

The simplest form of the standard 3-stage hierarchical model in the context of MEP size modeling can be described as follows. Let there be N_P exchange sequences $\{(x_i^p, y_i^p)_{i=1}^{n(p)} \mid j = 1, 2 \dots N_P\}$ of MEP size y_i^p recorded at stimulus intensity x_i^p from p^{th} participant, for a total of N_P participants.

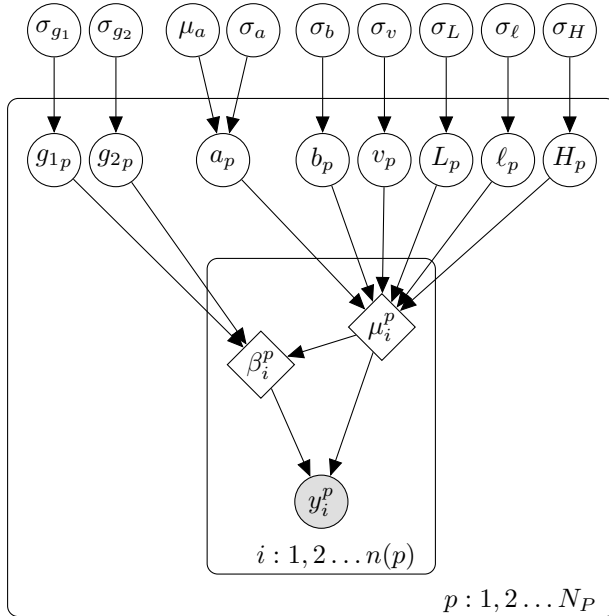
The first stage of hierarchy is the *participant-level* and specifies parametric model $P(y_i^p \mid x_i^p, \theta_p)$ for each of the N_P sequences and models the MEP size y_i^p as a function of stimulus intensity x_i^p and participant specific parameters θ_p . In the second stage, $\theta_1, \theta_2 \dots \theta_P$ are assumed to be exchangeable and generated from a common distribution $P(\theta_p \mid \gamma)$ with hyper-parameters γ . In the third stage, the hyper-parameters γ are assumed to be unknown and are assigned a non-informative prior, also called hyperprior, density $P(\gamma)$.

$$y_i^p \mid x_i^p, \theta_p, \gamma \sim P(y_i^p \mid x_i^p, \theta_p, \gamma) \quad (4.3.1)$$

$$\theta_p \mid \gamma \sim P(\theta_p \mid \gamma) \quad (4.3.2)$$

$$\gamma \sim P(\gamma) \quad (4.3.3)$$

Figure 4.3.1 shows the graphical representation of the basic hierarchical model implemented by hbmap (Section 5.1) for recruitment-curve fitting



$$y_i^p \mid x_i^p, \Omega_1^p \sim \mathcal{G}(\mu_i^p \cdot \beta_i^p, \beta_i^p)$$

$$\mu_i^p = \mathcal{F}(x_i^p \mid \Omega_1^p) \quad \beta_i^p = g_{1p} + \frac{\mu_i^p}{g_{2p}}$$

$$a_p \sim \mathcal{TN}(\mu_a, \sigma_a)$$

$$\theta_p \sim \mathcal{HN}(\sigma_{\theta_p}) \quad \forall \theta_p \in \Omega \setminus \{a_p\}$$

$$\mu_a \sim \mathcal{TN}(50, 20) \quad \sigma_a \sim \mathcal{HN}(30)$$

$$\sigma_{\sigma_L} \sim \mathcal{HN}(.05) \quad \sigma_{\sigma} \sim \mathcal{HN}(5) \quad \forall \sigma \in \Omega_2$$

$$\Omega_1^p = \{a_p, b_p, v_p, L_p, \ell_p, H_p, g_{1p}, g_{2p}\}$$

$$\Omega_2 = \{\sigma_L, \sigma_H, \sigma_v, \sigma_{\ell}, \sigma_{g_1}, \sigma_{g_2}\}$$