# INFERENCE ON HIGH DIMENSIONAL DISCRETE CHOICE MODEL

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#### Introduction

- Discrete choice models are presented as development and renovation of the classical choice theory. The model has been used extensively to discrete choice processes in different fields such as econometrics and transportation.
- Abe (1999) developed a methodology for discrete choice data using the generalized additive model (GAM).
- Giron (2010) assessed the feasibility and efficiency of using principal components of the predictors in the generalized additive mode (GAM) for discrete choice data and determine its capacity to manage drawbacks of highdimensional data.
- Torres (2013) extended the work of Giron (2010) by using generalized adaptive sparse-PCA (GAS-PCA) as a data reduction tool and used GAM for high dimensional discrete data.



his study aims to conduct variable selection process on the explanatory ples using two methods of sparse principal component analysis. The

## Methodology

• The methodology starts with the data generating process to create a high dimensional data for discrete choice model.

$$P(Y = y_j | \underline{X} = \underline{x}_j^*) = \frac{exp\left(\underline{X}_j^*\beta_p\right)}{\sum_{i=1}^J exp\left\{\underline{X}_i^*\beta_p + \epsilon_i\right\}} \ i = 1, 2, \dots J$$

Where,

- $X_j^*$  matrix of predictors/characteristics of individual j
- Y the response variables, with possible values,  $Y_1, Y_2, ..., Y_r$
- $\beta_p$  beta coefficients
- $\epsilon_i$  random error

The data consists of a single categorical response variable, Y, which can take on j possible alternatives, and p predictor variables,  $x_1, x_2, ..., x_p$ , measured on n subjects, with population covariance matrix S =The following notations will be used throughout this study:



- sample size
- number of dimensions/ predictors
  - number of categories of the response variable
  - number of sparse principal components selected

# Methodology

- The proposed estimation proposed procedure composed of the following:
  - Principal Component Analysis and Sparse Principal Component Analysis using
    - Penalized Matrix Decomposition Sparse Principal Component (PMD-SPC)
    - General Adaptive Sparse Principal Component Analysis (GAS-PCA)
  - Local Scoring Algorithm and Backfitting Algorithm.

The proposed estimation procedure is then followed by the hypothesis testing on significant predictors, controlling false discovery rate and power and size computation. The design of the simulation study is stated also including all settings for each scenario.



- The data generation process involves five hidden factors that will generate 5 groups of strongly correlated predictors. The latent factors L<sub>1</sub>, L<sub>2</sub>, L<sub>3</sub>, L<sub>4</sub> and L<sub>5</sub> represent the "important drivers" that predict the response variable. The latent factors were generated such that:
- $L_1 \sim N(50, 40^2), L_2 \sim N(50, 20^2), L_3 \sim N(50, 10^2), L_4 \sim N(50, 5^2), L_5 \sim N(50, 5^2)$  $\operatorname{cor}(L_1, L_2, L_4, L_5 | L_3) = \begin{bmatrix} 1 & 0 & 0.6 & 0 & 0 \\ 0 & 1 & -0.5 & 0 & 0 \\ 0.6 & -0.5 & 1 & 0 & 0 \\ 0 & 0 & 0 & 1 & 0 \\ 0 & 0 & 0 & 0 & 1 \end{bmatrix}$
- Given those specifications,  $L_1$  and  $L_2$  are independent of each other,  $L_3$  is moderately correlated to  $L_1$  and  $L_2$ . On the other hand,  $L_4$  and  $L_5$  are independent on the other hidden factors.



- For NHD scenarios, L<sub>1</sub>, L<sub>2</sub> and L<sub>3</sub> will each have 8 significant predictors while L<sub>4</sub> and L<sub>5</sub> will each have 4 significant predictors. For HD scenarios, all 5 groups will have 8 significant predictors each.
- We then generate 5000 observations from each predictor and compute for  $Y^* = \alpha + \sum_i f_i(X_i) + \epsilon$  where  $\epsilon$  is from a Gumbel distribution.



Case	Relative Contribution by Group						
	G 1	G 2	G 3	G 4	G 5		
200 –	10%	15%	15%	25%	35%		
200	(20)	(30)	(30)	(50)	(70)		
200 –	1%	1.5%	1.5%	48%	48%		
2000	(20)	(30)	(30)	(960)	(960)		

Table1. Scenario Settings for Dichotomous Response Simulation Study

Table 2. Scenario Settings for 3-category Response Simulation Study

Case	Relative Contribution by Group				
	G 1	G 2	G 3	G 4	G 5
210 –	10%	10%	20%	30%	30%
210	(21)	(21)	(42)	(63)	(63)
210 –	1%	1%	2%	48%	48%
2100	(21)	(21)	(42)	(1008)	(1008)

Balanced (50-50) Let  $Y^{*}{}_{(i)} = \begin{cases} 1, & \text{for } i = 1, 2, ..., 2400 \\ 2, & \text{for } i = 2601, 2602, ..., 5000 \end{cases}$ Moderately Unbalanced (70-30) Let  $Y^{*}{}_{(i)} = \begin{cases} 1, & \text{for } i = 1, 2, ..., 3360 \\ 2, & \text{for } i = 3561, 3562, ..., 5000 \end{cases}$ Severely Unbalanced (10-90) Let  $Y^{*}{}_{(i)} = \begin{cases} 1, & \text{for } i = 1, 2, ..., 480 \\ 2, & \text{for } i = 681, 682, ..., 5000 \end{cases}$ 

We trimmed the middle 200 observations to put some threshold between the boundary of the two categories inducing the dichotomy. The population size for each distribution of the dichotomous category is 4800.



#### Balanced (33-33-34)

$$Y^{*}_{(i)} = \begin{cases} 1, & \text{for } i = 1, 2, ..., 1500 \\ 2, & \text{for } i = 1751, 1752, ..., 3250 \\ 3, & \text{for } i = 3501, 3502, ..., 5000 \end{cases}$$

Moderately Unbalanced (60-20-20) Let

$$Y^{*}{}_{(i)} = \begin{cases} 1, & \text{for } i = 1, 2, ..., 2700 \\ 2, & \text{for } i = 2951, 2952, ..., 3850 \\ 3, & \text{for } i = 4101, 4102, ..., 5000 \end{cases}$$

Severely Unbalanced (70-20-10) Let  $Y^{*}{}_{(i)} = \begin{cases} 1, & \text{for } i = 1, 2, ..., 3150 \\ 2, & \text{for } i = 3401, 3402, ..., 4300 \\ 3, & \text{for } i = 4551, 4552, ..., 5000 \end{cases}$ 

We trimmed the middle 250 observations to put some threshold between the boundary of the 1<sup>st</sup> and 2<sup>nd</sup> categories and 250 trimmed observations between 2<sup>nd</sup> and 3<sup>rd</sup> category. The population size for each distribution of the 3-category is 4500.

Let

#### **Benchmark/ Baseline Test**

To further evaluate the performance of the proposed test, a benchmark or baseline test was also implemented together with the proposed test using the same set of generated data. The baseline test that will be used Bonferroni Test for family wise error rate. Power and size will be computed for each of the predictors simultaneously.



# Results and Discussions

The proposed method used two different transformations of the utility function: linear and exponential. The proposed method was not able to detect significant predictors when there is an exponential transformation. In all exponential transformation of the utility function, only the predictors from the  $L_1$ were retained after the dimension reduction process. However, the backfitting algorithm only produced very few (min: 0; max: 4) significant predictors and were all eventually rejected by the BH procedure in the end.

## **Results and Discussions**

- The proposed method also was not able to detect any significant predictors coming from  $L_4$  and  $L_5$  groups. All significant predictors selected by the proposed method came from  $L_1$  to  $L_3$  group most of the time. The most consistent group with the highest number of predictors selected by PMD-SPC and GAS-PCA is  $L_1$ .
- The p-values computed came from the linear component of the ANOVA table. The p-values in the nonparametric part are always nonsignificant.
- For the estimated power and size for dichotomous and 3-category response HD case using GAS-PCA, GAS-PCA failed to reduce the dimensionality of the predictors since HPCs available for use can only run a batch for 7 days but in that period, GAS-PCA failed to do so. Power and size cannot be computed for HD cases of GAS-PCA. This is recommended for further studies.



# **Results and Discussions**

- In all cases using PMD-SPC, the proposed test is not always correctly sized. In fact, the baseline test is also not correctly sized at 0.05 level sometimes. The following tables provide a summary of the size for each variable.
- In all cases using GAS-PCA, the proposed test is not always correctly sized. In most cases, the baseline and the proposed method have the same size. In the case of 3-category response where n=p, only 1 not truly significant variable was selected by the GAS-PCA at 95% variance explained.



	Balanced						
	N	HD	HD				
Va	Prop	Base	Prop	Base			
<i>X</i> 9	0.042	0.041	0.025	0.010			
<i>X</i> <sub>10</sub>	0.044	0.06	0.031	0.012			
<i>X</i> <sub>11</sub>	0.053	0.049	0.039	0.025			
<i>X</i> <sub>12</sub>	0.050	0.054	0.047	0.016			
<i>X</i> <sub>13</sub>	0.047	0.039	0.021	0.015			
<i>X</i> <sub>14</sub>	0.062	0.052	0.036	0.016			
<i>X</i> <sub>15</sub>	0.055	0.067	0.033	0.022			
<i>X</i> <sub>16</sub>	0.046	0.054	0.042	0.028			
<i>X</i> <sub>17</sub>	0.056	0.047	0.073	0.032			
<i>X</i> <sub>18</sub>	0.052	0.043	0.029	0.016			
<i>X</i> <sub>19</sub>	0.073	0.067	0.051	0.026			
<i>X</i> <sub>20</sub>	0.047	0.039	0.088	0.043			
Va	Prop	Base	Prop	Base			
<i>X</i> 9	0.044	0.015	0.030	0.018			
<i>X</i> <sub>10</sub>	0.041	0.012	0.063	0.030			
<i>X</i> <sub>11</sub>	0.078	0.036	0.022	0.005			
<i>X</i> <sub>12</sub>	0.027	0.013	0.057	0.021			
<i>X</i> <sub>13</sub>	0.045	0.010	0.015	0.008			
<i>X</i> <sub>14</sub>	0.039	0.020	0.054	0.030			
<i>X</i> <sub>15</sub>	0.053	0.031	0.056	0.023			
<i>X</i> <sub>16</sub>	0.05	0.020	0.021	0.005			
<i>X</i> <sub>17</sub>	0.064	0.023	0.028	0.008			
<i>X</i> <sub>18</sub>	0.051	0.023	0.058	0.022			
<i>X</i> <sub>19</sub>	0.076	0.041	0.093	0.041			
<i>X</i> <sub>20</sub>	0.074	0.032	0.098	0.067			
<i>X</i> <sub>21</sub>	0.071	0.055	0.121	0.043			

**Results and Discussions** 

- For the estimated power using PMD-SPC, the test is powerful in detecting the significant predictors in all scenarios compared to the baseline test. It should be noted that as distribution becomes more unbalanced, the power increases.
- In all cases using GAS-PCA n=p, the test is powerful in detecting the significant predictors in all scenarios compared to the baseline test.

# Summary and Conclusion

- The simulation showed that the proposed test is somehow correctly sized only in balanced cases. In all unbalanced cases, the proposed test is always incorrectly sized. Even though the test is incorrectly sized, the baseline test is also incorrectly sized and the proposed method showed a comparable size compared to the baseline test.
- The result of the simulation shows that the proposed test is found to be more powerful than the baseline test across different parameters and number of variables involved.



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