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# Correcting the Quasi-complete Separation Issue in Logistic Regression Models

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#### **ABSTRACT**

Quasi-complete separation is a commonly detected issue in logit/probit models. Quasi-complete separation occurs when the dependent variable separates an independent variable or a combination of several independent variables to a certain degree. In other words, levels in a categorical variable or values in numeric variable are separated by groups in a discrete outcome variable. Most of the time, it happens in categorical independent variable(s). Quasi-complete separation can cause convergence failures in the model, which consequently result in inflated coefficient estimates and standard errors. Therefore it potentially yields biased results.

This paper reviews various approaches for correcting Quasi-complete separation using a binary logistic regression model as the example. First it introduces the concept of Quasi-complete separation and how to diagnose the issue. Then it provides step-by-step guidelines for fixing the problem from straightforward data configuration approach to complicated statistical modeling approach, such as EXACT method and FIRTH method.

**Keywords:** Quasi-complete separation, logistic regression, Greenacre's method, FIRTH method and cluster analysis.

#### INTRODUCTION

Logistic regression is a statistical method used to measure the relationship between a dichotomous outcome variable and one or more independent variables. It is also called a logit model, because the log odds of the outcome variable is modeled as a linear combination of the predictor variables. Logistic regression is the special case of generalized linear model (GLM) with a logit function. Quasi-complete separation is a common issue in regression analysis when outcome variable is categorical variable, including logistic regression and probit regression. This paper describes several remedies ranging from simple data configuration to relatively complex statistical modeling.

# WHAT IS QUASI-COMPELET SEPARATION

Quasi-complete separation occurs when the dependent variable separates an independent variable or a combination of several independent variables to a certain degree. In other words, at least one group of the dependent variable has zero frequency for at least one category of an independent variable. It can happen with continuous variables as well. However, it most often occurs with categorical variables. Here is an example.

## **EXAMPLE DATA**

The following example is based on a randomly generated artificial dataset called DEMO and it does not refer to any business/company. The dataset (N=1,500) consists one ID variable, one dependent variable and three independent variables.

- Y: is the binary dependent variable (0 or 1)
- X1: is the continuous variable
- X2: is the nominal discrete variable (1-15)
- X3: is the ordinal discrete variable (1-8)

Table 1 shows the first 10 observations of the DEMO dataset.

ID	Υ	X1	X2	Х3
1	0	-0.043	2	8
2	0	0.648	2	3
3	0	-0.222	5	3
4	0	-0.731	14	6
5	0	-1.521	12	4
6	0	-0.430	11	6
7	0	0.176	5	1
8	0	-0.626	3	2
9	0	-0.644	15	3
10	0	2.387	2	6

# **DETECTION QUASI-COMPELET SEPARATION**

First, run logistic regression model with PROC LOGISTIC procedure.

```
proc logistic data=demo desc;
  class x2 x3;
  model y = x1 x2 x3;
run;
```

Some warning messages from LOG indicate that the model is not adequate and questionable.

WARNING: There is possibly a quasi-complete separation of data points. The maximum likelihood estimate may not exist.

WARNING: The LOGISTIC procedure continues in spite of the above warning. Results shown are based on the last maximum likelihood iteration. Validity of the model fit is questionable.

Ana	aly	sis	of Maxim	um Likeli	hood Estima	ates
				Standard	Wald	
Parameter	-	DF	Estimate	Error	Chi-Square	Pr > ChiSq
Intercept		1	-4.9729	27.8511	0.0319	0.8583
x1		1	-0.2316	0.1048	4.8794	0.0272
x2	1	1	1.3249	16.1333	0.0067	0.9346
x2	2	1	1.0359	16.1352	0.0041	0.9488
x2	3	1	0.5199	16.1350	0.0010	0.9743
x2	4	1	0.3724	16.1380	0.0005	0.9816
x2	5	1	0.5070	16.1366	0.0010	0.9749
x2	6	1	0.8538	16.1346	0.0028	0.9578
x2	7	1	1.1387	16.1342	0.0050	0.9437
x2	8	1	0.4046	16.1366	0.0006	0.9800
x2	9	1	0.8057	16.1351	0.0025	0.9602
x2	10	1	1.2411	16.1347	0.0059	0.9387
x2	11	1	1.3379	16.1337	0.0069	0.9339
x2	12	1	-12.0529	225.8	0.0028	0.9574
x2	13	1	0.8858	16.1351	0.0030	0.9562
x2	14	1	1.0258	16.1342	0.0040	0.9493
x3	1	1	1.7404	22.7056	0.0059	0.9389
x3	2	1	1.5500	22.7057	0.0047	0.9456
x3	3	1	2.1626	22.7053	0.0091	0.9241
x3	4	1	1.2059	22.7062	0.0028	0.9576
х3	5	1	1.7343	22.7055	0.0058	0.9391
х3	6	1	-11.4435		0.0052	0.9426
x3	7	1	1.7112	22.7057	0.0057	0.9399

From the estimate table, we can see that standard errors are extremely large when x2=12 and X3=6 compared to other levels in x2 and X3 variables. And the magnitudes of point estimate of these two levels are also very large. All of these abnormal signs convince us that the estimates of coefficients and standard errors from the model are not reliable. We can validate this assumption easily by running 2 by 2 frequency tables shown as below,

			X2													
		1	2	3	4	5	6	7	8	9	10	11	12	13	14	15
V	0	106	82	122	92	89	102	87	96	92	76	81	96	85	97	91
ĭ	1	13	7	7	4	5	8	9	5	7	8	11	0	7	9	6

				X3								
			1	2	3	4	5	6	7	8		
Ī	V	0	171	173	165	178	188	177	169	173		
	T	1	16	14	23	10	17	0	15	11		

When X2=12 or X3=6, the cell frequency of Y=1 are all 0. In a 2 by 2 table, at least one cell's frequency is zero, which is a typical example of a quasi-complete separation. A quasi-complete separation can cause infinite parameter estimation in the model and failure of convergence of maximum likelihood estimation (MLE).

## FIXING QUASI-COMPELET SEPARATION

## **COLLECTING MORE DATA**

Most of the time quasi-complete separation is associated with small sample sizes. Recruiting more samples in the development data could be a possible remedy by filling the empty cells. However, it is not practical because additional data may not be available.

#### **EXCLUDING CASES**

Excluding the level(s) causing the issue and applying the rest of data into the model is another way to fix quasi-complete separation.

```
proc logistic data=Demo desc;
  class x2 x3;
  model y = x1 x2 x3;
  where x2 not in (12) and x3 not in (6);
run;
```

Ana	Analysis of Maximum Likelihood Estimates										
				Standard	Wald						
Parameter		DF	Estimate	Error	Chi-Square	Pr > ChiSq					
Intercept		1	-2.4772	0.1122	487.6614	<.0001					
x1		1	-0.2316	0.1048	4.8794	0.0272					
x2	1	1	0.4639	0.2989	2.4097	0.1206					
x2	2	1	0.1749	0.3862	0.2051	0.6506					
x2	3	1	-0.3411	0.3789	0.8103	0.3680					
x2	4	1	-0.4884	0.4900	0.9936	0.3189					
x2	5	1	-0.3539	0.4423	0.6400	0.4237					
x2	6	1	-0.00711	0.3607	0.0004	0.9843					
x2	7	1	0.2778	0.3452	0.6476	0.4210					
x2	8	1	-0.4563	0.4411	1.0697	0.3010					
x2	9	1	-0.0552	0.3836	0.0207	0.8855					
x2	10	1	0.3802	0.3678	1.0683	0.3013					
x2	11	1	0.4770	0.3208	2.2114	0.1370					

Ana	Analysis of Maximum Likelihood Estimates										
				Standard	Wald						
Parameter	-	DF	Estimate	Error	Chi-Square	Pr > ChiSq					
x2	13	1	0.0248	0.3846	0.0042	0.9485					
x2	14	1	0.1648	0.3444	0.2290	0.6323					
x3	1	1	0.1056	0.2494	0.1792	0.6721					
x3	2	1	-0.0848	0.2606	0.1059	0.7448					
x3	3	1	0.5278	0.2191	5.8051	0.0160					
x3	4	1	-0.4288	0.2974	2.0789	0.1493					
х3	5	1	0.0995	0.2420	0.1690	0.6810					
х3	7	1	0.0765	0.2551	0.0899	0.7644					

After applying WHERE statement, there is no more warning message and all statistical inferences are in the normal range.

## **COLLAPSING THE LEVELS**

When categorical variable with quasi-complete is ordinal, one might think of redefining that categorical variable by collapsing the levels based on the order in that variable. For example, there is an ordinal categorical variable called age group, which contains five levels (0-18, 19-30, 30-50, 50-60 and 60+). Let's suppose all Y are equal to 1 when age group=30-50. In other words, the frequency cell of Y=0 and age group=30-50 is 0. As we already knew from prior example, quasi-complete will occur and model will fail to converge. What you can do is to combine age group of 30-50 with group 50-60 together and create a new age group, 30-60. You also may combine age group 30-50 with 19-30. In the demo example, X3 is the ordinal categorical independent variable. So level 6 of X3 can be combined with either level 5 or 7. In the next example, level 6 was combined with level 7.

```
Data demo2;
    set demo;
    if x3 = 6 then x3 = 7;
run;

proc logistic data=demo2 desc;
    class x3;
    model y = x1 x3;
run;
```

Ana	Analysis of Maximum Likelihood Estimates											
	Γ			Standard	Wald							
Parameter	-	DF	Estimate	Error	Chi-Square	Pr > ChiSq						
Intercept		1	-2.6087	0.1071	593.0077	<.0001						
x1		1	-0.2337	0.1025	5.1984	0.0226						
х3	1	1	0.2137	0.2448	0.7624	0.3826						
x3	2	1	0.0517	0.2574	0.0404	0.8407						
х3	3	1	0.6171	0.2156	8.1913	0.0042						
х3	4	1	-0.3264	0.2944	1.2295	0.2675						
х3	5	1	0.1903	0.2387	0.6359	0.4252						
х3	7	1	-0.5617	0.2463	5.1998	0.0226						

The demo data has been reconstructed and the result showed the quasi-complete separation issue has been solved.

So far, I have introduced multiple ways of fixing separation issue in non-statistical approaches. In the next section, several statistical approaches will be introduced.

#### **GREENACRE'S METHOD**

Greenacre's method is a data-driven method to collapse levels in the categorical variable. It is a repeatable iteration process, which means, in each iteration, the Chi-square statistics from all possible

combinations of two levels are produced and compared. The levels giving the least difference in Chisquare statistic will be merged together. This method is a fast and efficient way to reduce the dimension of the categorical input since only univariate association was considered.

In the sample dataset DEMO, independent variable X2 has 15 levels. First, frequencies and the proportions of event by each level in X2 were calculated. For example, X2=1, FREQ=119 and PROP=0.109, means there are total 119 cases when X2=1 and among them, about 10.9% (13/119) cases have the events(Y=1).

x2	FREQ	PROP
1	119	0.109
2	89	0.079
3	129	0.054
4	96	0.042
5	94	0.053
6	110	0.073
7	96	0.094
8	101	0.050
9	99	0.071
10	84	0.095
11	92	0.120
12	96	0.000
13	92	0.076
14	106	0.085
15	97	0.062

Next step is to apply Greenacre's method by using PROC CLUSTER with the WARD method. WARD method or Ward's minimum variance method is a commonly used hierarchical cluster technique that merges two clusters together based on smallest decrease in the ANOVA sum of squares or Chi-square statistic. Ward's method tends to join clusters with a small number of observations, and is strongly biased toward producing clusters with roughly the same number of observations. It is also very sensitive to outliers (Milligan 1980).

```
ods output clusterhistory=clus_history;
proc cluster data=demo_freq method=ward;
   freq _freq_;
   var propor;
   id x2;
   run;
ods listing;
proc print data=clus_history;
run;
```

Step	NumberOfClusters	ldj1	ldj2	FreqOfNewCluster	SemipartialRSq	RSquared
1	14	3	5	223	0.0001	1
2	13	7	10	180	0.0001	1
3	12	6	9	209	0.0002	1
4	11	2	13	181	0.0002	0.999
5	10	CL14	8	324	0.0011	0.998
6	9	CL11	CL12	390	0.0025	0.996
7	8	1	11	211	0.0046	0.991
8	7	CL13	14	286	0.005	0.986
9	6	CL10	15	421	0.0055	0.981

10	5	CL6	4	517	0.0109	0.97
11	4	CL9	CL7	676	0.0375	0.932
12	3	CL8	CL4	887	0.1399	0.793
13	2	CL5	12	613	0.1832	0.609
14	1	CL3	CL2	1500	0.6094	0

The NumberOfClusters field gives the total number of cluster after merging. They are in descending order and the last value is 1, because all levels will be grouped into one big cluster eventually. Idj1 and Idj2 are the name of the levels used in the clustering. Any level starting with CL means the previously collapsed levels and the numbers following CL is the resulting cluster. For example, at step 5, Idj1 is CL14, which represents the new cluster of level 3 and level 5 at step1; Idj2 is 8, the level 8 in variable X2. FreqOfNewCluster indicates the number of observation after merging. SemipartialRSq is the semi-partial R-squares, which is the disease in the proportion of variance accounted for by joining two clusters. The value of SemipartialRSq at step1 is 0.0001, which is difference between the overall Chi-square statistic and Chi\_square statistic after collapsing level 3 and 5. They are in increasing order, because WALD method merges two clusters with smallest variation from all possible combinations of any two clusters. The last column is squared multiple correlation (RSQ), representing the proportion of variance remaining after the levels are collapsed.

```
data cutoff;
  if _n_=1 then set chi;
  set clus_history;
  chisquare = _pchi_*rsquared;
  degfree = NumberOfClusters-1;
  logpvalue = logsdf('CHISQ',chisquare,degfree);
run;
```

The next step is to calculate overall Chi-square statistics.

```
proc freq data=demo;
  tables x2*y / chisq;
  output out=chi(keep=_pchi_) chisq;
run;
```

Result: PCHI =18.36

Then get the logarithm value by multiplying overall Chi-square statistic with proportion of variance remaining by individual level.

```
proc sql;
    create table cutoff as
    select a.*, b.*,
    logsdf('CHISQ',_pchi_*rsquared,numberofclusters-1) as logpvalue
    from chi a, clus_history b;
quit;
```

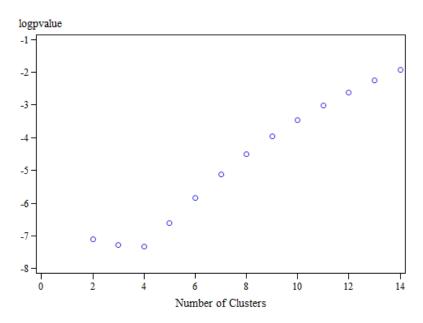


Figure 1. Plot of number of cluster vs. logPvalue

From the Figure 1, we can see that when clusters are 4, logvalue has the lowest value. So the optimum number of clusters is 4.

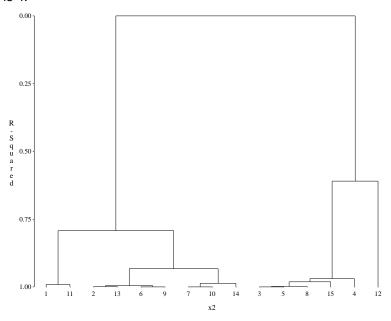


Figure 2. Dendrogram of clusters for variable X2.

The following summary table is based on dendrogram shown above.

CLUSTER	x2
	3
	5
1	8
	15
	4
	7
	10
	6
2	9
	2
	13
	14
2	1
3	11
4	12

## **EXACT METHOD**

Conventional logistic regression generates coefficient estimates through the maximum likelihood estimate (MLE) method. However, MLE is suitable for large sample estimation. If the sample is too small, standard error or P-value from MLE can be biased. Exact logistic regression can handle small sample size, skewed data or sample with quasi-complete separation. You may consider it as some generation format of Fisher's exact test. This method was originally introduced by Cox (1970), and the computation method has been employed in PROC LOGISTIC procedure since SAS® 8.1 version. Please keep in mind that logistic conditional exact methods is a time and memory consuming process and requires quite amount of computational time. Only 480 observations were randomly selected out of total 1500 from demo dataset.

```
proc logistic data=Demo_Sample desc;
  class x3 / param=ref;
  model y = x3;
  exact x3 / estimate=both;
run;
```

The following is the parameter estimate table from EXACT method.

	Exact Parameter Estimates										
Paramete	r Estimate	Standard Erro	r95% Confide	ence Limits	Two-sided p-Value						
х3	1 0.9617	0.8545	-0.8974	3.3517	0.4390						
х3	2-6.3E-15	1.0128	-2.6551	2.6551	1.0000						
х3	3 0.4194	0.9280	-1.7856	2.9362	1.0000						
х3	4 0.4194	0.9280	-1.7856	2.9362	1.0000						
х3	5 0.7224	0.8827	-1.2648	3.1619	0.6794						
х3	6 -0.8933 *		-Infinity	1.2410	0.4958						
x3	7 0.7224	0.8827	-1.2648	3.1619	0.6794						

Note: \* indicates a median unbiased estimate.

#### PENALIZED LIKELIHOOD METHOD OR FIRTH METHOD

Logistic regression with exact method can handle small sample size and quasi-complete separation data set. However, it can be very computationally demanding and not very efficient. An alternative method that can be used in large data set or quasi-complete separation is called penalized likelihood method or

FIRTH method, which was introduced by Firth (1993). In this method, Newton-Raphson algorithm was used to calculate the vector of first derivatives.

$$U(\beta) = \frac{\partial \log L}{\partial \beta} = \sum_{i} x_{i} y_{i} - \sum_{i} x_{i} y_{i} - \sum_{i} h_{i} x_{i} (.5 - y_{i})$$

Where,  $h_i$  is the i<sup>th</sup> diagonal element of the "hat" matrix H.

$$H = W^{1/2} X (X'WX)^{-1} X'W^{1/2} \text{ and } W = diag \{ \hat{y}_i (1 - \hat{y}_i) \}$$

FIRTH method is implemented in PROC LOGISTIC procedure by using "FIRTH" option and it is recommended to use it with CLPARM option as well. This is because conventional confidence interval from Wald test produces biased and misleading result. CLPARM=PL means computation of confident interval is based on profile likelihood; while CLPARM=WALD, means the confidence intervals are individual Wald test based.

```
proc logistic data=demo desc;
  class x3;
  model y = x1 x3 / firth clparm=pl;
run;
```

Analysis of Penalized Maximum Likelihood Estimates									
	Γ			Standard	Wald				
Parameter	-	DF	Estimate	Error	Chi-Square	Pr > ChiSq			
Intercept		1	-2.9000	0.2002	209.8496	<.0001			
x1		1	-0.2369	0.1013	5.4641	0.0194			
х3	1	1	0.5362	0.2994	3.2071	0.0733			
х3	2	1	0.3785	0.3097	1.4939	0.2216			
х3	3	1	0.9294	0.2762	11.3231	0.0008			
x3	4	- 1	0.0143	0.3400	0.0018	0.9666			
x3	5	1	0.5119	0.2944	3.0235	0.0821			
х3	6	1	-3.0067	1.2430	5.8509	0.0156			
х3	7	1	0.4856	0.3044	2.5445	0.1107			

Parameter Estimates and Profile-Likelihood										
Confidence Intervals										
Parameter		Estimate	95% Confid	lence Limits						
Intercept		-2.9000	-3.5218	-2.5875						
x1		-0.2369	-0.4385	-0.0361						
х3	1	0.5362	-0.0388	1.2498						
х3	2	0.3785	-0.2235	1.1027						
х3	3	0.9294	0.4153	1.6196						
х3	4	0.0143	-0.6679	0.7708						
x3	5	0.5119	-0.0505	1.2203						
х3	6	-3.0067	-7.2396	-1.3151						
x3	7	0.4856	-0.1023	1.2043						

# **CONCLUSION**

Quasi-complete separation is common in regression analyses where the outcome variable is categorical. It causes the failure of convergence in the model and yields unreliable inference and misleading results. This paper demonstrated multiple methods to remedy this issue ranging from data configuration to statistical approaches. Greenacre's method is good when the categorical variable(s) with problem has (have) a lot of levels; EXACT method is good when the sample is small and FIRTH method is suitable for small or big data.

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