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Supporting information for article:

Structural basis of self-assembling in the lipid-binding domain of mycobacterial polar growth factor Wag31

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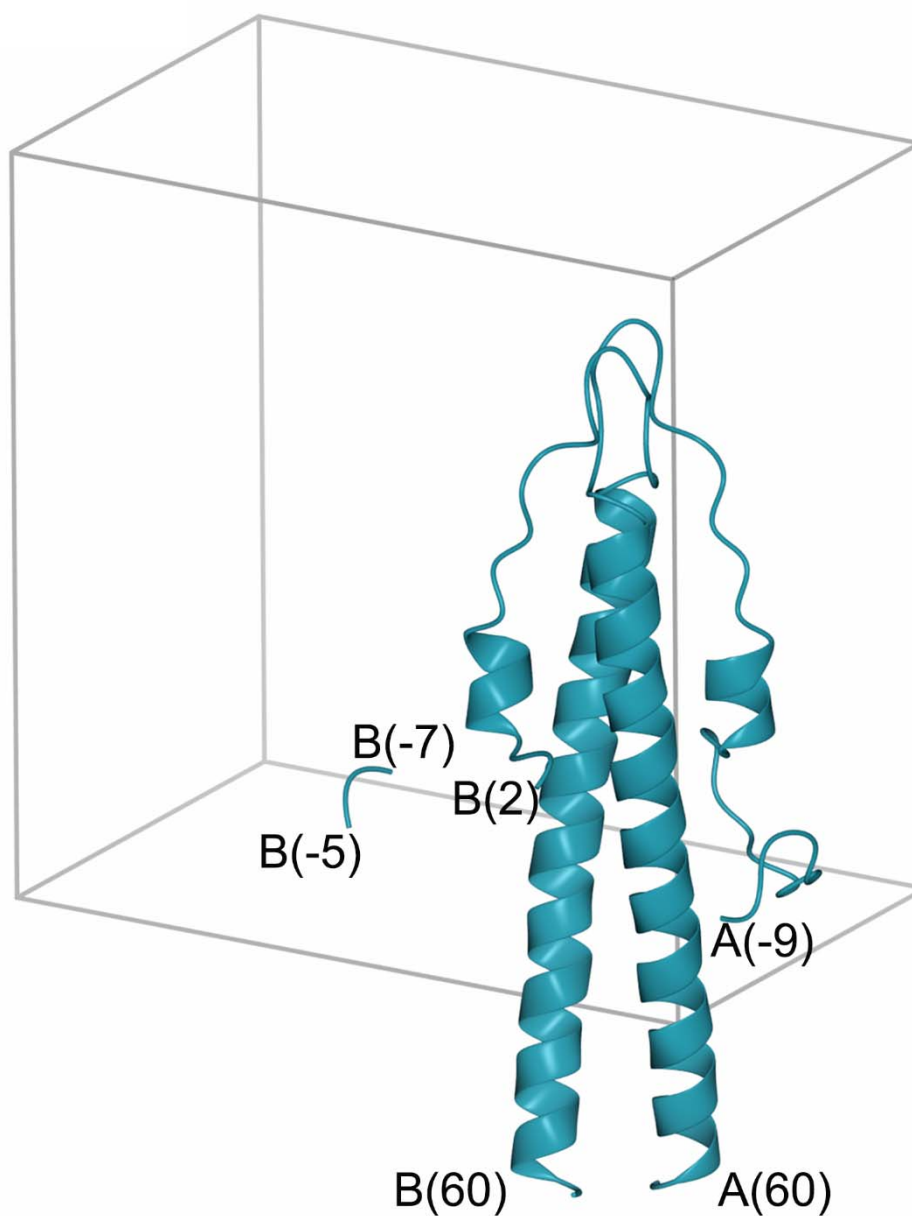


Figure S1 Structure of N-Wag31. The monoclinic unit cell is shown as a box. N-Wag31 dimer within the asymmetric unit, including the poly-histidine tag regions (see experimental section), is shown as a cartoon in blue.

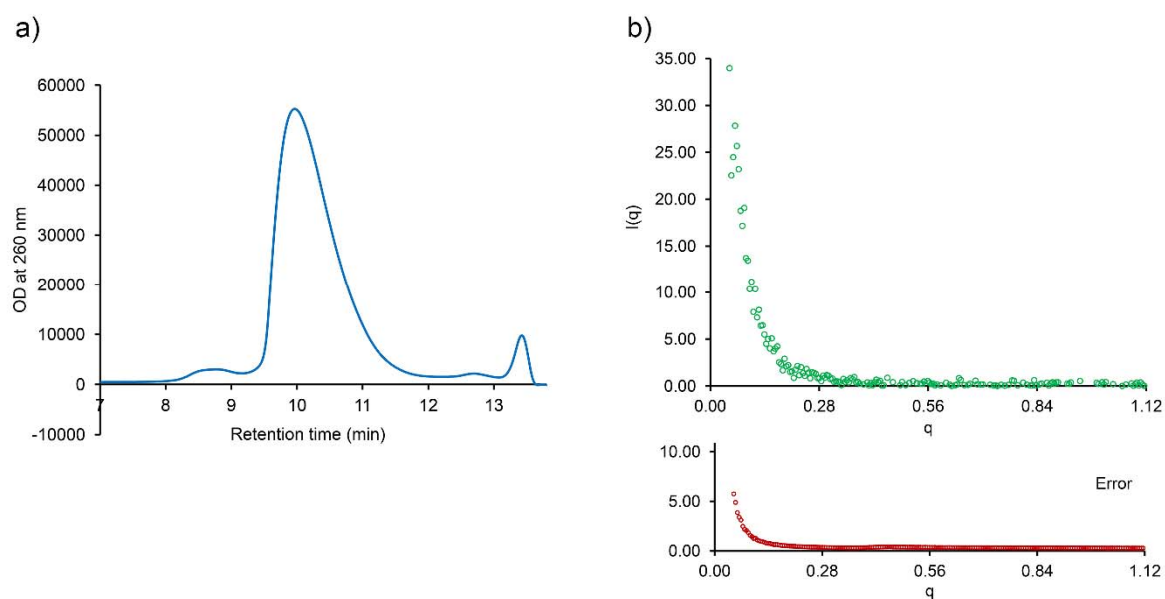


Figure S2 SEC-SAXS data of N-Wag31. (a) UV elution profile (UV absorbance in arbitrary unit at 260 nm wave-length *versus* retention time in minutes) of N-Wag31 during the SEC-SAXS experiment. (b) Intensity (I) *versus* momentum transfer (q in nm⁻¹) for averaged SAXS intensity profile, and associated error, obtained from the peak I region shown in figure 3a.