

# Joint Models with Multi-State Processes

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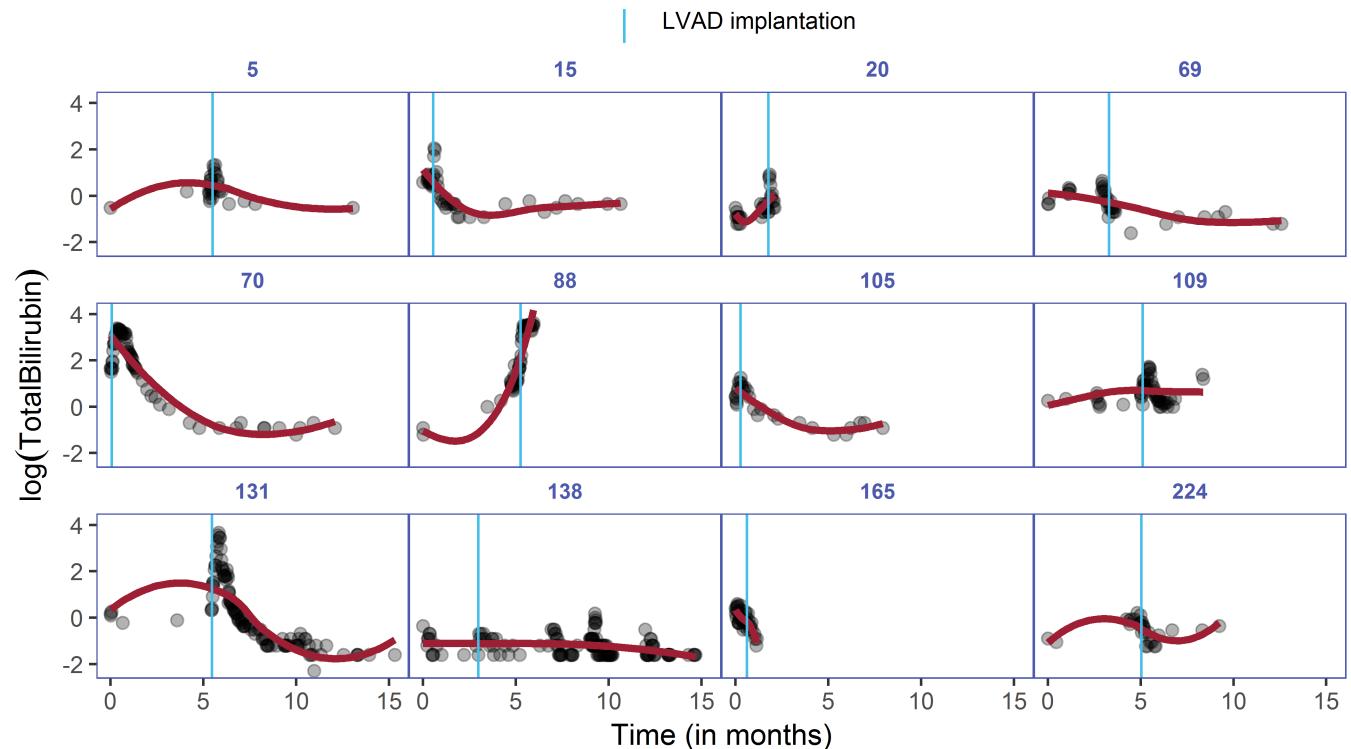
# Motivation

- Outcomes in follow-up studies
- Multiple longitudinal responses:
  - biomarkers, blood values
- Times of transitions between states of interest:
  - relapse, clinical complications, death
- Intermediate events
  - (re)intervention

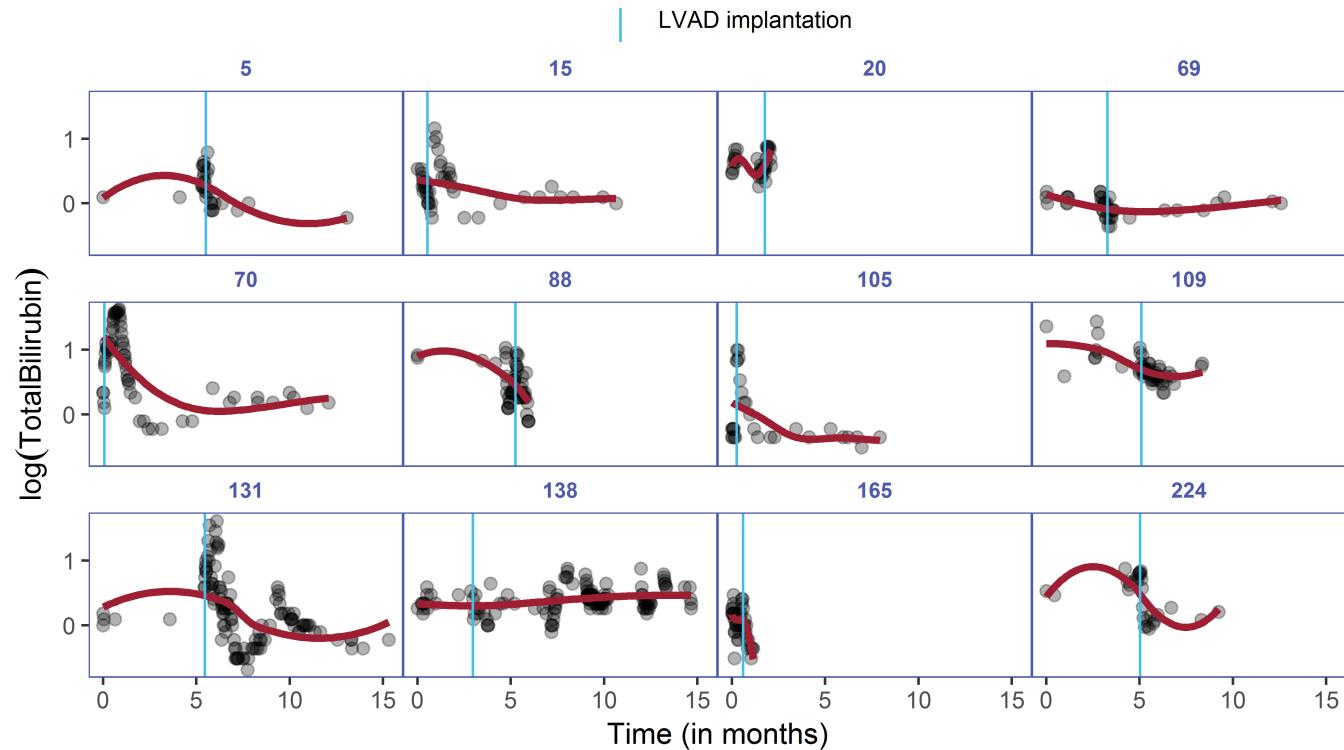
# Motivation

- 232 patients who were followed-up after heart failure
  - All patients received a **L**eft **V**entricular **A**ssist **D**evice during follow-up
- Longitudinal outcomes:
  - Total bilirubin (mg/dl)
  - Creatinine (mg/dl)
- Events of interest:
  - Complications: Thrombosis, Embolic Events, Dialysis
  - Transplantation/Death

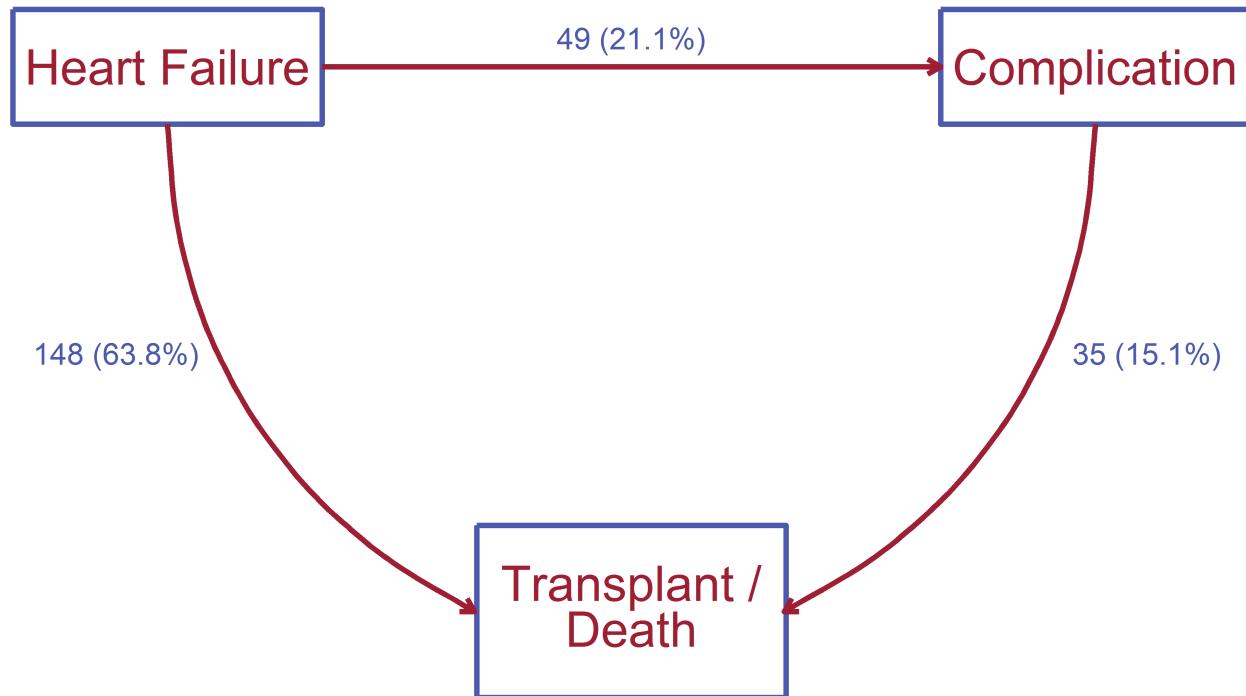
# Motivation



# Motivation



# Motivation



# Research Goals

- Investigate the association between the longitudinal and multi-state processes
  - Functional form of the association for **each transition**
  - Strength of the association for **each transition**
  - Selection of the association structure for **each transition**
- Investigate the impact of LVAD on the evolution of the markers

# Methods

## Multivariate generalized linear mixed-effects submodel:

$$g_k [E\{y_{ki}(t) \mid b_{ki}\}] = \eta_{ki}(t) = \begin{cases} x_{ki}^\top(t) \beta_k + z_{ki}^\top(t) b_{ki}, & 0 < t < \rho_i \\ x_{ki}^\top(t) \beta_k + z_{ki}^\top(t) b_{ki} + \tilde{x}_{ki}^\top(t) \tilde{\beta}_k + \tilde{z}_{ki}^\top(t) \tilde{b}_{ki}, & t \geq \rho_i, \end{cases}$$

- $y_{ki}$ : repeated measurements of the  $k^{th}$  outcome for the  $i^{th}$  subject
- $b^\top = (b_{1i}, \dots, b_{Ki}, \tilde{b}_{1i}, \dots, \tilde{b}_{Ki})^\top \sim \mathcal{N}(0, D)$
- $\rho_i$ : time of occurrence of the intermediate event
- $t_{i+} = \max(0, t_{ki} - \rho_i)$ : time relative to the occurrence of the intermediate event

# Methods

$$\eta_{ki}(t) = \begin{cases} x_{ki}^\top(t) \beta_k + z_{ki}^\top(t) b_{ki}, & 0 < t < \rho_i \\ x_{ki}^\top(t) \beta_k + z_{ki}^\top(t) b_{ki} + \tilde{x}_{ki}^\top(t) \tilde{\beta}_k + \tilde{z}_{ki}^\top(t) \tilde{b}_{ki}, & t \geq \rho_i, \end{cases}$$

- $\tilde{x}_{ki}^\top(t) \tilde{\beta}_k + \tilde{z}_{ki}^\top(t) \tilde{b}_{ki}$ : May be any function of  $t_{i+}$

# Methods

## Multi-state submodel:

- $S = \{1, \dots, M\}$ : state space
- $T_i = (T_{i1}, \dots, T_{im_i})^\top$ : vector of observed times for the  $i^{th}$  subject
- $\delta_i = (\delta_{i1}, \dots, \delta_{im_i})^\top$ : vector of observed transition indicators for the  $i^{th}$  subject

$$\lambda_{hl}^i(t) = \begin{cases} \lambda_{hl,0}(t) \exp \left[ W_{hl,i}^S {}^\top \gamma_{hl} + \sum_{k=1}^K \sum_{j=1}^J f_j \{ \eta_{ki}(t), \alpha_{kj} \} \right], & 0 < t < \rho_i, \\ \lambda_{hl,0}(t) \exp \left[ W_{hl,i}^S {}^\top \gamma_{hl} + \sum_{k=1}^K \sum_{j=1}^J f_j \{ \eta_{ki}(t), \alpha_{kj} \} \right], & t \geq \rho_i. \end{cases}$$



# Association Structures

Current value:  $f_j \{ \eta_{ki} (t), \alpha_{kj} \} = \eta_{ki} (t) \cdot \alpha_{kj}$

# Association Structures

Current slope:  $f_j \left\{ \eta_{ki} (t), \alpha_{kj} \right\} = \frac{d}{dt} \eta_{ki} (t) \cdot \alpha_{kj}$

# Association Structures

Cumulative effect:  $f_j \left\{ \eta_{ki} (t), \alpha_{kj} \right\} = \int_0^t \eta_{ki} (s) ds \cdot \alpha_{kj}$

# Feature Selection

- Which features of the longitudinal outcomes are associated with transition intensities?

#Biomarkers  $\times$  #Features  $\times$  #Transitions

- High dimensional parameter space
- (Potentially) high correlation among features from the same biomarker
- Feature selection  $\rightarrow$  difficult

# Bayesian Shrinkage

- Shrinkage priors for variable selection
- Priors that shrink towards zero:
  - Bayesian ridge
  - Bayesian horseshoe
  - Bayesian lasso
  - ...
- E.R. Andrinopoulou and D. Rizopoulos (2016) investigated the performance of local shrinkage priors

# Bayesian Shrinkage

- Local priors:

- $\alpha_{jk} \mid \tau_{jk}^2 \sim \mathcal{N}(0, \tau_{jk}^2)$

- $\tau_{jk}^2$ : Local shrinkage

- Global-local priors:

- $\alpha_{jk} \mid \tau_{jk}^2, \lambda^2 \sim \mathcal{N}(0, \tau_{jk}^2 \lambda^2)$

- $\tau_{jk}^2$ : Local shrinkage

- $\lambda^2$ : Global shrinkage

# Bayesian Shrinkage

- Global-local horseshoe prior
  - Double inverse-gamma prior leads to  $C^+(0, 1)$

$$\alpha_{jk} \mid \tau_{jk}^2, \lambda^2 \sim \mathcal{N}\left(0, \tau_{jk}^2 \lambda^2\right)$$

$$\tau_{jk}^2 \mid \nu_{jk}^2 \sim \mathcal{IG}\left(\frac{1}{2}, \frac{1}{\nu_{jk}}\right)$$

$$\lambda^2 \mid \xi \sim \mathcal{IG}\left(\frac{1}{2}, \frac{1}{\xi}\right)$$

$$\nu_{1k}, \dots, \nu_{JK}, \xi \sim \mathcal{IG}\left(\frac{1}{2}, 1\right)$$

- **Properties:**

- Strong spike that leads to severe shrinkage near 0
- Narrow tails that allow strong signals to remain strong

# Bayesian Shrinkage

- Global-local ridge prior

$$\begin{aligned}\alpha_{jk} \mid \tau_{jk}^2, \lambda^2 &\sim \mathcal{N}(0, \tau_{jk}^2 \lambda^2) \\ \tau_{jk}^2 &\sim \mathcal{IG}(\frac{1}{2}, 1) \\ \lambda^2 &\sim \mathcal{IG}(\frac{1}{2}, 1)\end{aligned}$$

- **Properties:**

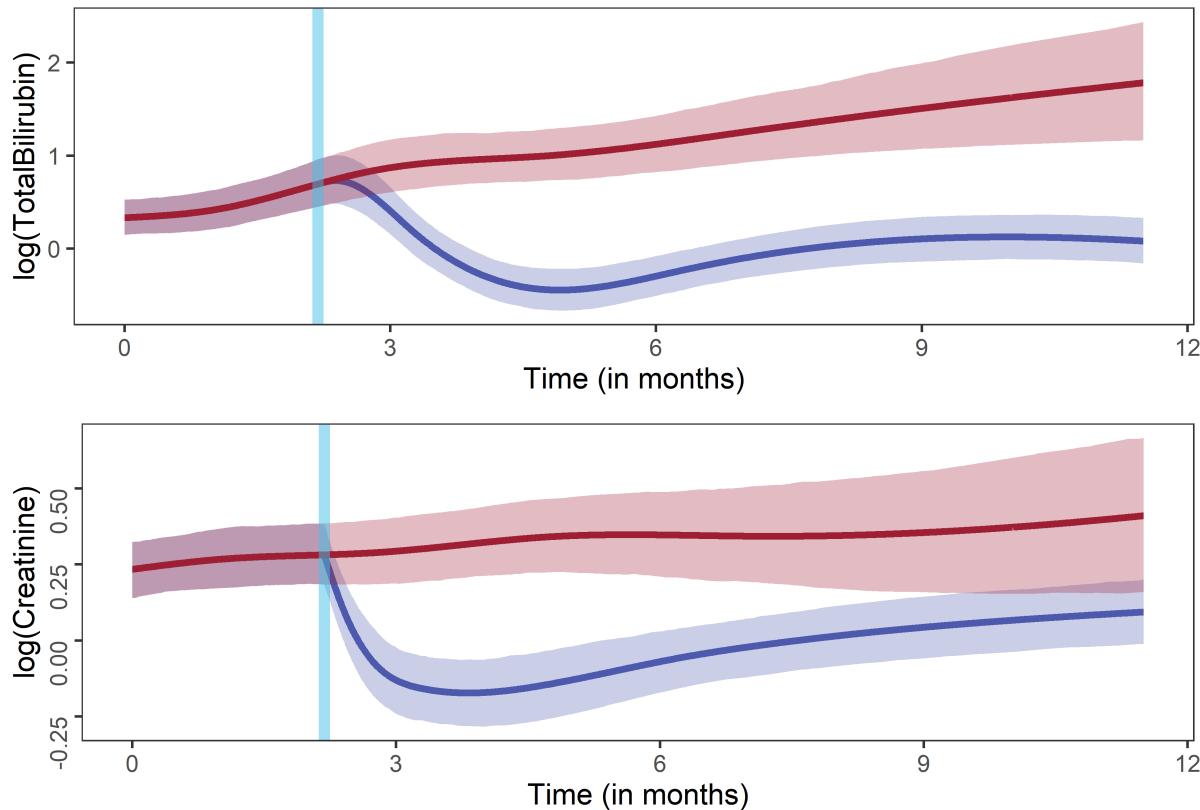
- Less shrinkage near 0
- Heavier tails than horseshoe

# Analysis

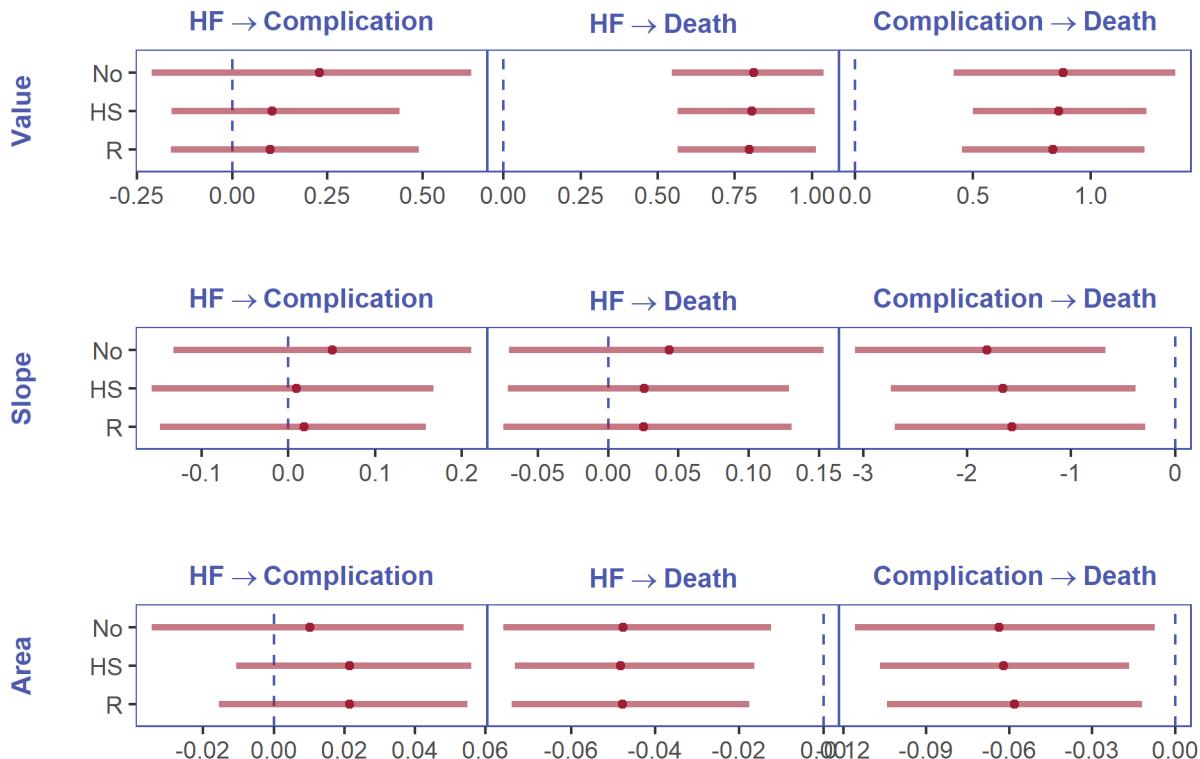
- **Longitudinal submodels:**

- **Fixed-effects:** natural cubic splines for time and time relative to **LVAD** implantation, adjusted for BMI, age, sex and etiology
- **Random-effects:** natural cubic splines for time and time relative to **LVAD** implantation
- **Multi-state submodel:**
  - **State-specific covariates:** BMI, age, sex and etiology
  - **Markers' features:** value, slope and cumulative effect association with **each transition**

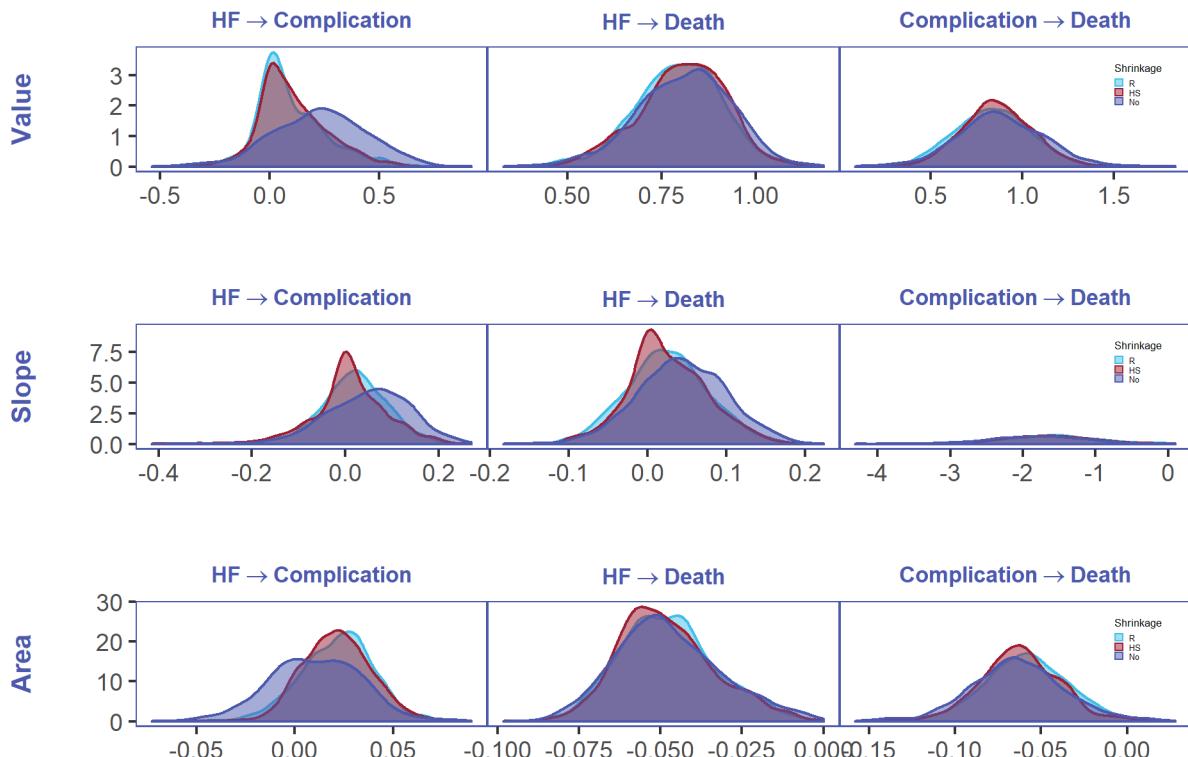
# Results



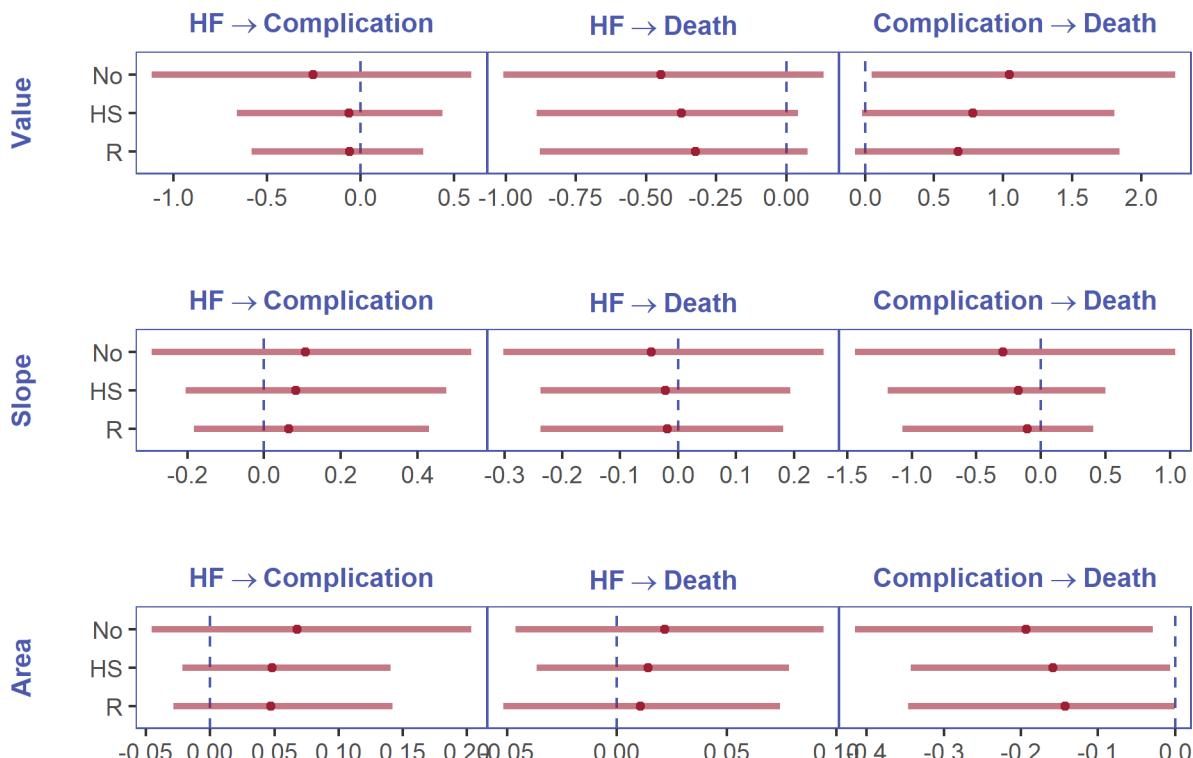
# Results: Bilirubin



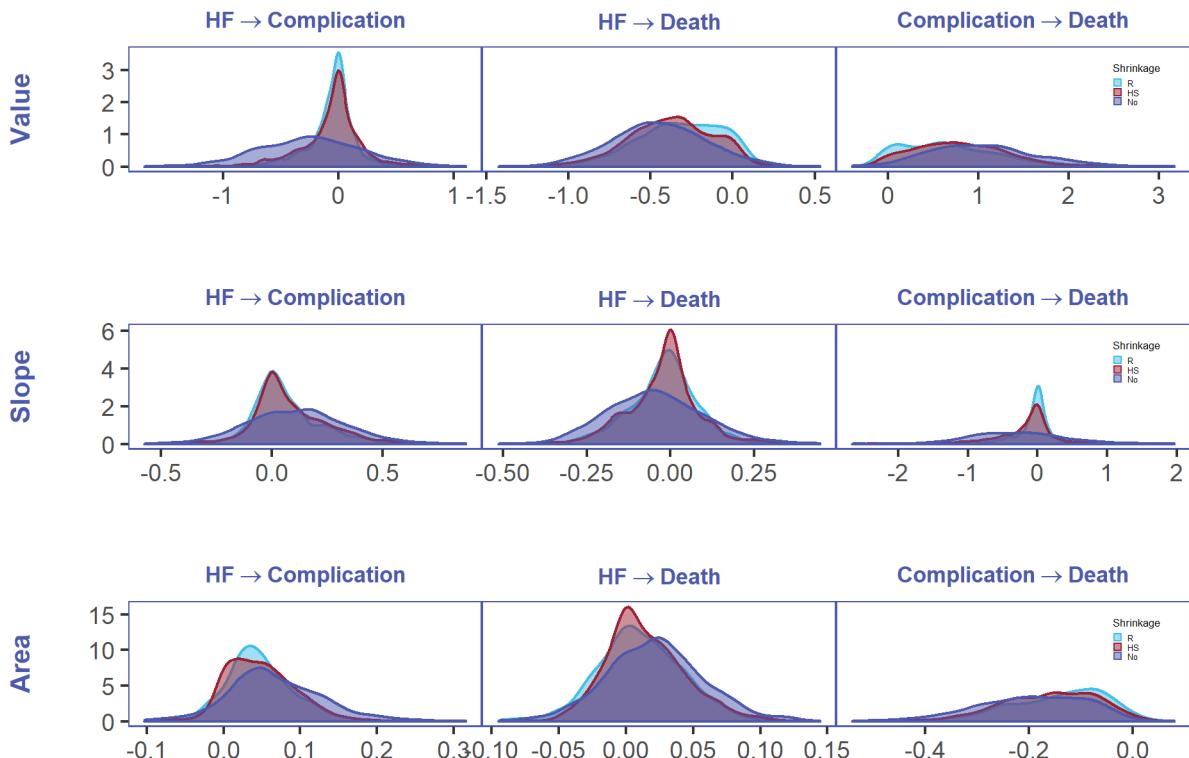
# Results: Bilirubin



# Results: Creatinine



# Results: Creatinine



# Software Implementation



Thank you

 @drizopoulos

 <https://github.com/drizopoulos>

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