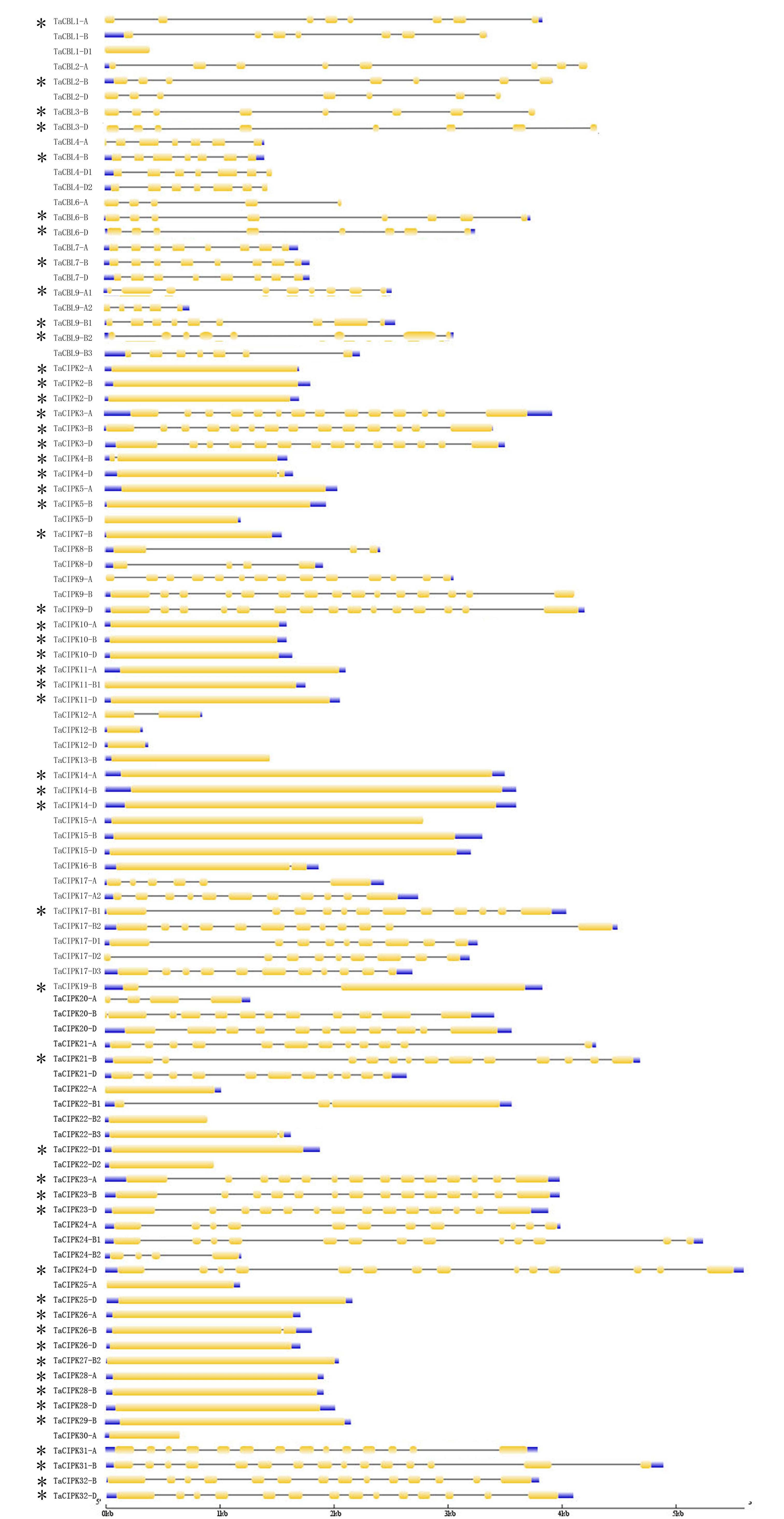




Additional files 4. Exon-intron analysis of *TaCBLs* and *TaCIPKs* in wheat. The boxes and solid lines represent “exon” and intron, respectively. The “upstream/downstream” indicates the flank sequence of “intron/exon” structure in genome sequence. The complete structures covering the full length ORF of genes were marked with “*”.