

Fixed effect threshold model for unbalanced panel data

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- Threshold model has been widely used in economics and finance, monetary policy, economic growth etc.
- Hansen (1999) proposed the threshold model for panel data. The model is only used for balanced panel.
- For unbalanced panel, the user need to transform it to balanced one.
 - ① potential sample selection bias.
 - ② subjectivity: longer period and less individual, or shorter period and more individual.



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- single threshold model

$$y_{it} = \mu + X_{it}(q_{it} < \gamma)\beta_1 + X_{it}(q_{it} \geq \gamma)\beta_2 + u_i + e_{it}. \quad (1)$$

with $i = 1, 2, \dots, G$, $t = 1, 2, \dots, T_i$. The independent variable z is regime independent and x is regime dependent.

- written as

$$y_{it} = \mu + X_{it}(q_{it}, \gamma)\beta + u_i + e_{it}. \quad (2)$$



- grid search to minimize $S_1(\gamma)$: $[\underline{\gamma}, \overline{\gamma}]$.

$$\hat{\gamma} = \underset{\gamma}{\operatorname{argmin}} S_1(\gamma) \quad (3)$$

- As proved by Hansen (1999), $\hat{\gamma}$ is consistent, and has non-standard distribution.



- threshold effect test:

$$H_0 : \beta_1 = \beta_2; \quad H_a : \beta_1 \neq \beta_2$$

- F -statistic

$$F_0 = (S_0 - S_1) / \hat{\sigma}_1^2.$$



- bootstrap p-value:
 - (1) estimate linear model, get \hat{e}_{it} ; estimate single threshold model, get F_0 .
 - (2) cluster resampling of \hat{e}_{it} , get bootstrap residual v_{it}^* .
 - (3) bootstrap dependent variable

$$y_{it}^* = X_{it}\hat{\beta}.$$

- (4) use y^* , X , and q to compute F -statistic.
- (5) (2) - (4) B times, get F -statistic (F_1, F_2, \dots, F_B) .

$$Prob = P(F_j > F_0).$$



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- DGP (data generating process)

$$y_{it} = \mu + \alpha z_{it} + \beta_1 x_{it} 1(q_{it} < \tau) + \beta_2 x_{it} 1(q_{it} \geq \tau) + u_i + e_{it}, \quad (4)$$

- $z_{it}, x_{it} \sim \chi^2(1) - 1$,
individual effect: $u_i \sim \chi^2(1) - 1$,
the idiosyncratic error $e_{it} \sim N(0, 1)$.
- parameter: $\mu = 1, \alpha = 1, \beta_1 = 1, \beta_2 = 2, \tau = 1$.

- allow cluster sizes to vary systematically, we allocate N observations among G clusters using the formula

$$N_i = \left\lfloor \frac{N \exp(\gamma i/G)}{\sum_{j=1}^G \exp(\gamma j/G)} \right\rfloor, \quad i = 1, 2, \dots, G - 1. \quad (5)$$

- $\lfloor \cdot \rfloor$ denotes the integer part of the argument, and $N_G = N - \sum_{j=1}^{G-1} N_j$. When $\gamma = 0$ and N/G is an integer, the panel is balanced and $N_i = N/G$ for all i . As γ increases, cluster size becomes more unequal.
For example, when $G = 50$ and $N = 500$, if $\gamma = 1$, the largest cluster size is 24, and the smallest cluster size is 6.
If $\gamma = 2$, the largest cluster size is 24, and the smallest cluster size is 3.

Table: Simulation result for balanced and unbalanced panel

G	\bar{T}	Obs	$\gamma = 0$		$\gamma = 1$		$\gamma = 2$	
			RMSE(10^3)	Rate	RMSE(10^3)	Rate	RMSE(10^3)	Rate
50	5	250	8.3	0.537	34.4	0.481	24.3	0.463
50	10	500	1.7	0.633	2.9	0.579	3.0	0.574
50	20	1,000	0.4	0.815	0.8	0.763	0.8	0.755
200	5	1,000	0.5	0.775	0.9	0.734	1.2	0.709
200	10	2,000	0.1	0.921	0.2	0.890	0.2	0.864
200	20	4,000	0.05	0.977	0.077	0.962	0.071	0.958
500	5	2,500	0.104	0.935	0.195	0.888	0.196	0.889
500	10	5,000	0.041	0.994	0.049	0.981	0.059	0.967
500	20	10,000	0.026	1.00	0.032	0.995	0.031	0.996

- Given the number of observations and the number of clusters, the more unbalanced the panel is, the higher the RMSE is.
- The RMSE decreases with a larger number of groups or a larger number of observations. This is true for both balanced and unbalanced panels. That implies the estimator is consistent in unbalanced panel data, so is the regression coefficient.
- The coverage rate increases with sample size. But the coverage rate gets higher than the nominal level when the sample size gets larger after some point. Too small sample tends to get a tighter interval and too large sample tends to get a wider interval.
- The RMSE increases and coverage rate decreases as the panel data become more unbalanced.



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- DGP:

$$y_{it} = 1 + z_{it} + x_{it} + u_i + e_{it}. \quad (6)$$

where $z_{it}, x_{it} \sim \chi^2(1) - 1$. $u_i \sim \chi^2(1) - 1$, $e_{it} \sim N(0, 1)$.

- (1) Draw a random sample. First estimate a fixed effect linear model under the null hypothesis, and get the estimate $\hat{\beta}$, the fitted value

$$\hat{y}_{it} = \hat{\mu} + x_{it}\hat{\beta} + \hat{u}_i,$$

the residual $\hat{e}_{it} = y_{it} - \hat{y}_{it}$, and the sum of squared residuals S_0 .

- (2) Estimate the fixed effect single threshold model under the alternative hypothesis, and get the sum of squared residuals S_1 . The F -statistic is computed as

$$F = \frac{S_0 - S_1}{\hat{\sigma}_1^2}. \quad (7)$$

- (3) Repeat step (1) - (2) S times, and get a series of probability values (P_1, P_2, \dots, P_S) . The test size is

$$Pr = \frac{\sum_{i=1}^S 1(P_i < \alpha)}{S}, \quad \alpha = 0.1, 0.05, 0.01. \quad (8)$$



- The probability of F -statistic: wild cluster bootstrap method.
(a) draw a random wild weight v_i^b from Rademacher distribution

$$v_i^b = \begin{cases} 1, & \text{with prob. } 1/2 \\ -1, & \text{with prob. } 1/2 \end{cases}.$$

The bootstrap error is

$$e_i^b = \hat{e}_i v_i^b$$

and the bootstrap dependent variable is

$$y_i^b = \hat{y}_i + \hat{e}_i v_i^b. \quad (9)$$

- (b) Repeat step (a) R times, get a series of F -statistic $F_j (j = 1, 2, \dots, R)$.

$$P = \frac{\sum_{i=1}^R 1(F_i > F)}{R}, \quad (10)$$



- We can also use $\tilde{y}_{it} = x_{it}\hat{\beta}$ in the bootstrap.
- The constant and individual effect will be dropped off in the within-deviation transformation.

- v_i^b can also be drawn from other distributions with mean 0 and variance 1. Some options for v_i^b include:

(a) Mammen (1993) two-point distribution

$$v_i^b = \begin{cases} 1 - \phi, & \text{with prob. } \phi/\sqrt{5} \\ \phi, & \text{with prob. } (1 + \sqrt{5})/2 \end{cases}$$

The golden ratio is $\phi = (1 + \sqrt{5})/2$.

(b) Webb (2014) six-point distribution which assigns probability $1/6$ to each of 6 points, namely, $\pm\sqrt{1/2}$, ± 1 , and $\pm\sqrt{3/2}$.

Rademacher and Mammen distribution can yield only 2^G distinct bootstrap samples. Webb distribution reduces, but not eliminate this problem.

(c) Standard normal distribution.

(d) Gamma distribution with shape parameter 4 and scale parameter $1/2$ as suggested by Liu(1988).



- As described in Mackinnon (2018), simulation studies suggest that wild bootstrap tests based on the Rademacher distribution perform better than ones based on other auxiliary distributions; see, among others, Davidson, Monticini, and Peel (2007); Davidson and Flachaire (2008); Finlay and Magnusson (2016). However, the Webb six-point distribution is preferred to the Rademacher when G is less than 10 or perhaps 12.



- which residual to use: restricted model (null hypothesis) or unrestricted model (alternative)?
- It is generally better use the wild cluster bootstrap under restrictions, MacKinnon et al.(2018).
- Intuitively, since inference involves estimating the probabilities of obtaining certain results under the assumption that the null is true, inference is improved by using bootstrap datasets in which the null in fact holds. Simulation evidence on this issue is presented in, among many others, Davidson and MacKinnon (1999) and Djogbenou, MacKinnon, and Nielsen (2018).

Table: Test size of balanced and unbalanced panel

G	\bar{T}	Obs	α	$\gamma = 0$	$\gamma = 0.5$	$\gamma = 1$	$\gamma = 1.5$	$\gamma = 2$
50	10	500	0.10	0.112	0.118	0.108	0.096	0.13
			0.05	0.062	0.05	0.04	0.048	0.058
			0.01	0.014	0.026	0.008	0.014	0.026
200	10	2,000	0.1	0.138	0.124	0.104	0.126	0.118
			0.05	0.078	0.004	0.054	0.062	0.066
			0.01	0.03	0.008	0.014	0.016	0.016



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- DGP:

$$y_{it} = \mu + \alpha z_{it} + \beta_1 x_{it} 1(q_{it} < \tau) + \beta_2 x_{it} 1(q_{it} \geq \tau) + u_i + e_{it}, \quad (11)$$

If $\beta_1 = \beta_2$, then the threshold effect disappears. The bigger difference between β_1 and β_2 , the higher the test power.

- Set $\beta_1 = 1, \beta_2 = (1.1, 1.2), \gamma = (0, 0.5, 1, 1.5, 2)$

Table: Test power of balanced and unbalanced panel ($\alpha = 0.05$)

G	\bar{T}	Obs	β_2	$\gamma = 0$	$\gamma = 0.5$	$\gamma = 1$	$\gamma = 1.5$	$\gamma = 2$
50	10	500	1.1	0.248	0.18	0.132	0.164	0.178
			1.2	0.722	0.49	0.492	0.486	0.492
200	10	2,000	1.1	0.748	0.498	0.466	0.454	0.446
			1.2	1.000	0.966	0.976	0.970	0.946



- The test power is relatively low for small sample with $G = 50$ and $\bar{T} = 10$. As G gets bigger, the test power is improved quite a lot.
- With the same number of observations, the test power for unbalanced panel is lower than that for balanced panel. But, the test power doesn't seem to be affected much by the extent of the unbalancedness.



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- Basically, `xthreg2` inherits the syntax from `xthreg`.

```
xthreg2 depvar [indepvars][if] [in] , rx(varlist) qx(varname) [  
thnum(integer) grid(integer) trim(numlist) bs(numlist) thlevel(#)  
gen(newvarname) noreg nob slog wc(string) options ]
```

- `xthreg2` will give the same point and confidence interval estimation with `xthreg`, but the threshold effect significance test may give different critical values and different probability value because of the different bootstrap design.

- Hansen (1999)

```
. use hansen1999, clear
. set seed 123
. xthreg2 i q1 q2 q3 d1 qd1, rx(c1) qx(d1) thnum(2) trim(0.01) bs(300)
There exist time-invariant individual(s) (maybe only one obs): d1 qd1
Estimating the threshold parameters: 1st ... 2nd ... Done
Bootstrap for single threshold
.....
..... + 300
Bootstrap for double threshold model:
.....
..... + 300
```

Threshold estimator (level = 95):

model	Threshold	Lower	Upper
Th-1	0.0158	0.0139	0.0177
Th-21	0.0158	0.0139	0.0177
Th-22	0.5394	0.5321	0.5470

Threshold effect test (bootstrap = 300):

Threshold	RSS	MSE	Fstat	Prob	Crit10	Crit5	Crit1
Single	16.7953	0.0024	30.25	0.047	26.942	29.982	37.916
Double	16.7457	0.0023	21.14	0.207	25.339	28.949	36.070

● unbalanced example

```

. use hansen1999,clear
. set seed 123
. drop if (year<=1977 | year>=1985) & inrange(runiform(0, 566), 1, 200) & rnormal(<0.1
(740 observations deleted)
. xtset
   id: 1, 2, ..., 565          n =          565
   year: 1974, 1975, ..., 1987      T =          14
      Delta(year) = 1 year
      Span(year)  = 14 periods
      (id*year uniquely identifies each observation)

```

```

Distribution of T_i:  min      5%    25%      50%      75%      95%      max
                   10      11     12        13        13        14        14

```

Freq.	Percent	Cum.	Pattern
138	24.42	24.42	1111111111111111
39	6.90	31.33	11.11111111111111
32	5.66	36.99	.1111111111111111
30	5.31	42.30	1.1111111111111111
29	5.13	47.43	111111111111.11
27	4.78	52.21	111.111111111111
26	4.60	56.81	11111111111111.
23	4.07	60.88	111111111111.1
10	1.77	62.65	..11111111111111
211	37.35	100.00	(other patterns)
565	100.00		XXXXXXXXXXXXXXXX

• triple threshold model

```
. set seed 123
. xthreg2 i q1 q2 q3 d1 qd1, rx(c1) qx(d1) thnum(3) trim(0.01) bs(300)
There exist time-invariant individual(s) (maybe only one obs): d1 qd1
Estimating the threshold parameters: 1st ... 2nd ... 3rd ... Done
Bootstrap for single threshold
..... + 300
Bootstrap for double threshold model:
..... + 300
Bootstrap for triple threshold model:
..... + 300
```

Threshold estimator (level = 95):

model	Threshold	Lower	Upper
Th-1	0.0154	0.0121	0.0173
Th-21	0.0154	0.0129	0.0173
Th-22	0.5421	0.5252	0.5487
Th-3	0.9230	.	.

Threshold effect test (bootstrap = 300):

Threshold	RSS	MSE	Fstat	Prob	Crit10	Crit5	Crit1
Single	17.3886	0.0027	36.65	0.010	24.212	27.861	36.592
Double	17.3201	0.0027	25.41	0.077	24.423	28.217	31.632
Triple	17.2933	0.0027	9.99	0.470	21.504	24.319	32.615

● double threshold reg. result

```
. xthreg2 i q1 q2 q3 d1 qd1, rx(c1) qx(d1) thnum(2) trim(0.01)
Fixed-effects (within) regression      Number of obs   =      6,992
Group variable: id                    Number of groups =      550
R-sq:                                  Obs per group:
    within = 0.1099                      min =          10
    between = 0.1166                     avg  =          12.7
    overall = 0.1064                     max  =          14

corr(u_i, Xb) = -0.1886                  F(8,6434)       =      99.29
                                          Prob > F        =      0.0000
```

i	Coef.	Std. Err.	t	P> t	[95% Conf. Interval]	
q1	.0115655	.0009494	12.18	0.000	.0097043	.0134267
q2	-.0248182	.0028863	-8.60	0.000	-.0304763	-.0191602
q3	.0013585	.0002162	6.28	0.000	.0009348	.0017823
d1	-.0249041	.0041553	-5.99	0.000	-.0330498	-.0167583
qd1	.0010209	.0013583	0.75	0.452	-.0016418	.0036837
_cat#c.c1						
0	.0580387	.0066704	8.70	0.000	.0449624	.071115
1	.1353776	.0176681	7.66	0.000	.1007422	.1700131
2	.0861913	.0054647	15.77	0.000	.0754787	.0969039
_cons	.0636492	.0017055	37.32	0.000	.0603059	.0669925
sigma_u	.03530003					
sigma_e	.04558565					
rho	.37486107	(fraction of variance due to u_i)				

F test that all u_i=0: F(549, 6434) = 6.88

Prob > F = 0.0000



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- A more unbalanced panel will result in more volatile threshold estimate and tighter coverage rate
- The test size is not affected by the unbalancedness.
- The test power is affected by the unbalancedness, but it doesn't deteriorate when the panel becomes more unbalanced.

- Our conclusions don't mean that it is be always better to use the unbalanced panel than to transform it into balanced one.
- Two choices for a unbalanced panel:
First, transform the panel data into balanced one by dropping some observations, then estimate the threshold model using the balanced panel.
Second, directly estimate the threshold model using the unbalanced panel.
- Both choices have pros and cons. The balanced panel has a less volatile estimate for the threshold, but less efficient estimate for regression coefficient due to less observations. The unbalanced panel has a more volatile estimate for the threshold, but more efficient estimate for regression coefficient beneficial from more observations.



- the effect of heteroscedasticity
- sample selection bias

Thank you