Erratum to: Mapping of a gene that confers short lateral branching (*slb*) in melon (*Cucumis melo* L.)

N. Fukino · T. Ohara · M. Sugiyama ·
N. Kubo · M. Hirai · Y. Sakata · S. Matsumoto

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Regrettably, an error was introduced in the last row of Table 3 of the above mentioned publication. The

position of the QTL on LG XI (flanking markers CMAGN89 and ECM149) should have read 53.8–88.9 (instead of 16.6–39.7). The correct representation of Table 3 is published on the next page and should be treated as definitive by the reader:

The online version of the original article can be found under doi:[10.1007/s10681-012-0667-3](https://doi.org/10.1007/s10681-012-0667-3).

N. Fukino (✉) · T. Ohara · M. Sugiyama ·
Y. Sakata · S. Matsumoto
NARO Institute of Vegetable and Tea Science,
Tsu, Mie 514-2392, Japan
e-mail: nbk@affrc.go.jp

N. Kubo · M. Hirai
Graduate School of Life and Environmental Sciences,
Kyoto Prefectural University, Seika,
Kyoto 619-0244, Japan

Present Address:

Y. Sakata
NARO Agricultural Research Center for Kyushu Okinawa
Region, Kurume, Fukuoka 839-8503, Japan

Table 3 QTLs for lateral branching in 94 F₂ plants derived from Nou-4 × Harukei-3, detected by using composite interval mapping

Linkage group	Flanking markers	Position ^a	Threshold LOD	Maximum LOD	R^2 ^b	Additive effect ^c	Dominance effect
III	ECTAMACA218 and TJ10	6.0–18.8	3.5	4.2	0.10	2.31	−8.00
XI	CMAGN89 and ECM147	53.8–88.9		12.5	0.51	−13.2	0.82

^a Expressed in Kosambi cM

^b R^2 is the proportion of the phenotypic variance explained by the QTL

^c Additive effects indicate an additive main effect of the parent contributing the higher-value allele. A positive value indicates that the higher-value allele (i.e., contributing to longer branches) came from Nou-4; a negative value indicate that the higher-value allele came from Harukei-3