
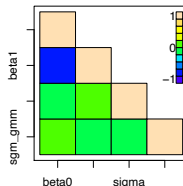


# Estimation

Thanks to MCMC algorithms, one can obtain a **Monte Carlo sample from the *posterior* distribution** for a **Bayesian model**

**Monte Carlo method** can then be used to get ***posterior* estimates** :

- Point estimates (*posterior* mean, *posterior* median, ...)
- Credibility interval (shortest: *Highest Density Interval* – *HDI* with  package `HDInterval`)
- Correlations between parameters
- ...



# Deviance Information Criterion (*DIC*)

Deviance is:  $D(\theta) = -2\log(p(\theta|\mathbf{y})) + C$  with  $C$  a constant

**Deviance Information Criterion** is then:

$$DIC = \overline{D(\theta)} + p_D$$

where  $p_D = \left( D(\bar{\theta}) - \overline{D(\theta)} \right)$  represents a penalty for the effective number of parameters

⇒ *DIC* allows to compare different models estimated on the same data  
*the smaller the DIC, the better the model !*

[M Plummer, Penalized loss functions for Bayesian model comparison, *Biostatistics*, 2008]

# Your turn !



**Practical:** exercise 4