
Weak identification in probit models with endogenous covariates

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Abstract Weak identification is a well-known issue in the context of linear structural models. However, for probit models with endogenous explanatory variables, this problem has been little explored. In this paper, we study by simulation the behavior of the usual z-test and the LR test in the presence of weak identification. We find that the usual asymptotic z-test exhibits large level distortions (over-rejections under the null hypothesis). The magnitude of the level distortions depends heavily on the parameter value tested. In contrast, asymptotic LR tests do not over-reject and appear to be robust to weak identification.

Keywords probit model · weak identification

JEL Classification C35

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1 Introduction

Probit models are widely used in applied econometrics; for some recent examples, see Abramitzky and Lavy (2014), Beck et al (2014), Bijsterbosch and Dahlhaus (2015), Bouoiyour et al (2016), Cornelli et al (2013), Croushore and Marsten (2016), Engelhardt et al (2010), Esaka (2010), Fitzenberger et al (2011), Haider and Jahangir (2017), Hao and Ng (2011), Hlaing and Pourjalali (2012), Horvath and Katuscakova (2016), Khanna et al (2015), Litchfield et al (2012), Massa and Zhang (2013), Wen and Gordon (2014). As in linear models, one or more explanatory variables can be endogenous. This problem can be solved by using instrumental variables; see Wilde (2008) for a comparison of different estimation methods using instrumental variables. The resulting estimates can be used to calculate test statistics for the parameters of the model.

In linear models, it is well known that weak instruments can cause considerable level (or size) distortions [see Dufour (2003) for an overview]. Wald-type tests like the usual t-tests and F-tests are especially vulnerable to this problem [see Dufour (1997)]. In probit models, a single parameter hypothesis is usually tested by the so-called z-test, i.e. the ratio of a consistent estimate and its asymptotic standard error. This is a Wald-type test. Therefore, large level distortions can be expected. Nevertheless, the topic seems to be largely a white spot in the literature. Exceptions are the recent theoretical papers of Andrews and Cheng (2013, 2014), who address the probit model as an example. However, Andrews and Cheng (2013) restrict their numerical analysis to a probit model with a nonlinear regression function and without endogeneity, and Andrews and Cheng (2014) do not analyze the probit model numerically.¹

This paper makes several contributions to this problem. First, large level distortions in probit models with endogeneity are demonstrated by a simulation study. Second, we show that level distortions depend heavily on which parameter value is tested: whereas level distortions are moderate for the problem of testing the null of a zero parameter, testing other values of the parameter yield large level distortions. Third, the behaviour of the classical likelihood ratio statistic in this case is analyzed. For the simulation design considered, no level distortions are observed. However, the probability of type I error can be notably lower than the nominal level of the test (undersizing). Fourth, some new insights concerning the estimation of probit models with endogenous covariates are provided.

Section 2 describes the econometric model and the test statistics. Section 3 explains the simulation design and the estimators used. Since a probit equation is part of the model, some formulae become more complicated than in the linear case. They are described in detail because textbook descriptions are missing so far. Section 4 presents the results of the simulation study, and Section 5 concludes. For ease of exposition, we focus on the binary probit model.

2 Model and classical tests

We study a structural probit model, where one of the explanatory variables is endogenous, and a reduced-form equation for this variable is specified. The specific model considered is:

¹ A further exception is Magnusson (2007), who considered in an early version of his paper the probit model with endogenous covariates as an example and found medium level distortions. However, in later versions of the working paper and in the published version Magnusson (2010) the probit example was deleted.

$$\begin{aligned} y_{1i}^* &= \gamma_1 y_{2i} + \beta_1 x_{1i} + u_{1i}, \\ y_{2i} &= \pi_{21} x_{1i} + \pi_{22} x_{2i} + v_{2i}, \end{aligned} \quad y_{1i} = \begin{cases} 1, & \text{if } y_{1i}^* > 0 \\ 0, & \text{otherwise} \end{cases}, \quad i = 1, \dots, N, \quad (1)$$

where y_{1i}^* is a latent variable, y_{1i} is its observable indicator, y_{2i} is an endogenous (observable) variable, x_{1i} and x_{2i} are $K_1 \times 1$ and $K_2 \times 1$ vectors of exogenous variables, γ_1 , β_1 , π_{21} , π_{22} are unknown parameter vectors of dimensions 1 , $1 \times K_1$, $1 \times K_1$ and $1 \times K_2$ respectively, and u_{1i} and v_{2i} are error terms with mean zero, variances $\sigma_{u_1}^2$ and $\sigma_{v_2}^2$ respectively, and $\text{Cov}(u_{1i}, u_{1j}) = \text{Cov}(v_{2i}, v_{2j}) = \text{Cov}(u_{1i}, v_{2j}) = 0$ for $i \neq j$. The probit model assumes that the u_{1i} 's are normally distributed. Whether the distribution of v_{2i} must also be specified depends on the estimation method. If a distribution is needed, we assume that u_{1i} and v_{2i} follow a joint normal distribution, i.e.

$$(u_{1i}, v_{2i}) | x_{1i}, x_{2i} \stackrel{\text{iid}}{\sim} \text{N} \left[\begin{pmatrix} 0 \\ 0 \end{pmatrix}, \begin{pmatrix} \sigma_{u_1}^2 & \sigma_{u_1 v_2} \\ \sigma_{u_1 v_2} & \sigma_{v_1}^2 \end{pmatrix} \right].$$

The parameter of special interest is γ_1 . It is not identified when π_{22} is equal to zero. Therefore, we say γ_1 is weakly identified if π_{22} is close to zero. Sometimes weak identification is quantified by the so-called concentration parameter; see Stock et al (2002), p. 519. However, this parameter grows with N , and hence it suggests that the problem of weak identification is reduced by enlarging the sample size. This is misleading, and therefore we do not use it as guideline in our study.

Testing the significance of γ_1 in empirical studies is usually done by the z-statistic (implemented in almost all econometric software packages):

$$z = \frac{\hat{\gamma}_1}{\sqrt{\hat{V}(\hat{\gamma}_1)}} \quad (2)$$

where $\hat{V}(\hat{\gamma}_1)$ is a consistent estimate (assuming identification) of the asymptotic variance of $\sqrt{N}(\hat{\gamma}_1 - \gamma_1)$. If $\hat{\gamma}_1$ is consistent with a standard normal asymptotic distribution, z is asymptotically standard normal under the assumption of strong identification. The parameter γ_1 can be estimated by two-step methods [see Blundell and Smith (1993) for an overview] or via joint GMM or maximum likelihood (ML) estimation of both equations [see Wilde (2008) for a comparison].

The z-test is a Wald-type test. The classical alternatives to it — the Likelihood Ratio (LR) and the Lagrange Multiplier (LM) tests — are based on ML estimation of the parameters. In linear models, the latter are less affected by weak identification than Wald-type tests. Here, we focus on the LR test. Given the estimates, the LR statistic is calculated easily, whereas the LM test requires the estimation of a complicated information matrix, so the results may depend on the estimation procedure chosen. The log-likelihood function of model (1) under the above standard assumptions is:

$$\begin{aligned} \ln l(\theta) &= \sum_{i=1}^N \left[-0.5 \ln(2\pi\sigma_{v_2}^2) - 0.5 \left(\frac{y_{2i} - \pi_{2i} x_i}{\sigma_{v_2}} \right)^2 \right. \\ &\quad \left. + y_{1i} \ln \Phi \left(\frac{1}{\sqrt{1-\rho_v^2}} \left(\frac{(\gamma_1 \pi_{21} + \beta_1)x_{1i} + \gamma_1 \pi_{22} x_{2i}}{\sigma_{v_1}} + \rho_v \left(\frac{y_{2i} - \pi_{2i} x_i}{\sigma_{v_2}} \right) \right) \right) \right. \\ &\quad \left. + (1 - y_{1i}) \ln \left(1 - \Phi \left(\frac{1}{\sqrt{1-\rho_v^2}} \left(\frac{(\gamma_1 \pi_{21} + \beta_1)x_{1i} + \gamma_1 \pi_{22} x_{2i}}{\sigma_{v_1}} + \rho_v \left(\frac{y_{2i} - \pi_{2i} x_i}{\sigma_{v_2}} \right) \right) \right) \right) \right] \quad (3) \end{aligned}$$

where $x_i = (x'_{1i}, x'_{2i})'$, $v_{1i} = u_{1i} + \gamma_1 v_{2i}$, $\theta = (\gamma_1, \beta_1, \pi_2, \sigma_{v_2}, \rho_v)'$, $\pi_2 = (\pi_{21}, \pi_{22})$, $\sigma_{v_1} = \sqrt{\text{Var}(v_{1i})}$, and $\rho_v = \text{Corr}(v_{1i}, v_{2i})$; see Wilde (2008), appendix 2. Since the structural parameters enter the likelihood only through ratios with a standard deviation and the latter does not appear separately, only these ratios are identifiable. Therefore, in our simulation study, σ_{v_1} is taken as known.

We consider the problem of testing

$$H_0 : \gamma_1 = \tilde{\gamma}_1 \quad \text{vs.} \quad H_1 : \gamma_1 \neq \tilde{\gamma}_1.$$

We denote by $\hat{\theta}_{\text{ML}}$ the unrestricted ML estimator of θ [based on (3)] and by $\hat{\theta}_{\text{RML}}$ the restricted ML estimator under the null hypothesis. The LR statistic has the form

$$\text{LR} = 2 [\ln l(\hat{\theta}_{\text{ML}}) - \ln l(\hat{\theta}_{\text{RML}})].$$

Under the usual assumptions (including strong identification) LR is asymptotically $\chi^2(1)$ under H_0 .

3 Simulation design

In order to avoid arbitrary choices of unnecessary nuisance parameters, we consider a simple simulation design with $K_1 \in \{0, 1\}$ and $K_2 = 1$. The value $K_1 = 0$ defines a model without constants, whereas $K_1 = 1$ is used to add a constant in both equations. With $K_1 = 0$, ML estimation causes numerical problems, i.e. many replications end with the message ‘‘maximum iterations reached’’ even after 200 and more iterations. Since GMM estimation is a well-known alternative to ML estimation for probit-type models and the efficiency loss in models like ours seems to be small [see Wilde (2008)], we used GMM estimation for the z-test. As will be shown below, the results under weak identification are nearly the same for models with and without constants. Therefore, we focus our comparison of z-test and LR test for models including constants in both equations.

In (1), the second equation is a reduced-form equation. Endogeneity of y_2 can occur for at least two reasons: correlation between the error terms of the structural equations and/or simultaneity between y_1 and y_2 . Both yield a correlation between u_1 and v_2 . In our simulation, we focus on simultaneity because it can be interpreted more easily. Nevertheless, all results can be reproduced by assuming correlation between the error terms of the structural equations. Therefore, our main data generating model with constants is:

$$\begin{aligned} y_{1i}^* &= \gamma_1 y_{2i} + \beta_{11} + u_{1i}, \\ y_{2i} &= \gamma_2 y_{1i}^* + \beta_{21} + \beta_{22} x_i + u_{2i}, \end{aligned} \quad y_{1i} = \begin{cases} 1, & \text{if } y_{1i}^* > 0 \\ 0, & \text{otherwise} \end{cases}, \quad i = 1, \dots, N. \quad (4)$$

The residuals u_{1i} and u_{2i} are drawn independently from the $N(0, 16)$ distribution, so the residual variances are equal for both equations. The exogenous variable x_i is drawn from the $N(0.5, 16)$ distribution, so the expected number of ones for y_{1i} differs from the expected number of zeros also for the model without constants. Alternatively, we draw the residuals independently from the $N(0, 1)$ distribution and x_i from an $N(0.5, 1)$ distribution. The constants are $\beta_{11} = 0.5$ and $\beta_{21} = 0.25$. Weak identification is equivalent to β_{22} being close to zero. In our simulations, we took $\beta_{22} = 0.0001$. Smaller values of β_{22} yield qualitatively similar results. The case of strong identification is simulated with $\beta_{22} = 1$. The simulations were also run with $\beta_{22} = 0.1$ to see whether weak identification also causes problems

for moderate parameter values. The sample sizes are $N = 400$ (medium sample size) and $N = 2000$ (large sample), and the number of replications is 5000.

The estimated model in these simulations is:

$$\begin{aligned} y_{1i}^* &= \gamma_1 y_{2i} + \beta_{11} + u_{1i} \\ y_{2i} &= \pi_{21} + \pi_{22} x_i + v_{2i} \end{aligned} \quad y_{1i} = \begin{cases} 1, & y_{1i}^* > 0 \\ 0, & \text{else} \end{cases}, \quad i = 1, \dots, N, \quad (5)$$

where

$$\pi_{21} = (\gamma_2 \beta_{11} + \beta_{21}) / (1 - \gamma_1 \gamma_2), \quad \pi_{22} = \beta_{22} / (1 - \gamma_1 \gamma_2), \quad v_{21} = (\gamma_2 u_{1i} + u_{2i}) / (1 - \gamma_1 \gamma_2).$$

In (5), γ_1 is exactly identified as long as π_{22} is different from zero. (5) is estimated by GMM using the ‘natural’ moment conditions [see Wilde (2008)]:

$$E \begin{bmatrix} x_i \left(y_{1i} - \Phi \left(\frac{\gamma_1 (\pi_{21} + \pi_{22} x_i) + \beta_{11}}{\sigma_{v_1}} \right) \right) \\ x_i (y_{2i} - \pi_{21} - \pi_{22} x_i) \\ 1 \left(y_{1i} - \Phi \left(\frac{\gamma_1 (\pi_{21} + \pi_{22} x_i) + \beta_{11}}{\sigma_{v_1}} \right) \right) \\ 1 (y_{2i} - \pi_{21} - \pi_{22} x_i) \end{bmatrix} = 0.$$

Setting

$$\theta = \begin{pmatrix} \gamma_1 \\ \pi_{22} \\ \beta_{11} \\ \pi_{21} \end{pmatrix}, \quad m_i(\theta) = \begin{bmatrix} x_i \left(y_{1i} - \Phi \left(\frac{\gamma_1 (\pi_{21} + \pi_{22} x_i) + \beta_{11}}{\sigma_{v_1}} \right) \right) \\ x_i (y_{2i} - \pi_{21} - \pi_{22} x_i) \\ 1 \left(y_{1i} - \Phi \left(\frac{\gamma_1 (\pi_{21} + \pi_{22} x_i) + \beta_{11}}{\sigma_{v_1}} \right) \right) \\ 1 (y_{2i} - \pi_{21} - \pi_{22} x_i) \end{bmatrix}, \quad \bar{m}(\theta) = \frac{1}{N} \sum_{i=1}^N m_i(\theta),$$

we calculate

$$\hat{\theta} = \hat{\theta}_{\text{GMM}} = \underset{\theta}{\operatorname{argmin}} \{ \bar{m}(\theta)' W_N \bar{m}(\theta) \}$$

where W_N is a weighting matrix. Since the number of moment conditions is equal to the number of parameters, the weighting matrix in the criterion function of the GMM estimator does not matter theoretically, and the same asymptotic covariance matrix of the estimator can be used for all choices of W_N [Harris and Mátyás (1999), p. 22]. To be more precise, the asymptotic covariance matrix is [see Greene (2008), p. 445]:

$$\operatorname{asyVar}(\hat{\theta}) = \frac{1}{N} [G' \Psi^{-1} G]^{-1},$$

$$\Psi = \operatorname{asyVar}(\sqrt{N} \bar{m}), \quad \varphi_i := \varphi \left(\frac{\gamma_1 (\pi_{21} + \pi_{22} x_i) + \beta_{11}}{\sigma_{v_1}} \right),$$

$$G = \frac{\partial \bar{m}}{\partial \theta'} = \begin{pmatrix} \frac{1}{N} \sum_{i=1}^N -\frac{\pi_{21} x_i + \pi_{22} x_i^2}{\sigma_{v_1}} \varphi_i & \frac{1}{N} \sum_{i=1}^N -\frac{\gamma_1 x_i^2}{\sigma_{v_1}} \varphi_i & \frac{1}{N} \sum_{i=1}^N -\frac{x_i}{\sigma_{v_1}} \varphi_i & \frac{1}{N} \sum_{i=1}^N -\frac{\gamma_1 x_i}{\sigma_{v_1}} \varphi_i \\ 0 & \frac{1}{N} \sum_{i=1}^N -x_i^2 & 0 & \frac{1}{N} \sum_{i=1}^N -x_i \\ \frac{1}{N} \sum_{i=1}^N -\frac{\pi_{21} + \pi_{22} x_i}{\sigma_{v_1}} \varphi_i & \frac{1}{N} \sum_{i=1}^N -\frac{\gamma_1 x_i}{\sigma_{v_1}} \varphi_i & \frac{1}{N} \sum_{i=1}^N -\frac{1}{\sigma_{v_1}} \varphi_i & \frac{1}{N} \sum_{i=1}^N -\frac{\gamma_1}{\sigma_{v_1}} \varphi_i \\ 0 & \frac{1}{N} \sum_{i=1}^N -x_i & 0 & -1 \end{pmatrix}.$$

Given the above assumptions, this matrix can be estimated consistently by

$$\text{estasyVar}(\hat{\theta}) = \frac{1}{N} [\hat{G}'\hat{\Psi}^{-1}\hat{G}]^{-1},$$

$$\hat{\Psi} = \frac{1}{N} \sum_{i=1}^N m_i(\hat{\theta})m_i(\hat{\theta})', \quad \hat{G} = G \text{ after substituting } \hat{\theta} \text{ for } \theta.$$

The square root of the first diagonal element is the denominator of the z-statistic (2). $\hat{\Psi}^{-1}$ is the optimal weighting matrix and is used to calculate the nominator of (2).² In the model without constants only the first two moment conditions are used (setting π_{21} and β_{11} equal to zero), and G reduces to a 2×2 matrix.

For ML estimation of the model without constants, everything works fine under strong identification, whereas under weak identification the algorithm did not find the maximum for nearly half of the replications. This is not surprising under weak identification. For example, in the model without constants, the second summand of (3) reduces to

$$y_{1i} \ln \Phi \left(\frac{1}{\sqrt{1-\rho_v^2}} \left(\frac{\gamma_1 \pi_{22} x_i}{\sigma_{v_1}} + \rho_v \left(\frac{y_{2i} - \pi_{22} x_i}{\sigma_{v_2}} \right) \right) \right)$$

where the parameter γ_1 affects the log-likelihood function only through the product $\gamma_1 \pi_{22}$. However, under weak identification, π_{22} is close to zero, so that it is very difficult to find the ‘true’ value of γ_1 : the log-likelihood function is relatively flat with respect to γ_1 . However, including constants avoids the numerical problems discussed above without changing the results concerning weak identification. The latter aspect was confirmed for those parameter values for which the optimum was also found in the model without constants.³

4 Simulation results

We distinguish two cases. First, we test the null hypothesis $\gamma_1 = 0$, i.e. the significance of y_2 . Second, we test the null hypothesis $\gamma_1 = c$, where c is a constant different from zero, and we present the results for $c = 2$. In both cases, the simulations consider different values of γ_2 . Small values of γ_2 correspond to a ‘small’ simultaneity problem, here $\gamma_2 = \pm 0.5$. A ‘medium’ problem of simultaneity is represented by $\gamma_2 = \pm 1.5$, and a ‘large’ problem of simultaneity is represented by $\gamma_2 = \pm 3$ and $\gamma_2 = \pm 6$.

4.1 Results for the z-test

Main model

The results for the models with and without constants are similar. Therefore, we focus on the model with constants. The results for the model without constants can be found in the appendix. In case of strong identification and testing $\gamma_1 = 0$, no level problems can be observed at the nominal levels 10%, 5%, and (to a lesser degree) 1%; see Table 1. Under weak

² All simulations were done using R; see R Core Team (2016). The GMM estimation was done using the package GMM, version 1.6-1. The case of iid observations can be implemented by the option `vcov="iid"`; see Chaussé (2010), p. 13. All R codes are available on request.

³ For the ML estimation the exogenous variable x_i was drawn from an $N(0, 16)$ distribution.

identification, the picture is mixed.⁴ The key result is highlighted in Figure 1 for $N = 2000$ and a nominal level of 5%. If there is only ‘weak’ simultaneity, an extreme undersizing phenomenon is observed. However, ‘strong’ simultaneity causes medium level distortions. Nevertheless, the level distortions are smaller than those in linear simultaneous equations. This is surprising because weak identification should cause similar problems in probit models. Furthermore, the results for $\beta_{22} = 0.1$ are close to the results for $\beta_{22} = 0.0001$, i.e. oversizing occurs even for moderate values of the parameter β_{22} . All results are similar for $N = 2000$ and $N = 400$.

Next, we consider the results concerning data simulated with $\gamma_1 = 2$ and testing $H_0 : \gamma_1 = 2$. Again, under strong identification, no level distortion appears at the nominal levels 10% and 5%; see Table 2. However, with weak identification and strong simultaneity, the level distortions become very large ($\gamma_2 = 0.5$ is omitted because γ_1 is no longer identified in this case); see Figure 2. The empirical level becomes more than tenfold as high as the nominal level. Thus, level distortions comparable to those in linear simultaneous equations models can be observed in probit models. With strong simultaneity, level distortions also occur for $\beta_{22} = 0.1$. They are stronger for $N = 400$, i.e. the sample size makes a notable difference in this case.

The differences between testing $\gamma_1 = 0$ and $\gamma_1 = 2$ demonstrate an important feature: in probit models, level distortions depend heavily on the parameter value tested. This property is not easily explained by the concentration parameter. Following the definition of Stock et al (2002), p. 519, the concentration parameter μ^2 in the model without constants is:

$$\mu^2 = \frac{\sum_{i=1}^N x_i^2 \pi_{22}^2}{\sigma_{v_2}^2} = \frac{\sum_{i=1}^N x_i^2 \left(\frac{\beta_{22}}{1 - \gamma_1 \gamma_2} \right)^2}{\frac{\sigma_{u_2}^2 + \gamma_2^2 \sigma_{u_1}^2}{(1 - \gamma_1 \gamma_2)^2}} = \frac{\sum_{i=1}^N x_i^2 \beta_{22}^2}{\sigma_{u_2}^2 + \gamma_2^2 \sigma_{u_1}^2}.$$

The concentration parameter does not depend on γ_1 . In the model with constants the formula is more bulky, but the message remains unchanged. Thus, concerning the concentration parameter the problem of weak identification should not depend on the value of γ_1 .

Sensitivity analysis

We performed several robustness checks on our results. First, we vary the estimation method, by iterating our two-step GMM estimator, and using I instead of Ψ^{-1} as weighting matrix.⁵ Second, we change the variances, i.e. we draw the u_i 's from the $N(0, 1)$ distribution and x_i from an $N(0.5, 1)$ distribution. Third, we change from exact identification of γ_1 to underidentification ($\beta_{22} = 0$) and overidentification (a second exogenous variable is added to the second equation). Except for overidentification, the different variants do not change the results. Thus, we present only examples of these variants.

Changing the estimation method does not affect the results, i.e. the rejection frequencies are the same as in Tables 1 and 2. Changing the variances in the sample design slightly modifies the results, but the pattern remains the same: we observe only moderate level distortions when testing $\gamma_1 = 0$ and large level distortions when testing $\gamma_1 = 2$ are observed (see Table 3). The results in the case of underidentification are very close to those of $\beta_2 = 0.0001$ (see Table 4).

⁴ This result is similar to that of Magnusson (2007).

⁵ We also tried the option "CUE" for the continuous updating estimator. However, we always got the error message "node stack overflow".

In contrast with the above results, overidentification leads to new insights. In the second equations of (4) and (5), there is an additional exogenous variable drawn from an $N(0, 16)$ distribution. The corresponding structural parameter β_{23} varies like β_{22} : for strong identification, we consider $\beta_{22} = \beta_{23} = 1$, and for weak identification $\beta_{22} = \beta_{23} = 0.0001$ (very weak identification) and $\beta_{22} = \beta_{23} = 0.1$. The second exogenous variable is used as an instrument in the same way as x_i and the ones in the previous simulations. The matrix G now becomes a (6×5) -matrix, i.e. the number of moment conditions exceeds the number of parameters. Thus, the choice of the weighting matrix should matter. Therefore, we compare the results using the identity matrix I with those based on the ‘optimal’ weighting matrix. We focus on a nominal level of $\alpha = 5\%$.

First, the sample size is important. Under strong (over-)identification, everything works fine with $N = 2000$ (see Table 5), but $N = 400$ is too small to meet the nominal level. For testing $\gamma_1 = 0$, the rejection frequencies are in the interval $[0.357, 0.3754]$; for testing $\gamma_1 = 2$, the rejection frequencies are in the interval $[0.2846, 0.3942]$. This is an interesting side result and a cautionary note if models like ours are used with macroeconomic or experimental data, where sample sizes can easily be small.

Second, testing $\gamma_1 = 0$ with $N = 2000$ leads to larger level distortions with weak identification and large simultaneity. Now, empirical rejection frequencies reach four times the nominal level (see Table 5). The results are slightly stronger if I is used as weighting matrix. With weak identification and weak simultaneity, *undersizing* is still observable, but it is now less pronounced. With strong simultaneity, the results for $\beta_{22} = \beta_{23} = 0.1$ are again close to those under weak identification, whereas with weak simultaneity, the results with $\beta_{22} = \beta_{23} = 0.1$ are close to those under strong identification.

Third, when testing $\gamma_1 = 2$ with $N = 2000$, the picture is mixed. Using the weighting matrix I strengthens the results in case of strong simultaneity in comparison with the results for model (5). The results are now close to the theoretical expectation. However, using the optimal weighting matrix yields smaller level distortions than under exact identification. The interpretation of this result is difficult: for some parameter combinations, the rejection frequencies for $\beta_{22} = \beta_{23} = 0.0001$ are even lower than for $\beta_{22} = \beta_{23} = 0.1$. This may be due to the fact that the asymptotic standard errors are a complicated function of γ_1 . If the identity matrix I is used as the weighting matrix, only $G'G$ must be inverted for computing the asymptotic variance-covariance matrix [see Greene (2008), p. 451]. In contrast, using the optimal weighting matrix requires the inversion of $G'\Psi^{-1}G$. Under weak identification, the latter calculation can lead to ‘bad’ results. However, further research is needed to clarify the reasons for this. Nevertheless, under strong and even medium simultaneity, noteworthy level distortions are observed.

4.2 Results for the LR test on the main model

We consider again the main model (5). We use the same parameter values as in Section 4.1, and we calculate the Maximum Likelihood estimator and the LR statistic. In case of strong identification, no level problems can be observed (see Table 6). However, the results under weak identification differ substantially from those for the z-test. If simultaneity is weak, the observed rejection frequencies are close to the nominal level; if simultaneity is medium or strong, *undersizing* is observed. Thus, the LR test may be a conservative alternative to the z-test.

For testing $\gamma_1 = 0$, the results with $\gamma_2 = 6$ are missing, because the program stopped with an error message for some replications. This can be explained as follows. Consider

for instance $\gamma_1 = 0$ and $\gamma_2 = 6$. This implies $\rho_v = 0.9864$, i.e. the bivariate normal distribution of v_1 and v_2 is close to being singular. In the log-likelihood function (3), we then have $1/\sqrt{1 - \rho_v^2} = 6.08$, $\Phi(\approx 6) = 1$, $1 - \Phi(\approx 6) = 0$, and $\ln(1 - \Phi(\approx 6))$ is not defined. Therefore, ML estimation is less robust against a high correlation of the reduced-form errors than GMM estimation. This is similar to the findings in Wilde (2008), p. 476, and an interesting side finding of this paper.

In contrast with the z-test, the results for the LR test under weak identification do *not* change if $\gamma_1 = 2$ is tested. Furthermore, $\gamma_2 = 6$ does not raise numerical problems. With $\beta_{22} = 0.1$ and $N = 2000$, only strong simultaneity ($\gamma_2 = \pm 6$) leads to undersizing (for $N = 2000$), i.e. the results are more stable for moderate values of β_{22} than those for the z-test.

5 Conclusion

The paper analyses weak identification in probit models with endogenous covariates. It shows remarkable level distortions concerning the usual z-test. However, further research is needed to clarify why the magnitude depends heavily on the parameter value tested. The likelihood ratio statistic seems to be a conservative alternative which is robust to weak identification. Further research is useful to clarify how advanced methods like those of Andrews and Cheng (2014), Dufour (2006) or Kleibergen (2005) will work for probit models with endogenous covariates.

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Table 1 Rejection frequencies of the z-test, $H_0 : \gamma_1 = 0$

γ_2	N	Nominal level								
		$\beta_{22} = 1$			$\beta_{22} = 0.1$			$\beta_{22} = 0.0001$		
		10%	5%	1%	10%	5%	1%	10%	5%	1%
-6	2000	0.0908	0.0484	0.0178	0.1634	0.1142	0.0512	0.1690	0.1114	0.0464
	400	0.0984	0.0688	0.0320	0.1764	0.1206	0.0500	0.1700	0.1196	0.0514
-3	2000	0.0950	0.0502	0.0118	0.1310	0.0880	0.0414	0.1424	0.0916	0.0350
	400	0.0868	0.0508	0.0206	0.1528	0.1018	0.0394	0.1452	0.0938	0.0382
-1.5	2000	0.0988	0.0506	0.0114	0.0942	0.0626	0.0242	0.0854	0.0508	0.0124
	400	0.0964	0.0482	0.0136	0.1044	0.0640	0.0196	0.0896	0.0528	0.0144
-0.5	2000	0.0984	0.0500	0.0102	0.0700	0.0340	0.0058	0.0120	0.0030	0.0000
	400	0.0996	0.0510	0.0116	0.0432	0.0202	0.0012	0.0148	0.0056	0.0006
0.5	2000	0.0986	0.0470	0.0110	0.0698	0.0320	0.0080	0.0128	0.0046	0.0004
	400	0.0974	0.0500	0.0102	0.0382	0.0154	0.0016	0.0126	0.0038	0.0006
1.5	2000	0.0970	0.0486	0.0100	0.0914	0.0626	0.0254	0.0836	0.0458	0.0136
	400	0.0966	0.0480	0.0102	0.0960	0.0566	0.0178	0.0896	0.0480	0.0128
3	2000	0.0960	0.0468	0.0108	0.1212	0.0866	0.0388	0.1430	0.0922	0.0342
	400	0.0864	0.0464	0.0160	0.1408	0.0914	0.0332	0.1488	0.0976	0.0340
6	2000	0.0886	0.0462	0.0160	0.1548	0.1076	0.0484	0.1674	0.1138	0.0458
	400	0.0930	0.0658	0.0276	0.1696	0.1130	0.0454	0.1768	0.1196	0.0482

Table 2 Rejection frequencies of the z-test, $H_0 : \gamma_1 = 2$

γ_2	N	Nominal level								
		$\beta_{22} = 1$			$\beta_{22} = 0.1$			$\beta_{22} = 0.0001$		
		10%	5%	1%	10%	5%	1%	10%	5%	1%
-6	2000	0.0890	0.0556	0.0234	0.2724	0.2280	0.1696	0.5586	0.5142	0.4366
	400	0.1038	0.0726	0.0382	0.4480	0.4078	0.3322	0.5732	0.5260	0.4540
-3	2000	0.0996	0.0518	0.0144	0.1586	0.1244	0.0810	0.3686	0.3100	0.2076
	400	0.0842	0.0480	0.0200	0.2772	0.2314	0.1634	0.3756	0.3150	0.2180
-1.5	2000	0.0970	0.0498	0.0108	0.1042	0.0756	0.0320	0.1562	0.0996	0.0404
	400	0.0932	0.0490	0.0116	0.1460	0.1002	0.0434	0.1632	0.1040	0.0432
-0.5	2000	0.0990	0.0468	0.0108	0.0684	0.0316	0.0056	0.0110	0.0026	0.0000
	400	0.0928	0.0490	0.0086	0.0418	0.0150	0.0012	0.0126	0.0032	0.0002
1.5	2000	0.0996	0.0494	0.0102	0.0864	0.0540	0.0218	0.0694	0.0358	0.0078
	400	0.0914	0.0424	0.0084	0.0856	0.0522	0.0166	0.0700	0.0400	0.0084
3	2000	0.0946	0.0472	0.0116	0.1576	0.1182	0.0708	0.3184	0.2530	0.1554
	400	0.0774	0.0426	0.0146	0.2496	0.1968	0.1246	0.3246	0.2608	0.1620
6	2000	0.0872	0.0482	0.0198	0.2640	0.2260	0.1674	0.5416	0.4966	0.4124
	400	0.1036	0.0696	0.0334	0.4446	0.4004	0.3256	0.5592	0.5152	0.4304

Table 3 Rejection frequencies of the z-test, sample design with variances =1

γ_1	γ_2	N	Nominal level								
			$\beta_{22} = 1$			$\beta_{22} = 0.1$			$\beta_{22} = 0.0001$		
			10%	5%	1%	10%	5%	1%	10%	5%	1%
0	6	2000	0.0908	0.0474	0.0162	0.1278	0.0834	0.0352	0.1424	0.0898	0.0330
		400	0.0954	0.0618	0.0252	0.1358	0.0852	0.0340	0.1480	0.0946	0.0332
2	6	2000	0.0808	0.0510	0.0176	0.3126	0.2448	0.1708	0.5718	0.5084	0.4078
		400	0.1054	0.0724	0.0352	0.4376	0.3902	0.3110	0.5464	0.5034	0.4194

Table 4 Rejection frequencies of the z-test in case of underidentification, i.e. $\beta_{22} = 0$

γ_2	N	Nominal level		
		10%	5%	1%
6	2000	0.1674	0.1138	0.0458
	400	0.1770	0.1194	0.0482
6	2000	0.5424	0.4964	0.4116
	400	0.5592	0.5150	0.4300

Table 5 Rejection frequencies of the z-test, nominal level 5%, $N = 2000$, with overidentification of γ_1 in the theoretical model

γ_1	γ_2	Weighing matrix					
		$\beta_{22} = \beta_{23} = 1$		$\beta_{22} = \beta_{23} = 0.1$		$\beta_{22} = \beta_{23} = 0.0001$	
		I	Optimal	I	Optimal	I	Optimal
0	-6	0.0526	0.0450	0.2396	0.1724	0.2682	0.2072
0	-3	0.0484	0.0464	0.1578	0.1100	0.2304	0.1758
0	-1.5	0.0466	0.0464	0.0818	0.0686	0.1350	0.1174
0	-0.5	0.0466	0.0464	0.0456	0.0488	0.0112	0.0122
0	0.5	0.0470	0.0456	0.0458	0.0456	0.0124	0.0150
0	1.5	0.0474	0.0460	0.0826	0.0710	0.1310	0.1232
0	3	0.0492	0.0464	0.1482	0.1062	0.2252	0.1904
0	6	0.0548	0.0496	0.2256	0.1692	0.2686	0.2120
2	-6	0.0376	0.0454	0.3580	0.1574	0.7844	0.0990
2	-3	0.0452	0.0486	0.1952	0.1130	0.5708	0.1126
2	-1.5	0.0470	0.0456	0.0888	0.0704	0.2416	0.1156
2	-0.5	0.0442	0.0458	0.0440	0.0446	0.0108	0.0154
2	1.5	0.0538	0.0548	0.0714	0.0598	0.1100	0.1104
2	3	0.0460	0.0504	0.2006	0.1106	0.4968	0.2102
2	6	0.0470	0.0542	0.3882	0.1606	0.7736	0.1344

Table 6 Rejection frequencies of the LR test

γ_1	γ_2	N	Nominal level								
			$\beta_{22} = 1$			$\beta_{22} = 0.1$			$\beta_{22} = 0.0001$		
			10%	5%	1%	10%	5%	1%	10%	5%	1%
0	-3	2000	0.1010	0.0494	0.0100	0.0960	0.0464	0.0082	0.0654	0.0283	0.0048
		400	0.0992	0.0518	0.0128	0.0798	0.0388	0.0090	0.0698	0.0324	0.0066
0	-1.5	2000	0.1000	0.0492	0.0106	0.0998	0.0490	0.0104	0.0828	0.0366	0.0062
		400	0.1000	0.0514	0.0118	0.0946	0.0476	0.0108	0.0834	0.0408	0.0072
0	-0.5	2000	0.0986	0.0488	0.0120	0.0986	0.0488	0.0120	0.0958	0.0478	0.0108
		400	0.0996	0.0522	0.0112	0.0996	0.0520	0.0110	0.0956	0.0510	0.0110
0	0.5	2000	0.1000	0.0488	0.0112	0.1000	0.0488	0.0112	0.0966	0.0484	0.0098
		400	0.0996	0.0534	0.0116	0.0992	0.0530	0.0114	0.0962	0.0518	0.0110
0	1.5	2000	0.0988	0.0538	0.0110	0.0986	0.0536	0.0106	0.0810	0.0410	0.0080
		400	0.0990	0.0510	0.0106	0.0948	0.0434	0.0086	0.0806	0.0414	0.0070
0	3	2000	0.0998	0.0548	0.0106	0.0956	0.0488	0.0076	0.0664	0.0340	0.0054
		400	0.0962	0.0496	0.0102	0.0732	0.0324	0.0042	0.0676	0.0298	0.0050
2	-6	2000	0.1014	0.0536	0.0100	0.0648	0.0312	0.0066	0.0510	0.0240	0.0042
		400	0.0956	0.0514	0.0114	0.0482	0.0244	0.0058	0.0456	0.0238	0.0044
2	-3	2000	0.0974	0.0520	0.0092	0.0942	0.0482	0.0092	0.0554	0.0240	0.0046
		400	0.0990	0.0504	0.0106	0.0620	0.0300	0.0068	0.0498	0.0248	0.0050
2	-1.5	2000	0.0940	0.0466	0.0096	0.0978	0.0514	0.0114	0.0740	0.0350	0.0056
		400	0.0972	0.0524	0.0078	0.0826	0.0416	0.0070	0.0682	0.0328	0.0066
2	0.5	2000	0.0974	0.0478	0.0104	0.0968	0.0478	0.0124	0.0936	0.0472	0.0102
		400	0.1010	0.0462	0.0078	0.0954	0.0486	0.0086	0.0950	0.0456	0.0092
2	1.5	2000	0.0990	0.0514	0.0100	0.0994	0.0522	0.0096	0.0808	0.0404	0.0084
		400	0.0922	0.0496	0.0092	0.0914	0.0474	0.0062	0.0814	0.0380	0.0072
2	3	2000	0.0996	0.0492	0.0104	0.0926	0.0454	0.0062	0.0588	0.0284	0.0056
		400	0.0940	0.0476	0.0074	0.0708	0.0302	0.0020	0.0538	0.0258	0.0046
2	6	2000	0.0994	0.0492	0.0106	0.0678	0.0276	0.0046	0.0492	0.0268	0.0062
		400	0.0894	0.0462	0.0084	0.0492	0.0218	0.0036	0.0462	0.0238	0.0052

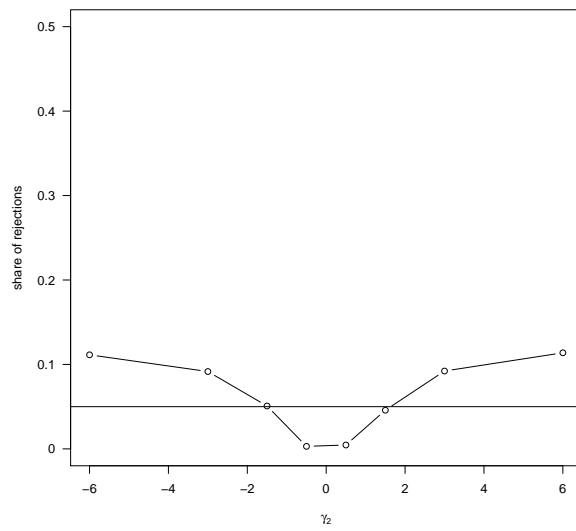


Fig. 1 Rejection frequencies of the z-test under weak identification, $H_0 : \gamma_1 = 0$, nominal level 5%, $N = 2000$

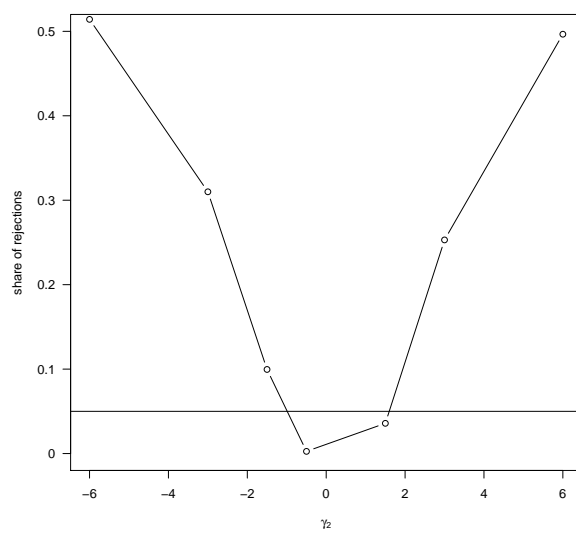


Fig. 2 Rejection frequencies of the z-test under weak identification, $H_0 : \gamma_1 = 2$, nominal level 5%, $N = 2000$

Appendix: Simulation results for model (5) with $\beta_{11} = \pi_{21} = 0$

Table 7 Rejection frequencies of the z-test, $H_0 : \gamma_1 = 0$

γ_2	N	Nominal level								
		$\beta_{22} = 1$			$\beta_{22} = 0.1$			$\beta_{22} = 0.0001$		
		10%	5%	1%	10%	5%	1%	10%	5%	1%
-6	2000	0.0888	0.0482	0.0178	0.1586	0.1118	0.0516	0.1724	0.1172	0.0474
	400	0.0982	0.0668	0.0308	0.1774	0.1188	0.0544	0.1776	0.1232	0.0498
-3	2000	0.0982	0.0470	0.0120	0.1248	0.0894	0.0408	0.1508	0.0942	0.0350
	400	0.0866	0.0478	0.0190	0.1516	0.1012	0.0396	0.1534	0.1010	0.0350
-1.5	2000	0.0998	0.0484	0.0090	0.0930	0.0636	0.0242	0.0892	0.0490	0.0140
	400	0.0960	0.0468	0.0126	0.1042	0.0608	0.0214	0.0940	0.0528	0.0132
-0.5	2000	0.1002	0.0502	0.0098	0.0714	0.0340	0.0078	0.0128	0.0030	0.0000
	400	0.0958	0.0470	0.0110	0.0456	0.0188	0.0018	0.0144	0.0050	0.0002
0.5	2000	0.1018	0.0510	0.0074	0.0712	0.0330	0.0096	0.0116	0.0042	0.0002
	400	0.0948	0.0478	0.0096	0.0398	0.0172	0.0026	0.0132	0.0046	0.0004
1.5	2000	0.0992	0.0492	0.0088	0.0956	0.0618	0.0242	0.0860	0.0488	0.0132
	400	0.0916	0.0482	0.0100	0.0972	0.0602	0.0194	0.0904	0.0508	0.0124
3	2000	0.0964	0.0492	0.0094	0.1264	0.0888	0.0392	0.1494	0.0942	0.0342
	400	0.0862	0.0464	0.0152	0.1438	0.0956	0.0370	0.1556	0.1004	0.0360
6	2000	0.0902	0.0482	0.0136	0.1596	0.1108	0.0504	0.1736	0.1192	0.0486
	400	0.0944	0.0626	0.0280	0.1706	0.1166	0.0486	0.1800	0.1236	0.0482

Table 8 Rejection frequencies of the z-test, $H_0 : \gamma_1 = 2$

γ_2	N	Nominal level								
		$\beta_{22} = 1$			$\beta_{22} = 0.1$			$\beta_{22} = 0.0001$		
		10%	5%	1%	10%	5%	1%	10%	5%	1%
-6	2000	0.0874	0.0542	0.0216	0.2668	0.2282	0.1668	0.5634	0.5166	0.4388
	400	0.1036	0.0702	0.0406	0.4466	0.4058	0.3358	0.5700	0.5276	0.4510
-3	2000	0.0968	0.0486	0.0128	0.1588	0.1248	0.0802	0.3724	0.3128	0.2164
	400	0.0826	0.0520	0.0198	0.2818	0.2362	0.1602	0.3756	0.3162	0.2214
-1.5	2000	0.0922	0.0462	0.0104	0.1058	0.0726	0.0330	0.1584	0.1020	0.0408
	400	0.0976	0.0500	0.0110	0.1422	0.0946	0.0416	0.1628	0.1046	0.0408
-0.5	2000	0.0938	0.0464	0.0100	0.0676	0.0318	0.0062	0.0124	0.0034	0.0000
	400	0.1004	0.0486	0.0078	0.0412	0.0148	0.0016	0.0142	0.0044	0.0002
1.5	2000	0.0994	0.0514	0.0102	0.0904	0.0580	0.0230	0.0712	0.0368	0.0078
	400	0.0956	0.0412	0.0096	0.0840	0.0510	0.0174	0.0736	0.0406	0.0092
3	2000	0.0986	0.0510	0.0106	0.1558	0.1214	0.0758	0.3184	0.2576	0.1604
	400	0.0830	0.0436	0.0154	0.2488	0.1966	0.1208	0.3266	0.2670	0.1630
6	2000	0.0922	0.0510	0.0188	0.2648	0.2252	0.1654	0.5490	0.5020	0.4210
	400	0.1044	0.0702	0.0324	0.4402	0.3968	0.3240	0.5604	0.5144	0.4304