

## Additional file 2

Table S1 contains genetic parameters for the mean and the variance of Box-Cox transformed harvest weight estimated using Double Hierarchical Generalized Linear Models.

**Table S1 Genetic parameters for the mean and the variance of Box-Cox transformed harvest weight**

Parameter	Mean	Variance
<sup>a</sup> $\sigma_A^2$	0.100 (0.019) <sup>b</sup>	0.239 (0.052)
$\sigma_e^2$	0.184 (0.004)	1.804 (0.035)
$\sigma_g^2$	0.050 (0.007)	0.035 (0.019)
$\sigma_k^2$	0.032 (0.004)	0.048 (0.025)
$\sigma_m^2$	0.008 (0.003)	0.014 (0.009)
$\sigma_P^2$	0.323 (0.011)	–
$h^2$	0.31 (0.05)	–
<sup>b</sup> $g^2$	0.15 (0.02)	–
<sup>c</sup> $k^2$	0.10 (0.02)	–
<sup>d</sup> $m^2$	0.02 (0.01)	–
<sup>e</sup> GCV	0.06	0.49

Standard errors are indicated between brackets

<sup>a</sup>Additive genetic variance was calculated as 4 times the sire-dam variance

<sup>b</sup>Group effect, calculated as  $g^2 = \sigma_g^2 / \sigma_P^2$

<sup>c</sup>Kin effect, calculated as  $k^2 = \sigma_k^2 / \sigma_P^2$

<sup>d</sup>Social maternal effect, calculated as  $m^2 = \sigma_m^2 / \sigma_P^2$

<sup>e</sup>Genetic coefficient of variation