Association of jacalin-related lectins with wheat responses to stresses revealed by transcriptional profiling

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Online Resource 6 Expression profiles of a selected set of TaJRL genes in Wangshuibai, as revealed by sqPCR. A, in shoot tissue after infection by Bgt and in inflorescence tissue after infection by *F. graminearum*. B, in seedling leaves and roots after 4°C, PEG, and NaCl treatments. C, in seedling leaves and roots after MeJA, SA, and ABA treatments. The results for genes that did not respond to the treatments and that are shown in Fig. 5 are not shown here.