² Derivations including the effects of epistasis between pairs of selected loci can be ³ found in a *Mathematica* notebook available as supplementary material. Although it is ⁴ possible to obtain general expressions valid for arbitrary selfing rate, these expressions ⁵ quickly become cumbersome when $\overline{\sigma} > 0$ and we thus only present results relative to ⁶ the initial spread of selfing in an outcrossing population ($\overline{\sigma} \approx 0$). In that case, the ⁷ change in mean selfing rate per generation can still be decomposed in a change during ⁸ selection $\Delta_{s}\overline{\sigma}$ and a change during reproduction $\Delta_{r}\overline{\sigma}$, with:

$$\Delta_{\rm r}\,\overline{\sigma} \approx \frac{1-\kappa}{2}\,V_{\sigma} \tag{D1}$$

⁹ representing direct selection for selfing (automatic transmission advantage), while
¹⁰ (from equation A1 in Supplementary File S1):

$$\Delta_{s} \overline{\sigma} = 2 \sum_{i,j} a_{j} \left(\tilde{D}_{ij} + \tilde{D}_{i,j} \right) + 2 \sum_{i,j} a_{j,j} \tilde{D}_{ij,j}$$

$$+ 2 \sum_{i,j < k} a_{jk} \left(\tilde{D}_{ijk} + \tilde{D}_{i,jk} \right) + 2 \sum_{i,j < k} a_{j,k} \left(\tilde{D}_{ij,k} + \tilde{D}_{ik,j} \right)$$

$$+ 2 \sum_{i,j < k} a_{jk,j} \left(\tilde{D}_{ijk,j} + \tilde{D}_{ij,jk} \right) + 2 \sum_{i,j < k} a_{jk,k} \left(\tilde{D}_{ijk,k} + \tilde{D}_{ik,jk} \right)$$

$$+ 2 \sum_{i,j < k} a_{jk,jk} \tilde{D}_{ijk,jk}.$$
(D2)

¹¹ Leading-order expressions fort the different genetic associations that appear in equa-¹² tion D2 can be obtained using the same methods as in Supplementary File S3 (see also ¹³ Mathematica notebook). The association $\tilde{D}_{ijk,jk}$ is generated by the effect of locus *i* ¹⁴ on the selfing rate, even in the absence of selection at loci *j* and *k* (representing the ¹⁵ fact that alleles increasing the selfing rate at locus *i* tend to be more often present in individuals which are homozygous at loci j and k). To leading order, it is given by:

$$\tilde{D}_{ijk,jk} \approx \frac{1}{2} \left[1 - 2\rho_{jk} \left(1 - \rho_{jk} \right) \right] \tilde{D}_{ii} pq_{jk} \tag{D3}$$

where $pq_{jk} = p_j q_j p_k q_k$. Associations $\tilde{D}_{ijk,j}$ and $\tilde{D}_{ij,jk}$ are given by:

$$\tilde{D}_{ijk,j} \approx \tilde{D}_{ij,jk} \approx \frac{1}{2} \left(1 - 2p_j \right) \left[\left(1 - \rho_{jk} \right) \tilde{D}'_{jk} + \rho_{jk} \, \tilde{D}'_{j,k} \right] \tilde{D}_{ii} \tag{D4}$$

¹⁸ where \tilde{D}'_{jk} and $\tilde{D}'_{j,k}$ are the associations \tilde{D}_{jk} and $\tilde{D}_{j,k}$ measured after selection. To the ¹⁹ first order in $a_{\mathbb{U},\mathbb{V}}$ coefficients, $\tilde{D}_{i,jk} = 0$ at QLE while $\tilde{D}_{ij,j}$ and $\tilde{D}_{ij,k}$ are given by:

$$\tilde{D}_{ij,j} \approx \frac{1}{2} p_j q_j \, \tilde{D}_{ii}, \quad \tilde{D}_{ij,k} \approx \frac{1}{2} \left(\tilde{D}'_{jk} + \tilde{D}'_{j,k} \right) \tilde{D}_{ii}.$$
 (D5)

Summing these expressions over all loci affecting the selfing rate, and using $V_{\sigma} = 2\sum_{i} \tilde{D}_{ii}$ and the expression for inbreeding depression measured after selection (δ') given by equation B9 in Supplementary File S2, equation D2 becomes:

$$\Delta_{\rm s}\,\overline{\sigma} = -\delta'\,V_{\sigma} + 2\sum_{i,j}a_j\left(\tilde{D}_{ij} + \tilde{D}_{i,j}\right) + 2\sum_{i,j< k}a_{jk}\,\tilde{D}_{ijk}\,.\tag{D6}$$

²³ To the first order in $a_{\mathbb{U},\mathbb{V}}$ coefficients, \tilde{D}_{ijk} at QLE is given by:

$$\tilde{D}_{ijk} \approx \frac{1}{\rho_{ijk} - (1 - \rho_{ijk}) [a_j (1 - 2p_j) + a_k (1 - 2p_k) + a_{jk} (1 - 2p_j) (1 - 2p_k)]} \times \left[\left[(1 - \rho_{jk}) a_{jk} + \rho_{jk} a_{j,k} + a_{jk,j} (1 - 2p_j) + a_{jk,k} (1 - 2p_k) + a_{jk,jk} (1 - 2p_j) (1 - 2p_j) \right] \tilde{D}_{ijk,jk} + \left[(1 - \rho_{jk}) a_{j,k} + \rho_{jk} a_{jk} + a_{jk,j} (1 - 2p_j) \right] p_k q_k \tilde{D}_{ij,j} + \left[(1 - \rho_{jk}) a_{j,k} + \rho_{jk} a_{jk} + a_{jk,k} (1 - 2p_k) \right] p_j q_j \tilde{D}_{ik,k} + \frac{1}{2} (a_{jk} + 2\rho_{jk} a_{j,k}) \tilde{D}_{ii} pq_{jk} \right]$$
(D7)

²⁴ where ρ_{ijk} is the probability that at least one recombination event occurs between ²⁵ the three loci *i*, *j*, *k* during meiosis, and where $\tilde{D}_{ijk,jk}$, $\tilde{D}_{ij,j}$ and $\tilde{D}_{ik,k}$ are given by

equations D3 and D5. Using equations A12 – A14 in Supplementary File S1, equation 26 D7 yields equation 45 in the main text for the case of uniformly deleterious alleles with 27 fixed epistasis (assuming that p_j , p_k are small). In the case of stabilizing selection on 28 quantitative traits, the terms in $a_{jk,j}$, $a_{jk,k}$ and $a_{jk,jk}$ in equation D7 all generate terms 29 in $(1 - 2p_j)(1 - 2p_k)$ (see equations A25 – A26 and A54 – A55 in Supplementary File 30 S1), that should vanish when summed over a large enough number of loci coding for 31 the traits (due to the fact that two symmetric equilibria exist for p_j , one with $p_j < 1/2$ 32 and the other with $p_j > 1/2$). This yields (using the fact that $a_{jk} = a_{j,k}$, and assuming 33 that recombination rates are large relative to selection coefficients): 34

$$\sum_{j,k} a_{jk} \tilde{D}_{ijk} \approx \sum_{j,k} \frac{2 + \rho_{jk}^2}{\rho_{ijk}} a_{jk}^2 p q_{jk} \tilde{D}_{ii} \,. \tag{D8}$$

Using the fact that $\sum_{j,k} a_{jk}^2 p q_{jk} \approx 4U^2/n$ independently of the parameter Q describing the shape of the fitness peak (see equation A49 in Supplementary File S1), equation D8 yields equation 50 in the main text.

Finally, we have $\tilde{D}_{i,j} \approx 0$ when $\overline{\sigma} \approx 0$, while \tilde{D}_{ij} is given by (to the first order in $a_{\mathbb{U},\mathbb{V}}$ coefficients):

$$\tilde{D}_{ij} \approx \frac{1}{\rho_{ij} - (1 - \rho_{ij}) a_j (1 - 2p_j)} \bigg[[a_j + a_{j,j} (1 - 2p_j)] \tilde{D}_{ij,j} + \sum_k a_{jk,k} p_j q_j \tilde{D}_{ik,k} + \sum_k [a_{jk,k} + a_{jk,jk} (1 - 2p_j)] \tilde{D}_{ijk,jk} \bigg].$$
(D9)

⁴⁰ Using equations D3 and D5, equation D9 yields equation 44 in the main text.