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dCAPS GENOTYPING FOR 2010 PROJECT

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GENE/ Allele	FORWARD PRIMER (5' to 3')	REVERSE PRIMER (5' to 3')	ENZYME	DESCRIPTION	SOURCE
MS1/ ms1-1	GCCATTTCATCGAC CCTTGTG	CGTGTGTGTTTGCG TGGTTGGACA	RsaI	2870 bp product WT cuts; *ms1-1* does not	Wilson et al., 2001. The Plant Journal 28(1): 27-39
WUS/ wus1-1	AGTAGCCATGTCTA TGGATC**C**ATG	GTGCATAGGGAAG AGAGGAAGC	NcoI	312 bp product WT cuts to form a product of 292 bp *wus-1* does not cut	R. S. Lamb
SPT/ spt-2	TGTTTATCTTTCTTG TAACAGA**CC**A	ATAGACTGGTTTT AGGATTTTGG	ScrFI	305 bp product WT cuts to form a prouct of 281 bp *spt-2* does not cut	R. S. Lamb
CAL/ cal-1	GAGAAGTAAGAAA ACGTACGATGAAAC TACTTG	TTGTTTTGTCCTTA GGATGTTTTCCCTC TC	MboII	318 bp product *cal-1* cuts to produce a 299 bp fragment WT doesn't cut	M. Yang
EGL3/ egl3-1	CGCAGTTTCAGTTC GAAACATTC**C**ATG	CCCCACTCCAACA CTCTTCGTTTAG	NcoI	272 bp product WT is cut by *NcoI*	M. Yang
GL3/ gl3-1	CCATCAGTTAATTC TCGGAC**T**G	GAAGAATCGATTA TATAAATATACCTT GTTG	PstI	322 bp product PstI cuts WT, not *gl3-1*	M. Yang
AP1/ ap1-1	GTATGGAGAAGATA CTTGAACGACGTC	GCAAGTCTTCCCC AAGATAA**G**GC	StuI	360 bp product WT product is cut by *StuI* to 338bp ap1-1 not cut	E. Chae Yale University

All primers were designed, if possible, to amplify a region that includes an intron or within an intron, to allow the differentiation between genomic copies and cDNA containing transgenes. Product lengths noted are always for genomic DNA. Residues in bold = introduced mismatches designed to introduce the restriction enzyme cut site.