	Genotype	Plant number	X ² (1:2:1)	Phenotype	Additive effect	Dominant effect
GNP	LT homozygotes	41		154.3		
	Heterozygotes	80		203.4		
	TQ homozygotes	42	0.07	229.6	37.7	11.5

S2 Table. The effect of *GNP1* revealed in the BC_5F_3 population derived from a near-isogenic line with BC_5F_2 heterozygous at the RM227–RM85 region and confirmed by BC_5F_4 family data.