

# Full Bayesian models to handle missing values in cost-effectiveness analysis from individual level data

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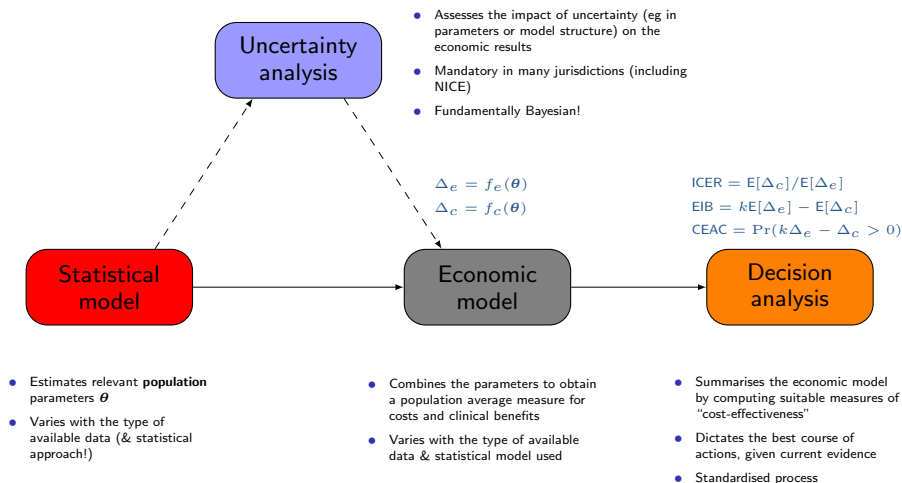
<http://www.statistica.it/gianluca>

<https://github.com/giabaio>

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**Objective:** Combine **costs** & **benefits** of a given intervention into a rational scheme for allocating resources, increasingly often under a Bayesian framework



- The available data usually look something like this:

ID	Trt	Demographics			HRQL data				Resource use data			
		Sex	Age	...	$u_0$	$u_1$	...	$u_J$	$c_0$	$c_1$	...	$c_J$
1	1	M	23	...	0.32	0.66	...	0.44	103	241	...	80
2	1	M	21	...	0.12	0.16	...	0.38	1204	1808	...	877
3	2	F	19	...	0.49	0.55	...	0.88	16	12	...	22
...	...	...	...	...	...	...	...	...	...	...	...	...

and the **typical** analysis is based on the following steps:

- 1 Compute individual QALYs and total costs as

$$e_i = \sum_{j=1}^J (u_{ij} + u_{i,j-1}) \frac{\delta_j}{2} \quad \text{and} \quad c_i = \sum_{j=0}^J c_{ij}, \quad \left[ \text{with: } \delta_j = \frac{\text{Time}_j - \text{Time}_{j-1}}{\text{Unit of time}} \right]$$

- 2 (Often implicitly) assume normality and linearity and model **independently** individual QALYs and total costs by controlling for baseline values

$$\begin{aligned} e_i &= \alpha_{e0} + \alpha_{e1}u_{0i} + \alpha_{e2}\text{Trt}_i + \varepsilon_{ei} [+ \dots], & \varepsilon_{ei} &\sim \text{Normal}(0, \sigma_e) \\ c_i &= \alpha_{c0} + \alpha_{c1}c_{0i} + \alpha_{c2}\text{Trt}_i + \varepsilon_{ci} [+ \dots], & \varepsilon_{ci} &\sim \text{Normal}(0, \sigma_c) \end{aligned}$$

- 3 Estimate population average cost and effectiveness differentials and use bootstrap to quantify uncertainty

# What's wrong with this?

- Potential correlation between costs & clinical benefits
  - Strong positive correlation — effective treatments are innovative and result from intensive and lengthy research  $\Rightarrow$  are associated with higher unit costs
  - Negative correlation — more effective treatments may reduce total care pathway costs e.g. by reducing hospitalisations, side effects, etc.
  - **NB:** In any case, the economic evaluation is based on both!
- Joint/marginal normality not realistic
  - Costs usually skewed and benefits may be bounded in  $[0; 1]$
  - Can use transformation (e.g. logs) — but care is needed when back transforming to the natural scale
  - Can use more suitable models (e.g. Gamma or log-Normal) — **especially under the Bayesian approach**
- ... and of course **Partially Observed** data
  - Can have item and/or unit non-response
  - Missingness may occur in either or both benefits/costs
  - The missingness mechanisms may also be correlated
  - Focus in decision-making — not inference!

# To be or not to be (Bayesians)?...

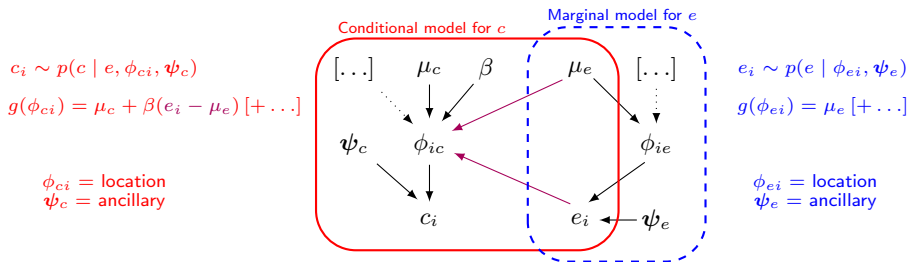
$$p(\theta | \mathbf{y}) \propto p(\theta)p(\mathbf{y} | \theta)?$$



# To be or not to be (Bayesians)?...

- In general, can represent a joint distribution as a **conditional regression**

$$p(e, c) = p(e)p(c | e)p(c | e) = p(c)p(e | c)$$



- For example:

$$e_i \sim \text{Beta}(\phi_{ei}\psi_e, (1 - \phi_{ei})\psi_e),$$

$$\text{logit}(\phi_{ei}) = \mu_e [+ \dots]$$

$$c_i | e_i \sim \text{Gamma}(\psi_c \phi_{ci}, \psi_c),$$

$$\text{log}(\phi_{ci}) = \mu_c + \beta(e_i - \mu_e) [+ \dots]$$

- Combining “modules” and fully characterising uncertainty about deterministic functions of random quantities is relatively straightforward using MCMC
- Prior information can help stabilise inference (especially with sparse data!), eg
  - Cancer patients are unlikely to survive as long as the general population
  - ORs are unlikely to be greater than  $\pm 5$

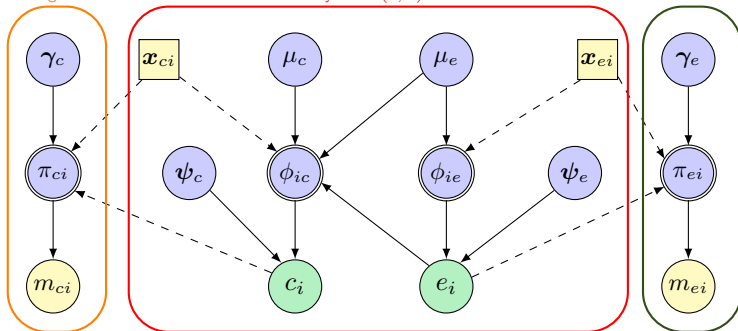
# Dealing with missing data — selection models

## MCAR ( $e, c$ )

Model of missingness for  $c$

Model of analysis for  $(c, e)$

Model of missingness for  $e$



- $m_{ei} \sim \text{Bernoulli}(\pi_{ei}); \quad \text{logit}(\pi_{ei}) = \gamma_{e0}$
- $m_{ci} \sim \text{Bernoulli}(\pi_{ci}); \quad \text{logit}(\pi_{ci}) = \gamma_{c0}$

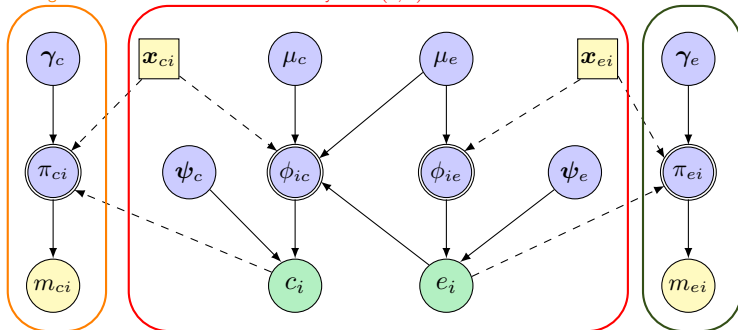
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Model of missingness for  $e$



- Partially observed data
- Unobservable parameters
- ◎ Deterministic function of random quantities
- Fully observed, unmodelled data
- Fully observed, modelled data

- $m_{ei} \sim \text{Bernoulli}(\pi_{ei}); \quad \text{logit}(\pi_{ei}) = \gamma_{e0} + \sum_{k=1}^K \gamma_{ek} x_{eik}$
- $m_{ci} \sim \text{Bernoulli}(\pi_{ci}); \quad \text{logit}(\pi_{ci}) = \gamma_{c0} + \sum_{h=1}^H \gamma_{ch} x_{cih}$



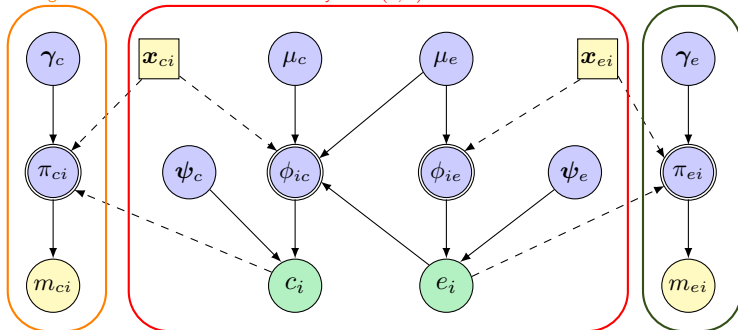
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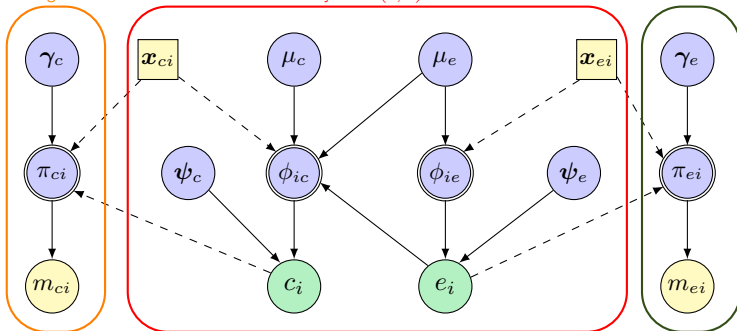
# Dealing with missing data — selection models

MNAR  $e$ ; MAR  $c$

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Model of missingness for  $e$



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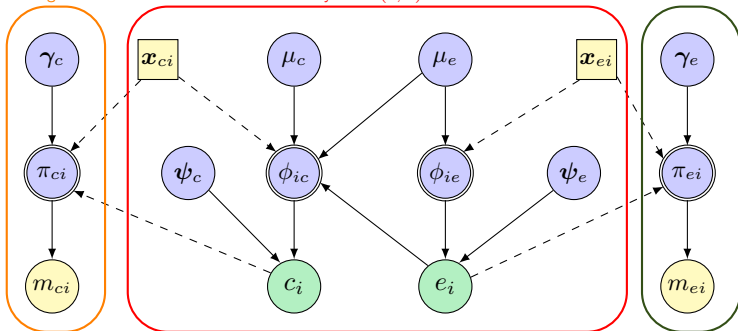
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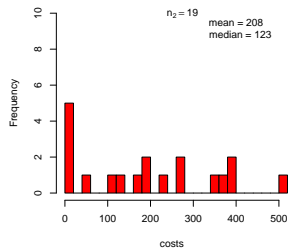
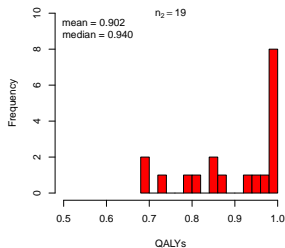
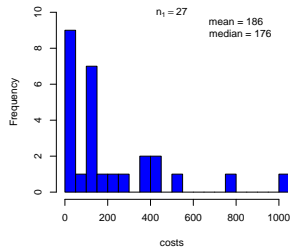
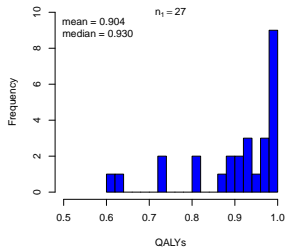
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$$\text{logit}(\pi_{ci}) = \gamma_{c0} + \sum_{h=1}^H \gamma_{ch} x_{cih} + \gamma_{cH+1} c_i$$

- The MenSS pilot RCT evaluates the cost-effectiveness of a new digital intervention to reduce the incidence of STI in young men with respect to the SOC
  - QALYs calculated from utilities (EQ-5D 3L)
  - Total costs calculated from different components (no baseline)

Time	Type of outcome	observed (%)	
		Control ( $n_1=75$ )	Intervention ( $n_2=84$ )
Baseline	utilities	72 (96%)	72 (86%)
3 months	utilities and costs	34 (45%)	23 (27%)
6 months	utilities and costs	35 (47%)	23 (27%)
12 months	utilities and costs	43 (57%)	36 (43%)
<b>Complete cases</b>	utilities and costs	27 (44%)	19 (23%)

- The MenSS pilot RCT evaluates the cost-effectiveness of a new digital intervention to reduce the incidence of STI in young men with respect to the SOC
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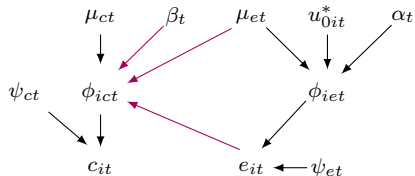
## 1 Bivariate Normal

- Simpler and closer to “standard” frequentist model
- Account for **correlation between QALYs and costs**

### Conditional model for $c | e$

$$c_{it} | e_{it} \sim \text{Normal}(\phi_{c_{it}}, \psi_{ct})$$

$$\phi_{c_{it}} = \mu_{ct} + \beta_t(e_{it} - \mu_{et})$$



### Marginal model for $e$

$$e_{it} \sim \text{Normal}(\phi_{e_{it}}, \psi_{et})$$

$$\phi_{e_{it}} = \mu_{et} + \alpha_t(u_{0it} - \bar{u}_{0t})$$

$$= \mu_{et} + \alpha_t u_{0it}^*$$

## 1 Bivariate Normal

- Simpler and closer to “standard” frequentist model
- Account for **correlation between QALYs and costs**

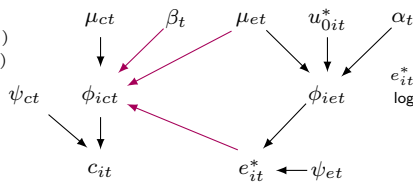
## 2 Beta-Gamma

- Account for **correlation between outcomes**
- Model the relevant ranges: QALYs  $\in (0, 1)$  and costs  $\in (0, \infty)$
- **But:** needs to rescale observed data  $e_{it}^* = (e_{it} - \epsilon)$  to avoid spikes at 1

### Conditional model for $c \mid e^*$

$$c_{it} \mid e_{it}^* \sim \text{Gamma}(\psi_{ct} \phi_{c_{it}}, \psi_{ct})$$

$$\log(\phi_{c_{it}}) = \mu_{ct} + \beta_t (e_{it}^* - \mu_{et})$$



### Marginal model for $e^*$

$$e_{it}^* \sim \text{Beta}(\phi_{eit} \psi_{et}, (1 - \phi_{eit}) \psi_{et})$$

$$\text{logit}(\phi_{eit}) = \mu_{et} + \alpha_t (u_{0it} - \bar{u}_{0t})$$

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## 1 Bivariate Normal

- Simpler and closer to “standard” frequentist model
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## 2 Beta-Gamma

- Account for
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- **But:** needs to rescale observed data  $e_{it}^* = (e_{it} - \epsilon)$  to avoid spikes at 1

## 3 Hurdle model

- Model  $e_{it}$  as a **mixture** to account for **correlation between outcomes**, model the relevant ranges and account for **structural values**
- May expand to account for partially observed baseline utility  $u_{0it}$

### Conditional model for $c \mid e^*$

$$c_{it} \mid e_{it}^* \sim \text{Gamma}(\psi_{ct} \phi_{c_{it}}, \psi_{ct})$$

$$\log(\phi_{c_{it}}) = \mu_{ct} + \beta_t (e_{it}^* - \mu_{et})$$

$$\psi_{ct}$$

$$\phi_{ict}$$

$$c_{it}$$

$$\mu_{ct}$$

$$\beta_t$$

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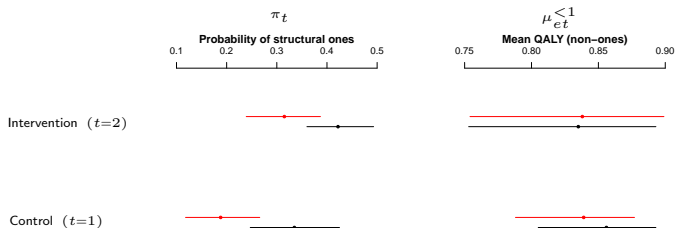
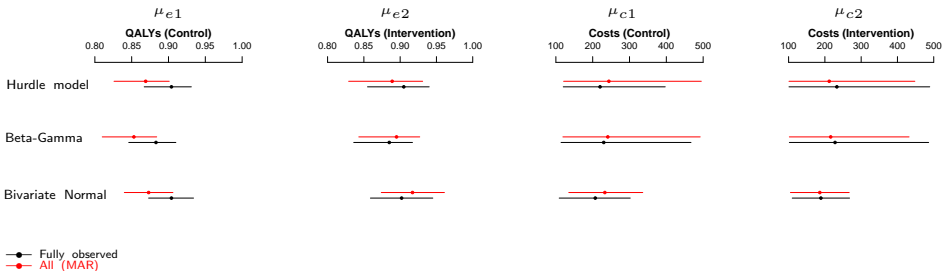
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$$\beta_t$$

$$\mu$$

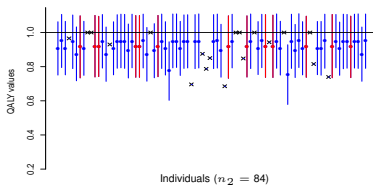
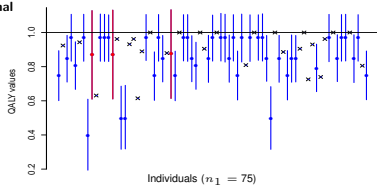


# Results — estimation of the main parameters (CCA + MAR)

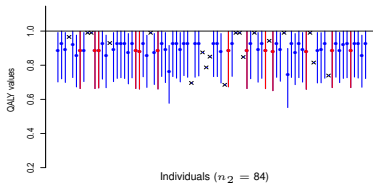
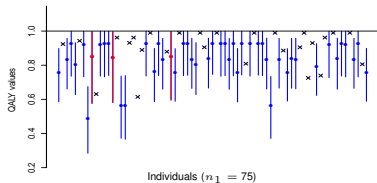


# Bayesian multiple imputation (under MAR)

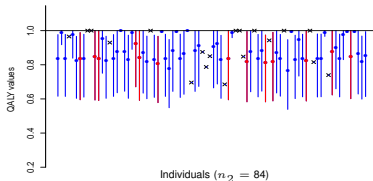
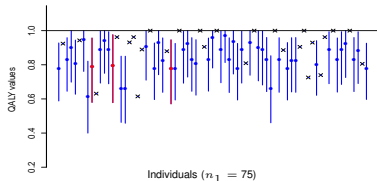
## Bivariate Normal



## Beta-Gamma



## Hurdle model



● Imputed, observed baseline  
● Imputed, missing baseline  
× Observed

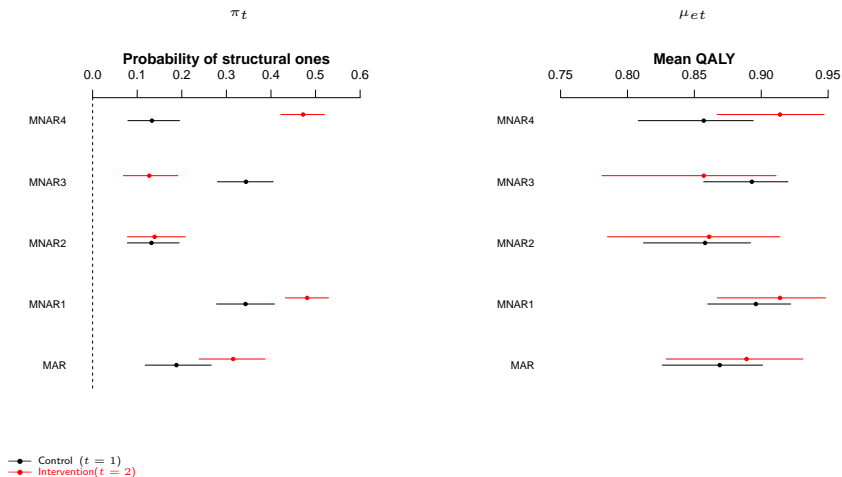
- We observe  $n_{01} = 13$  and  $n_{02} = 22$  individuals with  $u_{0it} = 1$  and  $u_{jit} = \text{NA}$ , for  $j > 1$
- For those individuals, we cannot compute directly the structural one indicator  $d_{it}$  and so need to make assumptions/model this
  - Sensitivity analysis to alternative MNAR departures from MAR

**MNAR1.** Set  $d_{it} = 1$  for all individuals with unit observed baseline utility

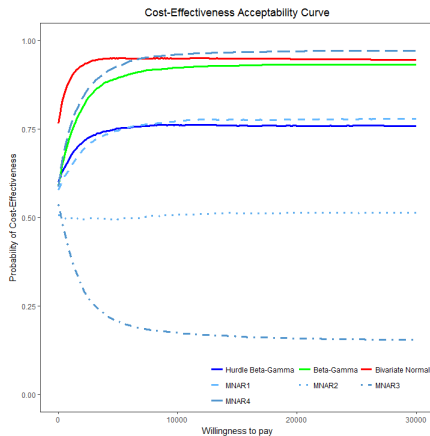
**MNAR2.** Set  $d_{it} = 0$  for all individuals with unit observed baseline utility

**MNAR3.** Set  $d_{it} = 1$  for the  $n_{01} = 13$  individuals with  $u_{0i1} = 1$  and  $d_{it} = 0$  for the  $n_{02} = 22$  individuals with  $u_{0i2} = 1$

**MNAR4.** Set  $d_{it} = 0$  for the  $n_{01} = 13$  individuals with  $u_{0i1} = 1$  and  $d_{it} = 1$  for the  $n_{02} = 22$  individuals with  $u_{0i2} = 1$



# Cost-effectiveness analysis



- A full Bayesian approach to handling missing data extends standard “imputation methods”
  - Can consider MAR and MNAR with relatively little expansion to the basic model
- Particularly helpful in cost-effectiveness analysis, to account for
  - Asymmetrical distributions for the main outcomes
  - Correlation between costs & benefits
  - Structural values (eg spikes at 1 for utilities or spikes at 0 for costs)
- Need specialised software + coding skills
  - R package `missingHE` under development to implement a set of general models
  - Preliminary work available at <https://github.com/giabaio/missingHE>
  - Eventually, will be able to combine with existing packages (eg `BCEA`: <http://www.statistica.it/gianluca/BCEA>; <https://github.com/giabaio/BCEA>) to perform the whole economic analysis

Thank you!