

Table S4. Heritability (h^2), additive genetic variance (σ^2_a) and phenotypic variance (σ^2_t) estimated considering the pedigree-based (A) relationship matrix. Standard errors are reported in parenthesis.

Trait*	h^2		σ^2_a		σ^2_t	
UTW	0.72	(± 0.08)	1.75	(± 0.26)	2.44	(± 0.13)
TTW	0.68	(± 0.09)	1.08	(± 0.19)	1.58	(± 0.10)

*UTW, Untrimmed thigh weight; TTW, Trimmed thigh weight.