Title

N-glycosylation of a cargo protein C-terminal domain recognized by the type IX secretion system in *Cytophaga hutchinsonii* affects protein secretion and localization

Running title

Protein *N*-glycosylation in *Cytophaga hutchinsonii*

Authors

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Supplemental Material

Fig. S1

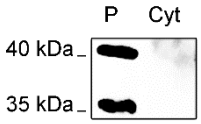

Fig. S1 The modification of the GFP-CTD_{CHU_2708} fusion protein occurred in the periplasm. The periplasmic and cytoplamic proteins (Cyt) of the $WT^{GFP\text{-}CTD}$ separated by SDS-PAGE, then identified by Western blotting with the anti-GFP antibody.

Fig. S2

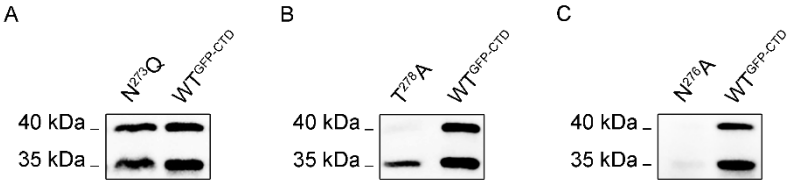

Fig. S2 Site-directed mutagenesis of CTD_{CHU_2708}. (A) The mutation of N273 in CTD_{CHU_2708} to glutamine. (B) The changing of T278 of CTD_{CHU_2708} to alanine. (C) The mutation of N276 of CTD_{CHU_2708} to alanine. The periplasmic space proteins of the $WT^{GFP\text{-}CTD}$, $N^{273}Q$, $T^{278}A$, and $N^{276}A$ strains were identified by Western blotting with the anti-GFP antibody.

Fig. S3

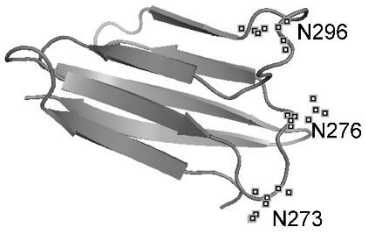

Fig. S3 The structural prediction of CTD_{CHU_2708}. The sites N273, N276, and N296

are shown.