Correction to: Utilization of a Wheat55K SNP array-derived high-density genetic map for high-resolution mapping of quantitative trait loci for important kernel-related traits in common wheat

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Unfortunately, the legends of Figures 1 and 2 were missed out in the original publication. The legends are given below.

Fig. 1 Chromosomal locations of the stable QTL associated with KWPS, SN, KDR, KW, KL, TKW and PH based on a 371 RIL population derived from a cross between CN18 and T1208. The QTL clusters were marked clearly on the chromosomes.

Fig. 2 Chromosomal locations of digenic epistatic QTL for PH, TKW, KL, KW, KDR, SN, and KWPS.

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