

$$\text{var}(y) = \text{var}(X\beta + Zg + e) = \underbrace{\text{var}(X\beta) + \text{var}(Zg) + \text{var}(e)}_{\text{const.}} + \underbrace{2\text{Cov}(X\beta, Zg) + 2\text{Cov}(X\beta, e) + 2\text{Cov}(Zg, e)}_{0}$$

$$= \text{var}(Zg) + \text{var}(e)$$

$$= Z \cdot \underbrace{\text{var}(g)}_{G \cdot \sigma_g^2} \cdot Z^T + \underbrace{\text{var}(e)}_{R = I \cdot \sigma_e^2}$$

$$= Z \cdot G \cdot Z^T \cdot \sigma_g^2 + I \cdot \sigma_e^2$$

$$* \text{Var}(c \cdot x) = \underline{c^2 \cdot \text{var}(x)} \quad \left\{ \begin{array}{l} \text{var}(5 \cdot x) = 25 \cdot \text{var}(x) \end{array} \right.$$

real scalar
constant

Matrix - Vector:

vector $g = \begin{bmatrix} g_1 \\ g_2 \\ \vdots \\ g_n \end{bmatrix}$ $\begin{matrix} \nearrow \text{var}(g_1) \\ \nearrow \text{var}(g_2) \end{matrix}$

$$\text{var}(g) = \begin{bmatrix} \text{var}(g_1) & \text{cov}(g_1, g_2) & \dots & \text{cov}(g_1, g_n) \\ \text{cov}(g_2, g_1) & \text{var}(g_2) & & \\ \vdots & & \ddots & \\ \text{cov}(g_n, g_1) & & & \text{var}(g_n) \end{bmatrix}$$

$$\text{var}(Zg) \neq Z \cdot \text{var}(g) \cdot Z^T$$

Sofar: Data sets where all animals had genotypes