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# Semi-nonparametric estimation of extended ordered probit models

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**Abstract.** This paper presents a semi-nonparametric estimator for a series of generalized models that nest the ordered probit model and thereby relax the distributional assumption in that model. It describes a new Stata command for fitting such models and presents an illustration of the approach.

**Keywords:** st0056, ordered response models, ordered probit, semi-nonparametric estimation

## 1 Introduction

The analysis of ordered response data is increasingly common in a growing number of research areas. The standard ordered probit (Aitchison and Silvey 1957) and ordered logit (McCullagh 1980) models are commonly used but require the researcher to make a specific distributional assumption.<sup>1</sup> This paper presents a semi-nonparametric estimator for a series of generalized models that nest the ordered probit model and relax the distributional assumption in that model.

## 2 A framework for ordered response models

As with many limited dependent variable models, it is useful to consider specification of a suitable model for ordered response data in terms of an underlying latent variable,

$$y_i^* = x_i' \beta + \varepsilon_i \quad (1)$$

for  $i = 1, \dots, N$ , where  $x_i$  is a vector of observations on a set of explanatory variables,  $\beta$  is a vector of unknown parameters, and  $\varepsilon_i$  is a random-error term independently distributed with distribution function  $F$ . If  $y_i^*$  was observed for all observations,  $\beta$  could be consistently estimated by ordinary least squares without requiring a distributional assumption on  $\varepsilon$ . For the type of data considered in this paper,  $y_i^*$  itself is not observed. Rather, the observed dependent variable,  $y_i$ , is discrete, taking one of the values  $\{1, 2, \dots, J\}$ , and is related to  $y_i^*$  as follows:

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<sup>1</sup>See Greene (2003) and Wooldridge (2002) for recent textbook treatments of such models.

$$y_i = \begin{cases} 1 & \text{if } y_i^* < \alpha_1 \\ 2 & \text{if } \alpha_1 \leq y_i^* < \alpha_2 \\ \vdots & \\ J & \text{if } \alpha_{J-1} \leq y_i^* \end{cases}$$

with the  $\alpha_j$  being additional parameters such that  $\alpha_1 < \alpha_2 < \dots < \alpha_{J-1}$ . Thus, the range of  $y^*$  is partitioned into  $J$  mutually exclusive and exhaustive intervals, and the variable  $y$  indicates the interval into which a particular observation falls. The dependent variable  $y$  is ordinal, and the  $\alpha_j$  are treated as parameters to be estimated. The probability of a particular observed outcome, for  $2 \leq j \leq J-1$ , is given by

$$\begin{aligned} \Pr[y_i = j] &= \Pr[\alpha_{j-1} \leq y_i^* < \alpha_j] \\ &= \Pr[\alpha_{j-1} - x_i'\beta \leq \varepsilon_i < \alpha_j - x_i'\beta] \\ &= F(\alpha_j - x_i'\beta) - F(\alpha_{j-1} - x_i'\beta) \end{aligned}$$

where  $F$  is the cumulative distribution function of  $\varepsilon_i$  and is assumed to contain no additional unknown parameters so that, for example,  $\varepsilon_i$  has a known variance. This assumption fixes the scale of the measurement of  $y^*$  but not the origin. Identification can be achieved by assuming a zero intercept (i.e., assuming that  $x_i$  does not contain a constant term) or by fixing one of the  $\alpha_j$ . The former is used here.

The full set of probabilities of the possible outcomes are

$$\Pr[y_i = j] = \begin{cases} F(\alpha_1 - x_i'\beta) & \text{if } j = 1 \\ F(\alpha_j - x_i'\beta) - F(\alpha_{j-1} - x_i'\beta) & \text{if } 2 \leq j \leq J-1 \\ 1 - F(\alpha_{J-1} - x_i'\beta) & \text{if } j = J \end{cases} \quad (2)$$

If we adopt the additional notation that  $\alpha_0 = -\infty$  and  $\alpha_J = +\infty$ , we can write these more compactly as

$$\Pr[y_i = j] = F(\alpha_j - x_i'\beta) - F(\alpha_{j-1} - x_i'\beta)$$

for all  $j$ . This defines a class of cumulative probability models in which a known transformation of the cumulative probabilities is taken to be a linear function of the  $x$  variables and only the intercept in this function differs across the categories:

$$F^{-1}\{\Pr[y_i \leq j]\} = \alpha_j - x_i'\beta$$

A natural estimator for this type of model is the maximum likelihood estimator. Define

$$y_{ij} = \begin{cases} 1 & \text{if } y_i = j \\ 0 & \text{else} \end{cases}$$

Then, the log likelihood for the model is given by

$$\log L = \sum_{i=1}^N \sum_{j=1}^J y_{ij} \log [F(\alpha_j - x_i'\beta) - F(\alpha_{j-1} - x_i'\beta)] \quad (3)$$

This is maximized with respect to  $(\beta, \alpha_1, \dots, \alpha_{J-1})$ , i.e.,  $M + J - 1$  parameters, where  $M$  is the number of exogenous variables, remembering that  $\beta$  (and hence  $M$ ) does not include an intercept.

By far the most commonly used models to date for analyzing ordered responses have been the ordered probit and ordered logit models (which take  $F$  to be standard normal and logistic, respectively). The ordered probit model, introduced by Aitchison and Silvey (1957), assumes that  $\varepsilon_i \sim N(0, \sigma^2)$ . Adopting the scale normalization  $\sigma = 1$  and imposing a zero intercept for identification, the probabilities are given by

$$\Pr[y_i = j] = \Phi(\alpha_j - x'_i \beta) - \Phi(\alpha_{j-1} - x'_i \beta)$$

where  $\Phi$  is the cumulative distribution function of a standard normal and the log likelihood by (3) with  $F$  replaced by  $\Phi$ .

### 3 Semi-nonparametric estimation

The “semi-nonparametric” series estimator of an unknown density, proposed by Gallant and Nychka (1987), approximates the density using a Hermite form. This approximation can be written as the product of a squared polynomial and a normal density. This gives a polynomial expansion with Gaussian leading term. In fact, an expansion around any density with a moment-generating function could be used, but the normal is a natural choice in the current context because the resulting model then nests the ordered probit. To ensure that the density approximation is proper (i.e., that it integrates to 1, in addition to the non-negativity ensured by the square), the approximation is specified as

$$f_K(\varepsilon) = \frac{1}{\theta} \left( \sum_{k=0}^K \gamma_k \varepsilon^k \right)^2 \phi(\varepsilon) \quad (4)$$

where  $\phi(\varepsilon)$  is the standard normal density function and where

$$\theta = \int_{-\infty}^{\infty} \left( \sum_{k=0}^K \gamma_k \varepsilon^k \right)^2 \phi(\varepsilon) d\varepsilon$$

However, this general specification of the density is invariant to multiplication of the vector  $\gamma = (\gamma_0, \gamma_1, \dots, \gamma_K)$  by a scalar, and a normalization is required. The restriction  $\gamma_0 = 1$  is a convenient choice.

The required distribution function is therefore specified as

$$F_K(u) = \frac{\int_{-\infty}^u \left( \sum_{k=0}^K \gamma_k \varepsilon^k \right)^2 \phi(\varepsilon) d\varepsilon}{\int_{-\infty}^{\infty} \left( \sum_{k=0}^K \gamma_k \varepsilon^k \right)^2 \phi(\varepsilon) d\varepsilon} \quad (5)$$

This defines a family of “semi-nonparametric” (SNP) distributions for increasing values of  $K$ .

Providing the unknown density satisfies certain smoothness conditions; it can be approximated arbitrarily closely by this Hermite series by increasing the choice of  $K$ , the degree of the polynomial. The class of densities that can be approximated by this form is very general. Any form of skewness, kurtosis, etc. is permitted, but “violently oscillatory” density functions are ruled out. In addition to the smoothness restriction, there is an upper bound on the tails of the densities. (See Gallant and Nychka (1987, 369) for a fuller discussion of the class of densities.)

Under these and other “mild” regularity conditions, and providing  $K$  increases with the sample size, the model parameters are estimated consistently (Gallant and Nychka 1987) by maximizing the pseudo-likelihood function obtained by replacing the unknown distribution function in (3) by that in (5).

A location normalization is also necessary for semi-parametric identification. That used by Gabler, Laisney, and Lechner (1993) in the binary response case to give the error term a zero mean is rather cumbersome. However, the restriction can be on either the error distribution or the systematic part of the model. Melenberg and van Soest (1996) use an alternative normalization, setting the constant term in their model equal to its probit estimate. This is far more convenient. The equivalent used here in the ordered response model context is to set the first threshold,  $\alpha_1$ , equal to its ordered probit estimate.

Note that in the  $K = 1$  case, the imposition of  $E(\varepsilon) = 0$  can be easily shown to imply  $\gamma_1 = 0$  so that the model in this case reduces to the ordered probit model. The imposition of the equivalent restriction used here has the same implication. An important additional feature of the model in the general case is that the score for  $\gamma_2$  is zero at the ordered probit estimates, which is equivalent to the result for the binary probit case proved in Appendix B of Gabler, Laisney, and Lechner (1993). This implies that the model for  $K = 2$  is also equivalent to the ordered probit model (as well as those for  $K = 0$  and  $K = 1$ ). The model with  $K = 3$  is therefore the first model in the series that is a generalization of the ordered probit model.

In practice, inference is conducted conditional on  $K$ , possibly for a range of alternative values of  $K$ , with the final specification of the model chosen by tests between them. Thus, in reality the model is treated as parametric for a given  $K$ , with the choice of  $K$  part of the model selection procedure. As Pagan and Ullah (1999) point out, the orientation is nonparametric, but the *modus operandi* is parametric. This latter characteristic is appealing because estimation can be conducted in a familiar maximum likelihood environment.

The SNP-ML approach can therefore be viewed as a series of polynomial densities and corresponding pseudo-likelihood functions and standard model selection procedures used to reduce the dimension of the parameter vector and improve efficiency in finite samples. This can, for example, be on the basis of Wald or likelihood-ratio tests on the  $\gamma$ -vector or using information criteria, such as those of Akaike, Schwarz, or Hannan and Quinn.

The polynomial form of (4) means that the cumulative probabilities given by (5), required for the pseudo-likelihood function, can be derived using the higher order truncated moments of a standard normal distribution. First, the SNP density can be written as

$$f_K(\varepsilon) = \frac{1}{\theta} \sum_{k=0}^{2K} \gamma_k^* \varepsilon^k \phi(\varepsilon)$$

where

$$\gamma_k^* = \sum_{i=a_k}^{b_k} \gamma_i \gamma_{k-i}$$

with  $a_k = \max(0, k - K)$  and  $b_k = \min(k, K)$ . The scaling factor to give a proper density is then given by

$$\theta = \int_{-\infty}^{\infty} \sum_{k=0}^{2K} \gamma_k^* \varepsilon^k \phi(\varepsilon) d\varepsilon = \sum_{k=0}^{2K} \gamma_k^* \mu_k \quad (6)$$

where  $\mu_k$  is the  $k$ th moment of a standard normal distribution. Given that  $\gamma_0^* = \gamma_0^2 = 1$  and that  $\mu_k = 0$  for  $k$  odd, this can be written as

$$\theta = 1 + \sum_{k=1}^K \gamma_{2k}^* \mu_{2k}$$

The cumulative probabilities in (5) required for the pseudo-likelihood function can be written as

$$F_K(u) = \frac{1}{\theta} \sum_{k=0}^{2K} \gamma_k^* \int_{-\infty}^u \varepsilon^k \phi(\varepsilon) d\varepsilon$$

This is a linear combination of the truncated moments of a standard normal distribution:

$$I_k(u) = \int_{-\infty}^u \varepsilon^k \phi(\varepsilon) d\varepsilon$$

Computation of the likelihood contributions for the maximum likelihood iterations can then be simplified by using the recursion

$$I_k(u) = (k-1)I_{k-2}(u) - u^{k-1}\phi(u) \quad (7)$$

with  $I_0(u) = \Phi(u)$  and  $I_1(u) = -\phi(u)$ . Each truncated moment can be written as a polynomial combination of the standard normal cumulative distribution and density functions. The formulation can be further simplified in the current context where each cumulative probability is a linear combination of truncated moments scaled by the corresponding combination of complete moments. It can be shown that the recursion need only involve the normal density function part of the expressions for these truncated moments. The cumulative probabilities in (5) can be written as

$$F_K(u) = \frac{1}{\theta} \sum_{k=0}^{2K} \gamma_k^* I_k(u)$$

with  $\theta$  given by the corresponding combination in (6). Using the recursion in (7), the truncated moments in this linear combination can also be given by

$$I_k(u) = \mu_k \Phi(u) - A_k(u) \phi(u)$$

where the  $A_k(u)$  are given by the recursion

$$A_k(u) = (k-1)A_{k-2}(u) + u^{k-1}$$

with  $A_0(u) = 0$  and  $A_1(u) = 1$ . Using this, the cumulative probabilities in (5) required for the pseudo-likelihood function can be written as

$$\begin{aligned} F_K(u) &= \frac{1}{\theta} \sum_{k=0}^{2K} \gamma_k^* [\mu_k \Phi(u) - A_k(u) \phi(u)] \\ &= \Phi(u) - \frac{1}{\theta} \left[ \sum_{k=0}^{2K} \gamma_k^* A_k(u) \right] \phi(u) \end{aligned} \quad (8)$$

The term in square brackets is a polynomial in  $u$  of order  $(2K-1)$ . Thus, the cumulative probabilities in the SNP framework have the standard normal cumulative distribution function as leading term and differ from this by the product of the standard normal density and this polynomial in  $u$ . The SNP estimator is then given by the maximization of the pseudo-likelihood function obtained by replacing the unknown distribution function in (3) by this final expression in (8).

## 4 The sneop command

### 4.1 Syntax

```
sneop depvar [varlist] [if exp] [in range] [weight] [, order(#) robust
      from(matname)]
```

`fweights`, `pweights`, and `iweights` are allowed; see [U] **14.1.6 weight**.

`sneop` shares the features of all estimation commands; see [U] **23 Estimation and post-estimation commands**.

### 4.2 Options

`order(#)` specifies the order of the Hermite polynomial to be used. The default is 3. Orders 1 and 2 give models equivalent to the ordered probit.

`robust` specifies that the Huber/White/sandwich estimator of variance is to be used in place of the traditional calculation; see [U] **23.11 Obtaining robust variance estimates**.

`from(matname)` specifies a matrix containing starting values. (This option can be used to investigate whether a global maximum has been found.) The default uses the ordered probit estimates as starting values.

### 4.3 Example using Stata Reference Manual data

This section gives an example of the use of the command and the output produced using the automobile data (<http://www.stata-press.com/data/r8/fullauto.dta>) used in [R] **oprobit** to model the 1977 repair records of 66 foreign and domestic cars. The variable `rep77` takes values poor, fair, average, good, and excellent (coded 1 to 5). The explanatory variables in the model are `foreign` (origin of manufacturer), `length` (a proxy for size), and `mpg`. All three have significant positive effects in the fitted ordered probit model. The output from the **sneop** program with  $K = 5$  is as follows:

```
. use http://www.stata-press.com/data/r8/fullauto.dta
(Automobile Models)
. sneop rep77 foreign length mpg, order(5)
Order of SNP polynomial = 5
Number of categories = 5, rep77 assumed coded 1,...,5
1st threshold set to Ordered Probit estimate = 10.158904
initial:      log likelihood = -78.020025
(output omitted)
Iteration 11: log likelihood = -77.099231
SNP Estimation of Extended Ordered Probit Model   Number of obs   =          66
                                                    Wald chi2(3)      =        333.83
Log likelihood = -77.099231                      Prob > chi2       =         0.0000
```

rep77	Coef.	Std. Err.	z	P> z	[95% Conf. Interval]	
foreign	1.311067	.3460665	3.79	0.000	.6327897	1.989345
length	.0485854	.0030438	15.96	0.000	.0426197	.0545511
mpg	.1550694	.0200241	7.74	0.000	.1158228	.1943159
Thresholds 1	10.1589	Fixed				
2	11.14811	.3948174	28.24	0.000	10.37428	11.92193
3	12.47549	.8385617	14.88	0.000	10.83194	14.11904
4	14.04136	1.023632	13.72	0.000	12.03508	16.04764
SNP coeffs: 1	-.3883669	.9299041	-0.42	0.676	-2.210946	1.434212
2	-.1782807	.1767086	-1.01	0.313	-.5246232	.1680619
3	-.1692988	.2396551	-0.71	0.480	-.6390143	.3004166
4	.0252651	.0580259	0.44	0.663	-.0884635	.1389938
5	.0189329	.0166693	1.14	0.256	-.0137383	.0516041

```
Likelihood ratio test of OP model against SNP extended model:
Chi2(3) statistic =      1.841588      (p-value = .6059271)
```

Estimated moments of error distribution:			
Variance =	1.39027	Standard Deviation =	1.179097
3rd moment =	2.120129	Skewness =	1.293343
4th moment =	13.94238	Kurtosis =	7.21338



The log likelihood is improved very little relative to the ordered probit model, and the likelihood-ratio test does not reject the ordered probit model: a  $\chi^2(3)$ -statistic of 1.84, giving a  $p$ -value of 0.61. This example is provided to illustrate the output produced by the program on the data and model used in [R] **oprobit**. It is, however, unlikely that we would be able to observe a significant difference between the fitted density and a standard normal on the basis of a sample as small as this.

## 5 Job satisfaction illustration

The use of the semi-nonparametric estimator is illustrated in more detail in this section on a model of self-declared job satisfaction, as specified by Clark and Oswald (1996), and using data from the 1991 wave of the British Household Panel Survey. Working respondents provided job satisfaction scores ranging from 1 (“not satisfied at all”) to 7 (“completely satisfied”). Clark and Oswald (1996) examine the hypothesis that job satisfaction depends on income relative to a “comparison” or reference level. Their main findings are that job satisfaction levels are positively related to own earnings and negatively related to “comparison” earnings. Other often-quoted features of their results are the higher job satisfaction levels of women and the U-shaped age profile.

Log-likelihood values and likelihood-ratio statistics for different values of  $K$  from 3 to 8 are given in table 1. The standard ordered probit (OP) model is rejected in all cases. The likelihood-ratio tests for  $K - 1$  against  $K$  reject the null, at conventional significance levels, for  $K \leq 5$  but accept it for  $K$  above 5, suggesting selection of the  $K = 5$  model. This use of likelihood-ratio tests runs into the usual problem with sequential testing procedures, which could be addressed with successive revision of the significance level.

Table 1: Likelihood-ratio tests for different values of  $K$

$K$	log likelihood	LR-test of OP	deg. of freedom	$p$ -value	LR-test of $K - 1$	$p$ -value
OP	-6174.25					
3	-6169.87	8.76	1	0.003	8.76	0.003
4	-6167.95	12.60	2	0.002	3.83	0.050
5	-6165.48	17.54	3	0.001	4.95	0.026
6	-6165.31	17.87	4	0.001	0.33	0.568
7	-6164.56	19.38	5	0.002	1.51	0.219
8	-6164.56	19.39	6	0.004	0.01	0.939

Note: 1 degree of freedom for LR test of  $K - 1$  in each row.

This is a standard model selection problem. There are a number of alternative model selection criteria available (Greene 2003). The best known is probably the Akaike information criterion (AIC), equal to the log likelihood minus the number of freely estimated parameters. However, in some cases, the AIC has been found to have a tendency towards selecting models with too many parameters, even asymptotically. A commonly used alternative is the Schwarz Bayesian criterion (SBC), based on a large sample approximation to the posterior odds ratio of the models. Under certain conditions, the SBC

selects the right model with probability 1 asymptotically (Amemiya 1985), but in some finite sample cases, it has been found to have a tendency to underfit. A third possibility is the Hannan–Quinn criterion (HQC), introduced in the context of autoregressive time-series models but more generally applicable. It gives values between the AIC and SBC and is sometimes viewed as a useful compromise. For the models in table 1, the AIC and the HQC both choose  $K = 5$ , while the SBC, which penalizes extra parameters more heavily, selects  $K = 3$ . For a sample of the size used here (3,895 observations), the SBC penalizes an extra parameter 4.13 on the log-likelihood scale, while the HQC and AIC penalize it 2.11 and 1, respectively.

Table 2: Job satisfaction models for different values of  $K$ 

	OP	SNP(3)	SNP(5)
	coef (s.e.)	coef (s.e.)	coef (s.e.)
log(earnings)	0.134 (.054)	0.096 (.050)	0.087 (.056)
log(comp. earn.)	−0.283 (.064)	−0.254 (.060)	−0.323 (.068)
Male	−0.156 (.044)	−0.147 (.044)	−0.130 (.046)
Age/10	−0.210 (.100)	−0.154 (.092)	−0.141 (.104)
Age <sup>2</sup> /100	0.038 (.013)	0.031 (.011)	0.030 (.013)
<u>Thresholds:</u>			
1	−4.125 (.377)	−4.125	−4.125
2	−3.917 (.376)	−3.879 (.043)	−3.847 (.047)
3	−3.562 (.375)	−3.476 (.093)	−3.390 (.075)
4	−2.995 (.375)	−2.877 (.161)	−2.603 (.125)
5	−2.416 (.374)	−2.319 (.211)	−1.889 (.164)
6	−1.664 (.374)	−1.657 (.254)	−1.151 (.177)
<u>Polynomial:</u>			
1		−0.050 (.193)	0.415 (.062)
2		−0.097 (.088)	0.366 (.252)
3		−0.051 (.021)	−0.073 (.171)
4			−0.087 (.034)
5			−0.002 (.016)
Log likelihood	−6174.25	−6169.87	−6165.48
Standard deviation	1	0.979	1.369
Skewness	0	0.034	0.064
Kurtosis	3	4.600	4.665
Test sum = 0 [ $\chi^2(1)$ ]	7.40	9.26	15.83
$p$ -value	[0.007]	[0.002]	[0.000]

Notes: (1) Sample size = 3895. (2) Models also contain 34 other variables.

The estimated parameters (coefficients on main variables only) for the ordered probit model and for the extended model with  $K = 3$  and  $K = 5$  are given in table 2.<sup>2</sup> A number of other statistics are also given.

<sup>2</sup>The other variables included in the model are as in table 3 of Clark and Oswald (1996). The data used here are from the current release of wave 1, while they used the original release. The ordered probit results are very close to theirs but not identical.

Estimates of  $\beta$  cannot be compared directly across models without adjustment, because the fitted densities differ. The  $\beta$  coefficients are only identified after the imposition of scale and origin restrictions (as is the case for the ordered probit model). Thus, even if considering effects on the latent dependent variable  $y_i^*$ , it is not appropriate to compare estimates of the  $\beta$  coefficients across the models directly.

However, there are a number of ways in which comparisons can be made, depending on the focus of the study. We might hypothesize that there is an underlying but unobservable continuous measure of job satisfaction (or utility or welfare, depending on one's perspective) that is an unknown but monotonic function of the variable  $y_i^*$  in (1). In this case, we would be interested in influences on this underlying variable and relations between them.

Consider two factors that influence job satisfaction (i.e., two elements of the vector  $x$  in (1)). We could plot “indifference curves” between them: combinations of the two factors that give the same level of job satisfaction. The specification in (1) means that they will be straight lines. The slopes of these (parallel) indifference curves between the two factors will be minus the ratio of their  $\beta$ -coefficients. Thus, ratios of coefficients are of interest and are compared below.

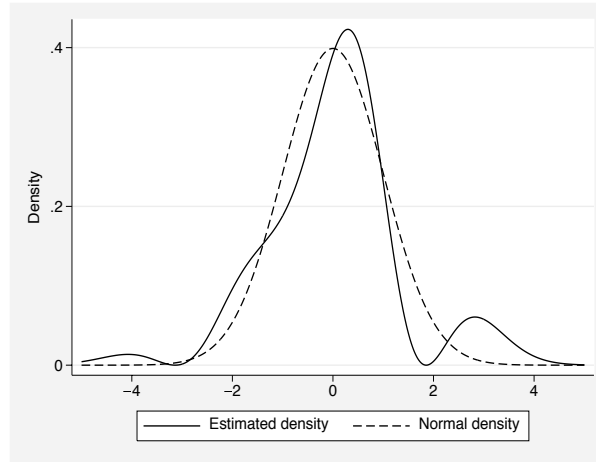
Ratios of  $\beta$  coefficients are also informative about the impacts of  $x$  variables on the probabilities of observed outcomes (or cumulations thereof). The partial derivatives of the probabilities of  $y$  with respect to  $x$ , the “marginal effects”, are proportional to  $\beta$  with the factor of proportionality being an evaluation of the density function (or a difference in densities). For example, from (2), the partial derivative of  $\Pr[y_i = J]$  with respect to  $x$  is  $\beta \cdot f(\alpha_{J-1} - x'_i \beta)$ . The “marginal effects” are therefore scaled versions of the  $\beta$  coefficients and the scaling factor is the same for each element of  $\beta$ . Hence, a ratio of “marginal effects” is equal to the corresponding ratio of  $\beta$  coefficients. Thus, ratios of coefficients are of interest from this perspective also.

Two such ratios are illustrated in table 3. The first is minus the ratio of the two earnings coefficients. In the ordered probit model, the ratio indicates that a 10% increase in comparison earnings would need to be compensated by a 21% increase in own earnings to give the same level of job satisfaction. This rises to 26% in the SNP model with  $K = 3$  and 37% in the SNP model with  $K = 5$ . The standard errors on these ratios, calculated by the “delta method” (using the Stata `nlcom` command), are also given in table 3. They indicate that these ratios are not very precisely estimated and that the differences in them between models, while appreciable, are small relative to the standard errors. The test of the hypothesis that the coefficients on the two earnings variables sum to zero, which would be the case if only relative earnings mattered, is rejected increasingly strongly (see table 2). The  $\chi^2(1)$  Wald statistic rises from 7.4 in the ordered probit model to 9.3 when  $K = 3$  is used and 15.8 when  $K = 5$  is used.

Table 3: Some derived statistics

	OP	SNP(3)	SNP(5)
	coef (s.e.)	coef (s.e.)	coef (s.e.)
log(earnings)	0.134 (.054)	0.096 (.050)	0.087 (.056)
log(comp. earn.)	-0.283 (.064)	-0.254 (.060)	-0.323 (.068)
Trade off	2.12 (0.70)	2.66 (1.15)	3.70 (2.04)
Age/10	-0.210 (.100)	-0.154 (.092)	-0.141 (.104)
Age <sup>2</sup> /100	0.038 (.013)	0.031 (.011)	0.030 (.013)
Age at minimum	27.6 (4.4)	24.8 (6.1)	23.1 (7.5)

Another coefficient ratio of interest is that involving the two coefficients of the quadratic in age. The minimum of the job satisfaction age profile is minus half the ratio of coefficients. It is estimated to be at about age 28 on the basis of the ordered probit model, but rather earlier with both the SNP models: age 25 if  $K = 3$  is used and age 23 if  $K = 5$  is used. These seem like important differences. However, the standard errors given in table 3 indicate that these ratios also are not very precisely determined. As a result, the differences across the models in the estimated age at the minimum of the age-satisfaction profile are small relative to the standard errors.

Figure 1: Estimated Density for  $K = 5$ 

Both SNP models exhibit positive skewness and indicate greater kurtosis than in the standard normal (table 2). Figure 1 gives a plot of the estimated density for the  $K = 5$  model. Both the positive skew and fatter tails than normal are evident. The central mode of the estimated density is at 0.30. Its height there is 0.423, compared with 0.399 for the standard normal density at its mode. The estimated density also has secondary modes in both tails. The larger of the two is at  $\varepsilon = 2.82$ , with density  $f_5(\varepsilon) = 0.061$ . There is also a smaller one at  $\varepsilon = -4.07$ ,  $f_5(\varepsilon) = 0.014$ .

## 6 Remarks on using sneop

1. The log likelihood of this model is not globally concave. It is therefore good practice to check for convergence to the global maximum rather than a local one by running the algorithm from different starting points using the `from` option. A natural alternative to the OP starting values is the SNP model for  $K - 1$ .
2. The program can be used to obtain SNP estimates of a binary response model. The binary response needs to be recoded to take the values 1 and 2.
3. It should be emphasized that this is a large sample technique. Small samples are unlikely to be able to detect divergencies from normality. Consequently, results on small samples can be expected to differ little from the `oprobit` estimates.
4. The desirable properties of the estimator, as well as requiring a large sample, also depend on  $K$  increasing with  $N$ . Experience with the estimator suggests that selecting  $K$  on the basis of either the criteria described above or likelihood-ratio tests does in fact do this.
5. Using `predict` with the `xb` option after `sneop` creates a new variable containing  $x'_i\hat{\beta}$ .
6. The program is much slower than `oprobit`, both because of the greater complexity of the likelihood function to be maximized and because it is written as an ado-file. The time required depends on the order of the polynomial used ( $K$ ), the number of observations ( $N$ ), and the number of categories ( $J$ ) (as well as the speed of the processor), but is little affected by the number of explanatory variables. Experiments with the program indicate a tight relationship for the time per iteration but only a rather loose one for the number of iterations required. As a guide, the combination  $N = 1000$ ,  $K = 3$ ,  $J = 7$  requires about 10 seconds per iteration on a 1.2 GHz Pentium III processor. This increases or decreases roughly proportional to  $N$  and roughly proportional to the squares of  $K$  and  $J$ , although in all three cases there are slight economies of scale. The number of iterations required is far more variable but seems roughly independent of  $N$ ,  $J$ , and the number of  $x$  variables. As a rough rule of thumb, convergence tends to require approximately  $5(K - 1)$  iterations.

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## 8 References

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