

Table S6. Alternative allele frequency (\pm St dev) in the deletion identified in chromosome 3 of ARR and NER

Sequencing	CLE	ARR	NER
Illumina ^a	0.48 \pm 0.1	0.81 \pm 0.1	0.71 \pm 0.1
Sanger ^b			
Young leaf			
Amplification	0.52 \pm 0.1	0.79 \pm 0.1	0.80 \pm 0.1
TA cloning	0.58 \pm 0.1	0.78 \pm 0.1	0.85 \pm 0.1
Fruit flavedo			
Amplification	0.54 \pm 0.0	0.78 \pm 0.1	0.75 \pm 0.1

^a Average allelic frequency based on Illumina reads (n > 10000)

^b Average allelic frequency based on after either direct amplification or after TA cloning