

Supplementary Data Material:

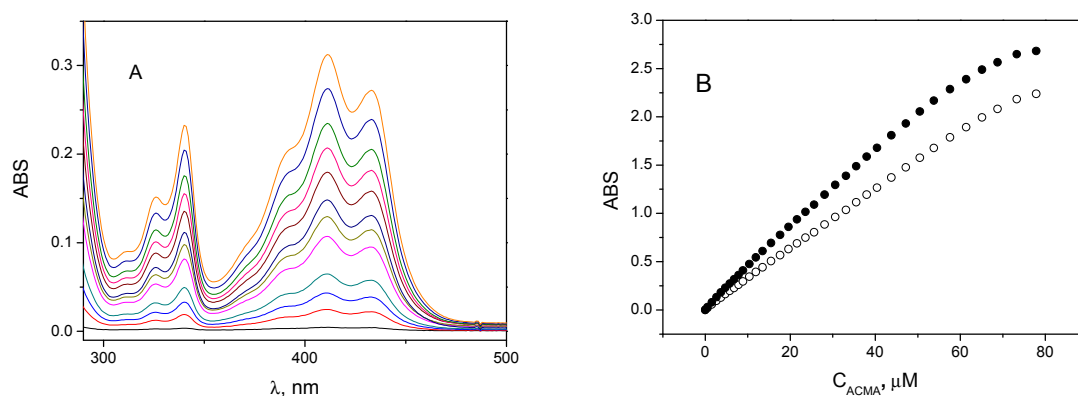


Fig. S1 A) Absorbance spectra of ACMA solutions, $C_{\text{ACMA}} = 0.5 - 80 \mu\text{M}$, $\text{pH} = 7.0$, $I = 0.1 \text{ M}$ and $T = 25^\circ\text{C}$. B) Representation of absorbance values vs ACMA concentration. ($\bullet$) $\lambda = 275 \text{ nm}$ and ($\circ$) $\lambda = 260 \text{ nm}$.

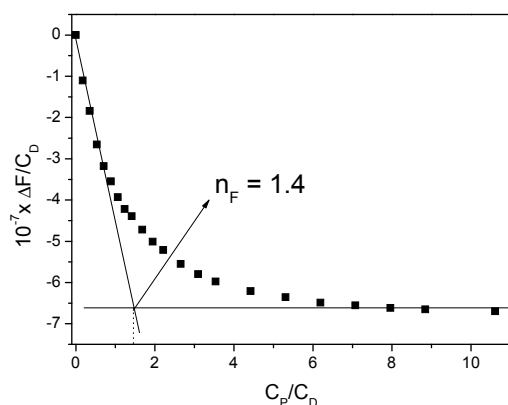


Fig. S2 Site Size determination for de CT-DNA/ACMA system by fluorescence titration. $\text{pH} = 7.0$, $I = 0.01 \text{ M}$, $T = 25^\circ\text{C}$, $\lambda_{\text{exc}} = 419 \text{ nm}$, $\lambda_{\text{em}} = 477 \text{ nm}$. The intersection of the straight lines yields n , the site size.

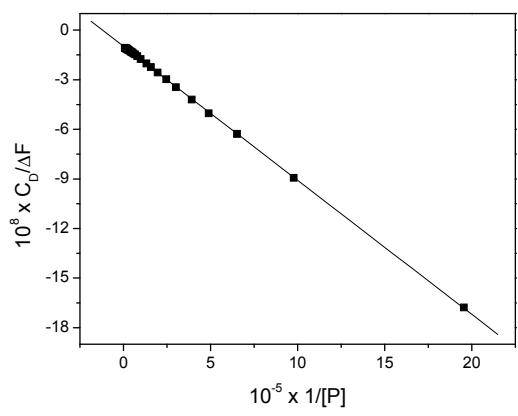


Fig. S3 Fit of a spectrofluorimetric titration for the CT-DNA/ACMA system to eqn 3. $C_D^0 = 1.50 \mu\text{M}$, $I = 0.1 \text{ M}$, $\text{pH} = 7.0$, $T = 25^\circ\text{C}$, $\lambda_{\text{exc}} = 419 \text{ nm}$ and $\lambda_{\text{em}} = 477 \text{ nm}$.

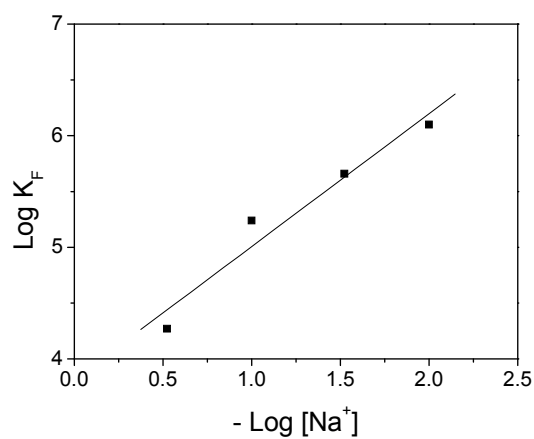


Fig. S4 Salt effect on the equilibrium constant, K , for ACMA binding to CT-DNA, $\text{pH} = 7.0$ and $T = 25^\circ\text{C}$.

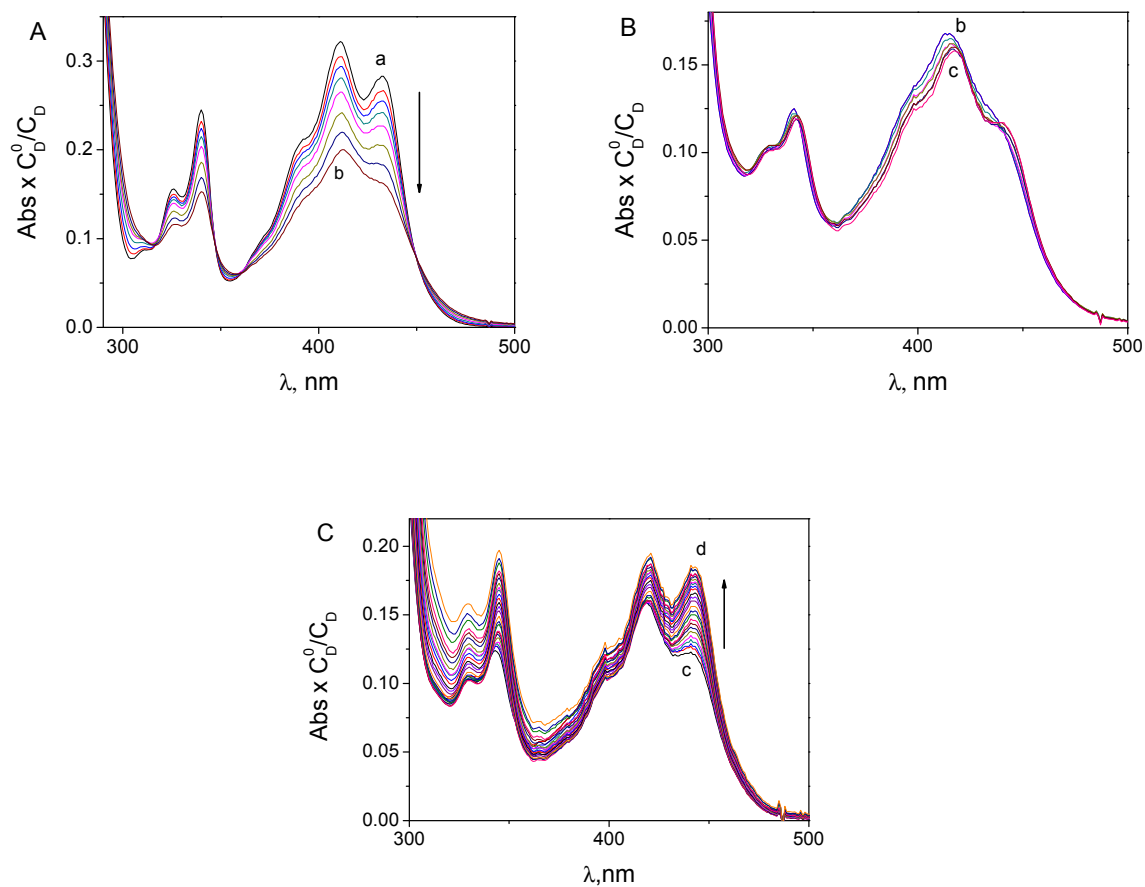


Fig. S5: Absorbance spectra of the CT-DNA/ACMA system recorded during titration at A) $0 \leq C_P/C_D \leq 0.9$; B) $0.9 \leq C_P/C_D \leq 2.2$; C) $C_P/C_D \geq 2.2$. $C_D^0 = 5.37 \times 10^{-5} \text{ M}$, $C_P = 7.41 \times 10^{-4} \text{ M}$. pH = 7.0, $I = 0.1 \text{ M}$ and $T = 25^\circ \text{C}$.

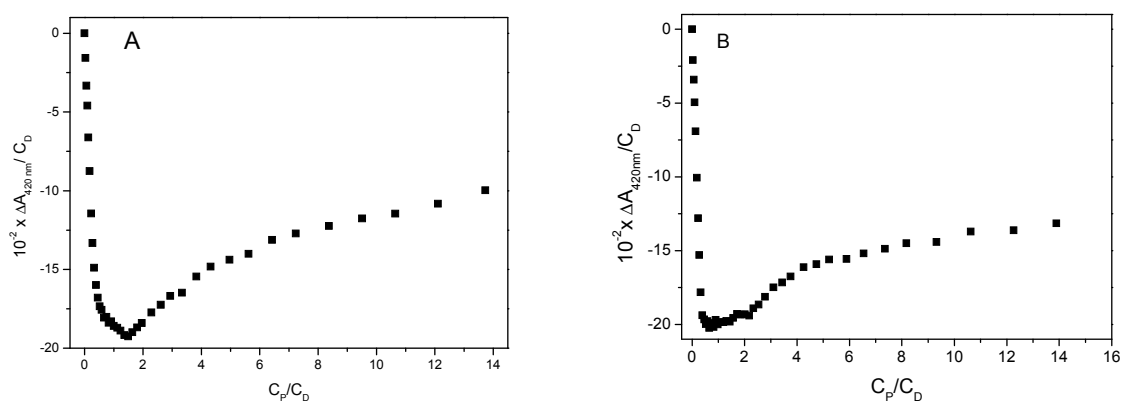


Fig. S6 Absorbance binding isotherm for the CT-DNA /ACMA system. $C_D^0 = 54 \mu\text{M}$, pH = 7.0, $T = 25^\circ \text{C}$. A) $I = 0.01 \text{ M}$; B) $I = 2.5 \text{ mM}$.

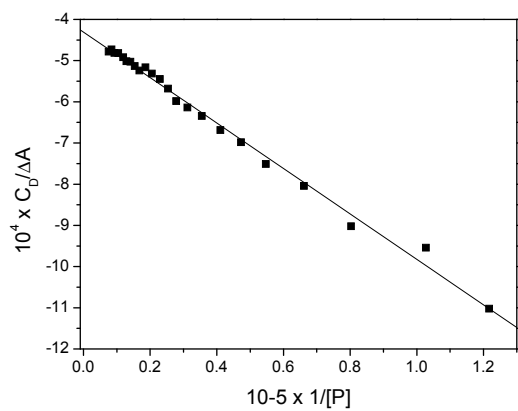


Fig. S7 Absorbance titration data fitted according to eqn 3

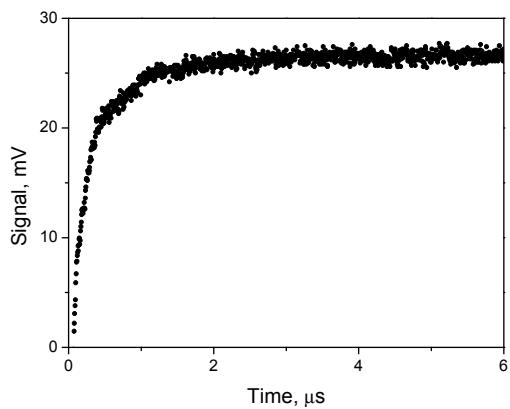


Fig. S8 T-jump relaxation curve for the CT-DNA/ACMA system, $I = 0.1$ M, $pH = 7.0$, $T = 25^\circ C$, $\lambda_{exc} = 420$ nm, $\lambda_{em} = 480$ nm, $C_D = 1.26 \times 10^{-5}$ M and $C_P = 1.50 \times 10^{-4}$ M.

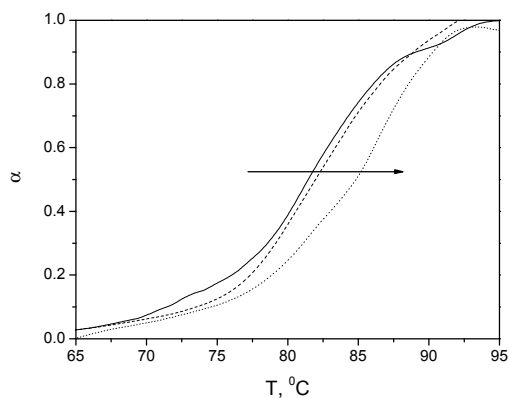


Fig. S9 Melting curves at different C_D/C_P ratios, $C_P = 1.19 \times 10^{-5}$ M, pH = 7.0, $T = 25^\circ\text{C}$, $I = 0.1$ M $\lambda = 260$ nm. Solid line in absence of ACMA, dash line at $C_D/C_P = 0.05$ and dot line at $C_D/C_P = 0.11$. The α value is calculated from the denaturation curve as $\alpha = (ABS - ABS_D)/(ABS_S - ABS_D)$, where ABS_D , ABS_S and ABS stand for the absorbance of the double, single strands and mixture, respectively.

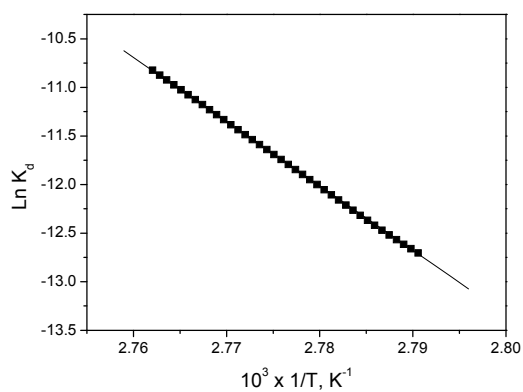


Fig. S10 van't Hoff plot for the denaturation process. $C_D/C_P = 0.11$, $C_P = 1.19 \times 10^{-5}$ M, pH = 7.0, $T = 25^\circ\text{C}$, $I = 0.1$ M. The denaturation constant, K_d , is defined as $K_d = 4C_0\alpha^2/(1 - \alpha)$.

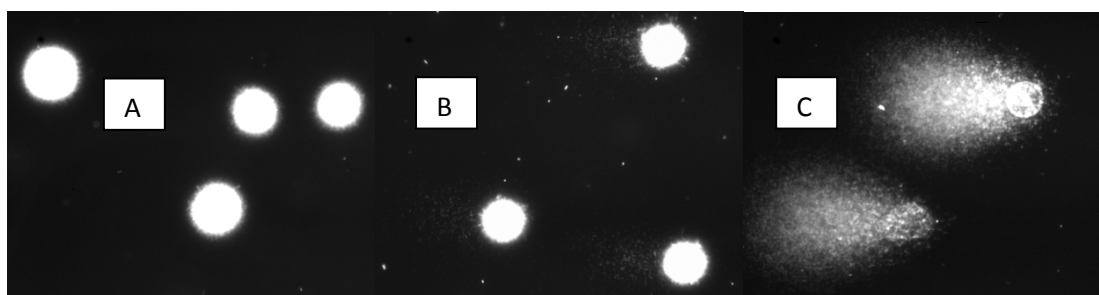


Fig. S11 Recorded images at 400 $\times$ magnification, (A) negative control, (B) ACMA 3.30 μ M and (C) positive control: H₂O₂ 250 μ M