

The Impact of Sexual Mixing By Age on Sexually Transmitted Infection Models

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Background

- Mathematical models of sexually transmitted infection (STI) dynamics are commonly used to evaluate interventions. For example, recently used to inform HPV vaccination policy recommendations.
- In STI models, a commonly used representation of sexual partner choice is the **assortative-proportionate (A-P) mixing structure**:
 - a single parameter defines the spectrum from assortative (only mixing with one's own subgroup) to proportionate (choosing a partner solely based on the proportion of the population in each subgroup and the number of partnerships they offer).
- Problems with A-P: not flexible enough to capture age mixing patterns
- We propose **empirical mixing**: use standard statistical methods to define probability distributions for mixing behavior.

Objectives

- Compare fit of A-P and empirical age mixing structures to survey data.
- Quantify effects of different mixing structures on model-predicted HPV vaccine benefits.

Methods

- We analyzed the British population probability survey National Survey on Sexual Attitudes and Lifestyles (Johnson, et al. 2015)
 - We used heterosexual age preference data, based on respondents' most recent sexual partners (up to 3).

Assortative-Proportionate Versus Empirical Mixing

Age Mixing Estimation: A-P

- Use maximum likelihood estimation to find optimal degree of assortativeness.

Age Mixing Estimation: Empirical

- Model the age preference data as a collection of probability distributions - one for each age group
- Partner age is heteroscedastic with respect to chooser age.
- Model variance:
 - Linear regression of chooser's age versus age of partner, predict mean partner age for each age group.
 - Model the squared residuals (and assorted transformations) as linear functions of chooser's age, and predict variance.

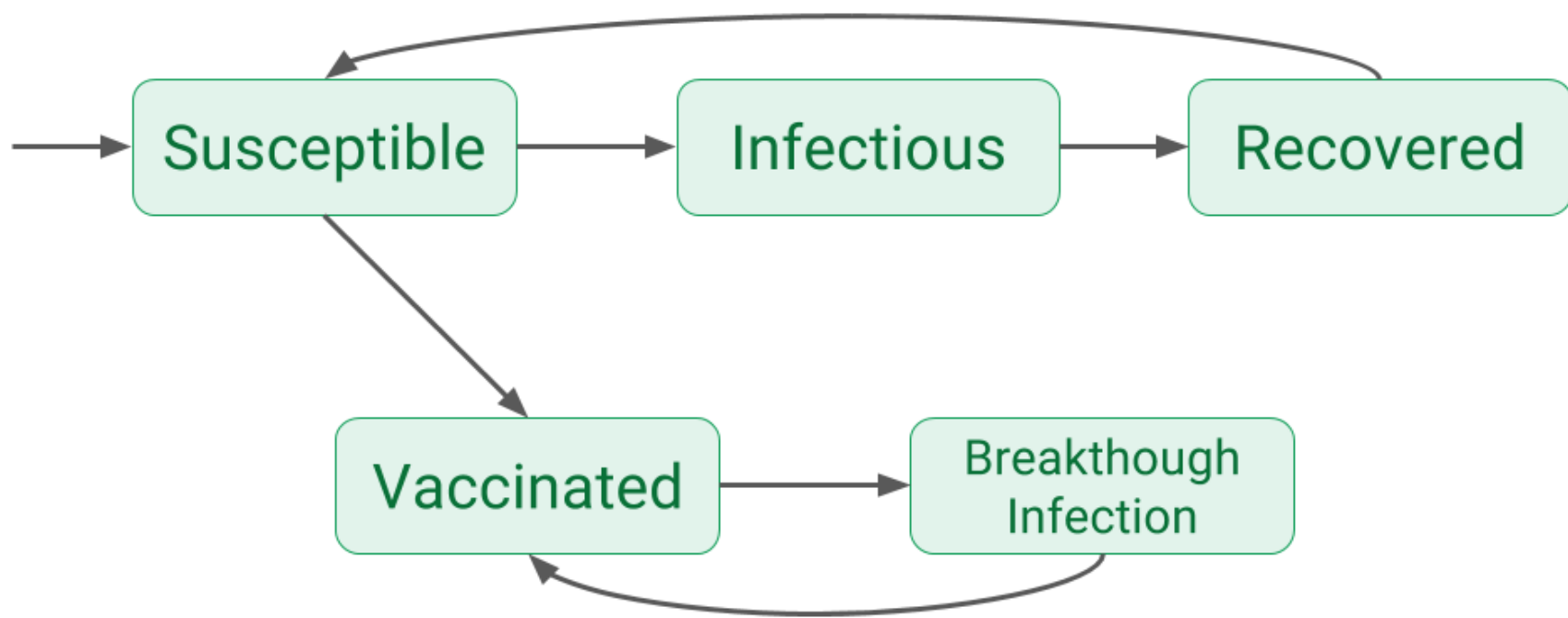
Likelihood Comparison

Calculate likelihood of Natsal-3 age mixing data under:

- A-P mixing
- Empirical mixing, with:
 - Linear: $\text{variance} \sim \text{chooser age}$
 - Square Root: $\sqrt{\text{variance}} \sim \text{chooser age}$
 - Log: $\log(\text{variance}) \sim \text{chooser age}$
 - Constant: $\text{variance} = \hat{\sigma}^2$

Mathematical Model of HPV

- Age- and sexual-activity-structured compartmental model
- Assume life-long vaccine protection.
- Model diagram on right, with age and sexual activity omitted for clarity.



Results: Age Mixing Structures

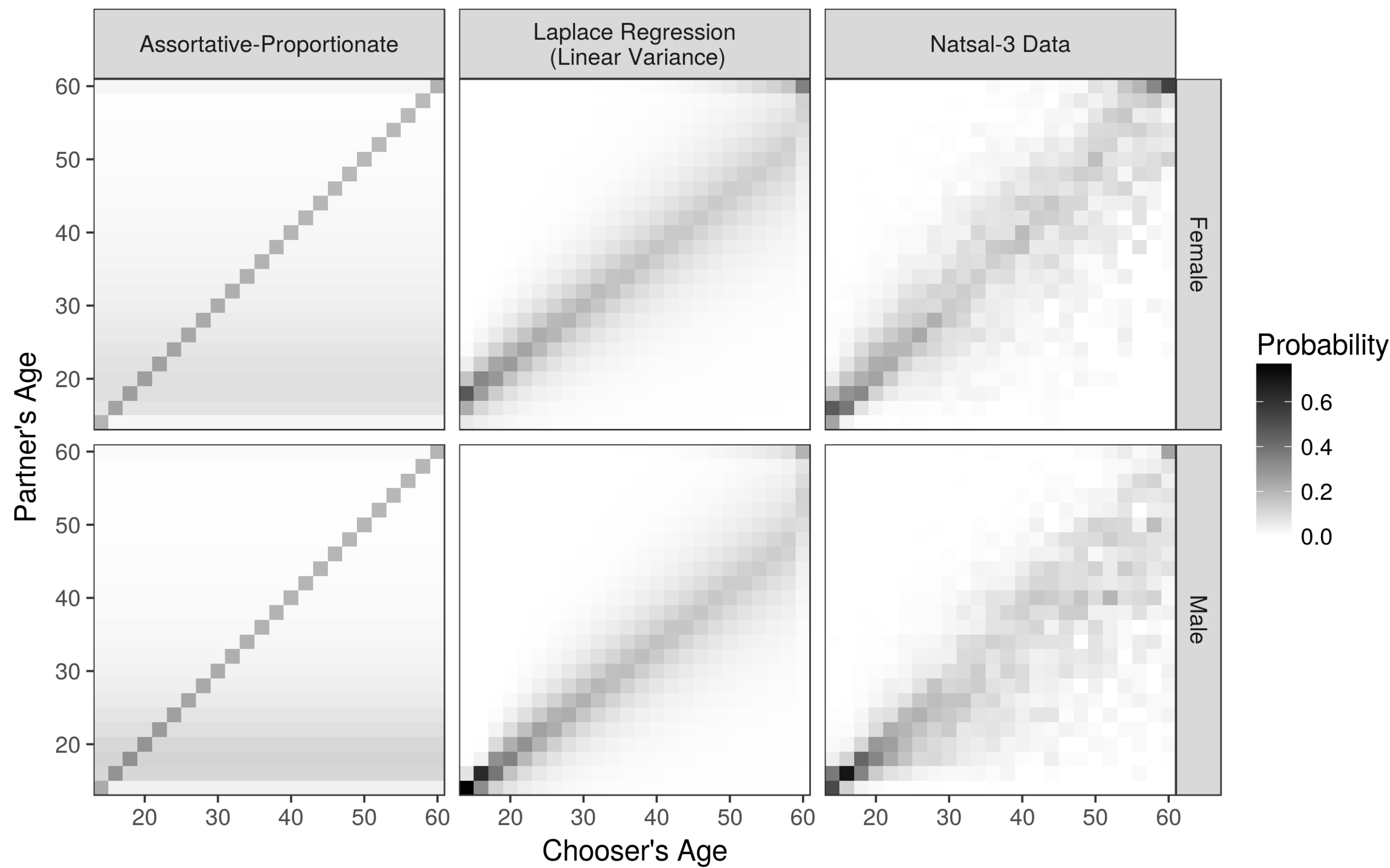


Figure 1: From left to right: the A-P structure, a regression model with Laplace distributed errors and variance as a linear function of age, and the Natsal data. Note the constant probabilities in rows of the A-P structure.

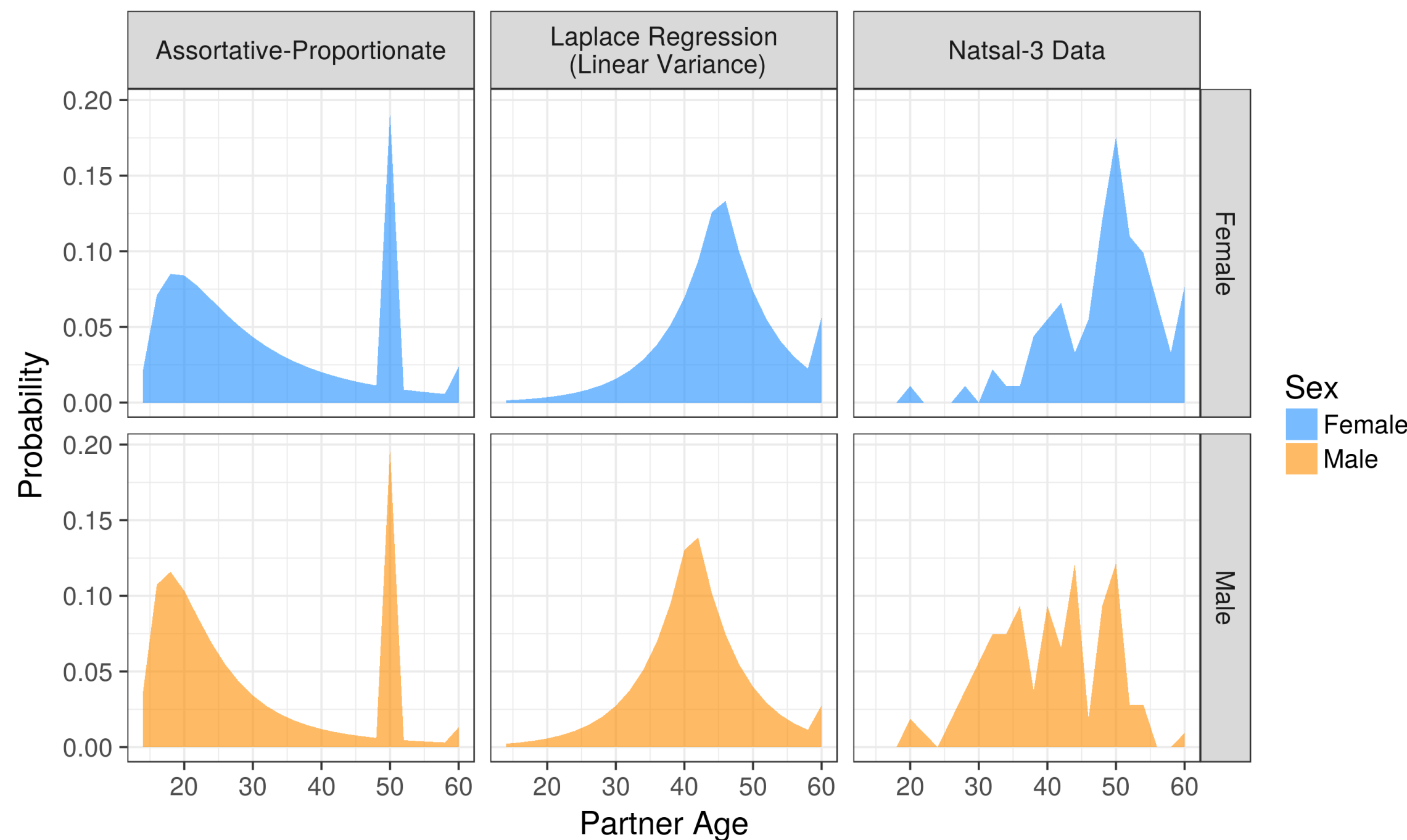


Figure 2: The partner age distributions for 50-year-old male and females.

Results: Likelihood Comparison

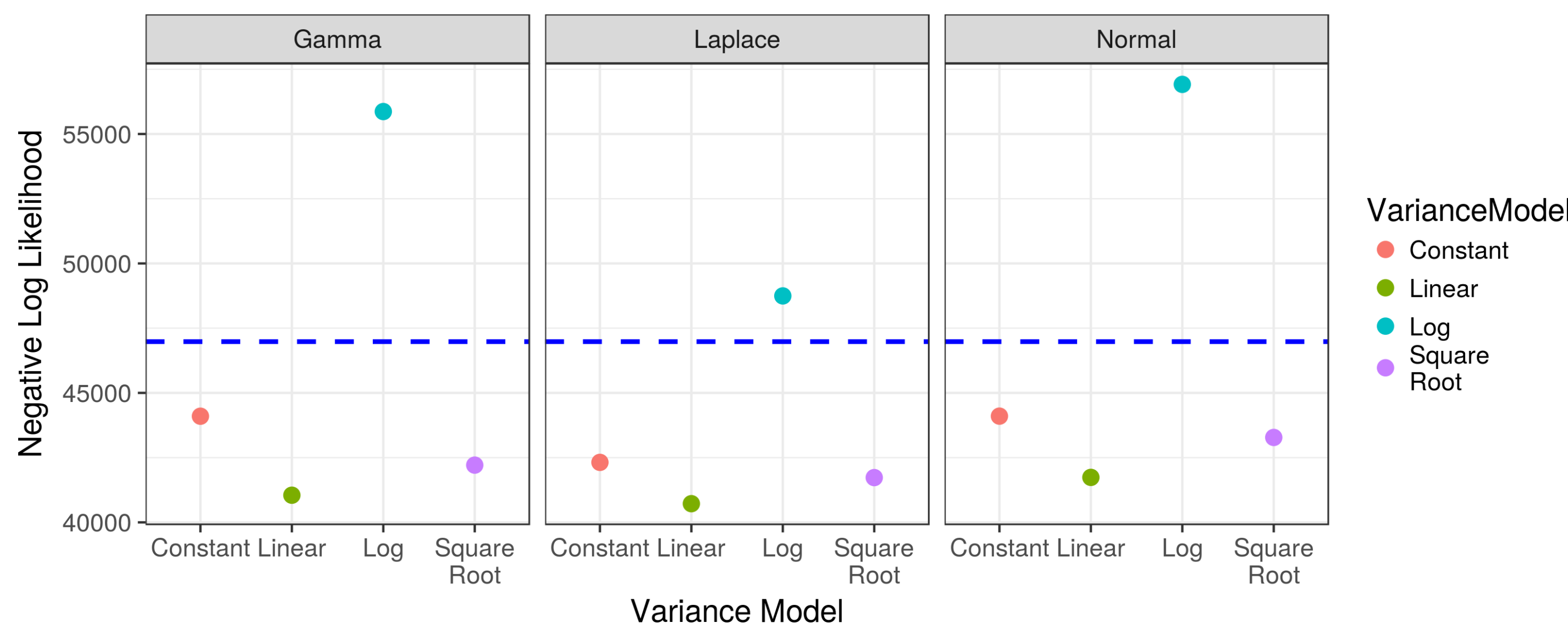


Figure 3: Likelihood of Natsal age mixing data under different mixing assumptions. The dotted blue line is the likelihood under A-P

Results: Extending Vaccination to 26-39-year-old Females

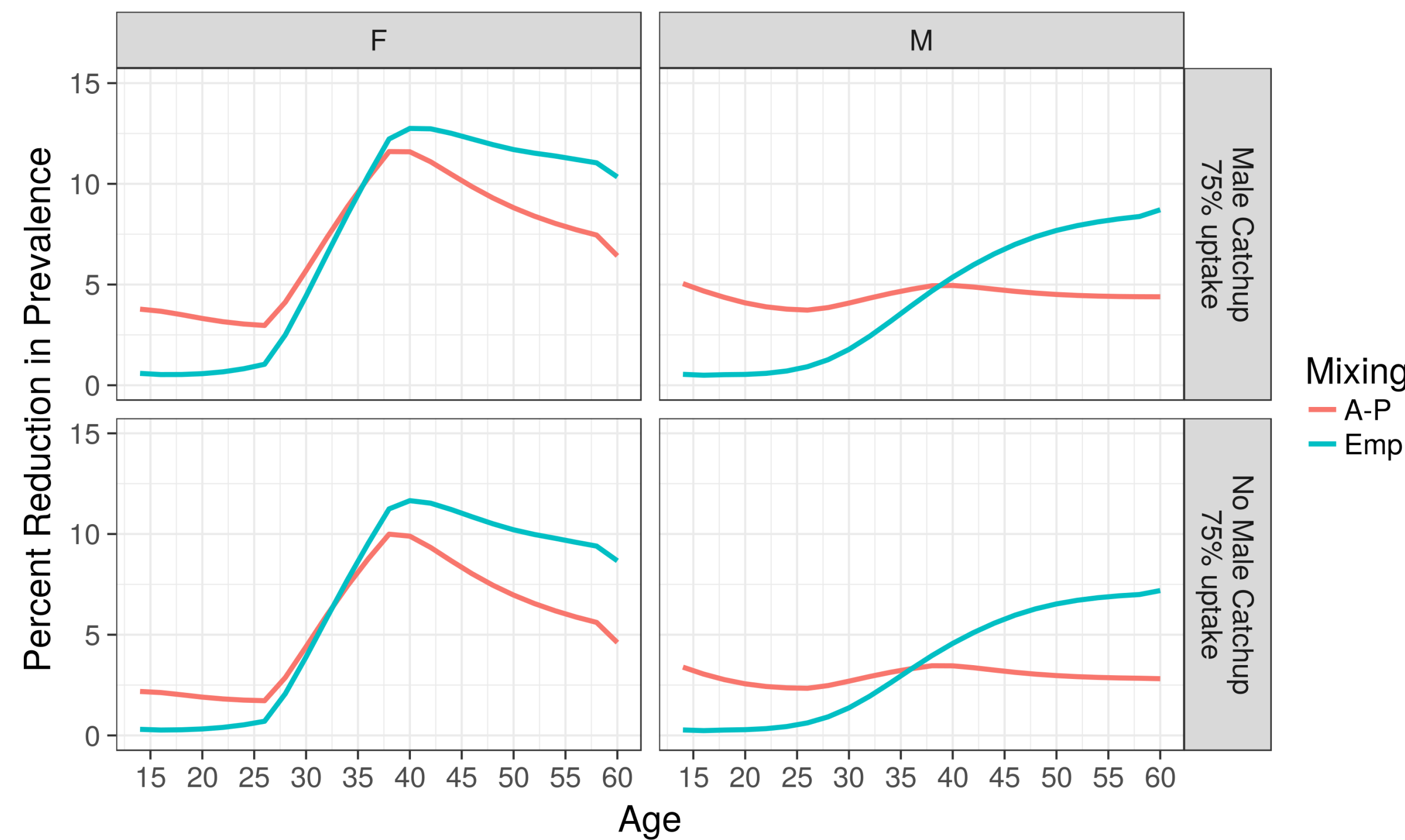


Figure 4: Relative reduction in prevalence due to extending vaccination, by sex, age, and male catch-up scenario.

Summary Measures

Table 1: Ratio of percentage reduction in prevalence predicted by A-P model to that predicted by empirical model.

Sex	<26 y/o >41 y/o	
	M	F
M	760.8	58.1
F	551.5	72.6

- The A-P model predicts a reduction in prevalence **7 times greater** (young males) and **5 times greater** (young females) than empirical model prediction.
- A-P model predicts slightly more than one half of the reduction for older males, and slightly less than 3/4 of the benefit for older females.

Conclusions

- Standard regression models fit Natsal-3 age mixing data better than the A-P mixing structure.
- The A-P mixing structure overestimates the sexual connection between those above and below 30.
- The choice of mixing structure impacts model-estimated vaccine benefits.
- A model with an A-P mixing structure:
 - Overestimates reduction in HPV prevalence for younger individuals
 - Underestimates reduction in prevalence for older individuals

Future Directions

- Analyze other sexual behavior surveys (National Survey of Family Growth, etc.) and compare results.
- Extend HPV model to include cervical cancer and other outcomes, for cost-effectiveness analysis.
- Calibrate model to HPV prevalence and incidence and examine degree to which calibration corrects for mixing structure differences.
- Develop empirical mixing estimates for non-heterosexual sexual mixing.