Erratum

Characterisation of microsatellites from *Actinidia chinensis*

Kurt Weising^{1,2}, Raymond W.M. Fung¹, D. Jeannette Keeling¹, Ross G. Atkinson³ & Richard C. Gardner¹, *

¹Centre for Gene Technology, School of Biological Sciences, University of Auckland, Private Bag 92019, Auckland, New Zealand (* author for correspondence); ²Plant Molecular Biology Group, Biozentrum, Johann Wolfgang Goethe University, Marie-Curie-Strasse 9, D-60439 Frankfurt, Germany; ³Horticulture and Food Research Institute of New Zealand Ltd, Private Bag 92169, Mt Albert, Auckland, New Zealand

Molecular Breeding **2**: 117–131, 1996.

Table 2 of the above article (p. 121) was printer incorrectly. The correct version is printed on the following page.

Table 2. Design of oligonucleotide primers for 18 cloned *Actinidia chinensis* microsatellites.

Prime pair	Plasmid repeat	Oligo sequence																		Size
714	(CA) ₁₀	5'	ATT	CCT	CGG	CCC	CTC	ACT	AG	3'	5'	GGT	TGC	AAC	TAT	TCT	TCA	CC	3'	100
717	(CT) ₁₀ :(CA) ₅ (AC) ₃	5'	CAA	GCC	CAT	TAG	TTC	AGC	CC	3'	5'	ATT	TGG	TAA	GTA	GAA	ATC	GG	3'	173
718	(CA) ₈	5'	CAG	CTT	CAT	TAT	GTT	CCA	CC	3'	5'	AAG	AGT	GCA	TGC	ATT	GGA	GG	3'	122
719	(CA) ₁₅	5'	CGA	GTT	CTA	TTA	AAT	TCA	CC	3'	5'	ATC	TGC	ATG	CCA	TCT	TCC	AG	3'	181
720	(CA) ₁₁	5'	ACT	CCA	ATT	CCA	CCA	TAT	CC	3'	5'	ATC	ATC	AAA	ACC	CAT	TTG	CC	3'	91
721	(CA) ₅ :(CA) ₁₇ :(CT) ₁₅	5'	TAC	CCA	TTT	ATG	GAA	AAA	GC	3'	5'	GTC	AAT	GTG	CTT	CAT	GAG	TC	3'	206
722	(CT) ₁₅	5'	TGT	CCA	TAT	GTA	GTA	AGT	GC	3'	5'	AGT	TTT	GGA	AAT	ATC	TCC	GC	3'	117
723	(CT) ₁₅	5'	GAT	CAC	ATA	CCA	CAA	<u>TCT</u>	<u>C</u>	3'	5'	TTA	TAA	CTC	AAT	GGA	CAC	CC	3'	147
735	(CT) ₉ :(CTAT) ₇ :(CA) ₁₃ :(GT) ₅	5'	TGC	CAA	TGA	AAC	CCA	AAA	AG	3'	5'	ATA	GTT	AGT	TTA	TGG	GTT	TG	3'	182
747	(CT) ₁₀	5'	CCT	CTA	CCG	CTT	CAT	AAG	TC	3'	5'	TCT	TGA	AGG	TTT	GGG	TTC	CC	3'	139
751	(CT) ₈	5'	GCA	AGC	CCT	AAA	TTC	AAT	AG	3'	5'	ATC	TCC	ACC	GAA	CTT	GGC	CG	3'	187
760	(CT) ₈	5'	GAT	CCA	TCC	AAA	ATC	<u>CT</u>		3'	5'	ACT	GTA	TGG	TGC	AAA	ATT	GG	3'	99
761	(CT) ₂ CC(CT) ₈	5'	GAT	TTG	CTG	TTT	GTA	AGG	CC	3'	5'	ATC	GCA	CTT	CCA	AGC	CTC	TC	3'	152
764	(CT) ₁₇	5'	TCG	AAA	TGT	CCC	TAA	TCG	CC	3'	5'	GCA	GTT	TCC	GTC	ATT	TAG	CC	3'	177
766	(CTT) ₄ C CTT CAT (CTT) ₂ : CAG(CGG) ₃	5'	CAT	CGA	AGC	ATA	TGA	AAC	AG	3'	5'	CTC	CTG	AGG	TTA	AGA	CGG	AG	3'	149
767	(CTT) ₅	5'	GAT	CAG	TTG	AGG	CAA	<u>GAA</u>	<u>G</u>	3'	5'	GGA	GTC	ATC	TAT	GGA	ACT	TG	3'	131
768	(CTT) ₅	5'	GAT	CCG	GGT	TTT	<u>AAG</u>	<u>AAG</u>		3'	5'	ATG	GAG	TTC	CTA	TGC	TTC	TG	3'	211
769	(CTT) ₈	5'	ACA	GCT	GAG	TTG	GCA	ATA	TG	3'	5'	AGG	AAT	CAC	CTG	TGT	TAG	TG	3'	118

Four of the primers include part of microsatellite (underlined). A colon is used in the repeat description to indicate non-contiguous microsatellites. The size is given for the predicted product of the PCR reaction using plasmid DNA as template, based on sequence information.