A Methodology for Selecting and Transforming Continuous Predictors for Logistic Regression

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ABSTRACT

This paper discusses the selection and transformation of continuous predictor variables for the fitting of binary logistic models. The paper has two parts: (1) A procedure and associated SAS® macro is presented which can screen hundreds of predictor variables and 10 transformations of these variables to determine their predictive power for a logistic regression. The SAS macro passes the training data set twice to prepare the transformations and one more time through PROC TTEST. (2) The FSP (function selection procedure) and a SAS implementation of FSP are discussed. The FSP tests all transformations from among a class of "FSP transformations" and finds the one with maximum likelihood when fitting the binary target. The FSP has been presented in full detail in a 2008 book by P. Royston and W. Sauerbrei.

INTRODUCTION¹

The setting for this discussion is direct marketing or credit scoring or other applications of binary logistic regression where sample sizes for model building are large (perhaps exceeding 1000 observations for each target value) and the emphasis is on building predictive models to be used for scoring future observations. In this setting:

The preparation of predictors for binary logistic regression includes the following phases:

- Screening predictors to detect predictive power
- Transforming the predictors to maximize the predictive power
- Other phases (not discussed) such as finding interaction of predictors and elimination of collinear predictors

Predictors fall into three broad categories:²

- (1) Nominal, ordinal, interval-scaled with only a few values
- (2) Counts
- (3) Continuous (e.g. distances, dollars, percents, time, quantities)

An effective and widely used transformation for category (1) is weight-of-evidence (WOE) coding. WOE is also often applied to count predictors. Optimal binning before WOE transformation is a key requirement.³

The meaning of "continuous" for a predictor is subjective. One definition might be "when it is hard to do binning". This definition applies when many of the values of X occur only on one observation.

The use of binning of continuous predictors followed by WOE coding (often performed in direct marketing or credit scoring) may over-fit and complicate these predictive models. Specifically, when a functional form can be accurately identified for a continuous predictor, the application of binning and WOE coding will lose predictive power.

FIGURE 1 provides an illustration. In this loose hypothetical case, the relationship between predictor X and log-odds of Y is linear. But the approximation to log-odds(Y) by X_cut3 creates three abrupt jumps in the prediction of log-odds(Y). These jumps are not related to underlying behavior of X and Y and create prediction error.

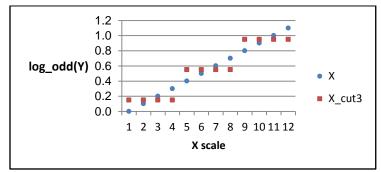


FIGURE 1: Hypothetical 3-cutpoint binning of a predictor X

¹ An earlier version of this paper appeared at MWSUG 2014, Lund (2014)

² To complicate this categorization some predictors are mixtures of types. Consider X: Ounces of alcohol consumed in past year.

Then X will have a spike at zero (nondrinkers).

³ A SAS macro for binning and WOE transformations is given by Lund and Brotherton (2013).

In this paper a statistical procedure and SAS macro %LOGIT_CONTINUOUS for this procedure are given for screening hundreds of continuous predictors for logistic regression. The goal is to identify predictor candidates that merit further study. This is done by measuring the predictive power of the original predictor and 10 transformations of the predictor. The entire procedure requires 3 passes of the original data set regardless of the number of predictors.

Once candidate predictors have passed screening by %LOGIT_CONTINUOUS, a final transformation for these predictors can be determined by the Function Selection Procedure (FSP) as described in Royston and Sauerbrei in *Multivariate Model-building* (2008). The FSP and a SAS implementation are discussed in Part II of this paper.

PART I: SCREENING CONTINUOUS PREDICTORS FOR PREDICTIVE POWER

Given dozens or hundreds of candidate continuous predictors, the "screening problem" is to test each predictor as well as a collection of transformations of the predictor for, at least, minimal predictive power in order to justify further investigation. If the number of candidate predictors is only a few, a brute force approach of fitting the predictor and transformations of the predictor by PROC LOGISTIC provides a simple, direct solution. However, each PROC LOGISTIC requires a pass of the training data set.

Instead, an alternative procedure is described which screens hundreds of predictors in a single run of PROC TTEST. In this paper a SAS macro to implement this procedure, called %LOGIT_CONTINUOUS, is discussed. %LOGIT_CONTINUOUS takes advantage of a connection between 2-group discriminant analysis, a t-test, and logistic regression. This connection is fully developed in Appendix A and only the key results are presented below.

THE CONNECTION BETWEEN 2-GROUP DISCRIMINANT ANALYSIS AND LOGISTIC REGRESSION

Let X be a predictor for a binary target Y. The values of Y identify two groups (j = 1, 2) of observations. It is assumed that X has a univariate normal distribution for each group with common standard deviation σ but differing means μ_1 and μ_2 . Then, just as in logistic regression, the 2-group discriminant analysis leads to the equation:

Log
$$(P(Y=1 | X=x) / P(Y=2 | X=x)) = \beta_0 + \beta_1 x$$

The key results are:

(1) When fitting the 2-group discriminant model to a sample, the coefficient β_1 is estimated by b_{1D} , where b_{1D} is found by substituting sample statistics from the two groups, \bar{x}_1 , \bar{x}_2 , and S_p^2 as shown in equation (A):

$$D_{1D} = (\bar{x}_1 - \bar{x}_2) / S_p^2 \dots (A)$$

Here, the pooled variance S_p^2 estimates σ^2 where S_j^2 are the sample variances for samples j = 1, 2 and

$$S_p^2 = {(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2}/(n_1 - 1 + n_2 - 1)$$

The "D" is added as a subscript to "b₁" to indicate that the method of fitting is by discriminant analysis. Additionally, "L" will be added as a subscript to "b₁" to give b_{1L} to indicate when the estimation of β_1 is by logistic regression maximum likelihood.

Both b_{1D} and b_{1L} are consistent estimators of β_1 . For large enough samples, b_{1D} will be close to b_{1L} .

(2) In order to test that $\beta_1 = 0$ vs. $\beta_1 \neq 0$ the discriminant analysis coefficient b_{1D} can be regarded as a t-statistic with $n_1 + n_2 - 2$ d.f. via this factorization: $b_{1D} = t$ (1/S_p) sqrt(1/n₁ + 1/n₂). The square of this t-statistic "t²" is a chi-square with 1 d.f. and it will be close to the Wald chi-square statistic for b_{1L} from logistic regression.⁴

COMPARISON OF DISCRIMINANT ANALYSIS AND LOGISTIC REGRESSION WITHOUT NORMALITY

Without the normality assumptions on X, there is not a theory that guarantees that b_{1D} and b_{1L} will be close in value. But for the purpose of screening predictor variables for logistic regression the closeness of b_{1D} and b_{1L} is not a requirement. Instead, the requirement is that the square of the t-statistic from discriminant analysis and the Wald chisquare from logistic regression concur in their measurement of the significance of X. This concurrence can occur without close agreement of b_{1D} and b_{1L} .

Pictorially, the comparison of the two significance measurements must produce very few examples that fall into cells "B" or "C". Especially troubling would be examples in cell "C", a false negative. See TABLE 1.

	Wald x2 significant	Wald x2 not significant	
t ² significant	A	B (false positive)	
t ² not significant	C (false negative)	D	

TABLE 1

⁴ Calculation of "t²" by Satterthwaite (instead of pooled) variance formula gives very similar values for examples to follow in this paper. The pooled variance is used because equal population variances are assumed when deriving the formula for t².

SIMULATIONS

Three predictors X1, X2, and X3 were generated by simulations. They will provide a test of the concurrence of t^2 and Wald chi-square in measuring the significance of a predictor. The test is given in Examples 1, 2, and 3 below:

EXAMPLE 1: Predictor X1

```
data Example1;
do i = 1 to 4000;
Y = (ranuni(12345) < .45);
if Y = 1 then X1 = rannor(12345) + 0.1; /* normal with mean=0.1, std dev = 1 */
else if Y = 0 then X1 = rannor(12345); /* normal with mean=0, std dev = 1 */
output;
end;
```

run;

The distributions of X1 for the two groups meet the assumptions of normality with equal standard deviations (=1). The coefficient estimates b_{1D} and b_{1L} of β_1 , as well as t^2 and Wald chi-square for X1 should be nearly identical.

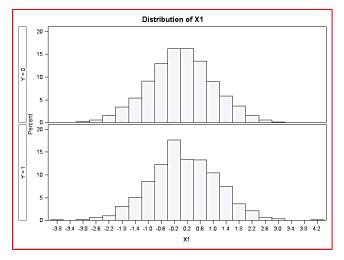


FIGURE 2: Histograms for Variable X1

TABLE 2A shows the very similar coefficient estimates.

TABLE 2A: Comparisor	of Coefficients from	Discriminant and Logistic
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Variable	b _{1D}	b _{1L}
X1	0.07420	0.07422

TABLES 2B and 2C show the almost perfect equality of the probabilities of the chi-squares

TABLE 2B: Chi-Square	Values for Coefficients from	Discriminant Analysis
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Variable	t Value	d.f.	ChiSq for b _{1D}	Prob ChiSq
X1	2.34	3998	5.467	0.0194

TABLE 2C: Chi-Square Values from Logistic

Variable	ChiSq for b _{1L}	Prob ChiSq
X1	5.455	0.0195

The SAS code that produced b_{1D} and the t-test values is shown below but SAS code for producing the logistic results is not included.

```
data example1;
do i = 1 to 4000;
    Y = (ranuni(12345) < .45);
    /* Normal distribution test */
    if Y = 1 then X1 = rannor(12345) + 0.1;
    else if Y = 0 then X1 = rannor(12345);
    output;
    end;
run;
ods listing;
ods output Statistics = TS;
ods output TTests = TT;
ods exclude ConfLimits; /* suppress unneeded print */
```

```
/* suppress unneeded print */
ods exclude Equality;
ods exclude EquivLimits; /* suppress unneeded print */
ods exclude EquivTests; /* suppress unneeded print */
proc ttest data=example1 plots=none;
  class Y; var X1;
data TT; set TT;
   tValue = -tValue; /* minus sign to model Y=1 as the response */
   chisq D = tValue**2;
  label Probt = "Prob ChiSq"; /* prob. for t-value = prob. for chi-sq. */
proc print data = TT label;
var Variable tValue DF Probt chisq_D;
where method = "Pooled"; /* Could consider alternative "Satterthwaite" */
run;
data TS 2; set TS;
retain mean1 mean2 n1 n2;
keep variable n1 n2 mean1 mean2 stddev b1D;
if class = "0" then do; mean1 = mean; n1 = n; end;
if class = "1" then do; mean2 = mean; n2 = n; end;
if class = "Diff (1-2)" /* this is the last row */
then
do;
  blD = - (mean1 - mean2) / (stddev **2); /* "-" to model Y=1 as response */
  output;
  end;
proc print data = TS_2 noobs; var Variable b1D;
run;
```

EXAMPLE 2: Predictor X2

```
data Example2;
do i = 1 to 500;
X2 = ranuni(12341);
Y = floor(X2 + ranuni(12341)); /* Y is more often equal to 1 when X2 is large */
output;
end;
run;
```

The distributions of X2 for the two groups strongly fail to meet the assumptions of normality as seen from the distributional histograms in FIGURE 3.

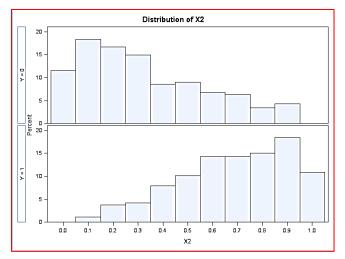


FIGURE 3: Histograms for Variable X2

The coefficient estimates b_{1D} and b_{1L} of $\beta 1$ are not close as seen in TABLE 3A.

TABLE 3A: Comparison of Coefficients from Discriminant and Logistic

Variable	b _{1D}	b _{1L}
X2	6.2743	5.3491

TABLES 3B and 3C show a wide divergence in value between the chi-squares, but both chi-squares are highly significant. As a screener of a predictor for a logistic model, both the t^2 from discriminant analysis and the Wald chi-square from logistic are highly significant.

TABLE 3B: Chi-Square Values for Coefficients from Discriminant Analysis

Variable	t Value	d.f.	ChiSq for b _{1D}	Prob ChiSq
X2	16.60	498	275.397	<.0001

TABLE 3C: Chi-Square Values from Logistic

Variable	ChiSq for b _{1L}	Prob ChiSq
X2	132.515	<.0001

EXAMPLE 3: Predictor X3

data Example3;

do i = 1 to 1000; X3 = ranuni(12345); /* Uniform on [0, 1] */ Y = (ranuni(12345) < .2); output; end; run;

Predictor X3 is uniform [0, 1] for both Y=0 or Y=1. The distributions of X3 for the two groups fail to meet the assumptions of normality as seen from the distributional histograms in FIGURE 4.

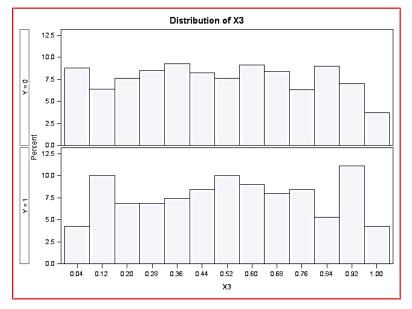


FIGURE 4: Histograms for Variable X3

The coefficient estimates b_{1D} and b_{1L} of $\beta 1$ are in close agreement as seen in Table 4A.

TABLE 4A: Comparison of Coefficients from Discriminant and Logistic

Variable	b _{1D}	b _{1L}
X3	0.35467	0.35550

Tables 4B and 4C show close agreement between the chi-squares. As a screener of X3 for a logistic model, both the t^2 from discriminant analysis and the Wald chi-square from logistic are strongly insignificant.

	•			•
Variable	t Value	d.f.	ChiSq for b _{1D}	Prob ChiSq
X3	1.25	998	1.554	0.2129

TABLE 4C: Chi-Square	Values	from	Logistic
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Variable	ChiSq for b _{1L}	Prob ChiSq
X3	1.552	0.2129

COUNTER-EXAMPLE

However, examples can be constructed where concurrence of the two methods in measuring significance is not attained. For predictor X4 in the data set below the t^2 is significant but the Wald chi-square is not significant at 5%.

```
data Example4;

Y = 0;

do i = 1 to 50;

X4 = rannor(12345);

output;

end;

do i = 1 to 2;

X4 = 50;

output;

end;

Y = 1;

do i = 1 to 500;

X4 = rannor(12345);

output;

end;
```

Predictor X4 is standard normal for Y=1 and X4 is also standard normal for Y=0 with the exception of two outliers at X4=50. Tables 5B and 5C show that chi-squares from discriminant analysis and from logistic disagree. The discriminant analysis chi-square is significant at < 0.01% while logistic chi-square fails a 5% significance test with probability value of 8.90%.

TABLE 5A: Comparison of Coefficients from Discriminant and Logistic

Variable	b _{1D}	b _{1L}
X4	-0.19914	-0.10782

TABLE 5B: Chi-Square Values for Coefficients from Discriminant Analysis

Variable	t Value	d.f.	ChiSq for b _{1D}	Prob ChiSq
X4	-4.25	550	18.0860	<.0001

TABLE 5C: Chi-Square	Values from	Logistic
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Variable	ChiSq for b _{1L}	Prob ChiSq
X4	2.892	0.0890

The relationship between X4 and Y would be highly unusual in an applied setting. It is likely that a modeler would research the two observations at X4=50 and correct them or eliminate them.

CONCLUSIONS

Examples 1-3 from the section on Simulations support the conclusion that the t-test approach can differentiate between significant and insignificant predictors for a logistic model.

Additional support is given by comments in *Applied Logistic Regression* by Hosmer, Lemeshow, and Sturdivant (2013 p. 91). These authors agree that if X has an approximately normal distribution for the two groups determined by values of Y, then the t-test is a good guide for screening a predictor for logistic regression.

The predictor X4 gives an example of a false positive (a "B" in TABLE 1). It was constructed by an extended trial and error process. I was unable to construct an example of a false negative (a "C" in TABLE 1) but, by no means, has this been ruled out. More simulation work is needed to understand the risk that such examples would present to the operation of the macro %LOGIT_CONTINUOUS.

%LOGIT_CONTINUOUS: MACRO FOR SCREENING HUNDREDS OF LOGISTIC PREDICTORS

The SAS macro %LOGIT_CONTINUOUS can screen hundreds of numeric predictors for logistic regression as well as 10 transformations of these predictors using the chi-square which is computed from a t-statistic. This t-statistic is mathematically derived from the coefficient of 2-group discriminant analysis as explained in Appendix A. The 10 transformations include 7 monotonic transformations and 3 quadratic transformations. Three passes of the data set are required for this computation:

- 1. PROC MEANS to determine the minimum value of each predictor
- 2. DATA STEP:
 - a. Predictors with minimum < 1 are shifted to have minimum value of 1.⁵
 - b. 10 transformations of the predictor are computed (such as LOG, X², and others).
- PROC TTEST to compute t-statistics whose square approximates the Wald chi-square from logistic regression.

⁵ In practice most continuous predictors have non-negative values domains (counts, distance, dollars, percents, time, quantity). Translation (except away from zero) is generally not needed.

The original X and the 10 transformations are:

- 8 monotonic: The "fractional polynomials" X^p where p is taken from S = {-2, -1, -0.5, 0, 1, 0.5, 2, 3} and where "0" denotes log(x). This list includes the original X.⁶
- 3 quadratic: (X-median)², (X-p25)², (X-p75)² where median, p25, and p75 are respectively the 50th, 25th, and 75th percentiles for X.

In practice, the relationship between a predictor X and the log-odds of Y⁷ is very often either monotonic or roughly quadratic (with a single maximum or minimum). Consequently, one (or more) of the 11 transformations is likely to have an approximate linear relationship to log-odds of Y, if, indeed, the predictor has any predictive power.

The parameters for %LOGIT_CONTINUOUS are:

DATASET: The input data set name.

Y: A numeric variable with two values. The larger value is the response that is modeled.

INPUT: A list of numeric predictor variables delineated by space. A predictor may have missing values.

EXAMPLE: %LOGIT_CONTINUOUS is run on X1 from Example1.

The macro call is %LOGIT_CONTINUOUS(example, Y, X1).

Variable	Transform	b _{1D}	ChiSq for b _{1D}	Prob ChiSq
X1_p7	x**3	0.0011	6.793	0.00919
X1_p11	(x-p25)**2	0.0404	6.424	0.01130
X1_p6	x**2	0.0082	6.268	0.01233
X1_p1	linear	0.0742	5.467	0.01943
X1_p5	x**0.5	0.3019	4.945	0.02622
X1_p8	log(x)	0.2954	4.325	0.03761
X1_p4	x**-0.5	-1.0970	3.593	0.05811
X1_p3	x**-1	-0.9437	2.739	0.09798
X1_p9	(x-p50)**2	0.0306	1.958	0.16179
X1_p2	x**-2	-0.8900	0.881	0.34809
X1_p10	(x-p75)**2	-0.0095	0.331	0.56510

TABLE 6: Results of %LOGIT_CONTINUOUS(example, Y, X1)

As shown in TABLE 6 the best transformations of X1 are X1³, (X1-p25)², and X1². These have similar chi-square values. The linear transform of X1 comes in at fourth place.

GUIDELINES FOR INTERPRETING THE RESULTS FROM %LOGIT_CONTINUOUS

In TABLE 6 the "Prob ChiSq" value should be viewed as a guide rather than a firm standard for significance. Due to by-chance over-fitting to the data by one or more of the 11 functions of X, the thresholds for the "Prob ChiSq" should be set conservatively. Here is one proposal for the threshold values for the best transformation:

- "Prob Chi-Square" less than 0.01 (1%)
- "ChiSq for b_{1D}" at least 10.0

Under these guidelines, X1 is a borderline case. Considering X1³, the best transform of X1, the first guideline is just barely met and the second guideline is not met.

- Guideline #1, "Prob Chi-Square" less than 0.01 (1%) vs. observed 0.919%
- Guideline #2, "ChiSq for b_{1D}" at least 10.0 vs. observed 6.793

TRANSLATIONS AND TRANSFORMATIONS OF X

Four transformations among the 11 are unaffected by translations: X, (X-median)², (X-p25)², (X-p75)². But could a "good" X be found that would otherwise be missed if translations were added to %LOGIT_CONTINUOUS to form translation-transformation combinations for the other 7?

SAS code below creates three predictors X5, X5_1, and X5_80. A translation of 1 unit was added to X5 to form X5_1 and a translation of 80 was added to form X5_80. Then %LOGIT_CONTINUOUS was run on X5, X5_1, and X5_80.

data Example5; do i = 1 to 500; X5 = ranuni(12341) + 1; Y = floor(X5 + ranuni(12341)); X5_1 = X5 + 1; X5_80 = X5 + 80;

⁶ These transformations are motivated by the FSP (function selection procedure) of Royston and Sauerbrei (2008).

⁷ Here, "Y" should be taken to mean the average of Y over a small interval of X.

output; end;

%LOGIT_CONTINUOUS(Example4, Y, X5 X5_1 X5_80);

The best transform of X5 is $X^{-0.5}$ while the best transform of X5_1 is X^{-1} . In this case the translation-transformation combination provided a very slightly better chi-square result than the transformation alone. Could a more extreme translation produce a translation-transformation that was even better? A translation of 80 units is used to create X5_80 but produced a lower chi-square of 275.88. See TABLE 7.

Variable	Best Transform	b _{1D}	ChiSq for b _{1D}	Prob ChiSq
X5_p4	x**-0.5	-22.270	281.24	<.0001
X5_1_p3	x**-1	-38.747	281.47	<.0001
X5_80_p2	x**-2	-1700484	275.88	<.0001

TABLE 7: Effect of a Translation Prior to running %LOGIT_CONTINUOUS

In general, the significance of the maximum value of the chi-squares for X and its 10 transformations is not greatly affected by translations. While a translation may result in a different transform being selected (as part of a translation-transformation), the effect on significance will not be decisive.

There is also a non-aesthetic aspect of a translation when the translation amount "c" places X+c outside the domain of observed or allowed values of X. Suppose a population consists of teenagers, the transformation is Log, X is age, and Y is buying a product or not. Then the coefficient b1 in the relationship log(Y/(1-Y)) = b0 + b1*log(X+10) is making a statement about the log-odds of buying for a population of non-teenagers.

SUMMARY

This paper shows that a process of screening a multitude of continuous predictors by %LOGIT_CONTINUOUS can efficiently and effectively identify predictors to retain for further study. But how should a final transformation be found for those predictors that pass the screening? This is the topic of Part II which follows below.

PART II: FUNCTION SELECTION PROCEDURE (FSP) FOR A CONTINUOUS PREDICTOR X

BACKGROUND

The Function Selection Procedure (FSP) is described by Royston and Sauerbrei (2008 p. 82) where a short history of its development is given. FSP uses transformations of X called fractional polynomials (FP). If X has negative values, then X must first be translated so that X is positive. Then the fractional polynomials of X can be defined by:

 X^{p} where p is taken from S = {-2, -1, -0.5, 0, 0.5, 1, 2, 3} where "0" denotes log(x)

FP1 refers to the collection of functions formed by the selection of one X^p. That is,

$$g(X,p) = \beta_0 + \beta_1 X^p$$

FP2 refers to the collection of functions formed by selection of two X^p. That is,

$$\begin{array}{ll} G(X,p_1,p_2) = \beta_0 + \beta_1 \; X^{p_1} + \beta_2 \; X^{p_2} & p_1 \neq p_2 \\ G(X,p_1,p_1) = \beta_0 + \beta_1 \; X^{p_1} + \beta_2 \; X^{p_1} \log(X) & p_1 = p_2 \end{array}$$

FP2 produces curves with a variety of non-monotonic shapes as shown by Royston and Sauerbrei (2008 p. 76). An example is given in FIGURE 5.

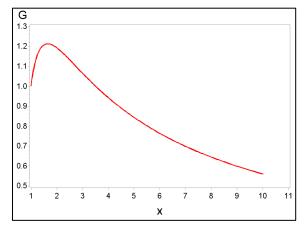


FIGURE 5: Graph of $G(X,-1,-1) = X^{-1} + 2X^{-1} \log(X)$

Royston and Sauerbrei (2008 p. 267) give links to software versions for performing FSP including Stata, R, and SAS. I downloaded the SAS version %mfp8 on 8/7/2014. It was written in SAS version 8. See notes in Appendix B which discuss a necessary change to the code in order to run %mfp8.⁶

FSP: SEARCHING FOR BEST TRANSFORMATIONS AND SIGNIFICANCE TESTING WITH %MFP8

Searching for best transformations: First, the FP1 and FP2 functions having the maximum likelihood are found by an exhaustive search in which 44 Logistic Models are run. Overall, PROC LOGISTIC is run 47 times.

Performing Significance Testing: Second, significance testing is performed. The following 3 steps for FSP significance testing are taken from documentation for the SAS macro implementation of FSP.

- Perform a 4 d.f. test at the α level of the best-fitting second-degree FP (i.e. FP2) against the null model. If the 1. test is not significant, drop X and stop, otherwise continue.
- Perform a 3 d.f. test at the α level of the best-fitting second-degree FP against a straight line. If the test is not 2. significant, stop (the final model is a straight line), otherwise continue.
- Perform a 2 d.f. test at the α level of the best-fitting second-degree FP against the best-fitting first-degree FP (i.e. 3 FP1). If the test is significant, the final model is the FP2, otherwise the FP1 is the final model.

The test-statistic for these three tests is the difference of deviances ¹⁰ as shown below:

Test-Statistic = (-2 Log Likelihood restricted model) - (-2 Log Likelihood full model)

For large samples, the Test-Statistic is approximately a chi-square. But the degrees of freedom for the three tests of the FSP need to be determined. The rationale for the degrees of freedom (4, 3, 2) used in the 3-step hypothesis tests of FSP is given by Royston and Sauerbrei (2008 p. 79).

EXAMPLE: RUNNING %MFP8 ON X2 FROM "EXAMPLE2" DATA SET

The FSP macro %mfp8 was run on predictor X2 from the Example2 data set. A translation was selected so that the minimum of the translated X2 would be 1 via this statement: X2 = X2 +1 - 0.001398486. See TABLE 8.

MFP8: Variable –X2-								
Best Functions for Different Degrees m								
Function m p1 p2 deviance diffra2 pdiffdev TEST:								
Null	-1			691.098	208.202	0.00000	FP2 v. Null: 4 d.f.	
Linear	0			489.001	6.106	0.10658	FP2 v. Linear: 3 d.f.	
First Degree	1	-2		483.540	0.645	0.72451	FP2 v. FP1: 2 d.f.	
Second Degree	2	-2	3	482.896	0.000	1.00000		

P Results from MFP8	R for Prodictor X2

For Step 1 the Test-Statistic is $\chi 2 = 691.098 - 482.896 = 208.202$

Predictor X2 passes Step 1 as seen from:

 $1 - \operatorname{Prob}(\chi^2(208.202,4)) < 0.0001$

For Step 2 the Test-Statistic is $\chi 2 = 489.001 - 482.896 = 6.106$

Predictor X2 fails Step 2 as seen from:

1- $Prob(\chi 2(6.106,3)) = 0.10658$ which is greater than 0.05 (5%)

On this basis FSP selects "Linear" as the transformation for X2.

Visualization of the Linear, FP1, and FP2 solutions:

The log-odds(Y) is plotted against the Linear, FP1, and FP2 solutions at the median value of X2 across 8 equal-sized ranks of X2 values. The log-odds(Y) "open-circles" give the mean of Y within the X2 rank.

Visually, both FP1 and FP2 give better fits than Linear. Since FP1 was approximately significant at 10% at Step 2, the FP1 solution has strong appeal.

See FIGURE 6.

⁸ FSP and %mfp8 also apply to ordinary least squares regression and Cox regression.

⁹ Meier-Hirmer, Ortseifen, and Sauerbrei (2003). Multivariable Fractional Polynomials in SAS, http://portal.uni-freiburg.de/imbi/mfp, beschreibung.pdf in SAS downloads. ¹⁰ The deviance is the -2 Log Likelihood value of a logistic model.

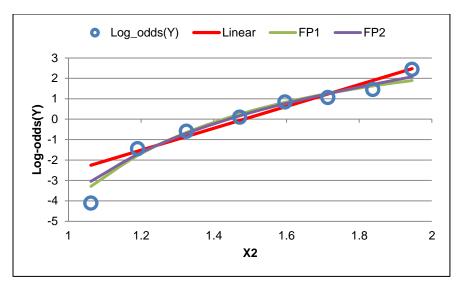


FIGURE 6: Linear = -7.9343 + 5.3491 X2; FP1 = 4.0901 - 8.3200 X2⁻²; FP2 = 2.9666 - 6.9522 X2⁻² + 0.1284 X2³

MORE SAS MACROS FOR FSP

The 47 PROC LOGISTIC's of %mfp8 can be reduced to 36. I wrote a macro that runs each of the 36 FP2 combinations through PROC LOGISTIC during which the log-likelihoods for the 8 FP1 functions are obtained.

Next, I tried to develop an approach that required only 8 PROC LOGISTIC runs. The idea is based on running PROC LOGISTIC with the options shown below on 8 sets of predictors called &Var1 to &Var8. See TABLE 9.

PROC LOGISTIC; MODEL Y = &Var<k> / SELECTION = FORWARD INCLUDE=1 START=1 STOP=2 SLE=1;

By this method all possible FP2 pairs have a chance to be selected. But selection of the second variable in a pair by FORWARD, to add to the first variable forced in by INCLUDE=1, may be sub-optimal. The reason is that the second variable is selected by the best Wald chi-square criterion, not by maximizing log likelihood of the two-variable model.

E.g. Consider &Var1.

- First, X⁻² is forced in.
- Now perhaps the best Wald chi-square criterion picks X⁻¹ to enter as the second variable.
- But the best log likelihood might be given by X³.

Such examples can be found.

Var1=	X ⁻²	X ⁻¹	X ⁵	X ^{.5}	Х	X ²	X ³	Log(X)	X ⁻² Log(X)
Var2=		X ⁻¹	X ⁵	X ^{.5}	Х	X ²	X ₃	Log(X)	X ⁻¹ Log(X)
Var3=			X ⁵	X ^{.5}	Х	X ²	X ³	Log(X)	X ⁵ Log(X)
Var4=				Х.5	Х	X ²	X3	Log(X)	X ^{.5} Log(X)
Var5=					Х	X ²	X ³	Log(X)	X Log(X)
Var6=						X ²	X ³	Log(X)	X ² Log(X)
Var7=							X ₃	Log(X)	X ³ Log(X)
Var8=								Log(X)	Log(X) Log(X)

TABLE 9: Eight sets of Predictors

The 8 PROC LOGISTIC approach may still have an advantage if there are hundreds of variables to be processed by FSP and the data set for these variables is relatively large, perhaps 50,000 observations. In this event, the 8 PROC LOGISTIC approach uses only about 17% (= 8 / 47) of the processing time versus the %mfp8 macro.

SUMMARY

This paper shows that a process of screening a multitude of continuous predictors by %LOGIT_CONTINUOUS can efficiently and effectively identify predictors to retain for further study. The FSP can then be focused on the surviving candidate predictors for either the selection of a final transformation or final elimination of the variable.

But there are open questions:

- What is the risk of false negatives (good predictors that are screened out) when running %LOGIT_CONTINUOUS?
- How often and to what degree would the 8 PROC LOGISTIC approach to FPS give sub-optimal FP2 solutions as measured by differences in log-likelihood?

SAS MACROS DISCUSSED IN THIS PAPER

The SAS macro %LOGIT_CONTINUOUS continues to be under development. Contact the author for a copy of the work-in-progress.

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APPENDIX A: TWO-GROUP DISCRIMINANT ANALYSIS AND LOGISTIC REGRESSION

In discriminant analysis it is assumed there are "p" populations G1, ..., Gp. The members of these p populations are characterized by predictor variables X_1 to X_K . The purpose of discriminant analysis is to develop for each population a linear combination of the predictors called the classification function (CF). Then an observation ($x_1 ..., x_K$) is assigned by the CF's to the population which has the largest CF value.

The development and application of the CF's follows two steps:

- (i) The "p" CF's are fitted using observations from G1, ..., Gp (i.e. observations where population membership is known).
- (ii) The CF's are applied to observations of X_1 to X_K where population membership is not known. An observation is assigned to the population with the largest CF value.

To support statistic inference, it is assumed the X₁... X_K follow a multivariate normal distribution.¹¹

THE CASE WHERE p = 2, K = 1, AND $\sigma_1 = \sigma_2$

The simplest case of discriminant analysis is when there are 2 populations, one predictor X, and the distributions of X for the two populations have a univariate normal distribution with common standard deviation σ but differing means μ_1 and μ_2 .

¹¹ See Huberty and Olejnik (2006, chapter 13) for discussion.

The formula for the distribution of X for population j = 1 or 2 is given by:

 $P(X=x | j) = (2\pi\sigma^2)^{-1/2} \exp(-0.5((x - \mu_j) / \sigma)^2)$... where μ_j is the mean for X for population j.

DEVELOPMENT OF THE CLASSIFICATION FUNCTIONS

Suppose a random sample for populations j = 1, 2 is taken and n_1 is the size from population 1 and n_2 is the size from population 2. The base-rate population probability of j is denoted by P(j). These probabilities are estimated by $q_j = n_j/(n_1 + n_2)$.

The probability that an observation with value X = x belongs to population j = 1, 2 is the conditional probability expressed by:

$$P(j | X=x)$$
 for $j = 1, 2$.

These are the probabilities which are needed to classify an observation x into a population. The classification rule is to assign x to j=1 if:

$$P(1 | X=x) > P(2 | X=x) \dots (A)$$

Otherwise assign x to j=2.

The P(j | X=x) probabilities can be calculated from the P(X=x | j) distributions using Bayes theorem as shown:

 $P(j | X=x) = P(X=x | j) P(j) / P(x) \dots (B)$

Substituting (B) into (A) gives:

$$P(X=x \mid 1) P(1) / P(x) > P(X=x \mid 2) P(2) / P(x) ... (C)$$

The P(x) will cancel when forming the classification rule (D) and there is no need to evaluate them. Equation (C) simplifies by cancelling the P(x) as well as the common factors in the normal distributions and then taking logarithms to produce:

$$-0.5 * ((x - \mu_1)/\sigma)^2 + \log(q_1) > -0.5 * ((x - \mu_2)/\sigma)^2 + \log(q_2) \dots (D)$$

The classification function CF_i for j = 1, 2 is:

$$CF_{i} = -0.5 * ((x - \mu_{i})/\sigma)^{2} + \log(q_{i}) \dots (E)$$

ESTIMATING CF FROM THE SAMPLES

The samples from populations 1 and 2 are used to estimate the parameters μ_j and σ . The sample means \bar{x}_j estimate μ_j . The pooled variance S_p^2 estimates σ^2 where S_j^2 is the sample variance for sample j and

$$S_p^2 = \{(n_1 - 1)S_1^2 + (n_2 - 1)S_2^2\}/(n_1 - 1 + n_2 - 1)$$

The sample CF is used in the sample based classification rule:

$$-0.5 * ((x - \bar{x}_1)/S_p)^2 + \log(q_1) > -0.5 * ((x - \bar{x}_2)/S_p)^2 + \log(q_2) \dots (F)$$

THE CLASSIFICATION FUNCTIONS AND LOG-ODDS-RATIO

Returning to equation (A), the odds ratio of membership in j=1 versus j=2 is given by:

$$P(1 | X=x) / P(2 | X=x) = (P(X=x | 1) P(1) / P(x)) / (P(X=x | 2) P(2) / P(x)) ... (G)$$

Logarithms are taken of equation G, and P(x) is cancelled to give equation (H):

Using equation (D), equation (H) becomes:

Log Odds-Ratio =
$$-0.5(-2x(\mu_1 - \mu_2) + {\mu_1}^2 - {\mu_2}^2) / \sigma^2 + \log(q1/q2) \dots (I)$$

Equation (I) shows that the Log-Odds-Ratio from 2-group discriminant analysis is a linear function of x

Log Odds-Ratio = $\beta_0 + \beta_1 x \dots (J)$

where $\beta_0 = -(\mu_1^2 - \mu_2^2) / 2\sigma^2 + \log(q1/q2)$ and $\beta_1 = (\mu_1 - \mu_2) / \sigma^2$

When fitting the 2-group discriminant analysis model to the sample, the estimates b_{0D} , b_{1D} for β_0 , β_1 are found by replacing μ_1 , μ_2 , σ with \bar{x}_1 , \bar{x}_2 , and S_p in equation (J)

$$b_{0D} = -(\bar{x}_1^2 + \bar{x}_2^2) / 2S_p^2 + \log(q1/q2)$$
 and $b_{1D} = (\bar{x}_1 - \bar{x}_2) / S_p^2 \dots (K)$

The "D" is added as a subscript to indicate that the method of fitting is by discriminant analysis.

If we make the assumption that $(1/S_p)$ is a constant, then, as shown by (L), the expression for b_{1D} is a linear transformation of a t-statistic with $n_1 + n_2 - 2$ d.f. The term K is the constant $(\mu_1 - \mu_2) / S_p^2$.

$$b_{1D} = \{(\bar{x}_1 - \bar{x}_2) - (\mu_1 - \mu_2)\} / S_p^2 + (\mu_1 - \mu_2) / S_p^2 = t (1/S_p) \text{ sqrt}(1/n_1 + 1/n_2) + K \dots (L)$$

The appropriateness of the assumption of constant S_p is justified by having large sample sizes n_1 and n_2 .

CONNECTION WITH LOGISTIC REGRESSION

The logistic regression model with predictor x also is formulated as:

Log Odds-Ratio = $\beta_0 + \beta_1 x$.

In the case of logistic regression the parameter estimates b_{0L}, b_{1L} are fitted by maximum likelihood.

The "L" is added as a subscript to indicate that the method of fitting is Logistic Regression with maximum likelihood.

CONNECTING b_{1D} AND b_{1L}

Under the assumptions of Appendix A, b_{1D} is essentially an unbiased estimator of β_1 . The actual expectation is given by $E(b_{1D}) = ((n_1 + n_2 - 2)/(n_1 + n_2 - 4))\beta_1$. Meanwhile, b_{1L} is asymptotically an unbiased estimator of β_1 .

The Wald chi-square for b_{1L} gives the significance for rejecting the null that $\beta_1 = 0$.

For discriminant analysis, the null hypothesis that $\beta_1 = 0$ makes $(\mu_1 - \mu_2) = 0$. The hypothesis H0: $\beta_1 = 0$ vs. H1: $\beta_1 \neq 0$ at significance " α " is tested by finding a critical value C so that the test statistic T = t (1/S_p) sqrt(1/n₁ + 1/n₂) satisfies P(|T| > C) = α . This test can be changed to finding a critical C' so that P(|t| > C') = α where C' = C / ((1/S_p) sqrt(1/n₁ + 1/n₂)). But then C' is the t-statistic value t_{$\alpha/2$}.

For large $n_1 + n_2$ the square of the t-statistic is essentially a chi-square with 1 d.f. The test above can be rephrased in terms of a chi-square.

APPENDIX B: NECESSARY CHANGES TO %MFP8

A. The "%then" must be removed in the sub-macro "fpmodels"

%macro fpmodels(model,y,x,pref,base,m,stvars);

%else /*%then*/

B. For purposes of running in Windows %mfp8 must be modified by replacing "/" with "\" as shown in the statements below:

%include "&MacPath.\boxtid.sas"; %include "&MacPath.\xtop.sas"; %include "&MacPath.\xvars.sas"; %include "&MacPath.\fpmodels.sas"; %include "&MacPath.\datasave.sas"; %include "&MacPath.\exlabb.sas"; %include "&MacPath.\exinc.sas"; %include "&MacPath.\labs.sas"; %include "&MacPath.\brename.sas"; %include "&MacPath.\funcfm.sas";