

Supplementary Table S1. Mapping of RNA-seq reads to carnation scaffold sequences

Sample	Input	Unique	Multiple	Unmapped
untouched rep1	21.2	15.24 (71.9%)	3.26 (15.36%)	2.7 (12.75%)
untouched rep2	23.3	16.56 (71.1%)	3.61 (15.49%)	3.13 (13.41%)
untouched rep3	21.08	14.98 (71.09%)	3.23 (15.32%)	2.86 (13.59%)
touched rep1	21.58	15.48 (71.74%)	3.33 (15.42%)	2.77 (12.84%)
touched rep2	24.73	17.66 (71.41%)	3.81 (15.39%)	3.27 (13.2%)
touched rep3	24.84	17.85 (71.85%)	3.84 (15.47%)	3.15 (12.68%)

Numbers indicate million reads, and ratio to input reads are shown in parenthesis.