

Bacterial Rotary Export ATPases are Allosterically Regulated by the Nucleotide Second Messenger
Cyclic-di-GMP

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SUPPLEMENTARY FIGURE

Figure S1 (attached .pdf file): Mass Spectrometry Peak Shift Analyses (MS-PSA) for treated (i.e. cross-linked to cdG) vs untreated FliI_{His} (S1A) and treated FliI_{His} only (S1B). The particularly dense clusters of related spectra in both analyses correspond to the most densely-modified peptide following cdG-capture compound crosslinking. In both cases, this peptide, (NVLLLMSLTR) comprises residues 259-269 of FliI.