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OPEN CACCESS QTL Mapping for chinese northern-style steamed bread specific					AS Subscription	
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Author(s) Peng Wu, Bin Liu, Tao Zhou, Zhuokun Li, Haiyun Du, Jiteng Wang, Jichun Tian ABSTRACT In this study, quantitative trait loci (QTLs) with additive effects, epistatic effects for CNSB specific volume in bread wheat (Triticum aestivum L.), were studied in cultivars Huapei 3 and Yumai 57 (Triticum aestivum L.). The DH population and the parents were planted in 2007 and 2008 in Tai' An and 2008 in Suzhou. QTL analyses were performed using the software of IciMapping v2.2 based on the mixed linear model. Five putative QTLs for CNSB specific volume were detected on 5 chromosomes where single QTLs explained 5.11% to 9.75% of phenotypic variations. All of them had negative effects on specific volume and were contributed by Yumai 57 alleles. Qsv-1B was detected in both environment 1 and 3 with 13.88% and 4.83% phenotypic variations which had positive effects and was transmitted by Huapei 3 alleles. Fourteen pairs of QTLs with epistatic effects were detected for specific volume. Seven major QTLs, Qsv-1B/Qsv-3A, Qsv- 2D/Qsv-3A, Qsv-3A/Qsv- 5B1, Qsv-1B/Qsv-6D, Qsv-2D/Qsv-4D, Qsv-4A/ Qsv-6B and Qsv-3A/Qsv-7D could account for 13.88%, 20.39%, 18.88%, 12.31%, 18.78%, 11.98%, and 17.05% of the phenotypic						
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variation of specific volume. The information obtained in this study will be useful for manipulating the QTLs for CNSB specific volume property by molecular marker-assisted selection (MAS).					Visits:	333,413
KEYWORDS Wheat; Doubled Haploid; Quantitative Trait Loci; CNSB; Specific Volume					Sponsors, Associates, and Links >>	
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