

# The formal demography of kinship: Two-sex models

Hal Caswell  
University of Amsterdam

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## Why kinship?



## The kinship model

The kin of Focal as a population

$$\mathbf{k}(x+1) = \mathbf{U}\mathbf{k}(x) + \beta(x)$$

$$\mathbf{k}(0) = \mathbf{k}_0$$

where

$\mathbf{k}(x)$  = age distribution of a kin at age  $x$  of Focal<sup>1</sup>

$\beta(x)$  = recruitment 'subsidy' at age  $x$  of Focal

$$= \begin{cases} \mathbf{0} & \text{no subsidy} \\ \mathbf{F}\mathbf{k}^*(x) & \text{subsidy from } \mathbf{k}^* \end{cases}$$

$\mathbf{k}_0$  = initial condition

<sup>1</sup>To be more precise, the expectation of the age distribution.

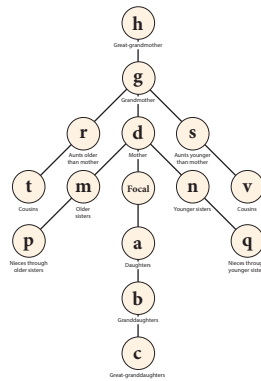


## Dynamics of kin of Focal at age $x$

Symbol	Kin	i.c. $\mathbf{k}_0$	$\beta(x)$
<b>a</b>	daughters	<b>0</b>	<b><math>\mathbf{F}e_x</math></b>
<b>b</b>	granddaughters	<b>0</b>	<b><math>\mathbf{F}a(x)</math></b>
<b>c</b>	great-granddaughters	<b>0</b>	<b><math>\mathbf{F}b(x)</math></b>
<b>d</b>	mothers	$\pi$	<b>0</b>
<b>g</b>	grandmothers	$\sum_i \pi_i \mathbf{d}(i)$	<b>0</b>
<b>h</b>	great-grandmothers	$\sum_i \pi_i \mathbf{g}(i)$	<b>0</b>
<b>m</b>	older sisters	$\sum_i \pi_i \mathbf{a}(i)$	<b>0</b>
<b>n</b>	younger sisters	<b>0</b>	<b><math>\mathbf{F}d(x)</math></b>
<b>p</b>	nieces via older sisters	$\sum_i \pi_i \mathbf{b}(i)$	<b><math>\mathbf{F}m(x)</math></b>
<b>q</b>	nieces via younger sisters	<b>0</b>	<b><math>\mathbf{F}n(x)</math></b>
<b>r</b>	aunts older than mother	$\sum_i \pi_i \mathbf{m}(i)$	<b>0</b>
<b>s</b>	aunts younger than mother	$\sum_i \pi_i \mathbf{n}(i)$	<b><math>\mathbf{F}g(x)</math></b>
<b>t</b>	cousins from aunts older than mother	$\sum_i \pi_i \mathbf{p}(i)$	<b><math>\mathbf{F}r(x)</math></b>
<b>v</b>	cousins from aunts younger than mother	$\sum_i \pi_i \mathbf{q}(i)$	<b><math>\mathbf{F}s(x)</math></b>



## Female kin through female lines of descent



We want both sexes through all lines of descent

## Two-sex kinship: demographic components

$U_f, U_m$  = female and male survival

$F_f, F_m$  = female and male fertility

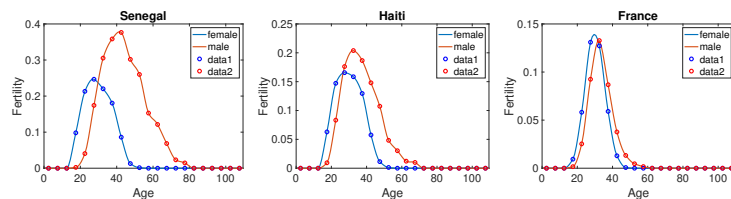
$\pi_f, \pi_m$  = female and male ages at birth

$\alpha$  = proportion male births

$\bar{\alpha} = 1 - \alpha$

## Female and male rates<sup>2</sup>

- mortality differences universal and well known
- fertility differences can be large
  - in timing
  - in amount



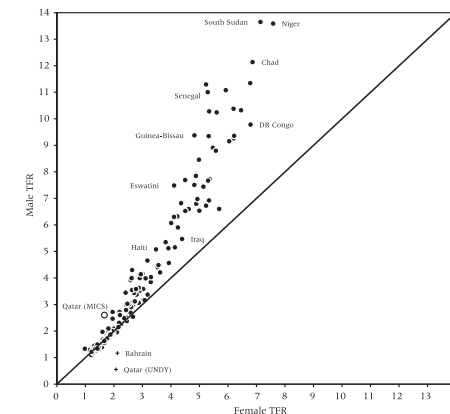
<sup>2</sup>Schoumaker, B. 2019. Male Fertility Around the World and Over Time: How Different is it from Female Fertility? Population and Development Review 45(3): 459-487

## Female and male fertility

BRUNO SCHOUMAKER

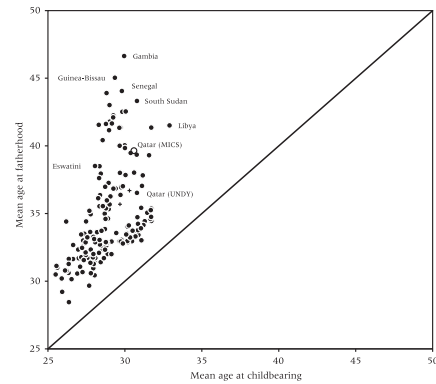
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FIGURE 2 Comparison of male and female total fertility rates, circa 2011 in 163 countries



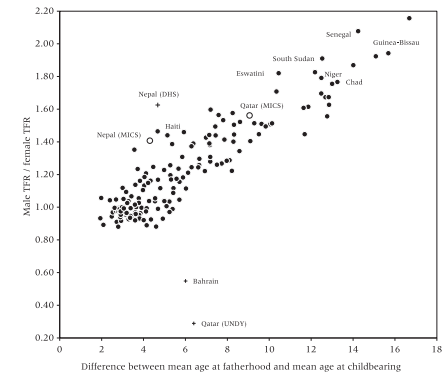
## Female and male fertility

**FIGURE 3** Comparison of mean age at childbearing (women) and mean age at fatherhood (men), circa 2011 in 163 countries



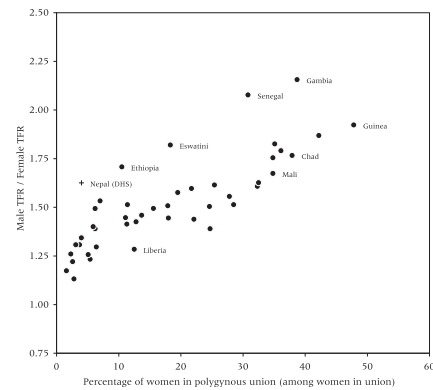
## Female and male fertility

**FIGURE 5** Relationship between the ratio of male TFR to female TFR and the difference between mean age at fatherhood and mean age at childbearing, circa 2011 in 163 countries

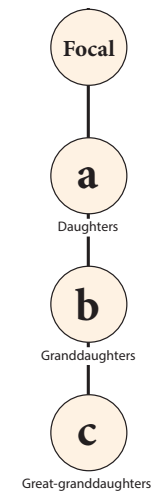


## Female and male fertility

**FIGURE 6** Relationship between male-female fertility ratio and polygyny, circa 2011 in 43 countries

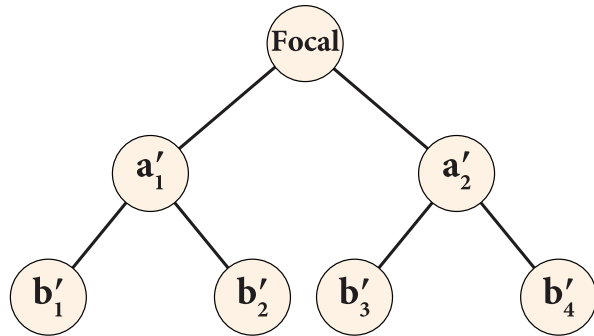


## Descendants: one-sex lineage



## Descendants of both sexes, both lineages

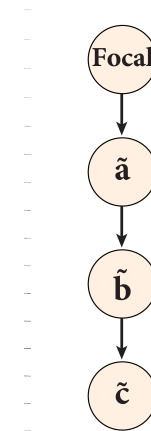
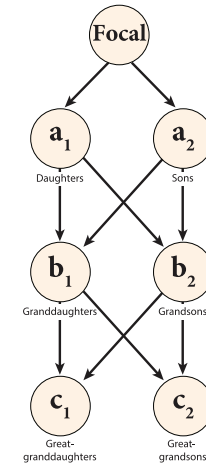
female and male lineages



## Descendants of both sexes

rates independent of lineage

block-structured vectors



## Two-sex models

Block-structured population vector

$$\tilde{\mathbf{k}} = \begin{pmatrix} \mathbf{k}_f \\ \mathbf{k}_m \end{pmatrix}$$

## Two-sex models

Block-structured matrices

$$\tilde{\mathbf{U}} = \left( \begin{array}{c|c} \mathbf{U}_f & \mathbf{0} \\ \hline \mathbf{0} & \mathbf{U}_m \end{array} \right)$$

$$\tilde{\mathbf{F}} = \left( \begin{array}{c|c} \bar{\alpha}\mathbf{F}_f & \bar{\alpha}\mathbf{F}_m \\ \hline \alpha\mathbf{F}_f & \alpha\mathbf{F}_m \end{array} \right)$$

$$\tilde{\mathbf{F}}^* = \left( \begin{array}{c|c} \bar{\alpha}\mathbf{F}_f & \mathbf{0} \\ \hline \alpha\mathbf{F}_f & \mathbf{0} \end{array} \right)$$

where

$\alpha$  = sex ratio (proportion male births)

$\bar{\alpha} = 1 - \alpha$

## Age at maternity and paternity

where

$$\tilde{\pi} = \left( \frac{\pi_f}{\pi_m} \right)$$

$$\|\boldsymbol{\pi}_f\| = \|\boldsymbol{\pi}_m\| = 1$$

The dynamics of  $\tilde{\mathbf{k}}(x)$

$$\left(\frac{\mathbf{k}_f}{\mathbf{k}_m}\right)(x+1) = \left(\frac{\mathbf{U}_f \mid \mathbf{0}}{\mathbf{0} \mid \mathbf{U}_m}\right) \left(\frac{\mathbf{k}_f}{\mathbf{k}_m}\right)(x) + \left(\frac{\beta_f}{\beta_m}\right)(x)$$

$$\tilde{\mathbf{k}}(x+1) = \tilde{\mathbf{U}}\tilde{\mathbf{k}}(x) + \tilde{\beta}(x).$$

Children, grandchildren, ...

## Children

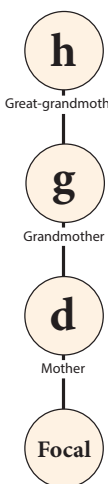
$$\tilde{\mathbf{a}}(x+1) = \tilde{\mathbf{U}}\tilde{\mathbf{a}}(x) + \tilde{\mathbf{F}}\tilde{\phi}(x)$$

## Grandchildren

$$\tilde{\mathbf{b}}(x+1) = \tilde{\mathbf{U}}\tilde{\mathbf{b}}(x) + \tilde{\mathbf{F}}\tilde{\mathbf{a}}(x)$$

etc.

## One-sex lineage of ancestry



Great-grandmother

Grandmother

Mother

(Focal

## Parents, grandparents, ...

Parents

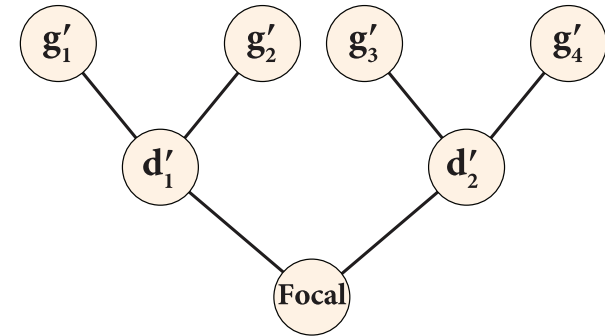
$$\begin{aligned}\tilde{\mathbf{d}}(x+1) &= \tilde{\mathbf{U}}\tilde{\mathbf{d}}(x) + \mathbf{0} \\ \tilde{\mathbf{d}}(0) &= \tilde{\pi}\end{aligned}$$

Grandparents

$$\begin{aligned}\tilde{\mathbf{g}}(x+1) &= \tilde{\mathbf{U}}\tilde{\mathbf{g}}(x) + \mathbf{0} \\ \tilde{\mathbf{g}}(0) &= \sum_i [\pi_f(i) + \pi_m(i)] \tilde{\mathbf{d}}(i)\end{aligned}$$

Navigation icons

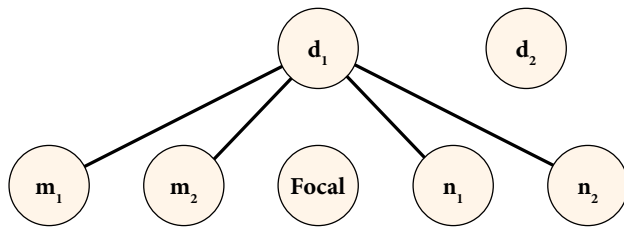
## Counting ancestors of both sexes



(and then combine to get block-structured vectors, as with descendants)

Navigation icons

## Ancestors don't reproduce independently



$$\begin{pmatrix} \mathbf{n}_f \\ \mathbf{n}_m \end{pmatrix}(x+1) = \tilde{\mathbf{U}} \begin{pmatrix} \mathbf{n}_f \\ \mathbf{n}_m \end{pmatrix}(x) + \begin{pmatrix} \bar{\alpha}\mathbf{F}_f & \mathbf{0} \\ \alpha\mathbf{F}_f & \mathbf{0} \end{pmatrix} \begin{pmatrix} \mathbf{d}_f \\ \mathbf{d}_m \end{pmatrix}(x)$$

$$\tilde{\mathbf{n}}(x+1) = \tilde{\mathbf{U}}\tilde{\mathbf{n}}(x) + \tilde{\mathbf{F}}^*\tilde{\mathbf{d}}(x)$$

Navigation icons

## Recruitment

1. by a direct ancestor of Focal

$$\begin{pmatrix} \beta_f \\ \beta_m \end{pmatrix}(x) = \begin{pmatrix} \bar{\alpha}\mathbf{F}_f & \mathbf{0} \\ \alpha\mathbf{F}_f & \mathbf{0} \end{pmatrix} \begin{pmatrix} \mathbf{k}_f^* \\ \mathbf{k}_m^* \end{pmatrix}(x)$$

$$\tilde{\beta}(x) = \tilde{\mathbf{F}}^*\tilde{\mathbf{k}}(x)$$

2. by any other kin type

$$\begin{pmatrix} \beta_f \\ \beta_m \end{pmatrix}(x) = \begin{pmatrix} \bar{\alpha}\mathbf{F}_f & \bar{\alpha}\mathbf{F}_m \\ \alpha\mathbf{F}_f & \alpha\mathbf{F}_m \end{pmatrix} \begin{pmatrix} \mathbf{k}_f^* \\ \mathbf{k}_m^* \end{pmatrix}(x)$$

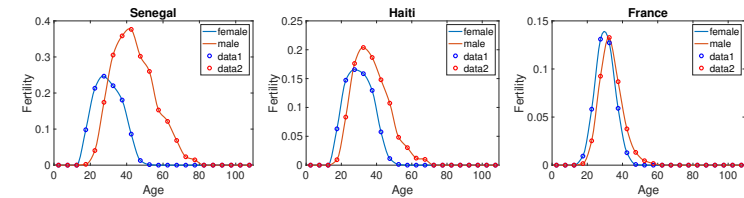
$$\tilde{\beta}(x) = \tilde{\mathbf{F}}\tilde{\mathbf{k}}(x)$$

3. no recruitment subsidy, then  $\tilde{\beta} = \mathbf{0}$ .

Navigation icons

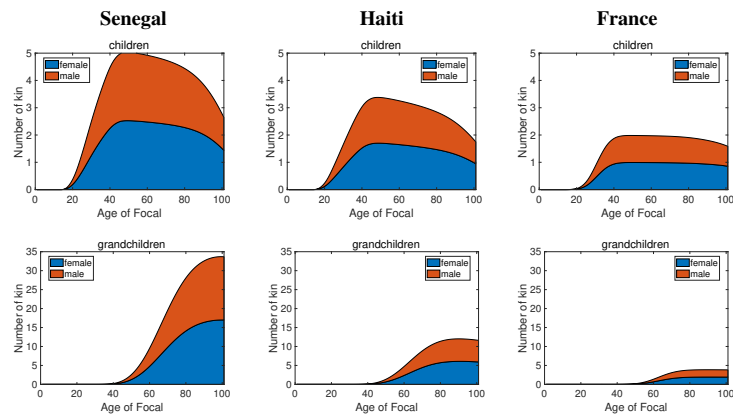
## The complete two-sex model

Symbol	Kin	initial condition $k_0$	Subsidy $\beta(x)$
$\tilde{a}$	children	0	$\tilde{F}\phi(x)$
$\tilde{b}$	grandchildren	0	$\tilde{F}\tilde{a}(x)$
$\tilde{c}$	great-grandchildren	0	$\tilde{F}\tilde{b}(x)$
$\tilde{d}$	parents	$\tilde{\pi}$	0
$\tilde{g}$	grandparents	$\sum_i \pi_{fm}(i) \tilde{d}(i)$	0
$\tilde{g}$	great-grandparents	$\sum_i \pi_{fm}(i) \tilde{g}(i)$	0
$\tilde{m}$	older siblings	$\sum_i \pi_f(i) \tilde{a}(i)$	0
$\tilde{n}$	younger siblings	0	$\tilde{F}^* \tilde{d}(x)$
$\tilde{p}$	nieces/nephews via older siblings	$\sum_i \pi_f(i) \tilde{b}(i)$	$\tilde{F}\tilde{m}(x)$
$\tilde{q}$	nieces/nephews via younger siblings	0	$\tilde{F}\tilde{n}(x)$
$\tilde{r}$	aunts/uncles older than mother	$\sum_i \pi_{fm}(i) \tilde{m}(i)$	0
$\tilde{s}$	aunts/uncles younger than mother	$\sum_i \pi_{fm}(i) \tilde{n}(i)$	$\tilde{F}^* \tilde{g}(x)$
$\tilde{t}$	cousins from aunts/uncles older than mother	$\sum_i \pi_{fm}(i) \tilde{p}(i)$	$\tilde{F}\tilde{r}(x)$
$\tilde{v}$	cousins from aunts/uncles younger than mother	$\sum_i \pi_{fm}(i) \tilde{q}(i)$	$\tilde{F}\tilde{s}(x)$

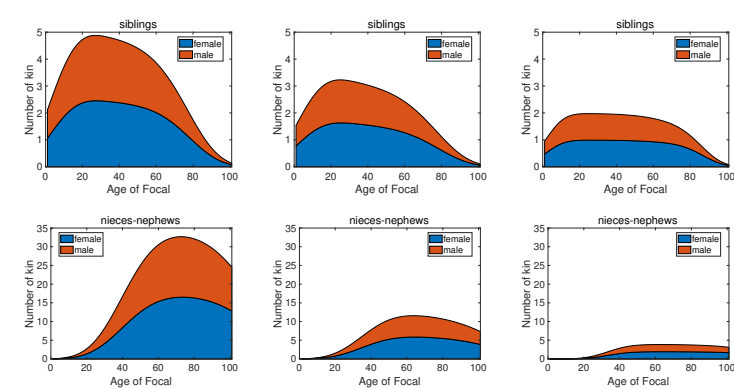


	Senegal	Haiti	France
TFR female	5.3	3.7	2.0
TFR male	11.0	5.1	2.0
age at maternity	29.8	30.0	30.1
age at paternity	44.1	37.0	33.5

## Female and male kin, through all lineages



## Female and male kin, through all lineages



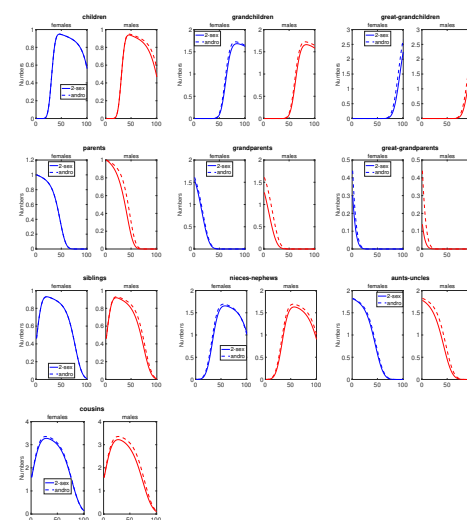
## Approximations to a two-sex model

$$\tilde{\mathbf{U}} = \left( \begin{array}{c|c} \mathbf{U}_f & \mathbf{0} \\ \hline \mathbf{0} & \mathbf{U}_m \end{array} \right) \quad \tilde{\mathbf{F}} = \left( \begin{array}{c|c} \bar{\alpha}\mathbf{F}_f & \bar{\alpha}\mathbf{F}_m \\ \hline \alpha\mathbf{F}_f & \alpha\mathbf{F}_m \end{array} \right)$$

- model 1: full two-sex model
- model 2: use female fertility for both sexes
- model 3: use female mortality for both sex
- model 4: males and females identical (the 'androgynous approximation')

Put appropriate survival and fertility matrices into  $\tilde{\mathbf{U}}$  and  $\tilde{\mathbf{F}}$

## The androgynous approximation<sup>3</sup>



<sup>3</sup>for Senegal

## Androgynous approximation: the GKP factors<sup>4</sup>

$$\left( \text{1-sex } \mathbf{k}(x) \right) \times \text{GKP factor} = \left( \text{androgynous 2-sex } \mathbf{k}(x) \right)$$

<sup>4</sup>Originally suggested by Goodman, Keyfitz, Pullum 1974

## Androgynous approximation: the GKP factors

