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## Randomized Response and the Binary Probit Model

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IAW-Discussion Paper



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## Randomized Response and the Binary Probit Model

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# Randomized response and the binary probit model

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JEL classification: C21, C25, C42, C81

## Summary

The paper analyzes effects of randomized response with respect to some binary dependent variable on the estimation of the probit model. This approach is used in interviews when asking sensitive questions. Alternatively randomization can be considered as a means of statistical disclosure control which has been termed post randomization method (PRAM). The paper shows that all properties concerning parameter estimation are maintained although there is a loss in (asymptotic) efficiency.

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# 1. Introduction

The binary probit model (see, for example, Greene 2003 chapter 21.3 or Ronning 1991 chapter 2.2.1) considers the effect of some explanatory variable  $x$  on a latent continuous variable  $Y^*$ , i.e. we assume that the model

$$Y^* = \alpha + \beta x + \varepsilon \quad (1)$$

holds where  $\varepsilon$  is a normally distributed random error with  $E(\varepsilon) = 0$  and  $V(\varepsilon) = 1$ . However  $Y^*$  is observed only as a binary or dichotomous variable  $Y$  which is defined by the threshold model

$$Y = \begin{cases} 1 & \text{if } Y^* > 0 \\ 0 & \text{if } Y^* \leq 0 \end{cases} \quad (2)$$

The sample information is given by  $n$  pairs  $(x_i, y_i)$  where  $y_i \in \{0, 1\}$  and  $x_i$  is an arbitrary real number. Maximum likelihood estimation of the two unknown parameters  $\alpha$  and  $\beta$  is straightforward; see some standard text as, for example, Greene (2003) or Ronning (1991). Note that we have already introduced the usual identifying restrictions, i.e. zero threshold and unit error variance. We confine ourselves to the case of just one regressor which is assumed to be continuous. The results in this paper however apply also to the more general case of an arbitrary number of explanatory variables after minor modifications. Here we consider randomization of the dichotomous variable  $y$  which switches its values with some prescribed transition probability (leaving the explanatory variable  $x$  in its original form). In the following section the method is described in some more detail. Section 3 then considers the effect on the estimation of the binary probit model. Some concluding remarks are added in section 4.

## 2. Randomized response and post randomization

Randomized response originally was introduced to avoid non-response in surveys containing sensitive questions on, e.g., drug consumption or AIDS disease. See Warner (1965). Särndal et. al. (1992 p. 573) suggested use of this method "to protect the anonymity of individuals". A good description of the difference between the two (formally equivalent) approaches is given by van den Hout and van der Heijden (2002): In the randomized response setting the stochastic model has to be defined in advance of data collection whereas in post randomization this method will be applied to the data already obtained.

For the case of a dichotomous variable the method can be described as follows: Consider a  $(2 \times 2)$  probability matrix describing transition probabilities for the two states "0" and "1" of a random variable  $Y$ :

$$P_y = \begin{pmatrix} p_{00} & p_{01} \\ p_{10} & p_{11} \end{pmatrix}$$

In the following we denote the dichotomous variable obtained from post randomization by  $Y^m$  which we call the 'masked' variable. Then the transition probabilities can

be defined by  $p_{jk} \equiv P(Y^m = j | Y = k)$  with  $j, k \in \{0, 1\}$  and  $p_{j0} + p_{j1} = 1$  for  $j = 0, 1$ . Since there is no argument not to treat the two states symmetrically, we define by  $\pi$  the probability of no change. Then the probability matrix can be written as follows:

$$P_y = \begin{pmatrix} \pi & 1 - \pi \\ 1 - \pi & \pi \end{pmatrix}$$

Note that this matrix is singular if  $\pi = \frac{1}{2}$  which will become important later on.

When the sample of the dependent variable has undergone randomization, we will have  $n$  observations  $y_i^m$  where  $y_i^m$  is the dichotomous variable obtained from  $y_i$  by the randomization procedure.

Randomization has the advantage that the original distribution of  $Y$  can be estimated from the masked observations  $y_i^m$ . See Kooiman et al (1997) for a detailed exposition. The sample of unmasked observations is completely characterized by  $n$ , the number of observations, and  $\theta = \sum_i y_i$ , the number of 'successes' which is the parameter of interest. Defining  $T^m = \sum_i Y_i^m$  an unbiased estimator is given by

$$\hat{\theta} = \frac{T^m - n(1 - \pi)}{(2\pi - 1)}$$

with variance

$$\text{var}(\hat{\theta}) = \frac{n \pi (1 - \pi)}{(2\pi - 1)^2} \quad (3)$$

which does not depend on  $\theta$ . However the coefficient of variation is inversely related to  $\theta$  and therefore distortion from post-randomization will be serious if this parameter is near to 0 or  $n$ . "This fits nicely into our general purpose to protect rare scores, since these are most vulnerable to disclosure." (Kooiman et al 1997 p. 4).

### 3. Estimation of the probit model under randomization

Let us now turn to the estimation of the probit model as given in (1) and (2). From figure 1 it is apparent that under randomization we have the following data generating process:

$$Y_i^m = \begin{cases} 1 & \text{with probability } \Phi_i \pi + (1 - \Phi_i)(1 - \pi) \\ 0 & \text{with probability } \Phi_i(1 - \pi) + (1 - \Phi_i)\pi \end{cases} \quad (4)$$

Here  $\Phi_i$  denotes the conditional probability under the normal distribution that the unmasked dependent variable  $Y_i$  takes on the value 1 for given  $x_i$ , i.e.  $\Phi_i \equiv \Phi(\alpha + \beta x_i) = P(Y_i^* > 0 | x_i)$ . See (1) and (2).

From (4) we obtain the following likelihood function:

$$\begin{aligned} \mathcal{L}(\alpha, \beta | (y_i^m, x_i), i = 1, \dots, n) \\ = \prod_{i=1}^n (\Phi_i \pi + (1 - \Phi_i)(1 - \pi))^{y_i^m} (\Phi_i(1 - \pi) + ((1 - \Phi_i)\pi)^{(1 - y_i^m)} \end{aligned}$$



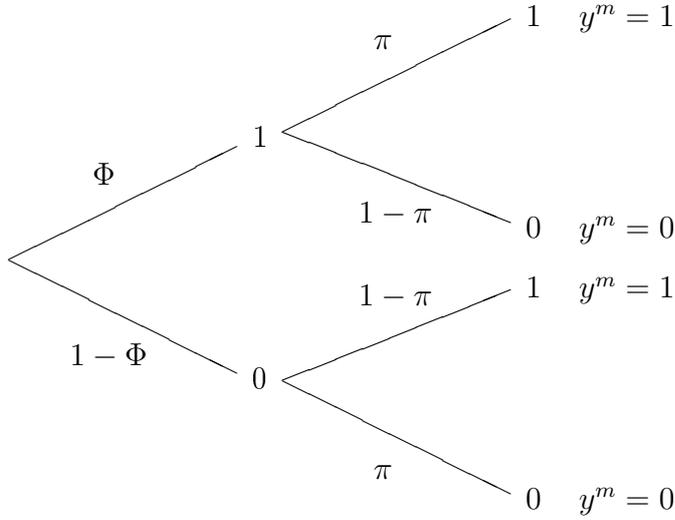


Figure 1: Data generating process under randomization

Global concavity of this function with respect to  $\alpha$  and  $\beta$  may be checked by deriving first and second (partial) derivatives of the log-likelihood function

$$L \equiv \log(\mathcal{L}) = \sum_i^n y_i^m \log[\Phi_i \pi + (1 - \Phi_i)(1 - \pi)] \\ + (1 - y_i^m) \log([\Phi_i(1 - \pi) + ((1 - \Phi_i)\pi)].$$

The partial first-order derivatives with respect to  $\alpha$  and  $\beta$  are given by

$$\begin{aligned} \frac{\partial L}{\partial \alpha} &= (2\pi - 1) \sum_i \left[ y_i^m \frac{\phi_i}{W_i} - (1 - y_i^m) \frac{\phi_i}{1 - W_i} \right] \\ &= (2\pi - 1) \sum_i \left[ \frac{(y_i^m - W_i) \phi_i}{W_i(1 - W_i)} \right] \\ \frac{\partial L}{\partial \beta} &= (2\pi - 1) \sum_i \left[ y_i^m \frac{\phi_i}{W_i} - (1 - y_i^m) \frac{\phi_i}{1 - W_i} \right] x_i \\ &= (2\pi - 1) \sum_i \left[ \frac{(y_i^m - W_i) \phi_i}{W_i(1 - W_i)} \right] x_i \end{aligned} \quad (5)$$

where we use the following definitions:

$$\phi_i \equiv \phi(\alpha + \beta x_i) = \frac{1}{\sqrt{2\pi}} \exp \left\{ -\frac{1}{2} (\alpha + \beta x_i)^2 \right\} \quad (\text{standard normal density})$$

and

$$W_i \equiv \pi \Phi_i + (1 - \pi)(1 - \Phi_i) \quad (\text{probability of observing } y_i^m = 1).$$

Note that the first order conditions would be fulfilled for  $\pi = \frac{1}{2}$  for *any*  $\alpha$  and  $\beta$  leaving these two parameters unidentified. Therefore we exclude this case in the following by assuming

$$\pi \neq \frac{1}{2} . \quad (6)$$

Moreover the two partial derivatives are similar (with  $W_i$  instead of  $\Phi_i$ ) in structure to those concerning the standard probit model disregarding the proportional factor  $(2\pi - 1)$ . See, e.g. , Ronning (1991 p. 45).

Unfortunately it turns out that the Hessian matrix formed from the second order partial derivatives is no longer negative definite contrary to the standard probit case which guarantees global concavity of the log-likelihood function. First note that

$$\begin{aligned} \frac{\partial W_i}{\partial \alpha} &= (2\pi - 1) \phi_i & \frac{\partial W_i}{\partial \beta} &= (2\pi - 1) \phi_i x_i \\ \frac{\partial (1-W_i)}{\partial \alpha} &= -(2\pi - 1) \phi_i & \frac{\partial (1-W_i)}{\partial \beta} &= -(2\pi - 1) \phi_i x_i \\ \frac{\partial W_i(1-W_i)}{\partial \alpha} &= (2\pi - 1) \phi_i (1 - 2W_i) & \frac{\partial W_i(1-W_i)}{\partial \beta} &= (2\pi - 1) \phi_i x_i (1 - 2W_i) \end{aligned}$$

and

$$\begin{aligned} \frac{\partial \phi_i}{\partial \alpha} &= -(\alpha + \beta x_i) \phi_i = -z_i \phi_i \\ \frac{\partial \phi_i}{\partial \beta} &= -(\alpha + \beta x_i) \phi_i x_i = -z_i \phi_i x_i \end{aligned}$$

using  $z_i \equiv \alpha + \beta x_i$ .

For the second-order partial derivative with respect to  $\alpha$  we obtain

$$\begin{aligned} \frac{\partial^2 L}{(\partial \alpha)^2} &= (2\pi - 1) \left\{ \sum_i \left[ \left( \frac{\partial}{\partial \alpha} \frac{y_i^m - W_i}{W_i(1 - W_i)} \right) \phi_i + \frac{y_i^m - W_i}{W_i(1 - W_i)} \left( \frac{\partial}{\partial \alpha} \phi_i \right) \right] \right\} \\ &= (2\pi - 1) \left\{ \sum_i \left[ \frac{\phi_i(2\pi - 1)(-W_i^2 - y_i^m(1 - 2W_i)) - W_i(1 - W_i)(y_i^m - W_i)z_i}{(W_i(1 - W_i))^2} \right] \phi_i \right\} \\ &= -(2\pi - 1) \left\{ \sum_i \left[ \frac{(2\pi - 1)(y_i^m - 2y_i^m W_i + W_i^2) \phi_i + (y_i^m - W_i)W_i(1 - W_i)z_i}{(W_i(1 - W_i))^2} \right] \phi_i \right\} \\ &= -(2\pi - 1) \left\{ \sum_i \frac{g_i \phi_i}{(W_i(1 - W_i))^2} \right\} \quad (7) \end{aligned}$$

where

$$g_i = (2\pi - 1)(y_i^m - 2y_i^m W_i + W_i^2) \phi_i + (y_i^m - W_i)W_i(1 - W_i)z_i . \quad (8)$$

Corresponding results are obtained for the two other second-order partial derivatives leading to the following Hessian matrix:

$$H^m = -(2\pi - 1) \sum_i \frac{g_i \phi_i}{(W_i(1 - W_i))^2} \mathbf{u}_i \mathbf{u}_i' \quad (9)$$

with  $\mathbf{u}_i' = (1 \ x_i)$ . Since  $2\pi - 1$  may be either positive or negative and the function  $g_i$  in (8) is more complex than in the standard case, the proof of negative definiteness used in the standard probit case does not go through here. See, for example, Amemiya(1985 p. 274) or Ronning (1991 p. 46).

However we obtain a simple formula for the information matrix from which it is immediately apparent that estimation under randomization implies an efficiency loss. First note that  $E[Y_i^m] = W_i$  and therefore the expected value of  $g_i$  as a function of  $Y_i^m$  is given by

$$\begin{aligned} E[g(Y_i^m)] &= E[(2\pi - 1)(Y_i^m - 2Y_i^m W_i + W_i^2) \phi_i + (Y_i^m - W_i)W_i(1 - W_i)z_i] \\ &= (2\pi - 1)(W_i - 2W_i^2 + W_i^2)\phi_i + (W_i - W_i)W_i(1 - W_i)z_i \\ &= (2\pi - 1)W_i(1 - W_i)\phi_i \end{aligned}$$

Therefore the information matrix (or the the expected value of the Hessian matrix multiplied by -1) in case of masked data is given by

$$\mathcal{I}^m = (2\pi - 1)^2 \sum_i \frac{\phi_i^2}{W_i(1 - W_i)} \mathbf{u}_i \mathbf{u}_i' \quad (10)$$

whereas in the case of unmasked data we obtain (e.g. Amemiya 1985 p. 272 or Ronning 1991 p. 46)

$$\mathcal{I} = \sum_i \frac{\phi_i^2}{\Phi_i(1 - \Phi_i)} \mathbf{u}_i \mathbf{u}_i' \quad (11)$$

We now want to show that the difference  $\mathcal{I} - \mathcal{I}^m$  is nonnegative definite. It is sufficient to show that for every  $i$

$$\frac{1}{\Phi_i(1 - \Phi_i)} > \frac{(2\pi - 1)^2}{W_i(1 - W_i)} \quad (12)$$

or  $W_i(1 - W_i) > (2\pi - 1)^2 \Phi_i(1 - \Phi_i)$  which follows immediately from

$$W_i(1 - W_i) = (2\pi - 1)^2 \Phi_i(1 - \Phi_i) + \pi(1 - \pi)$$

recalling the definition  $W_i = \pi\Phi_i + (1 - \pi)(1 - \Phi_i)$  from above.

Which values of  $\pi$  imply the largest efficiency loss ? For the 'weights' in (10) we can write

$$\begin{aligned} \frac{(2\pi - 1)^2}{W_i(1 - W_i)} &= \frac{(2\pi - 1)^2}{(2\pi - 1)^2 \Phi_i(1 - \Phi_i) + \pi(1 - \pi)} \\ &= \frac{1}{\Phi_i(1 - \Phi_i) + \frac{\pi(1 - \pi)}{(2\pi - 1)^2}} \end{aligned}$$

Since the function

$$h(\pi) = \frac{\pi(1 - \pi)}{(2\pi - 1)^2}$$

is symmetric, i.e.  $h(\pi) = h(1 - \pi)$ , monotonically increasing and tending towards infinity at  $\pi = 1/2$ , the weights in (10) tend towards zero and the information matrix  $\mathcal{I}^m$  tends towards a zero matrix when  $\pi$  (or  $(1 - \pi)$ ) tends towards  $1/2$ . Note that the function  $h(\pi)$  has already appeared in the variance of  $\hat{\theta}$  in section 2.

## 4. Concluding remarks

The paper has shown that randomization of the binary dependent variable involves an efficiency loss which is larger when the probability of switching tends towards  $1/2$  whereas this loss is small when this probability is near zero or one. We have not discussed performance of the maximum likelihood estimator since it becomes clear from the presented results that consistency and asymptotic normality still hold under randomization of the dependent variable. From the perspective of statistical disclosure control masking of the explanatory variable should also be taken into account. For example, the continuous explanatory variable could be masked by microaggregation or addition of noise. See, for example, Domingo-Ferrer (2002), in particular the section on microdata protection. However, this additional transformation seems to have no clear-cut effects on estimation but will be a topic of further research.

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