Estimating subnational populations of women of reproductive age in developing countries

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Introduction

Background

- Important to monitor demographic and health indicators at the subnational level
- Need population counts
 - of particular interest is women of reproductive age (15-49)
 - measures exposure to risk
 - denominator for maternal mortality, fertility rates, contraceptive prevalence..
- However, in many developing countries, population data are limited and/or of poor quality
- Develop statistical methods to overcome missing data and quality issues

Approach



Patrick Leslie





- Use a demographic model for estimation and projections to avoid implausible outcomes
- Estimation in a Bayesian framework to facilitate inclusion of various data sources and pooling of information across populations

Overview of project

We are developing a Bayesian demographic modeling approach to estimate and project

- female populations aged 15-49 (WRA)
- from \sim 1980 to 2020
- at the subnational level (admin2, i.e. county)
- using data sources that are commonly available in developing countries.

The model is applied to Kenya to estimate and project WRA populations from 1979-2020.

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Data

Data

Data

Administrative levels in Kenya



 Nationally, WRA increased from 3.4 million (1980) to 9.4 million (2009, last census year), projected to 13.2
million (2020)

• 8 provinces, 36 districts

Data sources used

Census:

- 10% microsamples are available for years 1979, 1989, 1999, 2009 (IPUMS)
- Subnational population counts (adjusted for age heaping)
- Data on internal migration in year before the census

National estimates from World Population Prospects (WPP):

- National estimates of population counts
- National estimates of mortality rates



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Model overview



Cohort component projection

Use a cohort component reconstruction/projection model:

$$\eta_{r,c,a+1} = \eta_{r,c,a} \cdot (1 - \rho_{r,c,a}) + \phi_{r,c,a},$$

with for region *r* and birth cohort *c*:

- $\eta_{r,c,a}$ the number of women of age $a = 15, 16, \ldots, 49$,
- $\rho_{r,c,a}$ the expected proportion of women who die between age a and a+1
- $\phi_{r,c,a}$ the expected number of net-migrants into/out of region r at age a

Cohort component projection

$$\eta_{r,c,a+1} = \eta_{r,c,a} \cdot (1 - \rho_{r,c,a}) + \phi_{r,c,a}$$

Observed census counts

$$y_{r,c,a} \sim N(\eta_{r,c,a}, s_{r,c,a}^2),$$

where sampling error s is due to 10% microsample.

• We constrain the sum of subnational populations to add up to national WPP estimates within lower and upper bounds (which is approximately 10%)

Mortality

$$\eta_{r,c,a+1} = \eta_{r,c,a} \cdot (1 - \rho_{r,c,a}) + \phi_{r,c,a}$$

Modeling the probability of death ρ by region, cohort and age:

Obtain age-specific 'principal component' vectors:

- mean mortality P_0
- mortality decline P_1
- impact of the AIDS epidemic P_2 using a principal component analysis of logit(national ρ 's).



Mortality

Use principal components as the basis of a regression model for $\rho_{r,c,a}$

$$logit(\rho_{r,c,a}) = P_{0,a} + \beta_{r,c,1} \cdot P_{1,a} + \beta_{r,c,2} \cdot P_{2,a}$$

Many different shapes of $\rho_{r,c,a}$ can be represented by different combinations of the *P*'s.

Mortality

$$logit(\rho_{r,c,a}) = P_{0,a} + \beta_{r,c,1} \cdot P_{1,a} + \beta_{r,c,2} \cdot P_{2,a}$$

Use hierarchical time series model on the β s:

$$\beta_{r,c,1} = \mu_c + \delta_{r,c}$$

with across-regions cohort mean μ_c and "cohort-specific deviations within a region" $\delta_{r,c}$ (modeled by an AR(1) process).



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Net migration

$$\eta_{r,c,a+1} = \eta_{r,c,a} \cdot (1 - \rho_{r,c,a}) + \phi_{r,c,a}$$

Data suggest constant age patterns in net-migration:



Net migration

We assume for net-migration numbers ϕ :

$$\phi_{r,c,a} = \eta_{r,c} \cdot x_{r,a} \cdot \pi_{r,c},$$

where

- $\eta_{r,c} = \sum_{a} \eta_{r,c,a}$ the total population of WRA,
- x_{r,a} = proportion of net-migration at age a, age pattern assumed constant across cohorts, obtained from census data,
- $\pi_{r,c}$ = proportion of WRA that is net-migration:
 - Modeled with a random walk, $\pi_{r,c} \sim N(\pi_{r,c-1}, \sigma_{\pi}^2)T(-0.2, 0.2)$,
 - Informed by data from census, $p_{r,c} \sim N(\pi_{r,c}, \sigma_p^2)$.

Model overview



Results

Population by district

Data and estimates for WRA by Province

Rift Valley Central Coast Eastern Nairobi Northeastern Nvanza Western 2500 2000 Population ('000) 1500 etch 🛨 estimate 1000 500 0 2020 1980 0661 2000 2010 2020 .086 .0661 2000 2010 2020 1980 .0661 2000 2010 2020 1980 0661 2000 2010 .086 .066 2000 2010 2020 2020 1980 .0661 2000 2010 2020 .086 .0661 2000 2010 2020 .0861 2000 2010 Year

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Population by district

Population (\cdot 1,000)



Age patterns by cohort



Summary

Summary

- Cohort component projection model embedded with a Bayesian hierarchical framework
- Robust estimates of WRA populations at the subnational level
- Incorporating different data sources that are widely available
- Future work will focus on investigating how to use other data sources
 - e.g. sibling death information; census household deaths

Thanks!

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