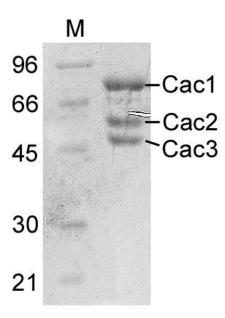
Supplementary Information

Yeast CAF-1 assembles histone (H3-H4)₂ tetramers prior to DNA deposition

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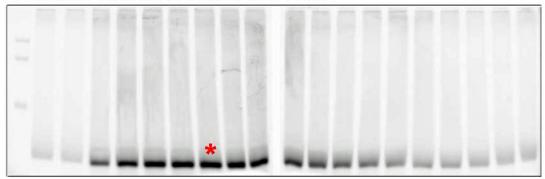


S. Figure 1 – SDS-PAGE analysis of recombinantly purified CAF-1. The three yeast CAF-1 subunits are shown in the right lane and correspond to Cac1 (70 kDa), Cac2 (51 kDa), and Cac3 (47 kDa). A low molecular weight marker (M) is shown the the left lane to provide a frame of reference.

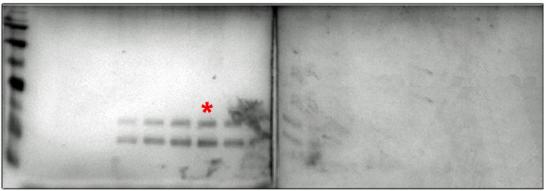
A. Elution Peak Fluorescence Scan

D5 D6 D7 D8 D9 D10 D11 D12 E1 E2 E3 E4 E5 E6 E7 E8 E9 E10 E11

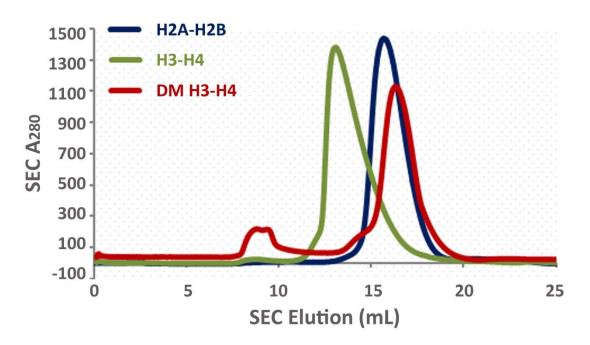
B. Elution Peak Fluorescence SDS-PAGE



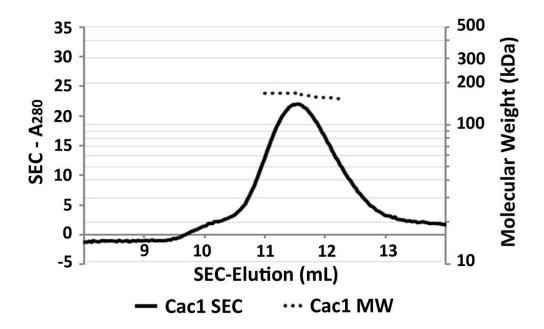
C. Elution Peak Silver Stain SDS-PAGE



S. Figure 2 – Analysis of the oligomeric state of H3-H4 using SEC. The elution fractions for each concentration of H3-H4 tested are aliquoted into a prepared 384-well clear-bottom microplate. The low concentration H3-H4 samples prohibit monitoring by 280 nm absorbance. The plate is then scanned for fluorescence of the H3-H4 conjugated fluorophore (panel A). The elution fractions are then run out on a 12% SDS-PAGE, scanned for fluorescence (panel B), and silver stained (panel C) to confirm overlapping of the fluorescence signal and H3-H4 polypeptides. The asterisks enable same sample (SEC elution fraction D11; highest fluorescence signal) comparison between panels A, B, and C.



S. Figure 3 – Size-exclusion chromatography peaks for high concentration (100 μM) histone complexes. Recombinantly expressed histones H2A and H2B; H3 and H4 are combined, refolded and then run over a Superdex 200 10/300 size exclusion chromatography column to ensure homogeneity and oligomeric state. Consistently, wild-type (H3-H4)₂ tetramers elute at a 14 mls (green); wild-type H2A-H2B dimer elution is delayed to 16 mls (blue). The elution profile of the H3 (L126R, I130R)-H4 variant (16.5 mls, red) is comparable to the H2A-H2B dimer which supports the inability of this histone complex to form tetramers even at high concentrations. The void volume for this column is near 8 mls.



S. Figure 4 – Determination of the apparent molecular weight of the Cac1 subunit using SEC-MALS. The Cac1 subunit elutes from the Superdex 200 GL column between 11 and 12 mls, which is slightly earlier than the complete CAF-1 complex. Light scattering confirms a molecular weight consistent with a Cac1 dimer (dotted line, right axis and Table 2). The asterisks denote same sample (SEC fraction D11; highest fluorescence signal fraction) for comparison across panels.