882
1518	FALSE 0.4863 -0.4863	-1.1542	-0.9730 0.038610* 3.318e-04	-0.036405
1524	FALSE 0.5722 -0.5722	-1.3032	-1.1566 0.024865* 3.405e-03	0.116667
1535	TRUE 0.6894 0.3106	0.8625	0.6713 0.026537* 1.788e-01	-0.861079*
1628	FALSE 0.5952 -0.5952	-1.3448	-1.2125 0.025310* 2.732e-04	-0.033035
1629	TRUE 0.6558 0.3442	0.9187	0.7245 0.031254* 7.783e-02	-0.561383*
1748	FALSE 0.3710 -0.3710	-0.9630	-0.7680 0.032507* 7.957e-03	0.178396
1775	FALSE 0.1171 -0.1171	-0.4991	-0.3642 0.022628* 1.879e-02	-0.274550
1776	TRUE 0.4444 0.5556	1.2736	1.1181 0.040658* 1.497e-03	0.077341
1777	FALSE 0.1137 -0.1137	-0.4913	-0.3582 0.022933* 1.880e-02	-0.274683
1780	FALSE 0.1486 -0.1486	-0.5672	-0.4178 0.025516* 1.562e-02	-0.250167
1781	TRUE 0.3834 0.6166	1.3847	1.2683 0.037391* 6.383e-03	0.159745
1782	FALSE 0.2802 -0.2802	-0.8109	-0.6240 0.030746* 3.558e-04	0.037698
1850	FALSE 0.4639 -0.4639	-1.1166	-0.9302 0.038909* 3.632e-04	0.038091
1904	TRUE 0.5667 0.4333	1.0658	0.8745 0.022855* 4.133e-03	0.128556
1935	TRUE 0.3890 0.6110	1.3741	1.2532 0.010635 3.819e-02	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.