Towards a New, Public Dataset for Studying Mortality Inequality

Matching the 1940 U.S. Census with Social Security death records, 1975-2005

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## Two quotations

The Human Mortality Database has launched thousands of papers, but we're at risk of falling behind. It's hard to study inequality without individual level data.

- PAA HMD workshop


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We lost our access to the SIPP \& SSA data file when the grant expired.

- a real researcher working on inequality and mortality


## The Challenges

- Protect individual privacy
- Easy access for all researchers
- Replication and scientific progress
(improving on others' work)
- Large sample sizes
- Lots of covariates
- Integrated estimation (matching and modeling)


## The Opportunity: Unrestricted data

- The 1940 U.S. Census with names and rich covariates
- Social Security Death Records with names and mortality


## 1940 Census



- 130 million people
- Rich individual level variables
(Full demographics \& income \& education \& housing)
- Street address level geography (Neighbors, blocks, ...)
- Already transcribed by Ancestry and MPC


## Social Security Death Index

## SSDI deaths / HMD counts

- 80 million deaths
- Nearly complete coverage (over age 65, 1975-2005)
- Name, SSN, DOB, DOD
- Public information



## Lexis diagram of CenSoc linkage



## Matching method

## Exact matching of unique keys

(first name, last name, birth year)

- About $75 \%$ of keys are unique


## Linkage

| Full 1940 census | 132 | million |
| :--- | ---: | :---: |
| and name and age available | 130 | million |
| and male | 64 | million |
| and aged 0-70 | 62 | million |
| and unique key | 46 | million |
| and expected death in interval | 14 | million |
| Matches | 6 | million |
| Match rate | 43 | $\%$ |

## Matching bias?



Matched also a bit whiter, more educated, and likely to be home-owners.

## Mortality rate validation



Key:
solid line $=$ HMD (Human Mortality Database) black dash $=$ reverse survival matched data
red dash $=$ reverse survival matched data, adjusted with HMD survivors

Mortality rates for matched US males,
by income quartile (1910 cohort)
Includes zero income


Mortality rates for matched US males,
by income quartile (1910 cohort)
Does not include zero income


## Smaller groups and new variables

OLS regressions of age-at-death for those aged 20-35 in 1940

| Intercept | 77.83*** | 74.70*** |
| :---: | :---: | :---: |
| Black/White | -0.90*** | 0.08 |
| Chinese/White | 1.01*** | 1.90*** |
| Filipino/White | 1.69*** | 2.40*** |
| Japanese/White | 1.90*** | $2.26 * * *$ |
| Other/White | $-1.27 * * *$ | -0.70*** |
| educ |  | 0.19*** |
| $\log$ (income) |  | $0.18 * * *$ |
| own/rent |  | 0.50*** |
| hh_head: Yes |  | -0.15 |
| N | 2 million | 1.1 mill |

## Validation: Middle Initials

|  | clean_key | ssn | fname.x | lname.x mname.x mname mi.match |  |  |  |
| ---: | ---: | ---: | ---: | ---: | ---: | ---: | ---: |
| 1: | AABERGEDWARD49 | 538074776 | EDWARD | AABERG | J |  |  |
| 2: | AABERGEELMER32 | 516228997 | ELMER | AABERGE |  |  |  |
| 3: | AABERGERIC34 | 521071090 | ERIC | AABERG | C | C | TRUE |
| 4: | AABERGLAWRENCE22 | 517169163 | LAWRENCE | AABERG | M | A | FALSE |
| 5: | AABERGRALPH30 | 522071496 | RALPH | AABERG |  | 0 |  |
| 6: | AABERGROBERT39 | 563033374 | ROBERT | AABERG | A | A | TRUE |
| 7: | AABERGSANDER43 | 535096685 | SANDER | AABERG | P |  |  |
| 8: | AABWILLIAM42 | 523147290 | WILLIAM | AAB |  |  |  |
| 9: | AABYCARLYLE20 | 473091698 | CARLYLE | AABY | P | P | TRUE |
| 10: | AABYELWIN18 | 517167623 | ELWIN | AABY |  |  |  |

Middle initial match rate $\approx 80 \%$

## Validation: Errors-in-variables framework

## Estimated coefficent of education on age at death by status of match



## Validation: Errors-in-variables framework

## Estimated coefficent of education on age at death by status of match



## Future Directions

- Public release of our 6-7 million linked deaths (with IPUMS, HMD)
- Other match methods (birthplace, probablistic matches?)
- Mortality estimation for linked data
- Parametric MLE for doubly-truncated cohorts
- Bayesian methods (Schmertmann et al.)
- Missing-at-random methods (Taylor, Sanders et al.)

