SUPPLEMENTARY MATERIAL

ATP-dependent Nucleosome Unwrapping Catalyzed by human RAD51

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Supplemental Table 1: PCR Primers

601D forward:	CTGGAGAATCCCGGTGCCGA
601D reverse:	ACAGGATGTATATATCTGACACGTGCCTGGAGACTAGGG
	AGTAATCCCCTTGGCGGTTAAAACGCGG(T-Cy3)GGACA
601E forward:	Cy3-CTGGAGAATCCCGGTGCCGA
601E reverse:	ACAGGATGTATATATCTGACACGTGCCTGGAGACTA
5SD forward:	GTACTAATACTGTATGAGCATACAGTACTTCCG
5SD reverse:	CTGGCATGGGGAGGAGCTGGGCCCCCCCAGAAGGCAGCA
	CAAGGGGAGGAAAAGTCAGCCTTGTGC(TCy3)CGC
5SE forward:	Cy3-GTACTAATACTGTATGAGCATACAGTACTTCCG
5SE reverse:	CTGGCATGGGGAGGAGCTGG
601-linker forward:	CTGGAGAATCCCGGTGCCGAGGC
601D-linker reverse:	GCACCCCACTGTCTATACACAGGATGTATATATCTGACACGTG
	CCTGGAGACTAGGGAGTAATCCCCTTGGCGGTTAAAACGCGG(T
	Cy3)GG
601E-linker reverse:	TCGCCTCCGCACCCCACTGTCTATACACA(T-Cy3)GATGTATA
linker forward:	CCTGTGTATAGACAGTGGGGTGCGGAGGCG
linker reverse:	CACGTGTGGGGGCCCGAGCTCGCGGC

SUPPLEMENTARY FIGURE LEGENDS

Supplementary Figure 1. Assembly of Nucleosomes. A) Agarose gel analysis of assembled nucleosomes using 601E, 601D, 5SE, 5SD, 601E-linker, and 601D-linker nucleosome positioning DNAs (see Fig. 1A). Nucleosome (Nuc) mobility and free DNA (DNA) mobility are marked on the left. B) Crystal structure of a nucleosome (Lugar et al., *Nature* 389:6648, 1997) showing the relative positions of the entry-exit (601E, left) and dyad (601D, right) Cy3 fluorophore relative to the Cy5 fluorophore linked to H2A(K119C) within the histone octamer.

Supplementary Figure 2. Nucleosome disassembly by HsRAD51(D316K). A)

Representative fluorescence emission spectra of 601E-linker nucleosomes (2 nM) excited at 510nM in the presence 130mM KCl, ATP (250 μ M) and 2mM MgCl2 (Mg+2) with HsRAD51(D316K) at 0nM (black), 100nM (blue), 200nM (green), 500nM (orange) and 900nM (purple). **D)** Representative fluorescence emission spectra of 601D-linker nucleosomes (2 nM) excited at 510 nM in the presence 130 mM KCl, ATP (250 μ M) and 2 mM CaCl2 (Ca+2) with HsRAD51 at 0 nM (black), 100 nM (blue), 200 nM (green), 500nM (orange) and 900nM (purple).





