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Directed evolution of RimJ For N-terminal acetylation with broad substrate specificity

Background

- N-terminal acetylation is essential for the stability, activity, and targeting of proteins in eukaryotes.
- However, most eukaryotic proteins are not acetylated when expressed in bacteria.
- RimJ is an N-terminal acetyltransferase known to acetylate many recombinant proteins with a narrow substrate specificity in E. coli, including the Z domain.
- N-terminal acetylation by RimJ has two requirements:
 - Initiator methionine cleavage
 - N-terminal amino acid sequence



Objective

- To increase the applicability of RimJ for the N-terminal acetylation of a broad range of recombinant proteins.
- Increase active site size to accommodate larger proteins, using a mutation to alanine.
- Determined from the Alpha Fold-predicted structure of E. coli RimJ, the following amino acids are of interest:

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Y35A E46A R49A Y106A Y170A L171A



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Methods

small neutral amino acid.

acetylate by wild type RimJ.

Purification of protein by Ni-NTA affinity chromatography.





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