

Correction to "Whole Genome Shotgun Sequences for Microsatellite Discovery and Application in Cultivated and Wild Macadamia (Proteaceae)"

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Erratum

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The authors discovered an error in a primer sequence reported in Table 1. The corrected primer sequence (locus Mac0010, reverse primer sequence) is listed below in boldface. The authors apologize for the error. The published article (full-text and PDF) has been corrected.

TABLE 1. Correction to: Characterization of 12 polymorphic microsatellite loci developed in Macadamia integrifolia.^a

Locus		Primer sequences $(5'-3')$	Repeat motif	Fluorescent label	Allele size range (bp)	$T_{\rm a}(^{\circ}{\rm C})$	GenBank accession no
Mac001	F:	GTGACTGGTGGACACCAAAACCCA	(AT) ₁₁	VIC	412-420	60	KF130888
	R:						
Mac002	F:	CCCAACTGGGTTTGCAAGGACCAA	(CT) ₈	NED	283–297	60	KF130889
	R:	AGTAGCCGCGAGCTGATCGAAGAT					
Mac003	F:	TGGACCATTGAGGAGTTGGACTGT	$(AT)_{9}$	FAM	258-276	60	KF130890
	R:	TCCACCGTTTCACTTTCGTCAGCC					
Mac004	F:		(AT) ₁₁	NED	224–240	60	KF130891
	R:	GGGAGACATCATACTTTTGACACATGCC					
Mac005	F:	011111001110110111100001111111	$(AAG)_{10}$	FAM	331–343	60	KF130892
	R:						
Mac006	F.:	TTTCATCATTGATCATCATAGGTACA	(AG) ₁₁	PET	322-360	55	KF130893
	R:						
Mac007	E.	AGGCCTTGGGATGTTCCAGTGTGA	$(CT)_{11}$	NED	368-390	60	KF130894
	F:	GCAATCAACACAAGCACCTGTGGC AACGGTTATGTCAAGTGCAACAGGA		E 434	200 200	(0)	WE120005
Mac008	r: R:		$(AT)_{10}$	FAM	388–398	60	KF130895
Mac009	F:		(AAG) ₁₃	VIC	241–244	60	KF130896
	P.	TAAATCTATGCCACATCACTAGGC					
Mac010	F:		(AG) ₁₁	PET	259–297	55	KF130897
	R:	TCCGATCATAGTCTTAGCATTTCA					
Mac011	F:	AGAGGGCGAGATCCCTGACTCTGA	(CT) ₉	FAM	175–199	60	KF130898
	R:	TGAAATTTGGCGTGGGGAAAGCGT					
Mac012	F:		(AC) ₁₀	VIC	309-321	60	KF130899
		GCCTGTTGTAGGTAAAGTGGAGAT	$(\Lambda C)_{10}$	VIC	507-521	00	Ki 150077

Note: T_a = annealing temperature used for all *Macadamia* species and cultivars.

^aValues based on 22 samples representing Macadamia cultivars located at Clunes Varietal Trial M2, New South Wales, Australia.

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