

Supplementary Material

Supplementary Table 2. Comparison of assembled sequences and those previously reported for *Fragaria chiloensis* in GenBank

Sequence	GenBank	RNAseq transcript id	% id	Alignment length	E-value
Fc_AAT1	FJ548610.1	comp6_c0_seq1	99.85	1353	0.0
Fc_MADSbox1	GQ398009.1	comp135_c0_seq2	99.81	1060	0.0
Fc_MYB1	GQ867222.1	comp1249_c0_seq1	99.65	564	0.0
Fc_PG1	EF441274.1	comp47_c0_seq1	99.84	1218	0.0
Fc_PL1	EF441273.1	comp34_c0_seq2	99.10	1217	0.0
Fc_XTH1	GQ280283.1	comp279_c0_seq2	98.98	886	0.0
Fc_XTH2	GQ280284.1	comp654_c0_seq1	99.56	919	0.0