# Additional file 1

## Details of the algebraic derivations of the formulas in the main text

# Algebraic derivation of expression (1)

Let  $K = 2^t$  be the number of possible haplotypes, at locus *i*, for a sliding window of *t* markers. Let  $\mathbb{1}_{\{u_{QTL,c_1,c_2}=1\}}$  and  $\mathbb{1}_{\{u_{QTL,c_1,c_2}=0\}}$  be the following indicator functions:

$$\begin{split} \mathbbm{1}_{\left\{u_{QTL,c_{1},c_{2}}=1\right\}} = \begin{cases} & 1 \text{ if } (c_{1},c_{2}) \text{ have identical alleles at the QTL} \\ & 0 \text{ else} \\ \\ \mathbbm{1}_{\left\{u_{QTL,c_{1},c_{2}}=0\right\}} = \begin{cases} & 1 \text{ if } (c_{1},c_{2}) \text{ have non-identical alleles at the QTL} \\ & 0 \text{ else} \end{cases} \end{split}$$

We have:

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \frac{1}{4n^{2}} \sum_{c_{1}=1}^{2n} \sum_{c_{2}=1}^{2n} |s_{i,c_{1},c_{2}}^{\mathcal{P}} - u_{QTL,c_{1},c_{2}}|$$
  
$$= \frac{1}{4n^{2}} \sum_{c_{1}=1}^{2n} \sum_{c_{2}=1}^{2n} \mathbb{1}_{\left\{u_{QTL,c_{1},c_{2}}=1\right\}} |s_{i,c_{1},c_{2}}^{\mathcal{P}} - 1| + \frac{1}{4n^{2}} \sum_{c_{1}=1}^{2n} \sum_{c_{2}=1}^{2n} \mathbb{1}_{\left\{u_{QTL,c_{1},c_{2}}=0\right\}} |s_{i,c_{1},c_{2}}^{\mathcal{P}} - 0|$$

Let  $E_{h_p}$  be the set of chromosome segments carrying haplotype  $h_p$   $(p \in \{1, .., K\})$  at locus *i*. We have:

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \frac{1}{4n^{2}} \sum_{c_{1}=1}^{2n} \sum_{c_{2}=1}^{2n} \sum_{p=1}^{K} \mathbb{1}_{\{c_{1}\in E_{h_{p}}\}} \sum_{q=1}^{K} \mathbb{1}_{\{c_{2}\in E_{h_{q}}\}} \mathbb{1}_{\{u_{QTL,c_{1},c_{2}}=1\}} |s_{i,c_{1},c_{2}}^{\mathcal{P}} - 1|$$
$$+ \frac{1}{4n^{2}} \sum_{c_{1}=1}^{2n} \sum_{c_{2}=1}^{2n} \sum_{p=1}^{K} \mathbb{1}_{\{c_{1}\in E_{h_{p}}\}} \sum_{q=1}^{K} \mathbb{1}_{\{c_{2}\in E_{h_{q}}\}} \mathbb{1}_{\{u_{QTL,c_{1},c_{2}}=0\}} |s_{i,c_{1},c_{2}}^{\mathcal{P}} - 0|$$

where  $\mathbb{1}_{\{c_1 \in E_{h_p}\}}$  and  $\mathbb{1}_{\{c_2 \in E_{h_q}\}}$  are the indicator functions of the events  $\{c_1 \in E_{h_p}\}$  and  $\{c_2 \in E_{h_q}\}$ respectively. Indeed  $\sum_{p=1}^{K} \mathbb{1}_{\{c_1 \in E_{h_p}\}}$  and  $\sum_{q=1}^{K} \mathbb{1}_{\{c_2 \in E_{h_q}\}}$  are always equal to one.  $d_1(\mathbf{M}^{\mathcal{P},i}, \mathbf{M}^{QTL})$  can thus be expressed as:

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \sum_{p=1}^{K} \sum_{q=1}^{K} \frac{1}{4n^{2}} \sum_{c_{1}=1}^{2n} \sum_{c_{2}=1}^{2n} \mathbb{1}_{\left\{u_{QTL,c_{1}\in E_{h_{p}},c_{2}\in E_{h_{q}}=1\right\}} |s_{i,h_{p},h_{q}}^{\mathcal{P}} - 1| + \sum_{p=1}^{K} \sum_{q=1}^{K} \frac{1}{4n^{2}} \sum_{c_{1}=1}^{2n} \sum_{c_{2}=1}^{2n} \mathbb{1}_{\left\{u_{QTL,c_{1}\in E_{h_{p}},c_{2}\in E_{h_{q}}=0\right\}} |s_{i,h_{p},h_{q}}^{\mathcal{P}} - 0|$$

where  $\frac{1}{4n^2} \sum_{c_1=1}^{2n} \sum_{c_2=1}^{2n} \mathbb{1}_{\left\{u_{QTL,c_1 \in E_{h_p}, c_2 \in E_{h_q}}=1\right\}} = f(c_1 \in E_{h_p}, c_2 \in E_{h_q}, \text{identical alleles at the QTL}) \text{ is the frequency of chromosome segments, at locus } i, carrying <math>h_p$  and  $h_q$  and having identical alleles at the

QTL. Since  $\{c_1 \in E_{h_p}\}$  and  $\{c_2 \in E_{h_q}\}$  are independent events we have:

$$f(c_1 \in E_{h_p}, c_2 \in E_{h_q}, \text{identical alleles at the QTL}) = p_{i,h_p,h_q}^{QTL} f_{i,h_p} f_{i,h_p} f_{i,h_q}$$

where  $p_{i,h_p,h_q}^{QTL}$  is the proportion of identical alleles shared at the QTL by the couples of chromosomes carrying  $h_p$  and  $h_q$  at position *i*.  $f_{i,h_p}$  and  $f_{i,h_q}$  are the frequencies of haplotypes  $h_p$  and  $h_q$  at position *i* respectively.

Similarly we have:

$$\begin{aligned} \frac{1}{4n^2} \sum_{c_1=1}^{2n} \sum_{c_2=1}^{2n} \mathbb{1}_{\left\{u_{QTL,c_1 \in E_{h_p}, c_2 \in E_{h_q}=0}\right\}} &= f(c_1 \in E_{h_p}, c_2 \in E_{h_q}, \text{non-identical alleles at the QTL}) \\ &= (1 - p_{i,h_p,h_q}^{QTL}) f_{i,h_p} f_{i,h_q} \end{aligned}$$

Consequently  $d_1(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL})$  can be written as (1), i.e.

$$d_1(\mathbf{M}^{\mathcal{P},i}, \mathbf{M}^{QTL}) = \sum_{p=1}^K \sum_{q=1}^K f_{i,h_p} f_{i,h_q} \left[ p_{i,h_p,h_q}^{QTL} (1 - s_{i,h_p,h_q}^{\mathcal{P}}) + (1 - p_{i,h_p,h_q}^{QTL}) s_{i,h_p,h_q}^{\mathcal{P}} \right]$$
(1)

Algebraic derivation of expression (2)

Let  $(n_{h_p})_{1 \le p \le K}$  be the counts of the possible haplotypes at a tested position *i*. And let  $(n_{h_p a_l})_{\substack{1 \le p \le K \\ 1 \le l \le 2}}$  be the counts of the 2K possible haplotypes defined between *i* and a QTL. Expression (1) can be rewritten as:

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \sum_{p=1}^{K} \sum_{q=1}^{K} \frac{n_{h_{p}}}{2n} \frac{n_{h_{q}}}{2n} \left[ \frac{[n_{h_{p}a_{1}}n_{h_{q}a_{1}} + n_{h_{p}a_{2}}n_{h_{q}a_{2}}]}{n_{h_{p}}n_{h_{q}}} (1 - s_{i,h_{p},h_{q}}^{\mathcal{P}}) \right. \\ \left. + \left( \frac{n_{h_{p}}n_{h_{q}} - [n_{h_{p}a_{1}}n_{h_{q}a_{1}} + n_{h_{p}a_{2}}n_{h_{q}a_{2}}]}{n_{h_{p}}n_{h_{q}}} \right) s_{i,h_{p},h_{q}}^{\mathcal{P}} \right] \\ = \sum_{p=1}^{K} \sum_{q=1}^{K} \left[ \left[ f_{i,h_{p}a_{1}}^{QTL} f_{i,h_{q}a_{1}}^{QTL} + f_{i,h_{p}a_{2}}^{QTL} f_{i,h_{q}a_{2}}^{QTL} \right] (1 - s_{i,h_{p},h_{q}}^{\mathcal{P}}) \right. \\ \left. + \left( \frac{(n_{h_{p}a_{1}} + n_{h_{p}a_{2}})(n_{h_{q}a_{1}} + n_{h_{q}a_{2}})}{4n^{2}} - f_{i,h_{p}a_{1}}^{QTL} f_{i,h_{q}a_{1}}^{QTL} - f_{i,h_{p}a_{2}}^{QTL} f_{i,h_{q}a_{2}}^{QTL} \right] s_{i,h_{p},h_{q}}^{\mathcal{P}} \right]$$

which simplifies to (2):

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \sum_{p=1}^{K} \sum_{q=1}^{K} \left[ \left[ f_{i,h_{p}a_{1}}^{QTL} f_{i,h_{q}a_{1}}^{QTL} + f_{i,h_{p}a_{2}}^{QTL} f_{i,h_{q}a_{2}}^{QTL} \right] (1 - s_{i,h_{p},h_{q}}^{\mathcal{P}}) + \left[ f_{i,h_{p}a_{2}}^{QTL} f_{i,h_{q}a_{1}}^{QTL} + f_{i,h_{p}a_{1}}^{QTL} f_{i,h_{q}a_{2}}^{QTL} \right] s_{i,h_{p},h_{q}}^{\mathcal{P}} \right]$$
(2)

Algebraic derivation of expression (3)

Let  $f_{i,h_pa_1}^{QTL} = f_{i,h_p}f_{a_1} + \Delta_{\mathbf{p}} = \alpha_p + \Delta_{\mathbf{p}}$  and  $f_{i,h_pa_2}^{QTL} = f_{i,h_p}f_{a_2} - \Delta_{\mathbf{p}} = \tilde{\alpha}_p - \Delta_{\mathbf{p}}$ . Note that we have  $\sum_{p=1}^{K} \Delta_{\mathbf{p}} = \sum_{p=1}^{K} (f_{i,h_pa_1}^{QTL} - f_{i,h_p}f_{a_1}) = f_{a_1} - f_{a_1} \sum_{p=1}^{K} f_{i,h_p} = 0$ . Replacing the haplotype frequencies in

expression (2) with the expressions including the  $\alpha_p$ ,  $\tilde{\alpha}_p$  and  $\Delta_p$  terms (same for the frequencies depending on the  $\alpha_q$ ,  $\tilde{\alpha}_q$  and  $\Delta_q$  terms) gives:

$$d(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \sum_{p=1}^{k} \left[ -4s_{i,h_{p},h_{p}}^{\mathcal{P}} \mathbf{\Delta}_{\mathbf{p}}^{2} + \left[ \alpha_{p} - \tilde{\alpha}_{p} + 4s_{i,h_{p},h_{p}}^{\mathcal{P}} (\tilde{\alpha}_{p} - \alpha_{p}) + \sum_{q \neq p}^{k} (\alpha_{q} - \tilde{\alpha}_{q}) \right] \mathbf{\Delta}_{\mathbf{p}} \right. \\ \left. + \alpha_{p}^{2} + \tilde{\alpha}_{p}^{2} + s_{i,h_{p},h_{p}}^{\mathcal{P}} (-\alpha_{p}^{2} - \tilde{\alpha}_{p}^{2} + 2\alpha_{p}\tilde{\alpha}_{p}) + \sum_{q \neq p}^{k} \alpha_{p}\alpha_{q} + \tilde{\alpha}_{p}\tilde{\alpha}_{q} \right. \\ \left. + \sum_{q \neq p}^{k} s_{i,h_{p},h_{q}}^{\mathcal{P}} \left( -4\mathbf{\Delta}_{\mathbf{p}}\mathbf{\Delta}_{\mathbf{q}} + 2(\tilde{\alpha}_{p} - \alpha_{p})\mathbf{\Delta}_{\mathbf{q}} + 2(\tilde{\alpha}_{q} - \alpha_{q})\mathbf{\Delta}_{\mathbf{p}} \right. \\ \left. - \alpha_{p}\alpha_{q} - \tilde{\alpha}_{p}\tilde{\alpha}_{q} + \tilde{\alpha}_{p}\alpha_{q} + \alpha_{p}\tilde{\alpha}_{q} \right) \right] \quad (*)$$

Replacing  $\Delta_{\mathbf{q}}$  with  $-\Delta_{\mathbf{p}} - \sum_{l \neq p,q}^{K} \Delta_{\mathbf{l}} \left( \text{ since } \sum_{p=1}^{k} \Delta_{\mathbf{p}} = 0 \right)$  in (\*) finally gives:

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{\mathcal{Q}^{TL}}) = \sum_{p=1}^{K} \left[ 4\left(\sum_{q\neq p}^{K} s_{i,h_{p},h_{q}}^{\mathcal{P}} - s_{i,h_{p},h_{p}}^{\mathcal{P}}\right) \mathbf{\Delta}_{\mathbf{p}}^{\mathbf{2}} + \left[\alpha_{p} - \tilde{\alpha}_{p} + 4s_{i,h_{p},h_{p}}^{\mathcal{P}}(\tilde{\alpha}_{p} - \alpha_{p}) + \sum_{q\neq p}^{K} \left(\alpha_{q} - \tilde{\alpha}_{q} + s_{i,h_{p},h_{q}}^{\mathcal{P}}\left(4\sum_{l\neq p,q}^{K} \mathbf{\Delta}_{\mathbf{l}} + 2(\alpha_{p} - \tilde{\alpha}_{p}) + 2(\tilde{\alpha}_{q} - \alpha_{q})\right)\right) \right] \mathbf{\Delta}_{\mathbf{p}} + \alpha_{p}^{2} + \tilde{\alpha}_{p}^{2} + s_{i,h_{p},h_{p}}^{\mathcal{P}}\left(-\alpha_{p}^{2} - \tilde{\alpha}_{p}^{2} + 2\alpha_{p}\tilde{\alpha}_{p}\right) + \sum_{q\neq p}^{K} \alpha_{p}\alpha_{q} + \tilde{\alpha}_{p}\tilde{\alpha}_{q} + \sum_{q\neq p}^{K} s_{i,h_{p},h_{q}}^{\mathcal{P}}\left(2(\alpha_{p} - \tilde{\alpha}_{p})\sum_{l\neq p,q}^{K} \mathbf{\Delta}_{\mathbf{l}} - \alpha_{p}\alpha_{q} - \tilde{\alpha}_{p}\tilde{\alpha}_{q} + \tilde{\alpha}_{p}\tilde{\alpha}_{q} + \alpha_{p}\tilde{\alpha}_{q}\right) \right]$$

Hence  $d_1(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL})$  can be expressed as (3):

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \sum_{p=1}^{K} \left[ 4 \left( \sum_{q \neq p}^{K} s_{i,h_{p},h_{q}}^{\mathcal{P}} - s_{i,h_{p},h_{p}}^{\mathcal{P}} \right) \mathbf{\Delta}_{\mathbf{p}}^{\mathbf{2}} + \Psi_{pq}^{\mathcal{P}}(\Delta_{l \neq p,q}) \mathbf{\Delta}_{\mathbf{p}} + \Phi_{pq}^{\mathcal{P}}(\Delta_{l \neq p,q}) \right]$$
$$= \xi^{\mathcal{P}}(\mathbf{\Delta}_{\mathbf{1}},..,\mathbf{\Delta}_{\mathbf{K}}) \qquad (3)$$

#### Algebraic derivation of expression (4)

Expression (4) is obtained directly from expression (3) for  $s_{i,h_p,h_p}^{\mathcal{P}} = 1$  and  $s_{i,h_p,h_q}^{\mathcal{P}} = 0$  when  $\mathcal{P} = \text{IBS}_{\text{hap}}.$ 

## Algebraic derivation of expressions (5) and (6)

Considering that only two haplotypes exist among the K possible ones is the same as setting K = 2. Expression (5) can be obtained directly by reducing expression (4) for the case where K = 2. However another derivation is given here so as to exhibit other properties, such as a lower bound, for the matrix distance. For K = 2 expression (2) becomes:

$$d_{1}(\mathbf{M}^{\mathcal{P},i}, \mathbf{M}^{QTL}) = (f_{i,h_{1}a_{1}}^{QTL-2} + f_{i,h_{1}a_{2}}^{QTL-2}) - s_{i,h_{1},h_{1}}^{\mathcal{P}} (f_{i,h_{1}a_{1}}^{QTL} - f_{i,h_{1}a_{2}}^{QTL})^{2} + 2(f_{i,h_{1}a_{1}}^{QTL} f_{i,h_{2}a_{1}}^{QTL} + f_{i,h_{1}a_{2}}^{QTL} f_{i,h_{2}a_{2}}^{QTL})(1 - s_{i,h_{1},h_{2}}^{\mathcal{P}}) + 2(f_{i,h_{1}a_{2}}^{QTL} f_{i,h_{2}a_{1}}^{QTL} + f_{i,h_{1}a_{1}}^{QTL} f_{i,h_{2}a_{2}}^{QTL})s_{i,h_{1},h_{2}}^{\mathcal{P}} - s_{i,h_{2},h_{2}}^{\mathcal{P}} (f_{i,h_{2}a_{1}}^{QTL} - f_{i,h_{2}a_{2}}^{QTL})^{2} + (f_{i,h_{2}a_{1}}^{QTL-2} + f_{i,h_{2}a_{2}}^{QTL-2})$$

and the frequencies in expression (2) can be written as:

$$f_{i,h_1a_1}^{QTL} = f_{i,h_1}f_{a_1} + \mathbf{\Delta}_1 = \alpha_1 + \mathbf{\Delta}_1$$
$$f_{i,h_1a_2}^{QTL} = f_{i,h_1}f_{a_2} - \mathbf{\Delta}_1 = \tilde{\alpha}_1 - \mathbf{\Delta}_1$$
$$f_{i,h_2a_1}^{QTL} = f_{i,h_2}f_{a_1} - \mathbf{\Delta}_1 = \alpha_2 - \mathbf{\Delta}_1$$
$$f_{i,h_2a_2}^{QTL} = f_{i,h_2}f_{a_2} + \mathbf{\Delta}_1 = \tilde{\alpha}_2 + \mathbf{\Delta}_1$$

Note that  $\Delta_1$ , in this case, can be expressed as  $\Delta_1 = f_{i,h_1a_1}^{QTL} f_{i,h_2a_2}^{QTL} - f_{i,h_1a_2}^{QTL} f_{i,h_2a_1}^{QTL}$  with its maximum and minimum value given by  $\frac{1}{4}$  and  $-\frac{1}{4}$  respectively. The maximum value of  $\Delta_1$  is given by  $f_{i,h_1a_1}^{QTL} = \frac{1}{4}$ 

 $f_{i,h_2a_2}^{QTL} = \frac{1}{2}$  and  $f_{i,h_1a_2}^{QTL} = f_{i,h_2a_1}^{QTL} = 0$ , and its minimum value is given by  $f_{i,h_1a_1}^{QTL} = f_{i,h_2a_2}^{QTL} = 0$  and  $f_{i,h_1a_2}^{QTL} = f_{i,h_2a_1}^{QTL} = \frac{1}{2}$ . Replacing the haplotype frequencies in  $d_1(\mathbf{M}^{\mathcal{P},i}, \mathbf{M}^{QTL})$  with the expressions including the  $\alpha_1, \tilde{\alpha}_1, \tilde{\alpha}_2, \alpha_2$  and  $\boldsymbol{\Delta}_1$  terms gives:

$$d_{1}(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = \left[-4s_{i,h_{1},h_{1}}^{\mathcal{P}} - 4s_{i,h_{2},h_{2}}^{\mathcal{P}} + 8s_{i,h_{1},h_{2}}^{\mathcal{P}}\right] \boldsymbol{\Delta}_{1}^{2} + \left[4s_{i,h_{1},h_{1}}^{\mathcal{P}}(\tilde{\alpha}_{1} - \alpha_{1}) + 4s_{i,h_{2},h_{2}}^{\mathcal{P}}(\alpha_{2} - \tilde{\alpha}_{2}) - 4s_{i,h_{1},h_{2}}^{\mathcal{P}}(\tilde{\alpha}_{1} + \alpha_{2} - (\alpha_{1} + \tilde{\alpha}_{2}))\right] \boldsymbol{\Delta}_{1} - s_{i,h_{1},h_{1}}^{\mathcal{P}}(\tilde{\alpha}_{1} - \alpha_{1})^{2} - s_{i,h_{2},h_{2}}^{\mathcal{P}}(\alpha_{2} - \tilde{\alpha}_{2})^{2} + 2s_{i,h_{1},h_{2}}^{\mathcal{P}}(\tilde{\alpha}_{1} - \alpha_{1})(\alpha_{2} - \tilde{\alpha}_{2}) + (\alpha_{1} + \alpha_{2})^{2} + (\tilde{\alpha}_{2} + \tilde{\alpha}_{1})^{2} = \xi^{\mathcal{P}}(\boldsymbol{\Delta}_{1})$$

Hence  $d_1(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL})$  can be expressed as:

$$\xi^{\mathcal{P}}(\mathbf{\Delta}_{\mathbf{1}}) = \left[-4s^{\mathcal{P}}_{i,h_1,h_1} - 4s^{\mathcal{P}}_{i,h_2,h_2} + 8s^{\mathcal{P}}_{i,h_1,h_2}\right] \mathbf{\Delta}_{\mathbf{1}}^{\mathbf{2}} + \Psi^{\mathcal{P}} \mathbf{\Delta}_{\mathbf{1}} + \Phi^{\mathcal{P}}$$
(5)

For the extreme values of  $\Delta_1$  we have:

$$\xi^{\mathcal{P}}\left(\frac{1}{4}\right) = \xi^{\mathcal{P}}\left(-\frac{1}{4}\right) = \frac{1}{2} + \frac{1}{2}s^{\mathcal{P}}_{i,h_1,h_2} - \frac{1}{4}(s^{\mathcal{P}}_{i,h_1,h_1} + s^{\mathcal{P}}_{i,h_2,h_2})$$

This quantity can also be obtained simply from expression (2), when K = 2, by replacing  $f_{i,h_1a_1}^{QTL}$ ,  $f_{i,h_2a_2}^{QTL}$ ,  $f_{i,h_1a_2}^{QTL}$  and  $f_{i,h_2a_1}^{QTL}$  by their corresponding values for the maximum and minimum value of  $\Delta_1$ . For  $\Delta_1 = \frac{1}{4}$  we have:

$$\begin{aligned} d_1(\mathbf{M}^{\mathcal{P},i},\mathbf{M}^{QTL}) = & \left(\left(\frac{1}{2}\right)^2 + 0^2\right) - s_{i,h_1,h_1}^{\mathcal{P}} \left(\frac{1}{2} - 0\right)^2 + 2\left(\frac{1}{2} \cdot 0 + 0 \cdot \frac{1}{2}\right) (1 - s_{i,h_1,h_2}^{\mathcal{P}}) \\ &+ 2\left(0 \cdot 0 + \frac{1}{2} \cdot \frac{1}{2}\right) s_{i,h_1,h_2}^{\mathcal{P}} - s_{i,h_2,h_2}^{\mathcal{P}} \left(0 - \frac{1}{2}\right)^2 + \left(0^2 + \left(\frac{1}{2}\right)^2\right) \\ &= \frac{1}{2} + \frac{1}{2} s_{i,h_1,h_2}^{\mathcal{P}} - \frac{1}{4} (s_{i,h_1,h_1}^{\mathcal{P}} + s_{i,h_2,h_2}^{\mathcal{P}}) \\ &= \xi^{\mathcal{P}} \left(\frac{1}{4}\right) \end{aligned}$$

In the same manner we can show that  $\xi^{\mathcal{P}}\left(-\frac{1}{4}\right) = \xi^{\mathcal{P}}\left(\frac{1}{4}\right)$  since the squares and the products of the frequencies in expression (2) when K = 2 are symmetric.  $\frac{1}{2} + \frac{1}{2}s_{i,h_1,h_2}^{\mathcal{P}} - \frac{1}{4}(s_{i,h_1,h_1}^{\mathcal{P}} + s_{i,h_2,h_2}^{\mathcal{P}})$  is greater or equal to  $\frac{1}{2}s_{i,h_1,h_2}^{\mathcal{P}}$  since the maximum possible value of  $s_{i,h_1,h_1}^{\mathcal{P}}$  and  $s_{i,h_2,h_2}^{\mathcal{P}}$  is equal to one. Hence if haplotypes  $h_1$  and  $h_2$  share allele similarity  $s_{i,h_1,h_2}^{\mathcal{P}}$  will be positive and  $\xi^{\mathcal{P}}\left(-\frac{1}{4}\right) = \xi^{\mathcal{P}}\left(\frac{1}{4}\right)$  $\in [\frac{1}{2}s_{i,h_1,h_2}^{\mathcal{P}}, 1]$ . For  $\mathcal{P} = \text{IBS}_{\text{hap}}$ , i.e.  $s_{i,h_1,h_1}^{\mathcal{P}} = s_{i,h_2,h_2}^{\mathcal{P}} = 1$  and  $s_{i,h_1,h_2}^{\mathcal{P}} = 0$ , we have  $\xi^{\mathcal{P}}\left(-\frac{1}{4}\right) = \xi^{\mathcal{P}}\left(\frac{1}{4}\right) = \xi^{\mathcal{P}}\left(\frac{1}{4}\right) = 0$ . Note that for  $\mathcal{P} = \text{IBS}_{\text{hap}}$   $\xi^{\mathcal{P}}(\Delta_1)$  becomes:

$$\xi^{\text{IBS}_{\text{hap}}}(\mathbf{\Delta}_{1}) = -8\mathbf{\Delta}_{1}^{2} + 4\left[\left(\tilde{\alpha}_{1} - \alpha_{1}\right) + \left(\alpha_{2} - \tilde{\alpha}_{2}\right)\right]\mathbf{\Delta}_{1} - \left(\tilde{\alpha}_{1} - \alpha_{1}\right)^{2} - \left(\alpha_{2} - \tilde{\alpha}_{2}\right)^{2} + \left(\alpha_{1} + \alpha_{2}\right)^{2} + \left(\tilde{\alpha}_{2} + \tilde{\alpha}_{1}\right)^{2}$$
(6)

Thus differentiating  $\xi^{\mathrm{IBS}_{\mathrm{hap}}}$  with respect to  $\Delta_1$  gives:

$$\mathbf{\Delta}_{1}^{*} = \frac{\tilde{\alpha}_{1} - \alpha_{1} + \alpha_{2} - \tilde{\alpha}_{2}}{4} = \frac{(2f_{a_{1}} - 1)(1 - 2f_{i,h_{1}})}{4}$$

The minimum of  $\Delta_1^*$ , which is equal to  $-\frac{1}{4}$ , is given by either  $f_{a_1} = f_{i,h_1} = 1$  or  $f_{a_1} = f_{i,h_1} = 0$ . Its maximum value, which is equal to  $\frac{1}{4}$ , is given by either  $f_{a_1} = 0$  and  $f_{i,h_1} = 1$  or  $f_{a_1} = 1$  and  $f_{i,h_1} = 0$ . Hence  $\Delta_1^*$  takes its minimum and maximum values when both the tested locus and the QTL are monomorphic.

#### Algebraic derivation of the matrix distance for a multiallelic QTL

For S distinct alleles at the QTL expression (2) generalizes to:

$$d_1(\mathbf{M}^{\mathcal{P},i}, \mathbf{M}^{QTL}) = \sum_{p=1}^K \sum_{q=1}^K \left[ (1 - s_{i,h_p,h_q}^{\mathcal{P}}) \sum_{l=1}^S f_{i,h_pa_l}^{QTL} f_{i,h_qa_l}^{QTL} + s_{i,h_p,h_q}^{\mathcal{P}} \sum_{l=1}^S \sum_{m\neq l}^S f_{i,h_pa_l}^{QTL} f_{i,h_qa_m}^{QTL} \right]$$
(7)

For  $\mathcal{P} = \text{IBS}_{\text{hap}}$  expression (7) becomes:

$$d_1(\mathbf{M}^{\mathrm{IBS_{hap}},i}, \mathbf{M}^{QTL}) = \sum_{p=1}^K \sum_{l=1}^S \sum_{m \neq l}^S f_{i,h_p a_l}^{QTL} f_{i,h_p a_m}^{QTL} + \sum_{p=1}^K \sum_{l=1}^S \sum_{q \neq p}^K f_{i,h_p a_l}^{QTL} f_{i,h_q a_l}^{QTL}$$
(7)

Let  $\mathbf{\Delta}_{\mathbf{pl}} = f_{i,h_pa_l}^{QTL} - f_{i,h_p}f_{a_l} = f_{i,h_pa_l}^{QTL} - \alpha_{pl}$ , which is equivalent to  $f_{i,h_pa_l}^{QTL} = \alpha_{pl} + \mathbf{\Delta}_{\mathbf{pl}}$ . Note that:

$$D_{i,QTL}^{2} = \sum_{p=1}^{K} \sum_{l=1}^{S} (f_{i,h_{p}a_{l}}^{QTL} - f_{i,h_{p}} \cdot f_{a_{l}})^{2} = \sum_{p=1}^{K} \sum_{l=1}^{S} \Delta_{\mathbf{pl}}^{2}$$

Since  $\sum_{p=1}^{K} \Delta_{\mathbf{pl}} = 0$  we have  $\sum_{q \neq p}^{K} \Delta_{\mathbf{ql}} = -\Delta_{\mathbf{pl}}$  and  $\sum_{m \neq l}^{S} \Delta_{\mathbf{pm}} = -\Delta_{\mathbf{pl}}$ . Replacing the haplotype fre-

quencies in expression (7) with the LD coefficients and the product of frequencies terms, and subsequently replacing  $\sum_{q \neq p}^{K} \Delta_{ql}$  and  $\sum_{m \neq l}^{S} \Delta_{pm}$  with  $-\Delta_{pl}$ , gives:

$$\begin{split} d_1(\mathbf{M}^{\mathrm{IBS}_{\mathrm{hap}},i},\mathbf{M}^{\scriptscriptstyle QTL}) &= \sum_{p=1}^K \sum_{l=1}^S \left[ -2\boldsymbol{\Delta}_{\mathbf{pl}}^2 + \left( \Psi_{pl}^{\mathrm{IBS}_{\mathrm{hap}},(1)} + \Psi_{pl}^{\mathrm{IBS}_{\mathrm{hap}},(2)} \right) \boldsymbol{\Delta}_{\mathbf{pl}} \right. \\ &+ \left( \Phi_{pl}^{\mathrm{IBS}_{\mathrm{hap}},(1)} + \Phi_{pl}^{\mathrm{IBS}_{\mathrm{hap}},(2)} \right) \right] \\ &= \xi^{\mathrm{IBS}_{\mathrm{hap}}}(\boldsymbol{\Delta}_{\mathbf{11}}, \boldsymbol{\Delta}_{\mathbf{12}}, ..., \boldsymbol{\Delta}_{\mathbf{KS}}) \end{split}$$

As for expression (3) the general behavior of the matrix distance for continuous predictors in [0, 1], as function of LD coefficients, is unspecifiable for the multiallelic QTL case. Hence we did not express the matrix distance, here in the multiallelic QTL case, for continuous predictors in [0, 1].