

1 **Table S1.** Primers used for REMAP, annealing temperatures, number of total and polymorphic bands scored for each primer combination used in
 2 the analysis.

Primer ID	Sequence (5'→3')	Ann. temp. (°C)	No. of bands scored/polymorphic
ISSR			
UBC-807	(AG) ₈ T	51	12/9
UBC-808	(AG) ₈ C	53	19/15
UBC-811	(GA) ₈ C	50	18/16
UBC-868	(GAA) ₆	55	17/15
9900F ^[a]	(GTG) ₇ A	58	21/16
LTR			
<i>Sukkula</i> ^[a]	ATAGGGTCGCATCTTGGGCGTGAC		

^[a] Described in Baumel *et al.* (2002)

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1 **Table S2.** Nuclear microsatellite markers used in this study, repeat motifs, position and primers sequence for PCR amplification.

Primer ID	Repeat	Position	Primers
Em133 ^[a]	(AC) ₁₃ (AT) ₄	Genomic	For: GCGGATCACCTGCAGTTACATTAC Rev: TCCTTTGACCTATAGTGGCACGTAGT
Em135 ^[a]	(CA) ₁₁ (GA) ₂₀	Genomic	For: ATCCTGTTGCTGCTCATTTTCCTC Rev: AGGAGGATCCAAGAGGTTTGTGTA
EEMS15 ^[b]	(C) ₁₂	Nuclear-EST	For: GGGACAAATCTGACCTTTGG Rev: CTGGTGGCAAATTCTTCGAT
EEMS49 ^[b]	(TA) ₁₂ (GA) ₇	Nuclear-EST	For: TGAAATTGATCAATACCTATAAATTTAG Rev: GAAAGCCAGGATAGCATTCG
Emg11A06 ^[c]	(AG) ₂₂	Genomic	For: AGTGCTAATATGCAAGGGGAATGG Rev: GTTTACGGTGATCTTCCGTATTCCAAA
Emg21A08 ^[c]	(TC) ₁₄ (TG) ₁₄ TCTGTCTG(TC) ₂₅ CA(TC) ₃	Genomic	For: ATGGCAAGGACTGAGGTATCACAA Rev: GTTTCCGCTTATTTGATGGATCTTGC
Emf11D18 ^[c]	(TA) ₁₀ (TG) ₅₁	Genomic	For: AGAGACAGGGAGAGTGCATTCTATG Rev: GTTTGCAGTTCATAAGGTTGCATCAATAC
Emf21C11 ^[c]	(TC) ₂₄	Genomic	For: AGGTTGGAGCCATGATTACTTGAA Rev: GTTTGCTACCTATCAAACAGGCGGAA
Emg11M09 ^[c]	(TC) ₇ TT(TC) ₃₁	Genomic	For: ATACATTGAAATTGGCTGAGCTTG Rev: GTTTGGATCTTCGCTAGAACTTTGGC

2 ^[a]Nunome *et al.*, 2003; ^[b]Stàgel *et al.*, 2008; ^[c]Nunome *et al.*, 2009

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