



附图 1 TA/Tetep 型细胞质雄性不育基因 *orf312*、RT102 型细胞质雄性不育基因 *orf352*、WA 型细胞质雄性不育基因 *WA352* 所编码氨基酸的序列比对<sup>[1,2]</sup>

Fig.S1 Multiple sequence alignment of amino acids encoded by the TA/Tetep-type CMS gene *orf312*, the RT102-type CMS gene *orf352*, and the WA-type CMS gene *WA352*<sup>[1,2]</sup>

- [1] Madeira F, Madhusoodanan N, Lee J, et al. The EMBL-EBI Job Dispatcher sequence analysis tools framework in 2024 [J]. Nucleic Acids Res, 2024, 52(W1): W521-W525.
- [2] Robert X, Gouet P. Deciphering key features in protein structures with the new ENDscript server [J]. Nucleic Acids Res, 2014, 42(W1): W320-W324.