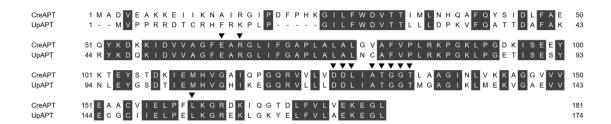
Title	Genome editing using a DNA-free clustered regularly interspaced short palindromic repeats-Cas9 system in green seaweed Ulva prolifera
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Appendix S2 Amino acid sequence alignment of the adenine phosphoribosyl transferase (*UpAPT*) gene. The deduced amino acid sequence of *U. prolifera* was compared with those of *Chlamydomonas reinhardtii* (CreAPT). Amino acid residues in common are highlighted and gaps are indicated by dashes. Arrowheads indicate the putative active site deduced by CDD/SPARCLE (Marchler-Bauer *et al.* 2017).