

Erratum to: A segregation distortion locus located on linkage group 4 of the chickpea genetic map

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Due to an unfortunate mistake in the production process, part of the heading of Table 2 is missing in

the above mentioned publication. The correct representation of Table 2 is published below and should be treated as definitive by the reader:

Table 2 Chi-square analysis of non-Mendelian segregation under three different hypotheses ($\chi^2_{(p=q)}$; $\chi^2_{(p^2:2pq:q^2)}$; $\chi^2_{(0.5p:0.5:0.5q)}$) in a chickpea F₂ population (F₂-1) derived from the intraspecific cross ILC3279 × WR315

Marker	Gametic selection in male and female				Gametic selection in male or female		
	Allelic frequency		χ^2_1		Allelic frequency		χ^2_1
	<i>p</i>	<i>q</i>	<i>p</i> = <i>q</i>	<i>p</i> ² :2 <i>pq</i> : <i>q</i> ²	<i>p</i>	<i>q</i>	0.5 <i>p</i> :0.5:0.5 <i>q</i>
GAA47	0.405	0.595	42.45***	0.00	0.315	0.684	0.68
TA72	0.285	0.715	215.59***	35.37***	0.062	0.937	0.14
SC-Y17	0.274	0.726	235.76***	34.99***	0.051	0.948	0.04

*, ** and *** are significant at $P < 0.05$, $P < 0.01$ and $P < 0.001$ levels, respectively

The online version of the original article can be found under doi:[10.1007/s10681-011-0356-7](https://doi.org/10.1007/s10681-011-0356-7).

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