



Fig. 1. Fragments of the aligned sequences of *DREB1* orthologous gene in *Thinopyrum bessarabicum* (lines 1-3), *Pseudoroegneria stipifolia* (line 4), *P. spicata* (lines 5-6), *Dasypyrum villosum* (line 7), *Th. intermedium* (lines 8-13), *Th. ponticum* (lines 14-29) and *TaDREB* of three subgenomes of bread wheat (three lower lines). The largest indels are highlighted in color: four indels in positions 299-327 (highlighted in yellow); region 682-689 (highlighted in purple); region 698-705 (highlighted in green); region 1266-1268 (highlighted in blue); region 1316-1324 (highlighted in red).