

## Erratum to: Assessment of genetic and epigenetic changes during cell culture ageing and relations with somaclonal variation in *Coffea arabica*

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There were errors in the methylation patterns shown in the bottom row of Table 2 in the original publication. The correct table is shown below.

**Table 2** MSAP DNA electrophoretic pattern as revealed by the specificity of the restriction enzymes *HpaII* and *MspI* to different methylation states of the symmetric sequence CCGG

Restriction enzymes	MSAP DNA electrophoretic patterns after enzymatic digestion			
	Pattern 1	Pattern 2	Pattern 3	Pattern 4
<i>HpaII</i>	1	1	0	0
<i>MspI</i>	1	0	1	0
Methylation state	Unmethylated	Hemi-methylated	Fully-methylated	Fully-methylated
Methylation position	None	External cytosine	Internal cytosines	External cytosines
Schematic representation	CCGG	CH <sub>3</sub>	CH <sub>3</sub>	CH <sub>3</sub>
	GGCC	CCGG	CCGG	CCGG
		GGCC	GGCC	GGCC
			CH <sub>3</sub>	CH <sub>3</sub>

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