



Correction: Multi-trait single-step genomic prediction accounting for heterogeneous (co)variances over the genome

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Correction to: Heredity

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In the original PDF version of this article, the below equation was incorrectly displayed.
Instead of:

$$\text{where } \mathbf{X} = [\mathbf{1}_L \mathbf{0} \mathbf{0} \mathbf{1}_H], \boldsymbol{\mu} = \begin{bmatrix} \mu_L \\ \mu_H \end{bmatrix}, \mathbf{B} = [\mathbf{B}_L \mathbf{B}_{LH} \mathbf{B}_{HL} \mathbf{B}_H]$$

The equation should be displayed as follows:

$$\text{where } \mathbf{X} = \begin{bmatrix} \mathbf{1}_L & \mathbf{0} \\ \mathbf{0} & \mathbf{1}_H \end{bmatrix}, \boldsymbol{\mu} = \begin{bmatrix} \mu_L \\ \mu_H \end{bmatrix}, \mathbf{B} = \begin{bmatrix} \mathbf{B}_L & \mathbf{B}_{LH} \\ \mathbf{B}_{HL} & \mathbf{B}_H \end{bmatrix}$$

This error has been corrected in the PDF version of the article. The HTML version of the article was correct at the time of publication.

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