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CORRECTION

Correction to: Development and characterization of a spring hexaploid wheat line with no functional *VRN2* genes

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In the original version of this article, PCR fragments and digestion product sizes for the *VRN-B2* and *VRN-D2* markers were not accurate. The corrected sizes are detailed below:

Primers SNF-B2-3p-F1 and SNFB2-3p-R2 for the *SNF-B2* gene tightly linked to *VRN-B2* amplified a 1281 bp product. Digestion of the amplified product with restriction enzyme *HpyCH4*IV yielded two fragments (1041 and 240 bp) for the allele linked to the *vrnB2*-null deletion and three fragments (633, 408 and 240 bp) for the wild type allele from the hexaploid winter wheat variety Goodstreak.

Primers for the *VRN-D2* gene amplified a 687-bp fragment from the functional allele in the hexaploid wheat Goodstreak and a 681-bp fragment from the non-functional allele from *Ae. tauschii* accession E1. Digestion of the amplified PCR products with the restriction enzyme *Mbo*II yielded three fragments (396, 196 and 95 bp) in hexaploid wheat and two fragments (586 and 95 bp) in *Ae. tauschii* E1.

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