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| **Article** | Differential expression of the *HvCslF6* gene late in grain development may explain quantitative differences in (1,3;1,4)-β-glucan concentration in barley |
| **Journal** | Molecular Breeding |
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**Online Resource 2** Annotated sequence of a 204-bp pre-mRNA 3' untranslated region of *HvCslF6* obtained by 3' RACE from cDNA of CDC Bold and TR251 barley

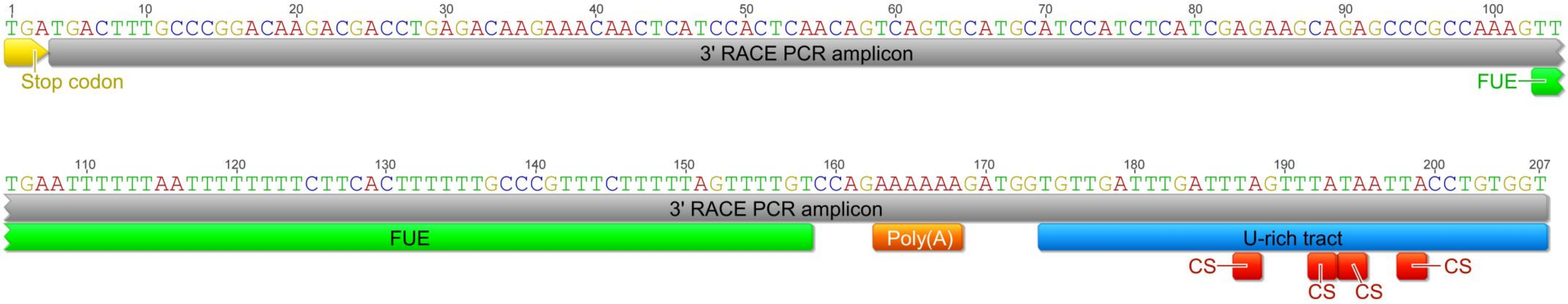

**Fig. S2** Annotated sequence of a 204-bp pre-mRNA 3' untranslated region of *HvCslF6* obtained by 3' RACE from cDNA of CDC Bold and TR251 barley. The far upstream element (FUE, green block) region contains multiple ambiguous UG motifs and is generally located 29 bp upstream of the mRNA cleavage site (CS, red blocks). The poly(A) signal (Poly(A), orange block) of *HvCslF6* is a AAAAAA hexamer. The U-rich tract (blue block) contains multiple putative mRNA cleavage sites which are denoted by TA di-nucleotide motifs at positions 187-188, 192-193, 194-195, or 198-199.