

Mixture Cure Rate Models for the Analysis of Survival Times in the COVID-19 Scenario

Bruna Wundervald, Rafael De Andrade Moral, Andrew Parnell, and the Hamilton Institute COVID-19 research group



**Maynooth
University**
National University
of Ireland Maynooth



Summary

- Introduction
- Methods
 - Mixture Cure Rate Models
 - GAMLSS Models
- Data
- Results
- Conclusions and Final Remarks



Introduction

- ❄ The pandemic of the new coronavirus has become an enormous health challenge worldwide
- ❄ The COVID-19 infection can be asymptomatic, but a significant proportion of the cases require hospitalization, including intensive care unit admission
- ❄ A few factors have been linked to the risk of more severe disease and death, such as age and sex

Survival modeling : helps us understand the correlation between human and environmental factors (age, sex) and the survival probabilities of each person



Methods

- ❄ Choice: motivated by the imbalance present in the data (deaths versus censored observations)
- ❄ Let T be the time to some specific event (death, for example), treated as a non-negative random variable
- ❄ In survival analysis, we commonly use the survival function $S(t) = P(T > t)$, that is nothing more than $S(t) = 1 - F(t)$, where $F(t)$ is the cumulative distribution function of T



Mixture Cure Rate Models

- ❄ Two sub-populations: one with a standard failure rate (**cured fraction**) and one with the same rate **added** with a subject-specific parametric failure rate
- ❄ The mixture survival function has the form

$$S_{pop}(t|\mathbf{x}, \mathbf{z}, \mathbf{n}) = (1 - \pi(\mathbf{z}))S(t|\mathbf{x}, \mathbf{n} = 1) + \pi(\mathbf{z}),$$

where $\pi(\mathbf{z})$ is the cured proportion, $S(t|\mathbf{x}, \mathbf{n} = 1)$ is the survival function of susceptible (uncured) individuals, and \mathbf{x} and \mathbf{z} are the covariates associated with π and S , respectively.



GAMLSS Models

- ❄ GAMLSS: generalized additive models for location, scale, and shape [2]
- ❄ Semi-parametric *Weibullcr* [1] distribution, for which the survival function will be

$$S_{pop}(t|\mu, \sigma, \nu) = \nu + (1 - \nu) \exp \left\{ - \left[\frac{t\Gamma(1/\sigma + 1)}{\mu} \right]^\sigma \right\}.$$

- ❄ The parameters $\theta = (\mu, \sigma, \nu)$ = (location, scale, shape) are estimated through a regression model that allows for complex regression structures



Data

→ Time until death or the end of the study (censoring)

Explanatory variables:

→ Age

→ Sex

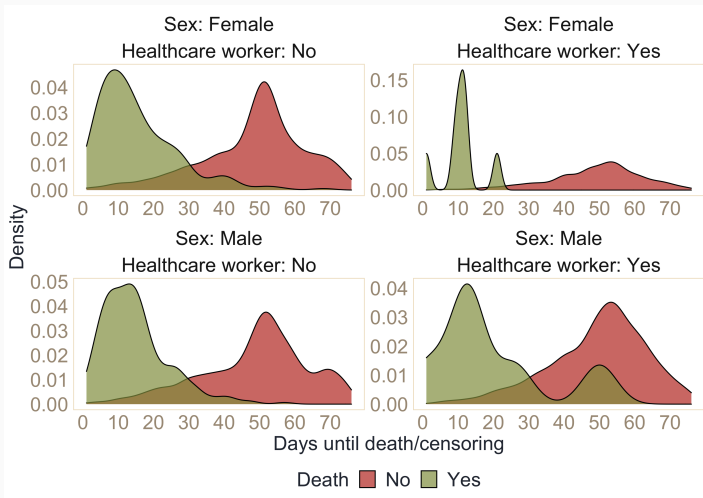
→ ICU status (yes or no)

→ Healthcare worker status (yes or no)

Note: the results presented here are from 'faked data', because the true Irish cases data is confidential



Data





Results

- ✧ Best model: selected by minimising the AIC
- ✧ Final model: does not use the same covariates in all parameters, and includes polynomial and interaction terms

$$\log(\mu_i) = \beta_{01} + \beta_{11}\text{age} + \beta_{21}\text{age}^2 + \beta_{31}\text{sex} + \beta_{41}\text{HW} + \beta_{51}\text{ICU} + \beta_{61}(\text{age} \times \text{sex}),$$

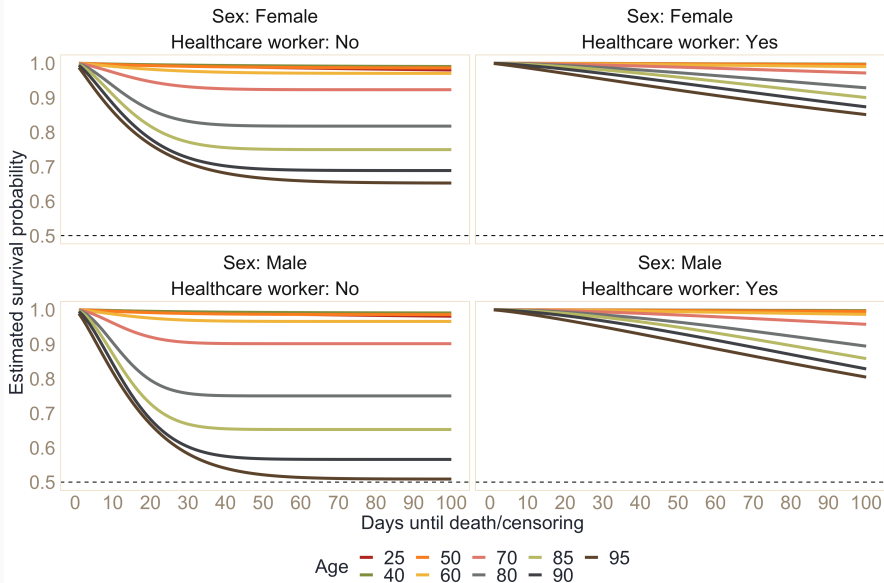
$$\log(\sigma_i) = \beta_{02} + \beta_{12}\text{sex} + \beta_{22}\text{age} + \beta_{32}\text{age}^2 + \beta_{42}\text{age}^3,$$

$$\text{logit}(\nu_i) = \beta_{03} + \beta_{13}\text{age} + \beta_{23}\text{age}^2 + \beta_{33}\text{age}^3 + \beta_{43}\text{ICU} + \beta_{53}\text{sex} + \beta_{63}(\text{age} \times \text{sex}),$$

where $\text{logit}(x) = \log(x/(1+x))$.

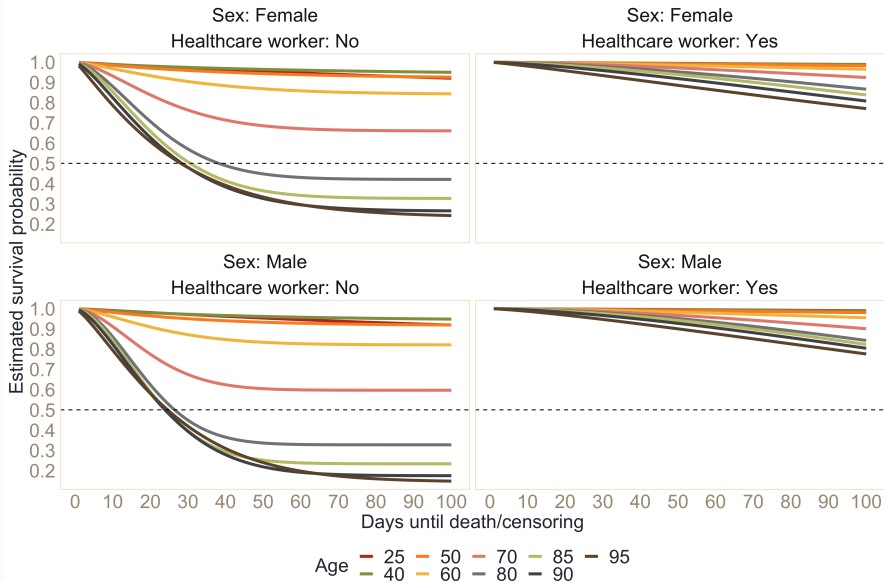
Model results per sex and healthcare worker status, non ICU patients

-- Dashed line represents the survival probability equal to 0.5

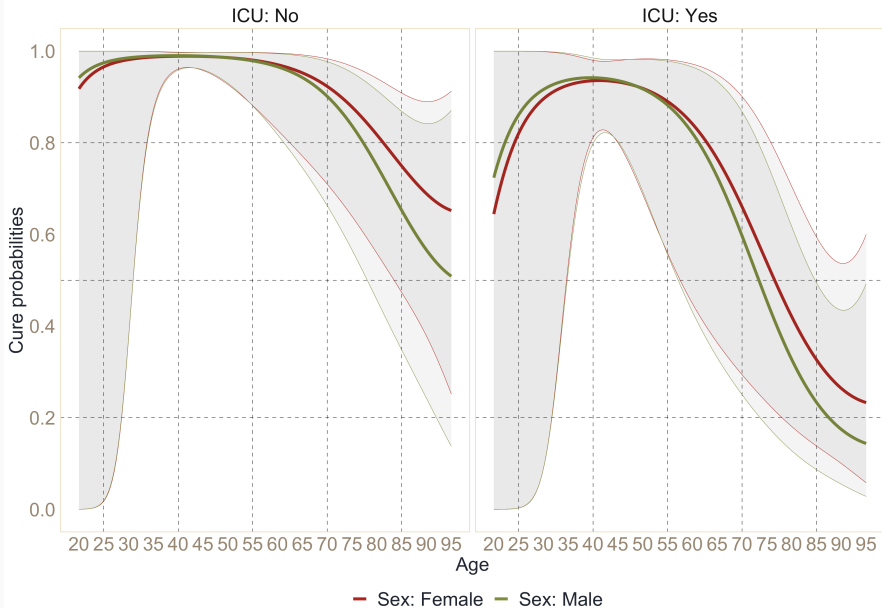


Model results per sex and healthcare worker status, ICU patients

-- Dashed line represents the survival probability equal to 0.5



Cure probabilities per age and sex





Conclusions and Final Remarks

- Age and sex seem to play the most important roles in the decrease of survival chances
- ICU patients, particularly males, have the lowest survival probabilities
- Healthcare workers appear to have a low mortality for COVID-19
- Many factors were not captured here, such as time and spatially variant dynamics and other influential covariates





Acknowledgements

This work was supported by Science Foundation Ireland and their emergency COVID-19 funding portal via grant number 20/COV/0081



<https://www.hamilton.ie/covid19/>

References

-  Thiago G Ramires et al. “A new semiparametric Weibull cure rate model: fitting different behaviors within GAMLSS”. In: *Journal of Applied Statistics* 46.15 (2019), pp. 2744–2760.
-  D Mikis Stasinopoulos, Robert A Rigby, et al. “Generalized additive models for location scale and shape (GAMLSS) in R”. In: *Journal of Statistical Software* 23.7 (2007), pp. 1–46.

Thanks!