

Additional file 2: Description of the system of equations of ssSNPBLUP_MS for a bivariate maternal model

A standard bivariate maternal model associated with the ssSNPBLUP_MS system of equations can be written as (ignoring the permanent environmental effects):

$$\mathbf{y} = \mathbf{X}\boldsymbol{\beta} + \begin{bmatrix} \mathbf{W}_n & \mathbf{0} \\ \mathbf{0} & \mathbf{W}_g \end{bmatrix} \begin{bmatrix} \mathbf{u}_n \\ \mathbf{a}_g \end{bmatrix} + \begin{bmatrix} \mathbf{0} \\ \mathbf{W}_g \mathbf{M}_z \end{bmatrix} \mathbf{g} + \begin{bmatrix} \mathbf{S}_n & \mathbf{0} \\ \mathbf{0} & \mathbf{S}_g \end{bmatrix} \begin{bmatrix} \mathbf{m}_n \\ \mathbf{q}_g \end{bmatrix} + \begin{bmatrix} \mathbf{0} \\ \mathbf{S}_g \mathbf{M}_z \end{bmatrix} \mathbf{g}_m + \mathbf{e}$$

where the subscripts g and n refer to genotyped and non-genotyped animals, respectively, $\mathbf{y} = \begin{bmatrix} \mathbf{y}_{n,1} \\ \mathbf{y}_{n,2} \\ \mathbf{y}_{g,1} \\ \mathbf{y}_{g,2} \end{bmatrix}$

is the vector of records (with the subscripts 1 and 2 referring to the first and the second trait, respectively), $\boldsymbol{\beta} = \begin{bmatrix} \boldsymbol{\beta}_{n,1} \\ \boldsymbol{\beta}_{n,2} \\ \boldsymbol{\beta}_{g,1} \\ \boldsymbol{\beta}_{g,2} \end{bmatrix}$ is the vector of fixed effects, $\mathbf{u}_n = \begin{bmatrix} \mathbf{u}_{n,1} \\ \mathbf{u}_{n,2} \end{bmatrix}$ is the vector of direct genetic effects

for non-genotyped animals, $\mathbf{a}_g = \begin{bmatrix} \mathbf{a}_{g,1} \\ \mathbf{a}_{g,2} \end{bmatrix}$ is the vector of direct residual polygenic effects for genotyped animals, $\mathbf{g} = \begin{bmatrix} \mathbf{g}_1 \\ \mathbf{g}_2 \end{bmatrix}$ is the vector of SNP effects for the direct genetic effects, $\mathbf{m}_n = \begin{bmatrix} \mathbf{m}_{n,1} \\ \mathbf{m}_{n,2} \end{bmatrix}$ is the vector of maternal genetic effects for non-genotyped animals, $\mathbf{q}_g = \begin{bmatrix} \mathbf{q}_{g,1} \\ \mathbf{q}_{g,2} \end{bmatrix}$ is the vector of maternal residual polygenic effects for genotyped animals, $\mathbf{g}_m = \begin{bmatrix} \mathbf{g}_{m,1} \\ \mathbf{g}_{m,2} \end{bmatrix}$ is the vector of SNP effects for the maternal genetic effects, and \mathbf{e} is the vector of residuals.

The matrices $\mathbf{X} = \begin{bmatrix} \mathbf{X}_{n,1} & \mathbf{0} \\ \mathbf{0} & \mathbf{X}_{n,2} \\ \mathbf{X}_{g,1} & \mathbf{0} \\ \mathbf{0} & \mathbf{X}_{g,2} \end{bmatrix}$, $\mathbf{W}_n = \begin{bmatrix} \mathbf{W}_{n,1} & \mathbf{0} \\ \mathbf{0} & \mathbf{W}_{n,2} \end{bmatrix}$, $\mathbf{W}_g = \begin{bmatrix} \mathbf{W}_{g,1} & \mathbf{0} \\ \mathbf{0} & \mathbf{W}_{g,2} \end{bmatrix}$, $\mathbf{S}_n = \begin{bmatrix} \mathbf{S}_{n,1} & \mathbf{0} \\ \mathbf{0} & \mathbf{S}_{n,2} \end{bmatrix}$

and $\mathbf{S}_g = \begin{bmatrix} \mathbf{S}_{g,1} & \mathbf{0} \\ \mathbf{0} & \mathbf{S}_{g,2} \end{bmatrix}$ are incidence matrices relating records to their corresponding effects.

The matrix \mathbf{M}_z is equal to $\mathbf{M}_z = \mathbf{I}_2 \otimes \mathbf{Z}$, with \mathbf{I}_2 being an identity matrix with size equal to the number of traits (i.e. 2) and the matrix \mathbf{Z} containing the SNP genotypes (coded as 0 for one homozygous genotype, 1 for the heterozygous genotype, or 2 for the alternate homozygous genotype) centred by their observed means.

Following the Eq. (2) in the main text, the matrix \mathbf{T} becomes $\mathbf{T} = \begin{bmatrix} \mathbf{I} & & & & & & & \\ \mathbf{0} & \mathbf{I} & & & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{I} & & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{I} & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{I} & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{M}_z & & \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{M}_z \end{bmatrix}$.

It follows that the matrix \mathbf{C}_{MSLS} in the Eq. (2) is equal to

$$\mathbf{C}_{MSLS} = \begin{bmatrix} \mathbf{X}'\mathbf{R}^{-1}\mathbf{X} & & & & & & & & \\ \mathbf{W}'_n\mathbf{R}_n^{-1}\mathbf{X}_n & \mathbf{W}'_n\mathbf{R}_n^{-1}\mathbf{W}_n & & & & & & & \\ \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{X}_g & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & & & & & & & \\ \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{X}_n & \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{W}_n & \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{W}_g & \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{S}_n & & & & & \\ \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{X}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{W}_n & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{S}_n & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{S}_g & & & & \\ \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{X}_g & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{W}_n & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{S}_n & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{S}_g & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & & & \\ \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{X}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{W}_n & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{S}_n & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{S}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{S}_g & & \end{bmatrix},$$

that the matrix \mathbf{C}_{MSR1} in the Eq. (2) is equal to

$$\mathbf{C}_{MSR1} = \begin{bmatrix} \mathbf{0} & & & & & & & & & & \\ \mathbf{0} & \mathbf{0} & & & & & & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{G}^{uu} \otimes \left(\left(1 - \frac{1}{w} \right) \mathbf{Q} \right) & & & & & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & & & & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{G}^{mu} \otimes \left(\left(1 - \frac{1}{w} \right) \mathbf{Q} \right) & \mathbf{0} & \mathbf{G}^{mm} \otimes \left(\left(1 - \frac{1}{w} \right) \mathbf{Q} \right) & & & & & & \\ \mathbf{0} & \mathbf{G}^{uu} \otimes \mathbf{A}^{gn} & \mathbf{G}^{uu} \otimes \mathbf{Q} & \mathbf{G}^{um} \otimes \mathbf{A}^{gn} & \mathbf{G}^{um} \otimes \mathbf{Q} & & \mathbf{G}^{uu} \otimes \mathbf{Q} & & & & \\ \mathbf{0} & \mathbf{G}^{mu} \otimes \mathbf{A}^{gn} & \mathbf{G}^{mu} \otimes \mathbf{Q} & \mathbf{G}^{mm} \otimes \mathbf{A}^{gn} & \mathbf{G}^{mm} \otimes \mathbf{Q} & & \mathbf{G}^{mu} \otimes \mathbf{Q} & \mathbf{G}^{mm} \otimes \mathbf{Q} & & & \end{bmatrix}$$

and that the matrix \mathbf{C}_{MSR2} in the Eq. (2) is equal to

$$\mathbf{C}_{MSR2} = \begin{bmatrix} \mathbf{0} & & & & & & & & & & \\ \mathbf{0} & \mathbf{G}^{uu} \otimes \mathbf{A}^{nn} & & & & & & & & & \\ \mathbf{0} & \mathbf{G}^{uu} \otimes \mathbf{A}^{gn} & \mathbf{G}^{uu} \otimes \frac{1}{w} \mathbf{A}^{gg} & & & & & & & & \\ \mathbf{0} & \mathbf{G}^{mu} \otimes \mathbf{A}^{nn} & \mathbf{G}^{mu} \otimes \mathbf{A}^{ng} & \mathbf{G}^{mm} \otimes \mathbf{A}^{nn} & & & & & & & \\ \mathbf{0} & \mathbf{G}^{mu} \otimes \mathbf{A}^{gn} & \mathbf{G}^{mu} \otimes \frac{1}{w} \mathbf{A}^{gg} & \mathbf{G}^{mm} \otimes \mathbf{A}^{gn} & \mathbf{G}^{mm} \otimes \frac{1}{w} \mathbf{A}^{gg} & & & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{G}^{uu} \otimes \left(\frac{m}{1-w} \mathbf{I} \right) & & & & \\ \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{0} & \mathbf{G}^{mu} \otimes \left(\frac{m}{1-w} \mathbf{I} \right) & \mathbf{G}^{mm} \otimes \left(\frac{m}{1-w} \mathbf{I} \right) & & & \end{bmatrix}$$

With the matrices \mathbf{C}_{MSLS} , \mathbf{C}_{MSR1} , and \mathbf{C}_{MSR2} being defined, the multiplication of the coefficient matrix \mathbf{C} , associated with a bivariate maternal ssSNPBLUP_MS, by a vector can be performed as proposed in the main text. For example, the multiplication of \mathbf{C}_{MSLS} by a vector can be performed with approaches already developed in animal breeding, such as iteration-on-data approaches [16-19]. Indeed,

the block matrix $\begin{bmatrix} \mathbf{X}'\mathbf{R}^{-1}\mathbf{X} & & & & & & & & \\ \mathbf{W}'_n\mathbf{R}_n^{-1}\mathbf{X}_n & \mathbf{W}'_n\mathbf{R}_n^{-1}\mathbf{W}_n & & & & & & & \\ \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{X}_g & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{W}_n & \mathbf{W}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & & & & & & \\ \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{X}_n & \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{W}_n & \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{W}_g & \mathbf{S}'_n\mathbf{R}_n^{-1}\mathbf{S}_n & & & & & \\ \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{X}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{W}_n & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{W}_g & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{S}_n & \mathbf{S}'_g\mathbf{R}_g^{-1}\mathbf{S}_g & & & & \end{bmatrix}$ extracted from

\mathbf{C}_{MSLS} is equal to the least-squares part of the coefficient matrix of a system of equations associated with a traditional pedigree BLUP based on the same model and datasets [19].