



## Correction to: Both overlapping and independent loci underlie seed number per pod and seed weight in *Brassica napus* by comparative quantitative trait loci analysis

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**Correction to: Mol Breeding (2021) 41:41**

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The original version of this article unfortunately contained incorrect Fig. 3b. The authors of this publication apologize for any inconvenience caused by this mistake. The corrected Fig. 3b should be.

Linkage disequilibrium analysis and haplotype analysis of 5 associated regions (a–e). Top, the horizontal axis represents haplotypes and the vertical axis represents the phenotypic values of

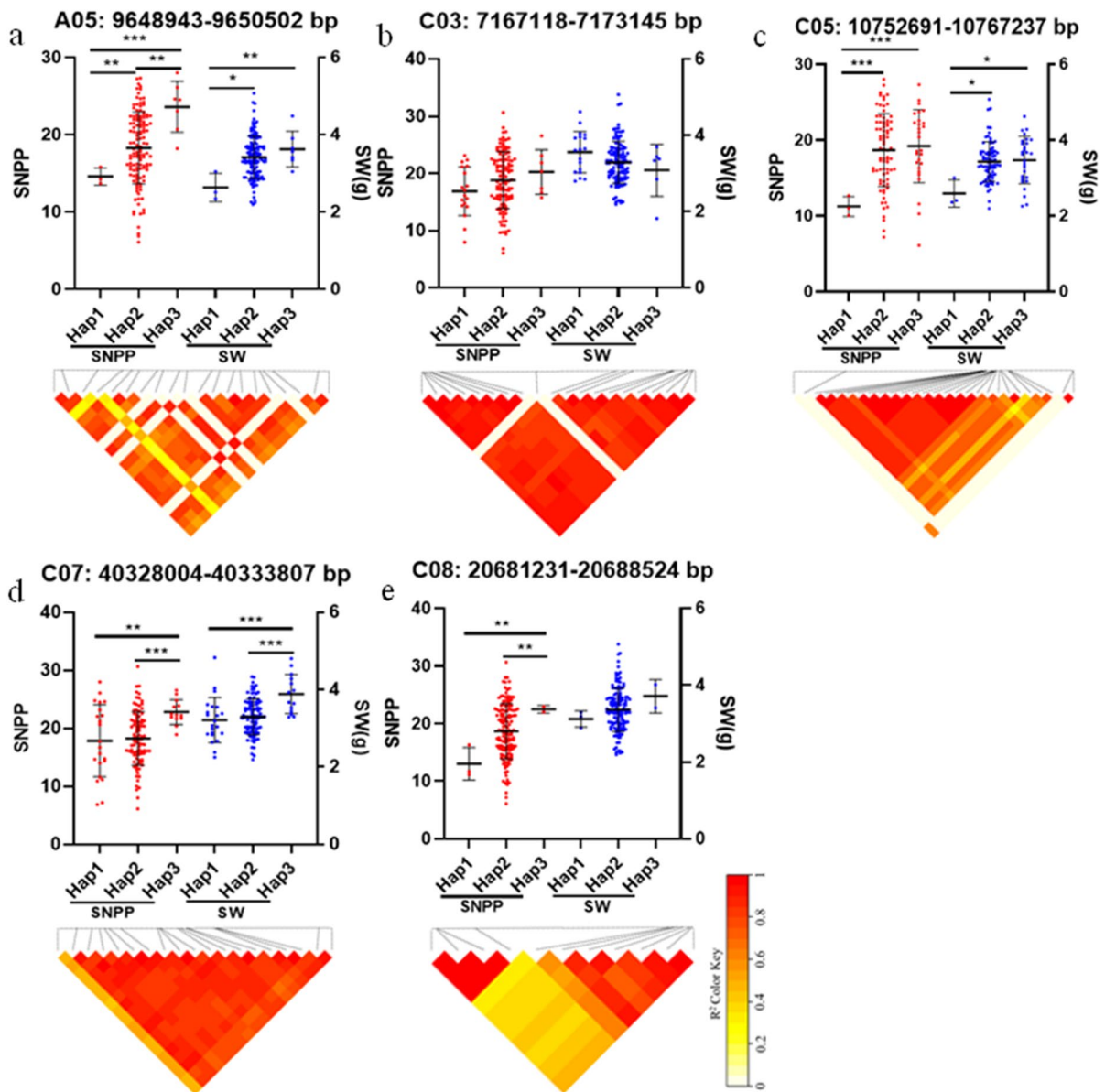
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**Fig. 3** Genetic effects on seed number per pod and seed weight among haplotypes in overlapping association regions in a natural population of rapeseed

seed number per pod and seed weight. The red and blue dots represent the average performance of SNPP and SW across years among 157 accessions in a natural population. \*\*\*, \*\*, \*. Significance at  $P < 0.001$ ,  $P < 0.01$ , and  $P < 0.05$ , respectively.

Bottom, pairwise LD estimates in the different haplotype blocks.

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