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Supporting information for article:

Crystal structure of the ATPase-C domain of the chromatin remodeller Fun30 from *Saccharomyces cerevisiae*

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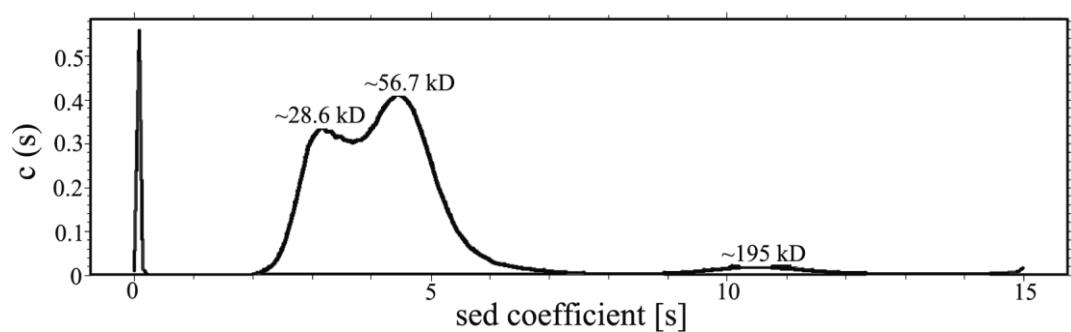


Figure S1 Analytic ultracentrifugation of Fun30 ATPase-N (residues 572-794) exhibits a mix of monomer and dimer.