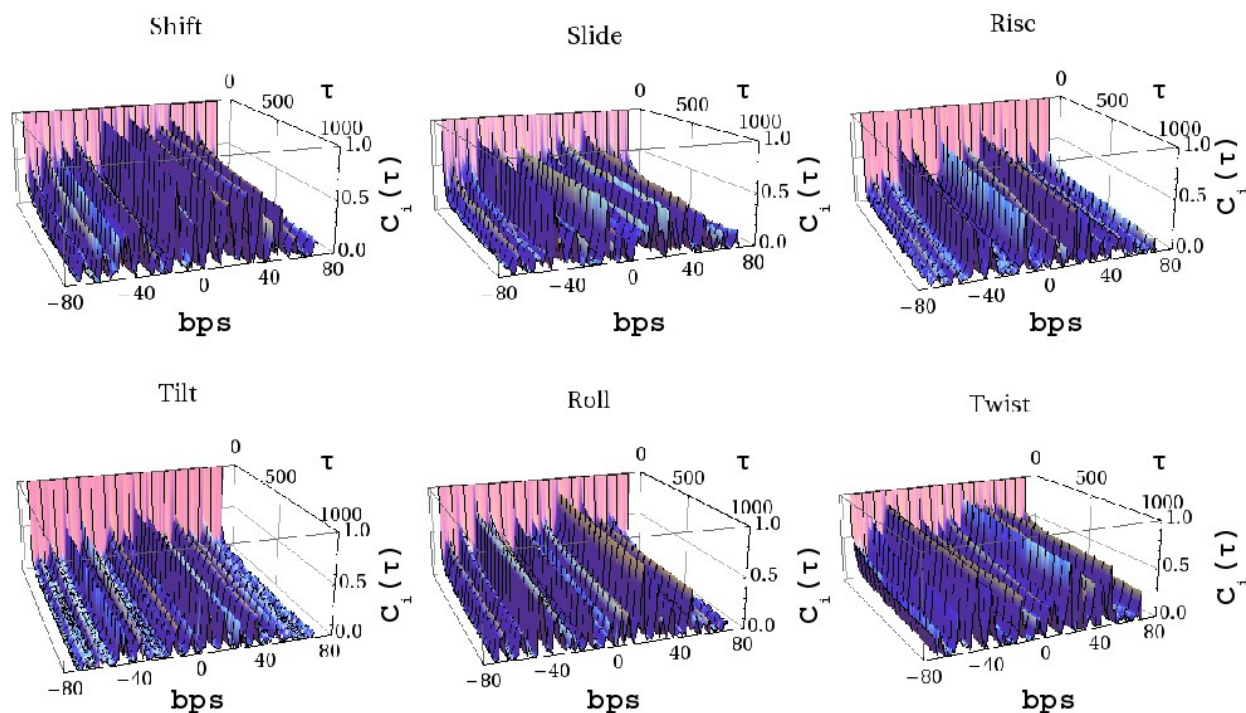
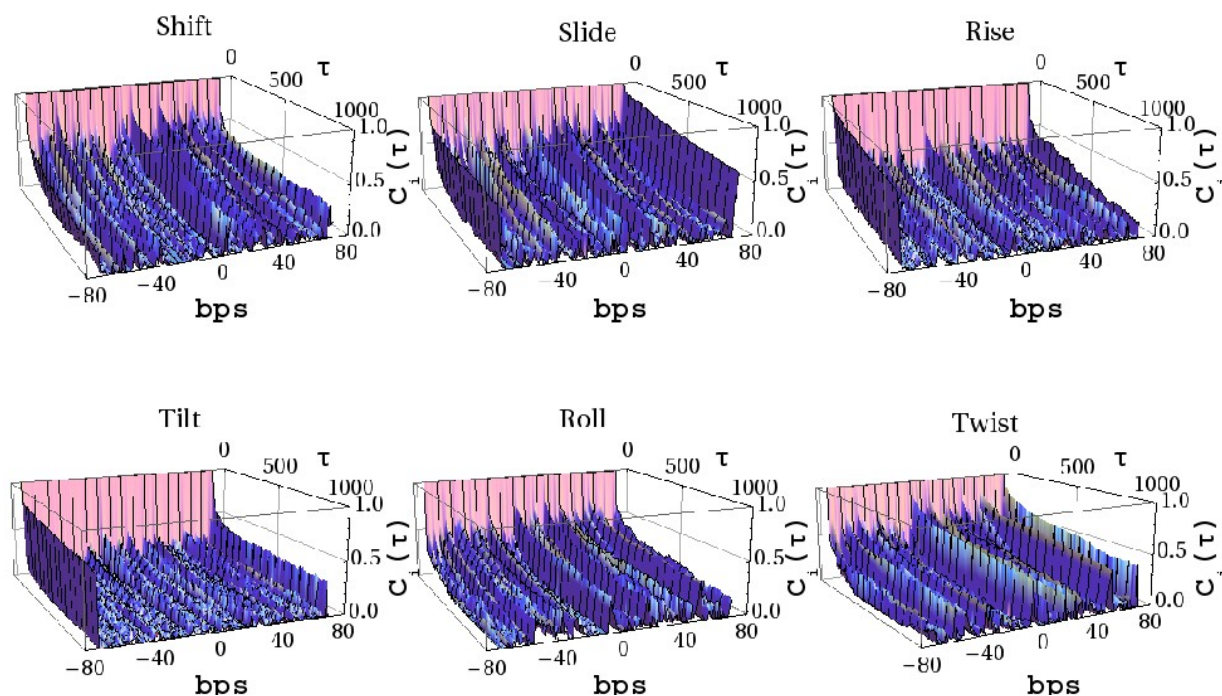


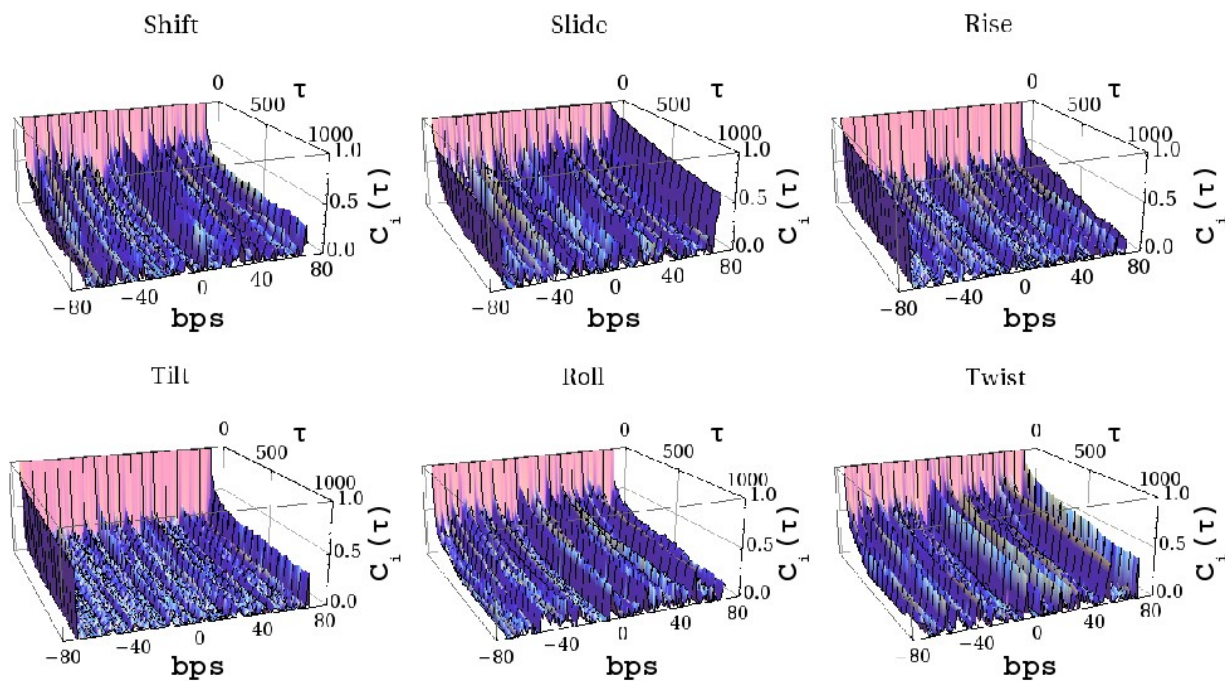
Supplementary Material



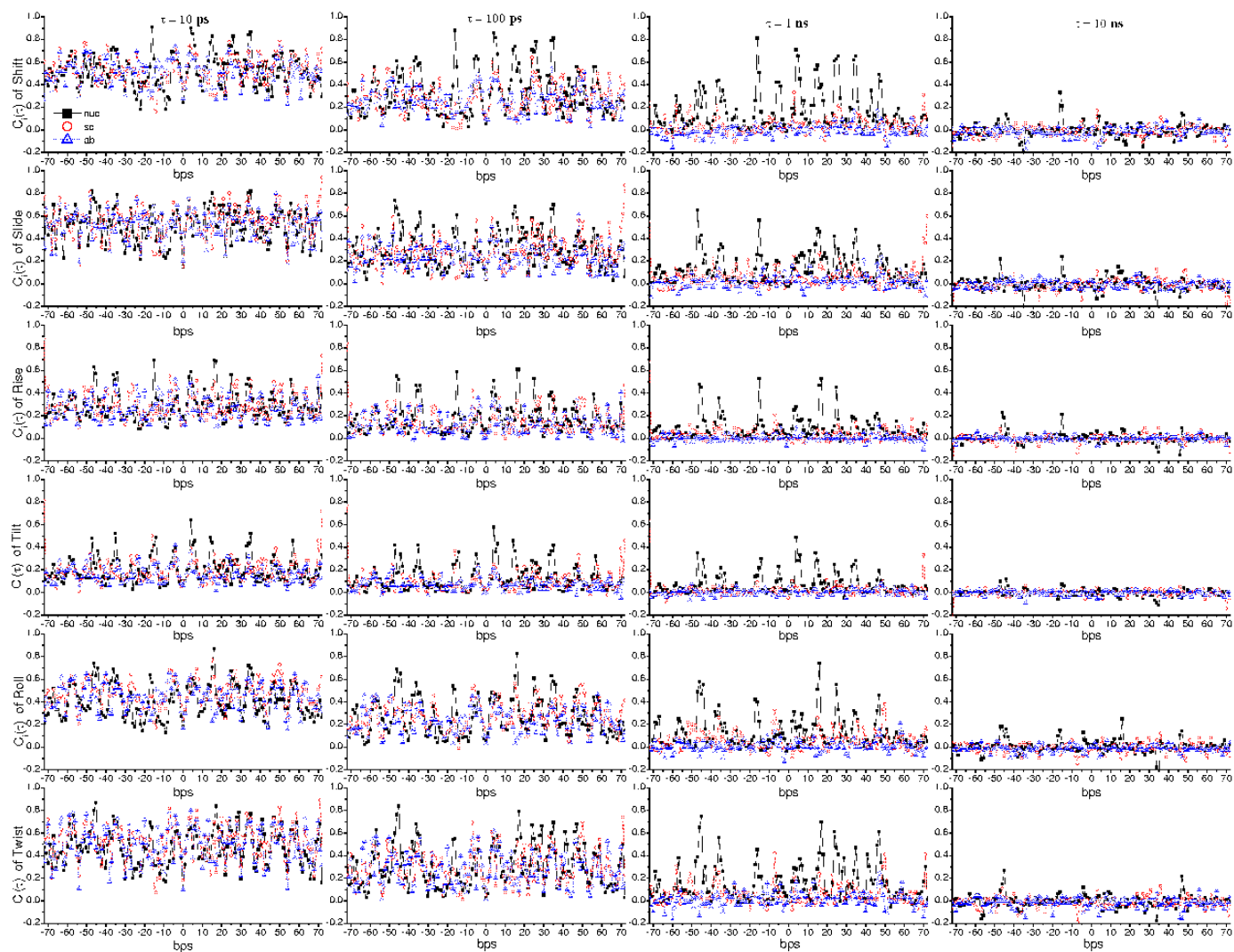
S1. 3D autocorrelation functions $C(\tau)$ of helix parameters for nuc-lid3, where τ is lag time, ps, and bps is DNA base pair number.



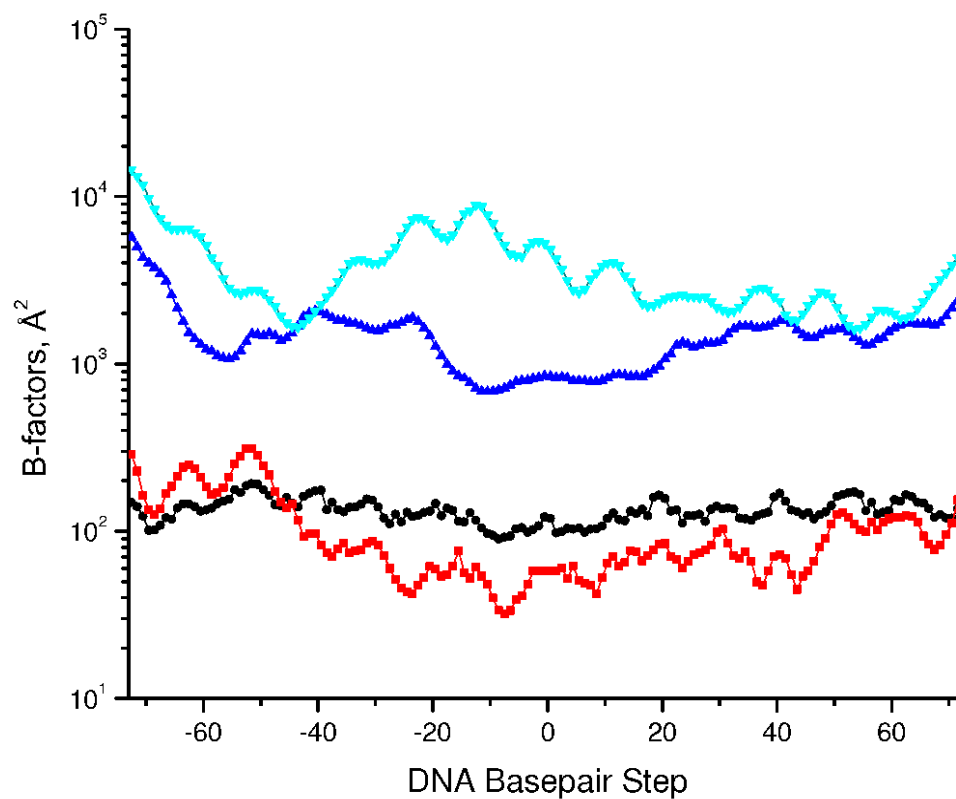
S2. 3D autocorrelation functions $C(\tau)$ of helix parameters for sc-DNA, where τ is lag time, ps, and bps is DNA base pair number.



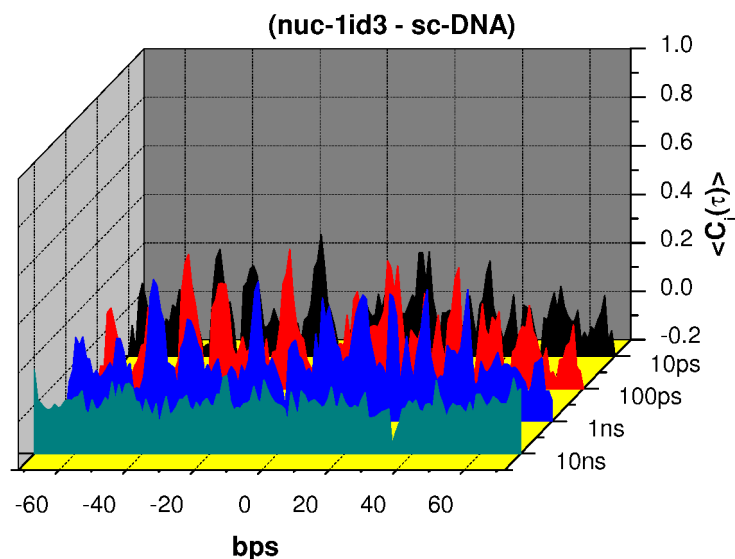
S3. 3D autocorrelation functions $C(\tau)$ of helix parameters for AB-DNA, where τ is lag time, ps, and bps is DNA base pair number.



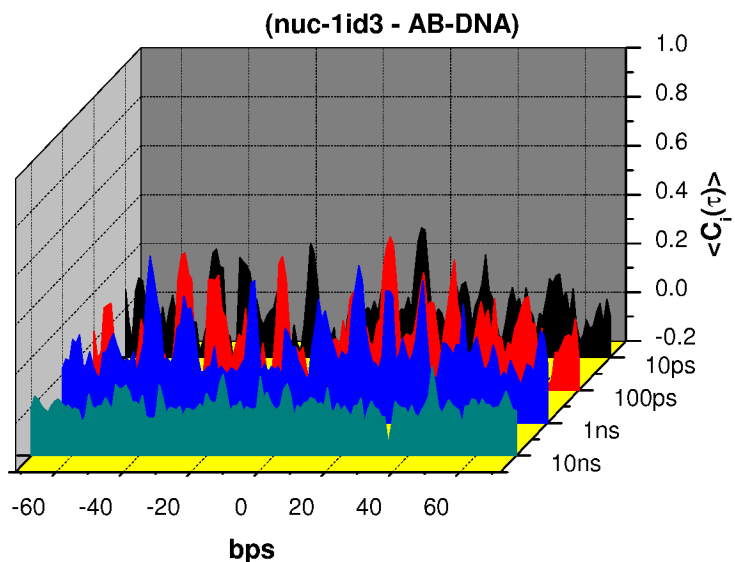
S4. 2D cross sections of 3D autocorrelation data of shift, slide, rise, tilt, roll and twist for nuc-lid3, sc-DNA and AB-DNA at times $\tau = 10$ ps, 100 ps, 1 ns, 10 ns.



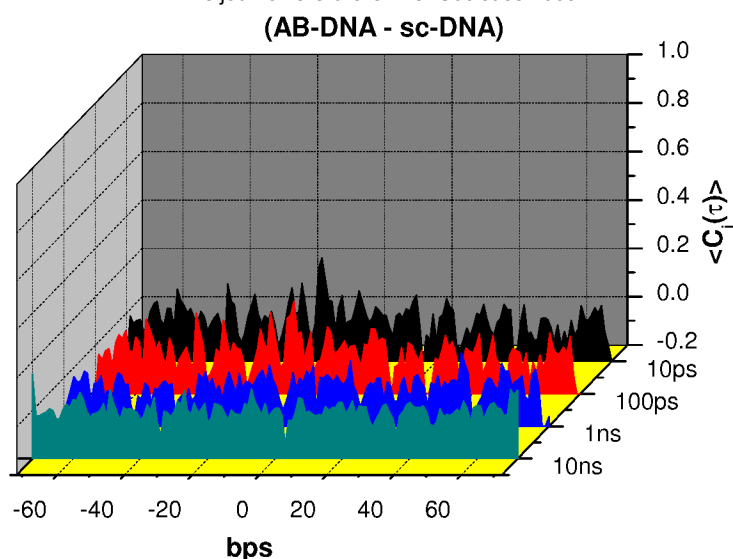
S5. B-factors from lid3 x-ray structure (black circles) and simulations of nuc-lid3 (red squares), sc-DNA (blue triangles) and AB-DNA (cyan down triangles) as a function of DNA basepair.



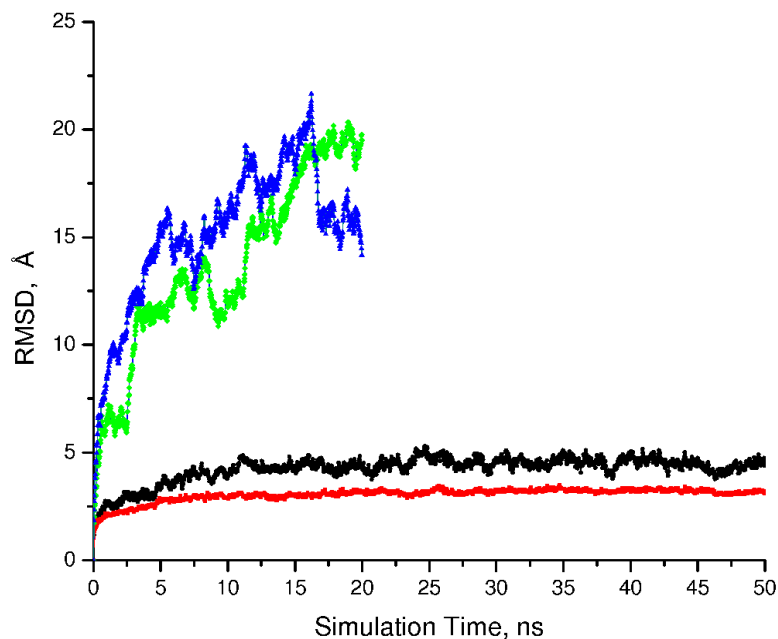
S6. Differential cross sections of 3D autocorrelation functions, $C_{\text{nuc-1id3}}(\tau) - C_{\text{sc-DNA}}(\tau)$, of helix parameters along the DNA length for nuc-1id3 at time instances $\tau = 10\text{ps}, 100\text{ps}, 1\text{ns}, 10\text{ns}$. Original cross sections are provided in Figure 2.



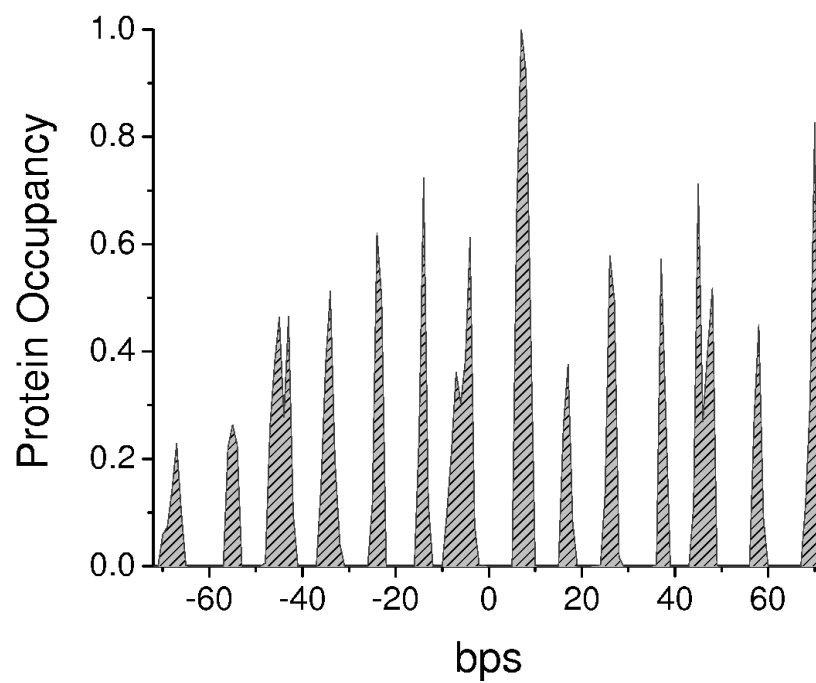
S7. Differential cross sections of 3D autocorrelation functions, $C_{\text{nuc-1id3}}(\tau) - C_{\text{AB-DNA}}(\tau)$, of helix parameters along the DNA length for nuc-1id3 at time instances $\tau = 10\text{ps}, 100\text{ps}, 1\text{ns}, 10\text{ns}$. Original cross sections are provided in Figure 2.



S8. Differential cross sections of 3D autocorrelation functions, $C_{\text{AB-DNA}}(\tau)$ - $C_{\text{sc-DNA}}(\tau)$, of helix parameters along the DNA length for AB-DNA at time instances $\tau = 10\text{ps}, 100\text{ps}, 1\text{ns}, 10\text{ns}$. Original cross sections are provided in Figure 2.



S9. Root mean square deviations of nuc-1ld3 (black circles - dna; red squares - protein), sc-DNA (green diamonds) and AB-DNA (blue triangles) with respect to their starting structures.



S10. Normalized protein occupancy as a function of DNA base pair step, bps,
in nuc-1id3.