

Computation of b_i :

$$\text{cov}(u_i, \tilde{y}_i) = \text{cov}\left(u_i, \left[\mu + u_i + p_i + \frac{1}{k} \sum_{j=1}^k t_{ij}\right]\right)$$

$$\begin{aligned} &= \frac{1}{k} \sum_{j=1}^k \text{cov}\left(u_i, \left[\mu + u_i + p_i + t_{ij}\right]\right) \\ &\quad \downarrow \text{genetic model} \\ &= \frac{1}{k} \sum_{j=1}^k \left[\underbrace{\text{cov}(u_i, \mu)}_{\frac{1}{k} \cdot k \cdot \mu = \mu} + \underbrace{\text{cov}(u_i, u_i)}_{u_i} + \underbrace{\text{cov}(u_i, p_i)}_{p_i} + \underbrace{\text{cov}(u_i, t_{ij})}_{\frac{1}{k} \sum_{j=1}^k t_{ij}} \right] \end{aligned}$$

$$\begin{aligned} \text{cov}(u_i, \tilde{y}_i) &= \overbrace{\text{cov}(u_i, \mu)}^{=0} + \underbrace{\text{cov}(u_i, u_i)}_k + \underbrace{\text{cov}(u_i, p_i)}_{=0} \\ &\quad + \underbrace{\text{cov}\left(u_i, \frac{1}{k} \sum_{j=1}^k t_{ij}\right)}_{=0} \end{aligned}$$

$$= \text{cov}(u_i, u_i) = \text{var}(u_i) = \sigma_u^2$$

genetic additive
variance