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How does the Persons per Household Affect Population Estimation and how to Measure it

Xiaomin Ruan



University of New Mexico BBER Population Estimation & Projection Program For the Data User Conference, Nov. 2008

Outlines

- Concepts related to persons per household (PPH)
- Effects of PPH in housing unit method
- Components that affect PPH
- PPH estimation model selection and validation
- Exploration of forecasting PPH

Concepts Related to Persons per Household

Persons per Household (PPH)

the number of persons in one household. Also named as
Household Size by Census Bureau.

PPH = Total Population / # Households (or # Housing Units)

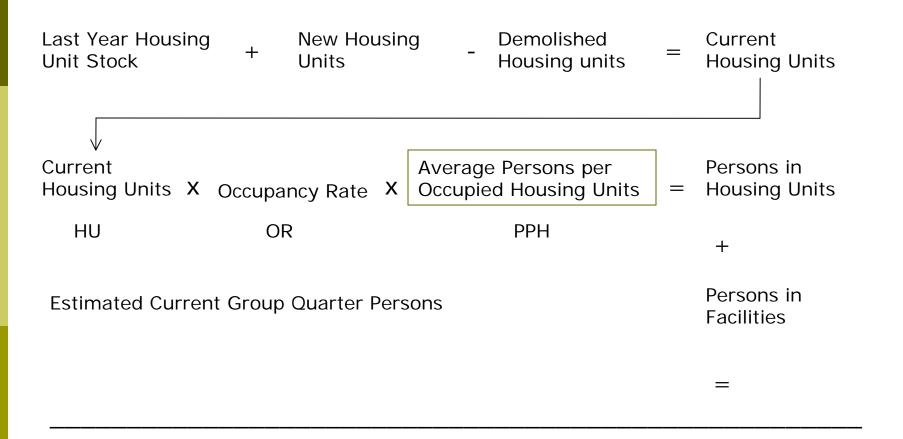
Household

- A household includes all persons who occupy a housing unit.

Housing Unit (HU)

— A housing unit is a house, an apartment, a mobile home, a group of rooms, or a single room that is **occupied** as a separate living quarters.

PPH in Population Estimation

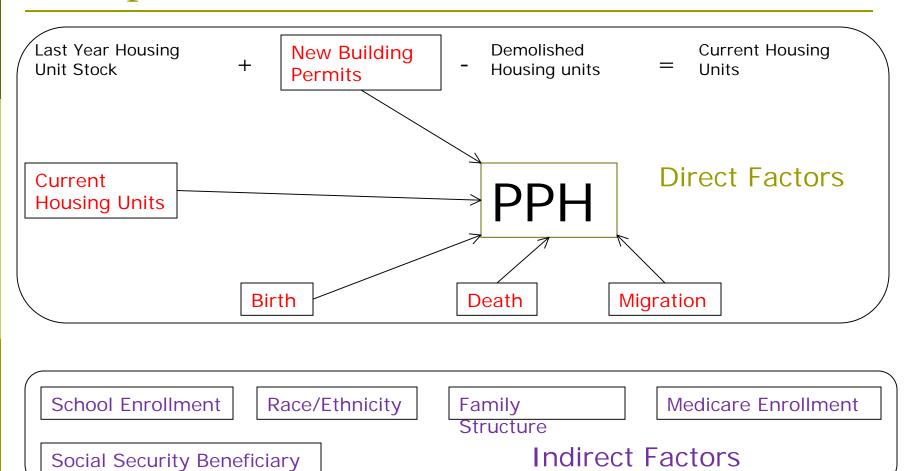


Total Persons

Sensitivity of Population Estimation to PPH

| Geographic Level | | | PPH Variation | | | | |
|------------------|------------------|-----------|---------------|-----------|-----------|-----------|--|
| Geogra | Geographic Level | | -1% | -2% | -5% | -10% | |
| | РРН | 2.34 | 2.32 | 2.27 | 2.16 | 1.94 | |
| Tract | Occupied HU | 1,183 | - | - | - | - | |
| nact | Population | 2,771 | 2,743 | 2,688 | 2,554 | 2,299 | |
| | Absolute Diff. | - | -28 | -83 | -217 | -472 | |
| | РРН | 2.52 | 2.49 | 2.47 | 2.39 | 2.27 | |
| County | Occupied HU | 220,936 | - | - | - | - | |
| county | Population | 556,678 | 551,111 | 545,544 | 528,844 | 501,010 | |
| | Absolute Diff. | - | -5,567 | -11,134 | -27,834 | -55,668 | |
| | РРН | 2.68 | 2.66 | 2.63 | 2.55 | 2.41 | |
| State | Occupied HU | 677,971 | - | - | - | - | |
| State | Population | 1,819,046 | 1,800,856 | 1,782,665 | 1,728,094 | 1,637,141 | |
| | Absolute Diff. | - | -18,190 | -36,381 | -90,952 | -181,905 | |

Components that affect PPH



Available Source Data

| Data Source | Time Span | Note |
|----------------------|-----------|--|
| ACS | 2002-2007 | Only MSA available |
| OASDI | 1984-2006 | Cibola missed 84-87 data, while Catron and Harding have imcomplete 86 and 87 data. |
| Medicare | 1998-2006 | Complete |
| Birth | 1990-2006 | Complete |
| Death | 1990-2006 | Complete |
| IRS Migration | 1981-2007 | Missed year 82, 83, 90, 91, 92 data |
| School Enrollment | 1986-2006 | Complete grade 1 to 12 |
| Building Permit | 2000-2006 | Complete |
| Employment | 1995-2006 | Complete |
| Race from Birth | 1990-2005 | Complete |
| Race from Death | 1990-2005 | Complete |
| Race from Sch_Enroll | 1989-2004 | Complete |

Regression Method

- Regression method uses the variation in independent variables to explain the variation in PPH
- Once the regression model is validated, the independent variable values can be plugged in inter decennial years to estimate the missing PPH and construct a PPH time series to support population estimation.
- Structural test (Chow Test) is critical to the regression model. Once the test shows the variance of residuals of Year 2000's PPH is indifferent from that of 1990, the estimated coefficients can then be applied to other years. Combining with validating process, the structural test can safely demonstrate the predicting power of the chosen model.

Stepwise Selection using SAS

- Including too many explanatory variables will decrease the degree of freedom, but including fewer important variables may reduce the explanatory power of the model. So, a stepwise selection combining forward selection and backward elimination is applied.
- The rule of forward selection is, adding variables to the model once at a time until no significant variable can be found to increase R-square of the previous model (Default significant level for individual variable is 50%).
- The rule of backward elimination is, deleting unimportant variables (decrease R-square the least or has the highest p-value) from the full model until any variable left is significant at default level (10%).
- Stepwise regression combines features of forward selection and backward elimination, which means, every time we add a variable to the model, we ask whether any of the variables added earlier can be omitted. (Default individual variable's significance level is 15%)

Data Transformation

- Knowing that PPH is a population count average by housing units, taking log of the count variables may reduce the model variation and increase model stability.
- Moreover, since PPH is a count variable weighted by housing units, weighting independent variables also by housing units may also increase the explanatory power of regression models.
- Weighting variables by housing units may cause another issue. Since housing unit stock data for non-census years are estimates by BBER, it may be inconvenient to use by outside data users, comparing to the access of administrative data.

Stepwise Selected Models

| | Count Model | Log Model | HU Weighted Model |
|-----------|-------------|-------------|-------------------|
| Dep. Var. | РРН | РРН | РРН |
| Ind. Var. | AI | - | HUW_AI |
| | - | Ln_OASDI_rw | HUW_OASDI_rw |
| | - | Ln_Birth | HUW_Birth |
| | - | Ln_G7_9 | HUW_G10_12 |
| | - | Ln_EmpEND | HUW_EmpAVE |
| | - | Ln_HispSE | HUW_HispSE |

F-Test, AIC/BIC screening

| Model | Count Model | Log Model | HU Weighted Model |
|-----------------|-------------|-----------|-------------------|
| Pseudo | | | |
| Chow Test | -13.15 | 0.85 | 3.06** |
| | | | |
| AIC Test | -2.97 | -4.38 | -4.87 |
| BIC Test | -79.47 | -172.32 | -204.22 |

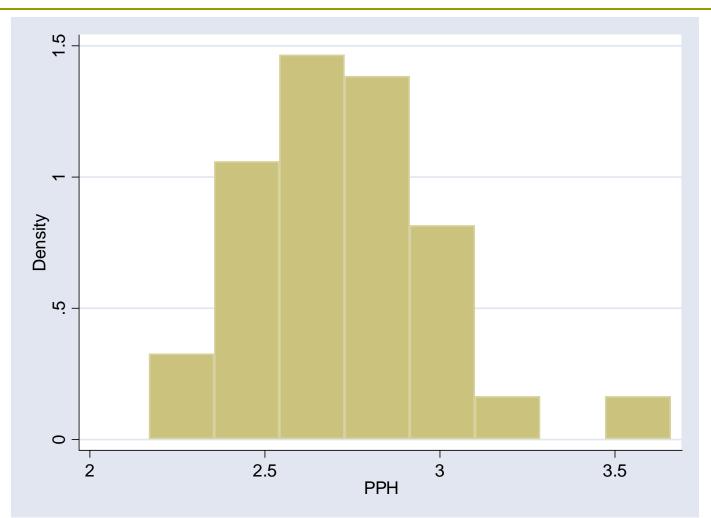
AIC = 2k + n[ln(RSS/n) + 1]

BIC = n*In(RSS/n) + k*In(n)

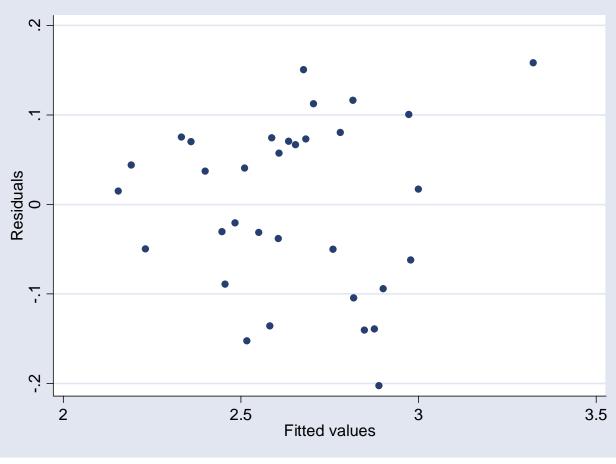
Descriptive Statistics

| | Var. Name | Obs | Mean | S.D. |
|-------------|--------------|-----|------|------|
| Dep. Var. | РРН | 33 | 2.65 | 0.28 |
| | Ln_OASDI_rw | 33 | 7.81 | 1.19 |
| | Ln_Birth | 33 | 5.72 | 1.58 |
| Log Model | Ln_G7_9 | 33 | 6.97 | 1.37 |
| | Ln_EmpEND | 33 | 8.86 | 1.54 |
| | Ln_HispSE | 33 | 7.60 | 1.43 |
| | | | | |
| | HUW_Indian | 33 | 0.04 | 0.09 |
| | HUW_OASDI_rw | 33 | 0.22 | 0.05 |
| HU Weighted | HUW_Birth | 33 | 0.03 | 0.01 |
| Model | HUW_G10_12 | 33 | 0.09 | 0.03 |
| | HUW_EmpAVE | 33 | 0.63 | 0.23 |
| | HUW_HispSE | 33 | 0.20 | 0.10 |

PPH Histogram

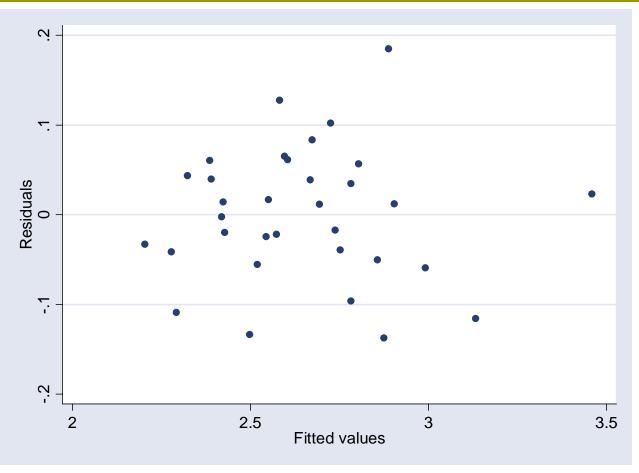


Heteroscedasticity Check Model 1



Log Model Residual Plot

Heteroscedasticity Check Model 2



HU Weighted Model Residual Plot

Estimated Equation

Stepwise Selected Log Model

PPH = 4.26386-0.36158*Ln_OASDI_rw+0.54668*Ln_Birth+0.19909*Ln_G7_9

(0.257)*** (0.070)*** (0.084)*** (0.085)**

-0.28872*Ln_EmpEND-0.09882*Ln_HispSE

(0.083)*** (0.039)**

 $R^2 = 0.8795$, $F(5,27) = 39.40^{***}$

Stepwise Selected HU Weighted Model

 $PPH = 2.25202-0.75932*HUW_OASDI_rw+15.49605*HUW_Birth+1.52901*HUW_G10_12$ (0.098)***(0.336)** (2.409)*** (0.644)** $-0.22723*HUW_EmpAVE+1.00188*HUW_Indian+0.40881*HUW_HispSE$ (0.080)*** (0.254)*** (0.207)* $R^2 = 0.9306, F(6,26) = 58.15***$

Validation

- The comparison of the fitted values against original data can tell us how good the model fits the data. But it may not reveal the ability to predict other place or other time spot.
- The model's ability to predict depends on both the model specification and the quality of the data in validating years.
- As we have to use ACS PPH estimates instead of other solid administrative records to validate our models, the differences between the predicted PPH and the ACS estimates may not reveal the true distance to the real PPH values.

Validation Continued

| | Log M | odel | HUW M | HUW Model | | |
|----------|--------|--------|--------|-----------|--|--|
| | 2000 | 2005 | 2000 | 2005 | | |
| # County | 33 | 7 | 33 | 7 | | |
| ME | 0.0001 | 0.0512 | 0.0000 | -0.0955 | | |
| MPE | 0.0011 | 0.0248 | 0.0008 | -0.0267 | | |
| MAPE | 0.0304 | 0.0809 | 0.0215 | 0.0657 | | |

Bayesian Basic Concepts for Simulation

- The basic idea behind Bayesian Modeling is a simple statistic idea: conditional probability.
- Conditional Probability Example: Flu test for 151 students

| | Flu | No Flu | | | Flu | No Flu | |
|-------|-----|--------|-----|--------|------|--------|---|
| st + | 46 | 15 | 61 | Test + | 0.30 | 0.10 | 0 |
| est - | 4 | 86 | 90 | Test - | 0.03 | 0.57 | C |
| | 50 | 101 | 151 | | 0.33 | 0.67 | 1 |

 $P(T^+/F) = 46/50 = 0.30/0.33 = P(T^+&F) / P(F)$ = P(F/T^+)P(T^+)/P(F) = (0.3/0.4)*0.4/0.33

Question: If one student was tested positive, what is his probability to have a real flu?

Parallel Bayesian Question in our Case

If the predictors are estimated like the equation below, with such a specified variance, what would they really look like once we have more than enough observations?

 $PPH = 4.26386-0.36158*Ln_OASDI_rw+0.54668*Ln_Birth+0.19909*Ln_G7_9$ (0.257)*** (0.070)*** (0.084)*** (0.085)** $-0.28872*Ln_EmpEND-0.09882*Ln_HispSE$ (0.083)*** (0.039)** $R^2 = 0.8795, F(5,27) = 39.40***$

WinBUGS do Bayesian Automatically

- What one needs to do is to specify the model first,
- Input data second, and then
- **D** Specify the prior information and let the program run simulation

The OPEN SOURCE WinBUGS website has a demonstration movie http://www.mrc-bsu.cam.ac.uk/bugs/winbugs/contents.shtml

First Try of Bayesian using Prior Estimates

Bayesian Simulated Log Model

PPH = 4.267-0.3656*Ln_OASDI_rw+0.5434*Ln_Birth+0.2015*Ln_G7_9

(0.269)** (0.072)**

(0.088)**

-0.2874*Ln_EmpEND-0.09627*Ln_HispSE

(0.089)**

(0.086)**

(0.040)**

Bayesian Simulated HU Weighted Model

PPH = 2.248-0.7529*HUW_OASDI_rw+15.44*HUW_Birth+1.528*HUW_G10_12

(0.102)** (0.352)** (2.512)** (0.678)**

-0.2209*HUW_EmpAVE+0.9972*HUW_Indian+0.4116*HUW_HispSE

(0.084)** (0.268)** (0.218)**

Bayesian Model Validation

| | Log M | odel | HUW | HUW Model | | |
|----------|--------|-----------|--------|-----------|--|--|
| | 2000 | 2000 2005 | | 2005 | | |
| # County | 33 | 7 | 33 | 7 | | |
| ME | 0.0009 | 0.1199 | 0.0001 | 0.0372 | | |
| MPE | 0.0015 | 0.0496 | 0.0008 | 0.0192 | | |
| MAPE | 0.0304 | 0.0890 | 0.0215 | 0.0904 | | |

2005 PPH Sensitivity Analysis

| | Fitted 2005 | ACS 2005 | | Pop Based on | Pop based on | |
|------------|-------------|----------|-------|--------------|--------------|---------|
| County | PPH | PPH | Diff. | Fitted | ACS | Pop Gap |
| Bernalillo | 2.55 | 2.37 | 0.18 | 635745 | 591860 | 43885 |
| Dona Ana | 2.76 | 2.75 | 0.01 | 185340 | 184377 | 964 |
| McKinley | 3.38 | 2.59 | 0.79 | 66919 | 51308 | 15611 |
| Sandoval | 2.63 | 2.75 | -0.12 | 102224 | 106692 | -4468 |
| San Juan | 3.01 | 3.29 | -0.28 | 114344 | 125106 | -10761 |
| Santa Fe | 2.35 | 2.61 | -0.26 | 124332 | 137805 | -13473 |
| Valencia | 2.68 | 2.74 | -0.06 | 65523 | 67108 | -1585 |

Total 30173

Future Work for Model Specification

- Look for previous census year data that can be used in regression models
- Compare fitted values to BBER Pop estimates.
- Use only those 7 MSA counties to do a structural change test.
- Use ACS 2006 data as reference instead of 2005 data.
- Try simultaneous equations

Appendix

SAS code

WinBUGS Bayesian Modeling Code

Thank you!

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