



Figure S2 Summary of cross-linking study on wild-type recombinant *S. cerevisiae* Ndc80 complex. (A) After cross-linking and trypsin digestion, four possible peptides were identified by mass spectrometry. (B-D) Diagram representation of cross-linking results for Ndc80, Nuf2, His₆-Spc24, and Spc25 (black bars). (B) Peptide sequence coverage (colored boxes) and mono-links (dotted vertical lines and circles). (C) Loop-links (dotted lines) and self cross-links (solid lines, cross-links between two peptides from the same protein). (D) Cross-links between different proteins (solid line). Vertical yellow lines denote positions of lysines. Full lists of cross-linking results are shown in Tables S4 and S5.