

Kinship models

An introduction to the analysis of population-level kinship structures

Diego Alburez-Gutiérrez[†]

[†]Kinship Inequalities Research Group,
Max Planck Institute for Demographic Research

International Institute for Population Sciences,
India, November 24-27, 2025



MAX PLANCK INSTITUTE
FOR DEMOGRAPHIC
RESEARCH

MAX-PLANCK-INSTITUT
FÜR DEMOGRAFISCHE
FORSCHUNG



Agenda

1. The Goodman-Keyfitz-Pullum kinship equations
2. Matrix kinship models
3. Implementations

What are kinship models?

- ① Kinship is an *emergent property* of demographic systems
- ② Simplified representation of interaction between reproduction and death
- ③ Not restricted to humans¹

- What are emergent properties?
- Can you think of other emergent properties in nature, society, or demography?

¹Coste, C. F. D., Bienvenu, F., Ronget, V., Ramirez-Loza, J.-P., Cubaynes, S., & Pavard, S. (2021). The kinship matrix: Inferring the kinship structure of a population from its demography (T. Coulson, Ed.). *Ecology Letters*, 24(12), 2750–2762. <https://doi.org/10.1111/ele.13854>

Formal models of kinship

Given a set of:

- ▶ age-specific fertility rates
- ▶ survival probabilities
- ▶ simplifying assumptions

The models produce:

- ① Number of (living/dead) kin
- ② Age distribution of relatives
- ③ From the point of view of an average member of the population ('Focal')

Focal: an average member of the population



Typology of kinship models

No	time	sex	state	reference
1	invariant	female	age	2
2	variant	female	age	3
3	invariant	two	age	4
4	invariant	female	multiple	5
5	variant	two	multiple	6

²Caswell, H. (2019). The formal demography of kinship: A matrix formulation. *Demographic Research*, 41, 679–712

³Caswell, H., & Song, X. (2021). The formal demography of kinship. III. kinship dynamics with time-varying demographic rates. *Demographic Research*, 45, 517–546

⁴Caswell, H. (2022). The formal demography of kinship IV: Two-sex models and their approximations. *Demographic Research*, 47, 359–396

⁵Caswell, H. (2020). The formal demography of kinship II: Multistate models, parity, and sibship. *Demographic Research*, 42, 1097–1146

⁶Williams, I., Alburez-Gutierrez, D., & DemoKin Team. (2023). *DemoKin*: 1.0.3. <https://CRAN.R-project.org/package=DemoKin>

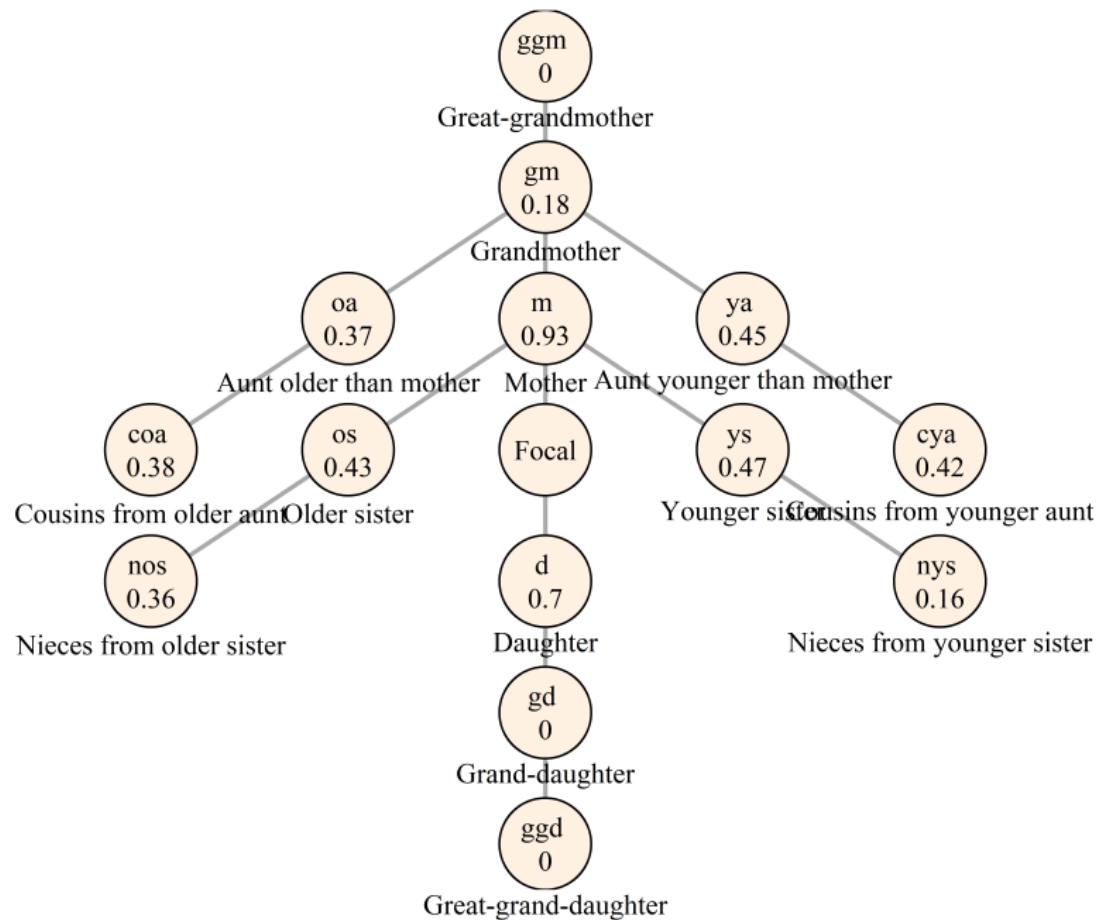
Model characteristics

Define the following model characteristics:

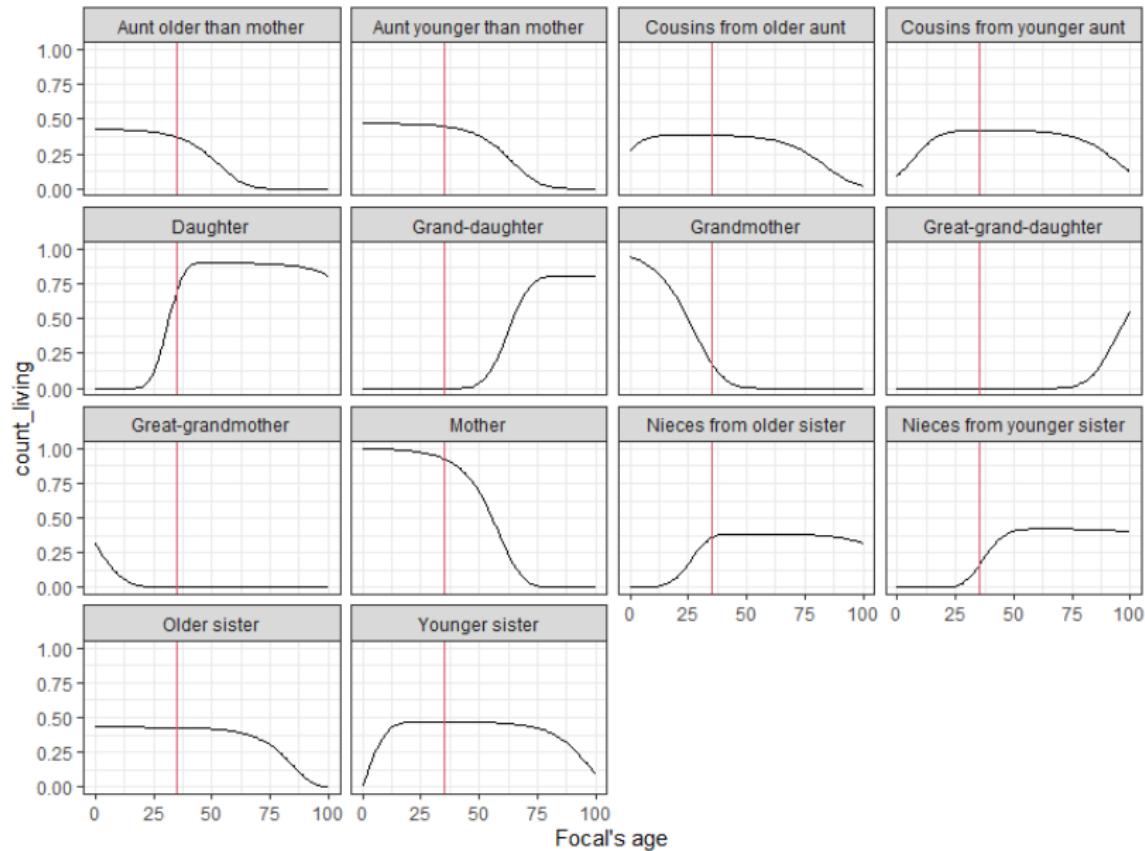
- ① Time-(in)variance
- ② One/two-sex models
- ③ (Multi)state models

The Goodman-Keyfitz-Pullum kinship equations

The tree of life



Expected number of kin



Daughters

$B_1(a)$ is the expected number of living daughters in a time-invariant female-only population⁷:

$$B_1(a) = \int_{\alpha}^a m(x)l(a-x) dx \quad (1)$$

where:

- ▶ $m(x)$ are fertility rates of mothers
- ▶ $l(a-x)$ are survival probabilities of daughters

⁷ Goodman, L. A. (1974). Family Formation and the Frequency of Various Kinship Relationships. *Theoretical Population Biology*, 27.

Daughters

If $a = 20$ and $\alpha = 15$; then:

$$B_1(20) \approx \sum_{15}^{20} m(x)/(20 - x)$$

So...

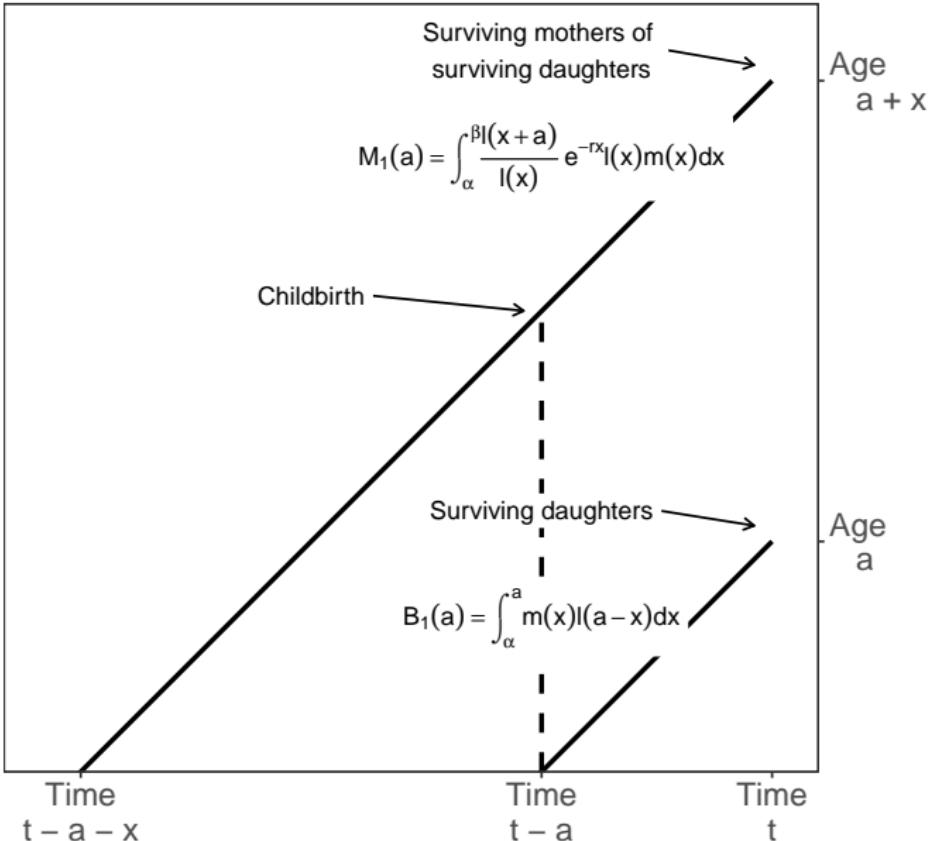
$$B_1(20) \approx m(15)/(5) + m(16)/(4) + m(17)/(3) \dots$$

Granddaughters

$B_1(a)$ is the expected number of living granddaughters in a time-invariant female-only population⁸:

$$B_2(a) = \int_{\alpha}^a m(x) \int_{\alpha}^{a-x} l(y)m(y)l(a-x-y) dy dx \quad (2)$$

⁸ Goodman, L. A. (1974). Family Formation and the Frequency of Various Kinship Relationships. *Theoretical Population Biology*, 27.



Mothers

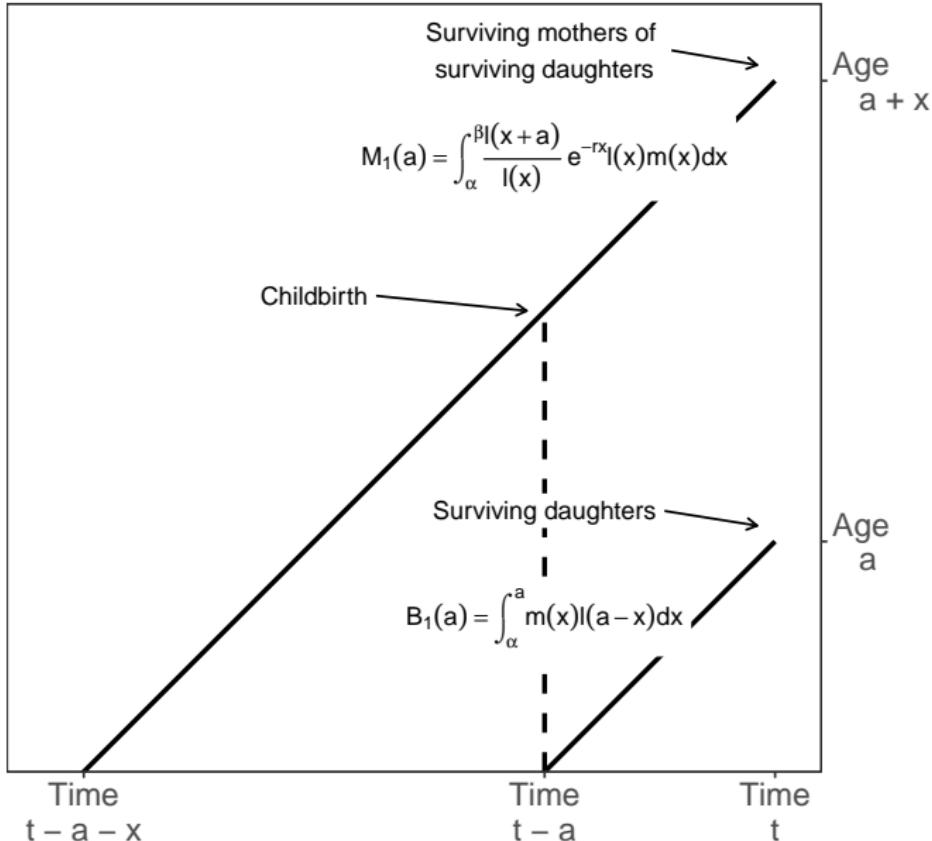
$M_1(a)$ is the probability of having a living mother in a time-invariant female-only population⁹:

$$M_1(a) = \int_{\alpha}^{\beta} \underbrace{\frac{I(x+a)}{I(x)}}_{\text{prob of surviving from } x \text{ to } a+x} \times \underbrace{W(x)}_{\text{age distribution of mothers}} dx. \quad (3)$$

where:

- ▶ $W(x) = e^{-rx} I(x) m(x)$ is the age distribution of mothers
- ▶ $I(x)$ are survival probabilities
- ▶ $m(x)$ are fertility rates
- ▶ r is the population growth rate
- ▶ $\alpha-\beta$ is the reproductive period

⁹ Goodman, L. A. (1974). Family Formation and the Frequency of Various Kinship Relationships. *Theoretical Population Biology*, 27.



Grandmothers

$M_2(a)$ is the expected number of living grandmothers in a time-invariant female-only population¹⁰:

$$M_2(a) = \int_{\alpha}^{\beta} \underbrace{M_1(a)}_{\text{prob of having living mother}} \times \underbrace{W(x)}_{\text{age distribution of mothers}} dx. \quad (4)$$

¹⁰ Goodman, L. A. (1974). Family Formation and the Frequency of Various Kinship Relationships. *Theoretical Population Biology*, 27.

Great-grandmothers

$M_3(a)$ is the expected number of living great-grandmothers in a time-invariant female-only population¹¹:

$$M_3(a) = \int_{\alpha}^{\beta} \underbrace{M_2(a)}_{\text{number of grandmother}} \times \underbrace{W(x)}_{\text{age distribution of mothers}} dx. \quad (5)$$

¹¹ Goodman, L. A. (1974). Family Formation and the Frequency of Various Kinship Relationships. *Theoretical Population Biology*, 27.

Sisters

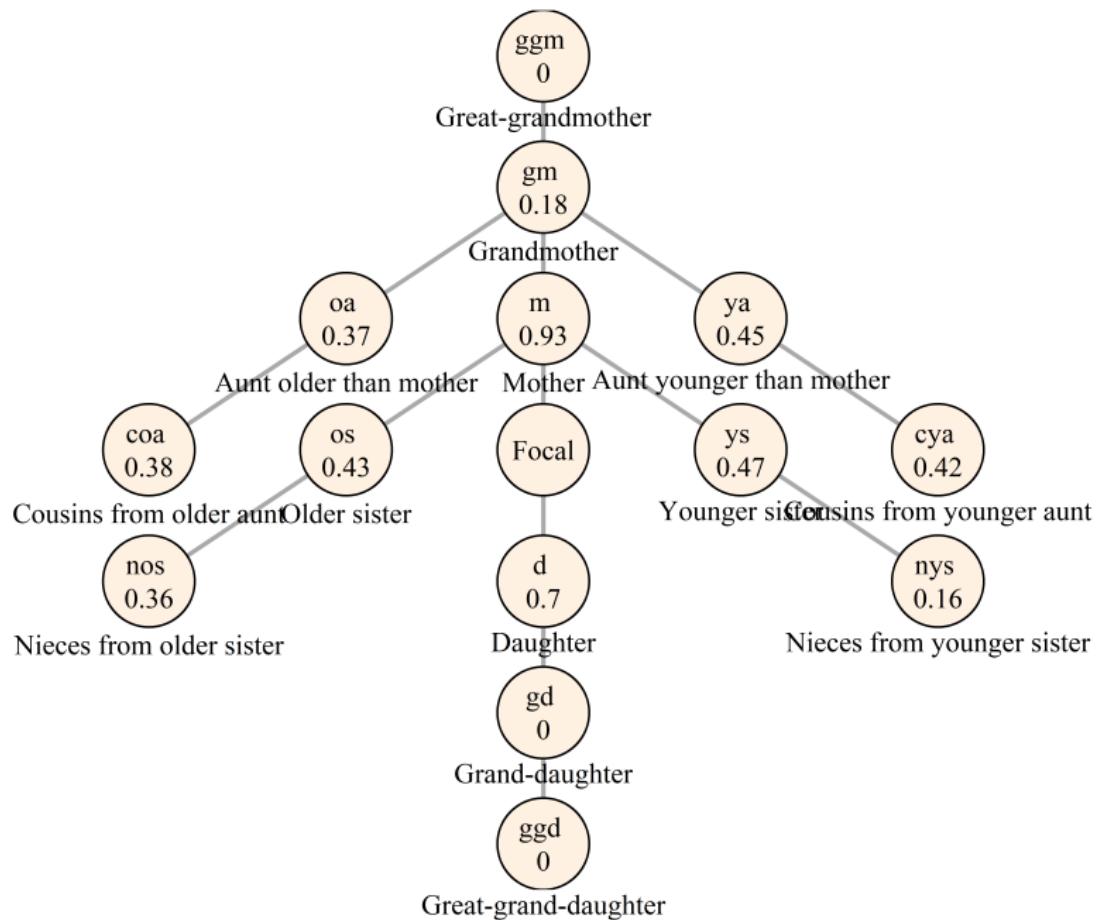
$B_1(a)$ is the expected number of living older sisters in a time-invariant female-only population¹²:

$$S^{old}(a) = \int_{\alpha}^{\beta} \int_{\alpha}^x m(y) I(a + x - y) W(x) dy dx \quad (6)$$

$$S^{young}(a) = \int_{\alpha}^{\beta} \int_0^a \left[\frac{I(x+u)}{I(x)} \right] m(x+u) I(a-u) du W(x) dx \quad (7)$$

¹² Goodman, L. A. (1974). Family Formation and the Frequency of Various Kinship Relationships. *Theoretical Population Biology*, 27.

Why do we model younger and older sisters/kin separately?



Demographic subsidy

"New members of the population arise not from reproduction of current members, but from elsewhere"¹³

- ▶ Can you think of other instances of 'subsidy' in demography?

¹³Caswell, H. (2019). The formal demography of kinship: A matrix formulation. *Demographic Research*, 41, 679–712

Break

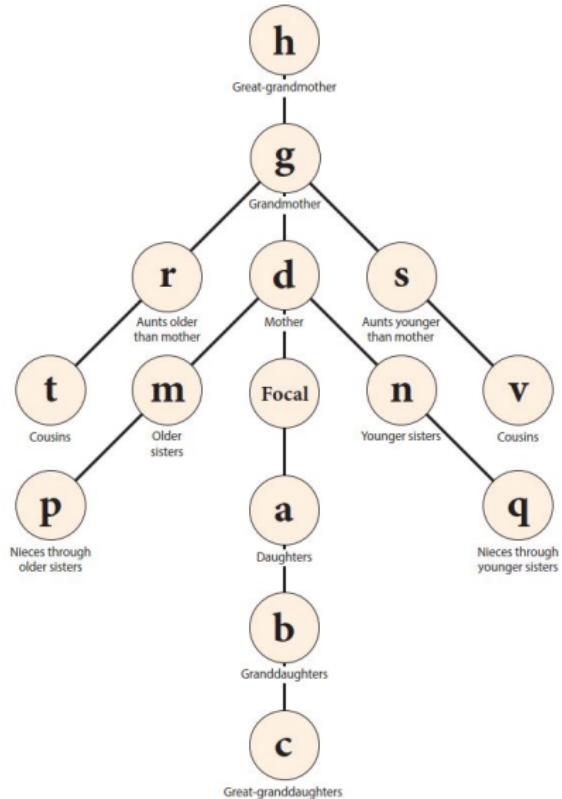
Matrix kinship models

From recursive equations to matrix operations

- ① The relatives of Focal constitute a population
- ② They can be modelled using traditional projection methods
- ③ Matrix operations provide an efficient implementation



The tree of life (2)



Implementation: time-invariant, one-sex models¹⁴

The models are of the general form:

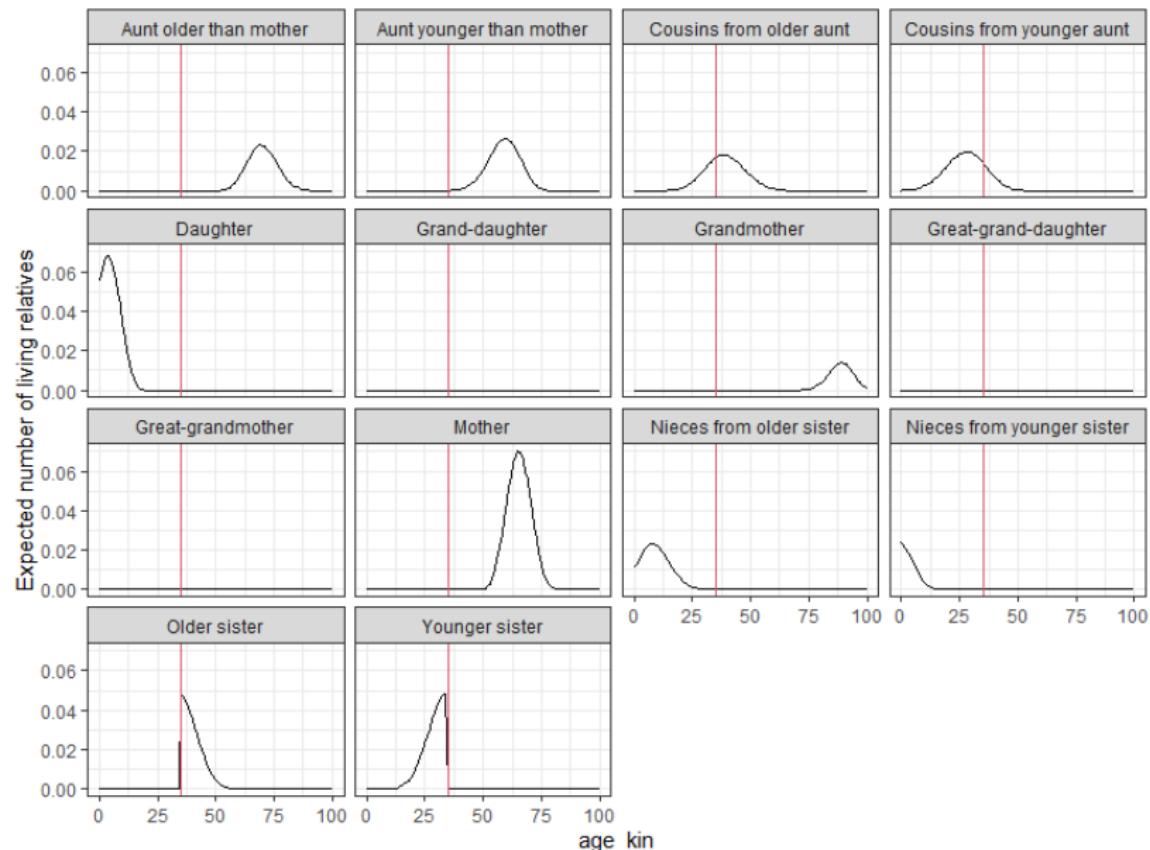
$$\underbrace{\mathbf{k}(x+1)}_{\substack{\text{age structure of kin} \\ \text{at Focal's age } x+1}} = \underbrace{\mathbf{U} \mathbf{k}(x)}_{\substack{\text{ageing and survival} \\ \text{of existing kin}}} + \underbrace{\begin{cases} \mathbf{0} \\ \mathbf{F} \mathbf{k}^*(x) \end{cases}}_{\substack{\text{new kin members} \\ \text{added to the population}}}.$$

where:

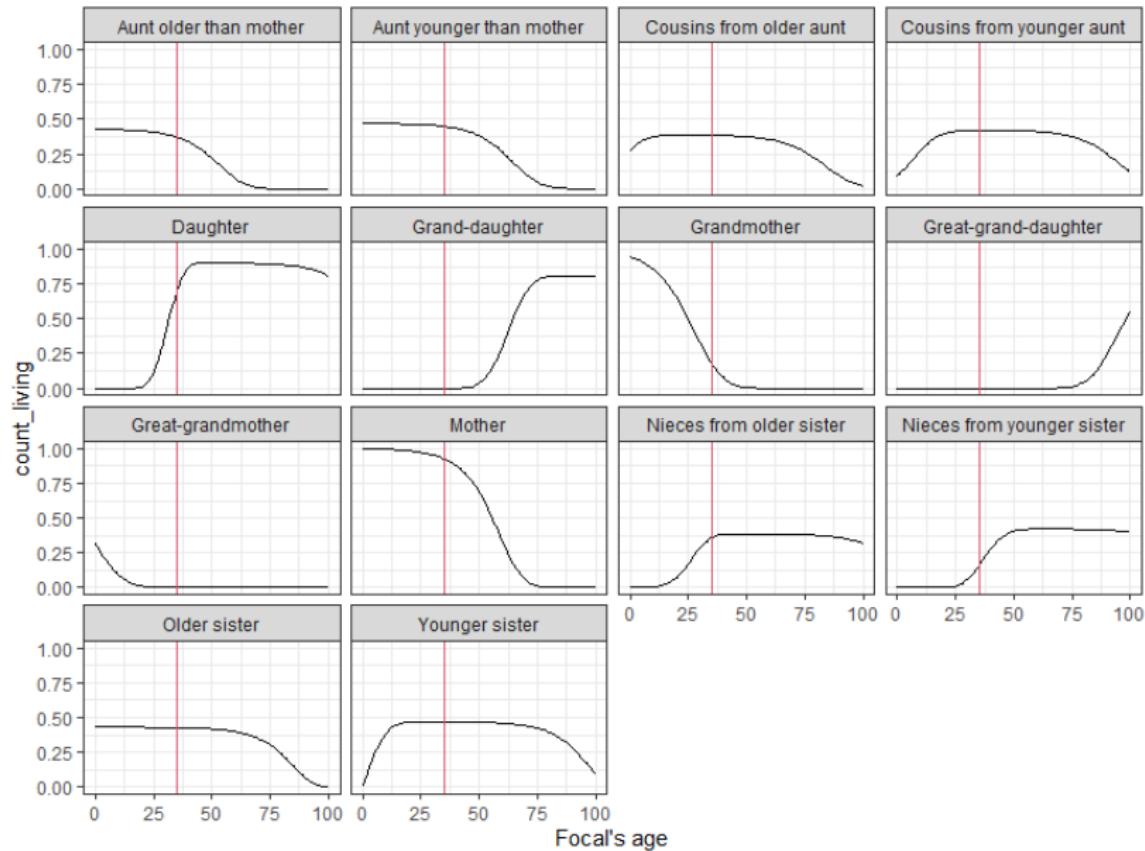
- ▶ **U** a matrix with survival probabilities in the subdiagonal
- ▶ **F** a matrix with fertility rates in the first row

¹⁴Caswell, H. (2019). The formal demography of kinship: A matrix formulation. *Demographic Research*, 41, 679–712

Age distributions of kin



Expected number of kin



Daughters

Daughters (**a**) are the result of the reproduction of Focal:

$$\underbrace{\mathbf{a}(x+1)}_{\text{age structure of daughters at Focal's age } x+1} = \underbrace{\mathbf{U} \mathbf{a}(x)}_{\text{ageing and survival of existing daughters}} + \underbrace{\mathbf{F} \mathbf{e}_x}_{\text{new daughters (subsidy)}}. \quad (8)$$

$$\mathbf{a}(0) = \mathbf{0}.$$

where:

- ▶ **U** is a matrix with survival probabilities in the subdiagonal
- ▶ **F** is a matrix with fertility rates in the first row
- ▶ **F e_x** is the subsidy vector
- ▶ **e_x** is the unit vector for age x
- ▶ $\mathbf{a}(0)$ is the distribution of daughters at Focal's birth

Mothers

The population of mothers (**d**) of Focal consists of at most a single individual:

$$\underbrace{\mathbf{d}(x+1)}_{\text{age structure of mothers at Focal's age } x+1} = \underbrace{\mathbf{U d}(x)}_{\text{ageing and survival of existing mothers}} + \underbrace{0.}_{\text{new mothers (subsidy)}} \quad (9)$$

$$d(0) = \pi.$$

where:

- ▶ $b(0)$ is the distribution of mothers at Focal's birth
- ▶ π is the distribution of ages of mothers in the population

All models¹⁵

Table 1: Summary of the components of the kin model given in equations (4) and (5)

Symbol	Kin	Initial condition	Subsidy $\beta(x)$
a	daughters	0	$F_a x$
b	granddaughters	0	$F_a(x)$
c	great-granddaughters	0	$F_b(x)$
d	mothers	π	0
g	grandmothers	$\sum_i \pi_i d(i)$	0
h	great-grandmothers	$\sum_i \pi_i g(i)$	0
m	older sisters	$\sum_i \pi_i a(i)$	0
n	younger sisters	0	$F_d(x)$
p	nieces via older sisters	$\sum_i \pi_i b(i)$	$F_m(x)$
q	nieces via younger sisters	0	$F_n(x)$
r	aunts older than mother	$\sum_i \pi_i m(i)$	0
s	aunts younger than mother	$\sum_i \pi_i n(i)$	$F_g(x)$
t	cousins from aunts older than mother	$\sum_i \pi_i p(i)$	$F_r(x)$
v	cousins from aunts younger than mother	$\sum_i \pi_i q(i)$	$F_s(x)$

¹⁵ Caswell, H. (2019). The formal demography of kinship: A matrix formulation. *Demographic Research*, 41, 679–712

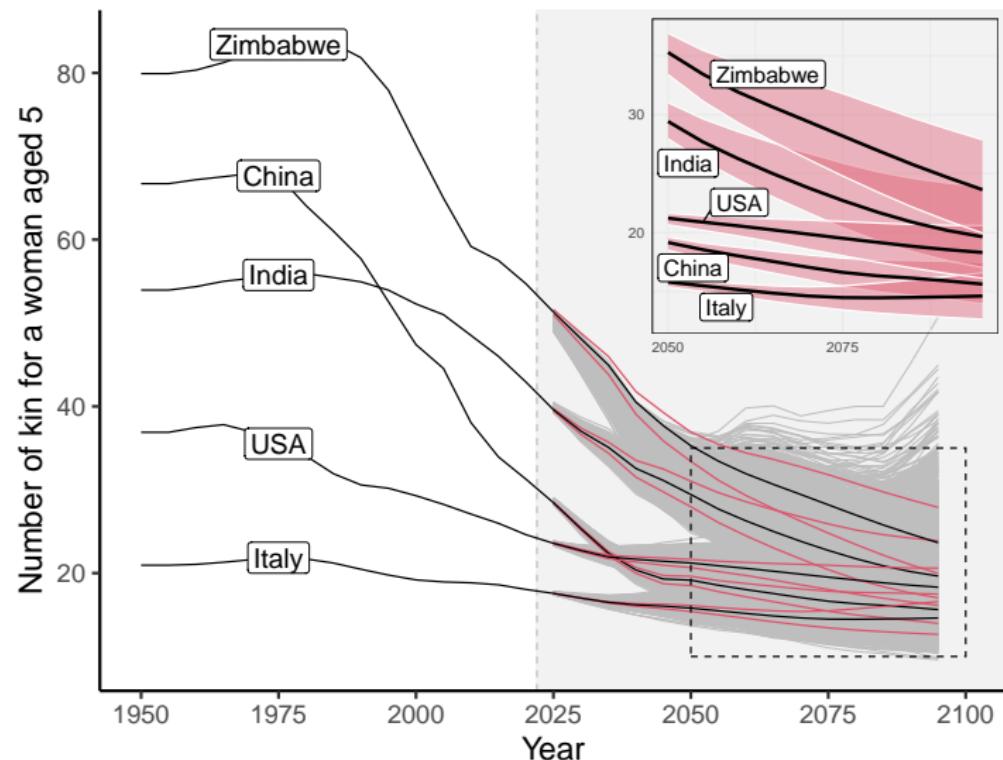
Consider a baby born in Spain in 1950...

- ① How old were her grandparents when she was born, on average?
- ② How many living children did she have on her 70th birthday?
- ③ How many grandchildren?

Break

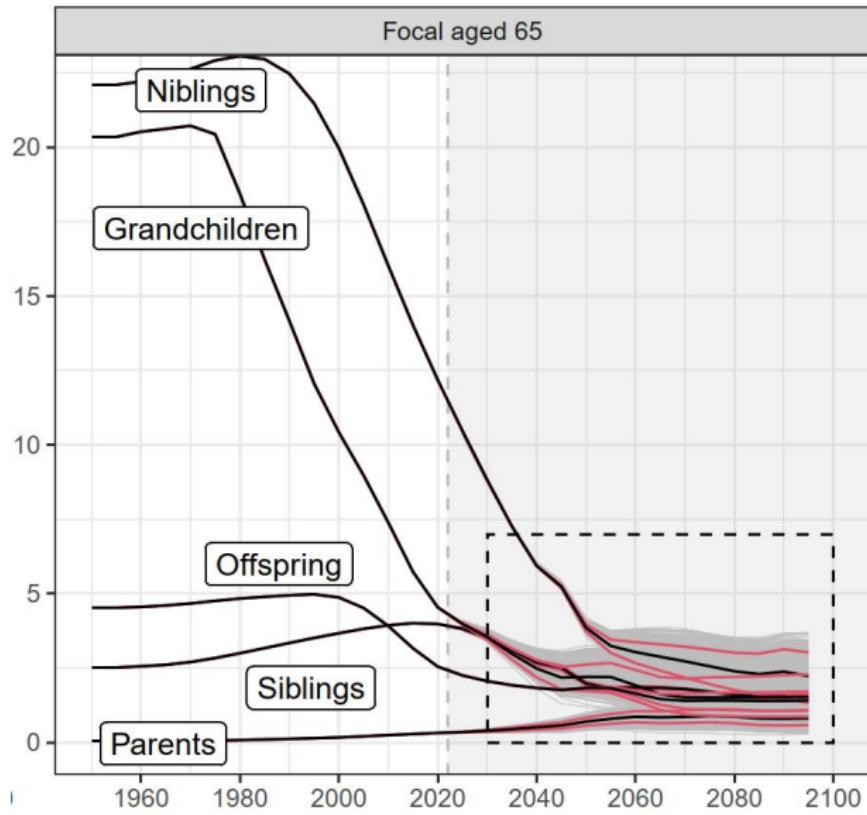
Implementations

Total number of kin (all kin combined) for a 5yo woman¹⁶



¹⁶ Alburez-Gutierrez, D., Williams, I., & Caswell, H. (2023). Projections of human kinship for all countries. *Proceedings of the National Academy of Sciences*, 120(52), e2315720120. <https://doi.org/10.1073/pnas.2315720120>

Number of living kin for a 65yo in China



Models vs reality

Discuss:

- ① What is the relationship between demographic models and reality?
- ② Would we expect kinship models to agree with 'empirical' observations of kinship?
- ③ Where can we find empirical data on kinship availability?