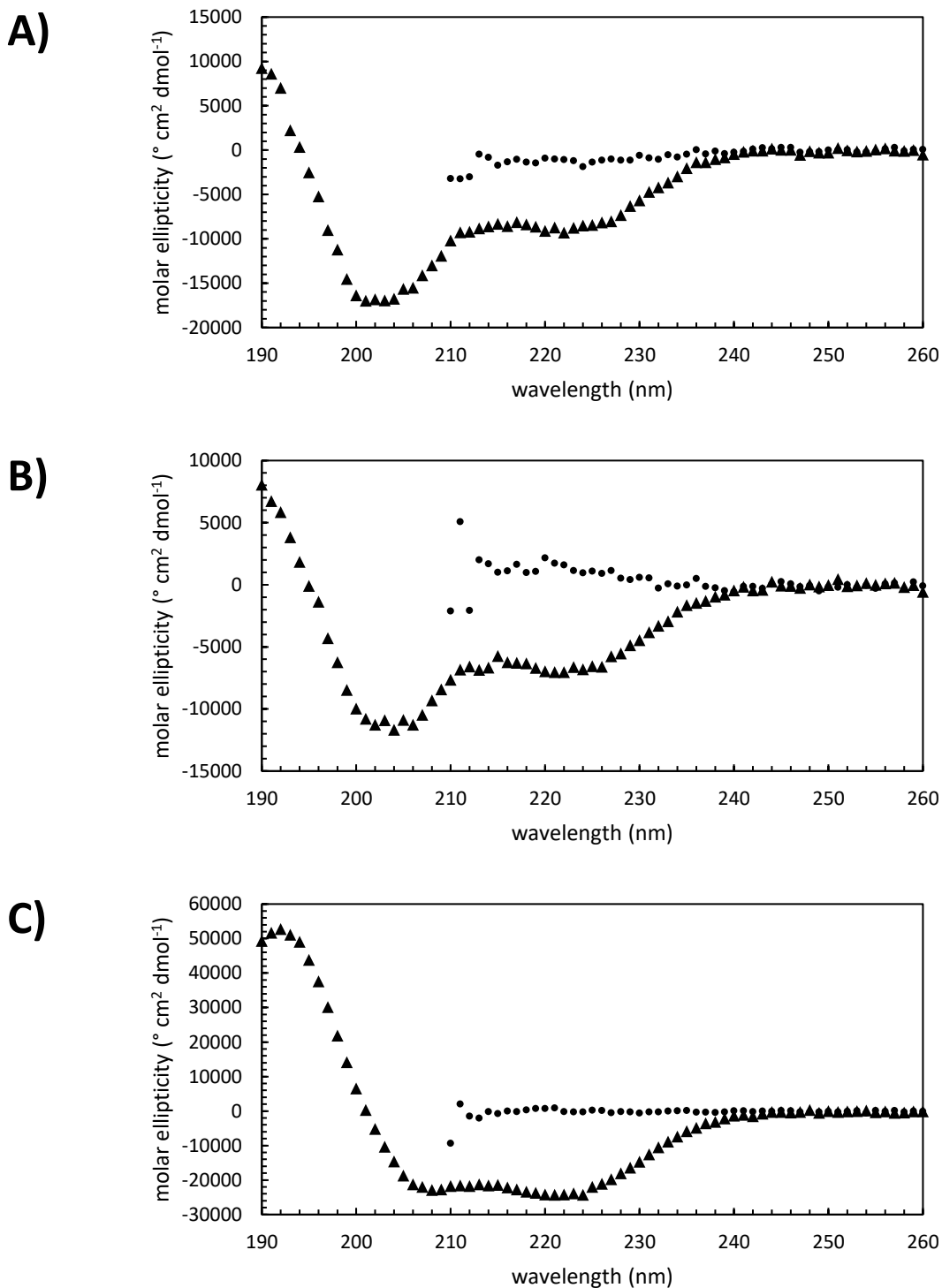


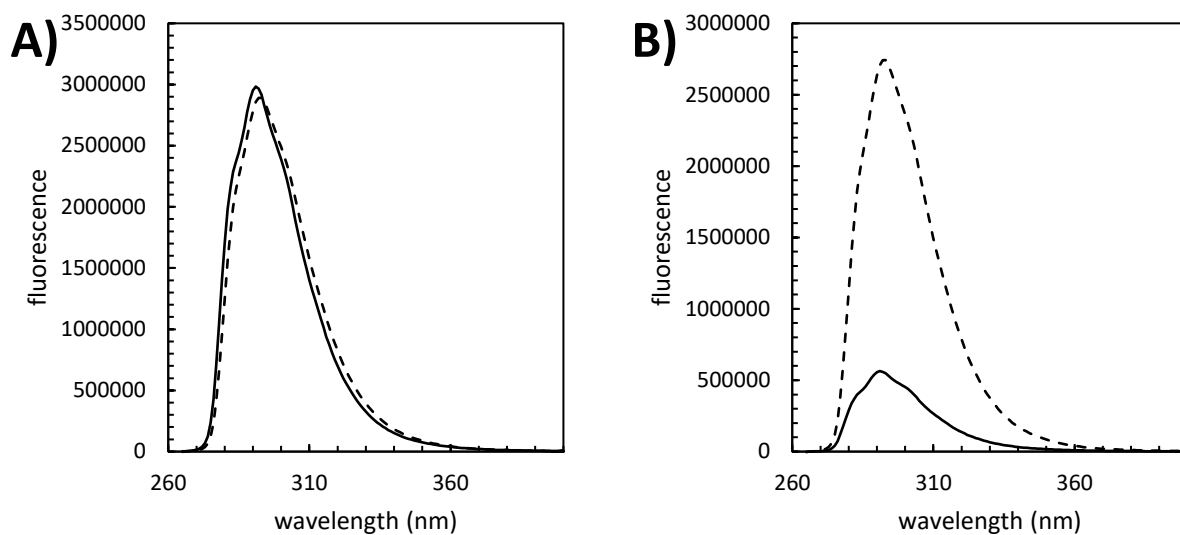
Supporting Information For

**A Non-perturbing Probe of Coiled Coil Formation Based on Electron Transfer Mediated  
Fluorescence Quenching**

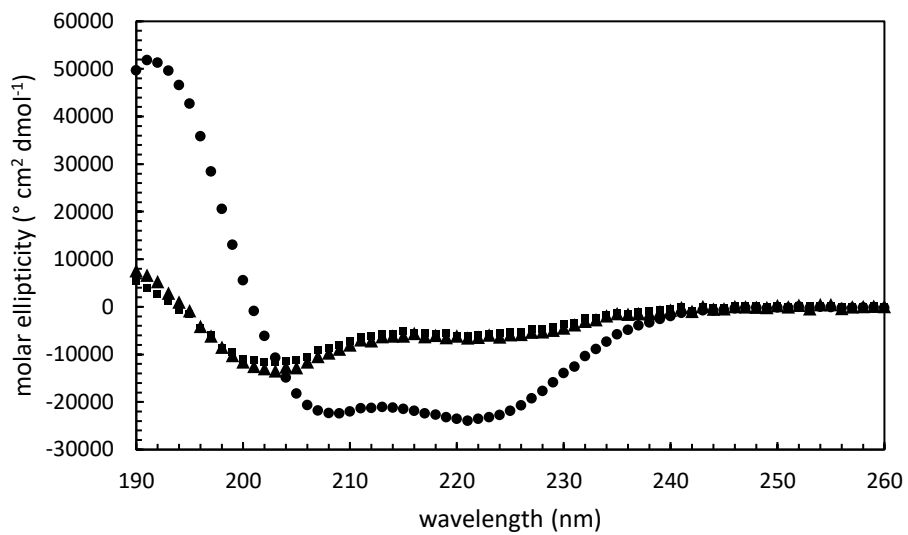
Matthew D. Watson, Ivan Peran, Daniel P. Raleigh



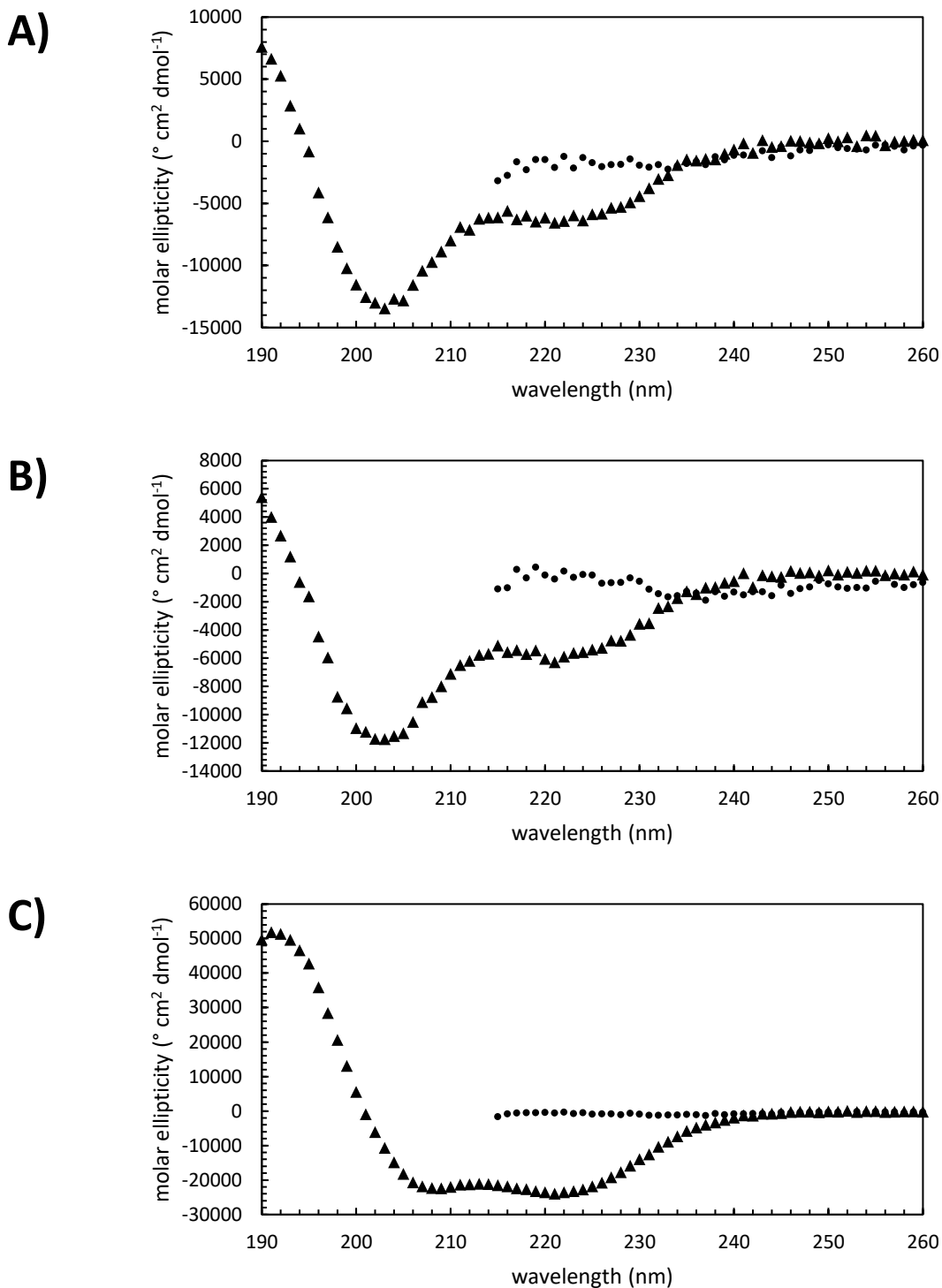
**Figure S1.** (A) Circular dichroism spectra of CC-19FCNA<sup>3.5</sup> (35  $\mu\text{M}$ ) at 20  $^{\circ}\text{C}$  in phosphate buffer (10 mM, pH 7.4, triangles) and in urea (10 M, circles). (B) Circular dichroism spectra of CC-20M<sub>se</sub>B<sub>N</sub><sup>3.5</sup> (35  $\mu\text{M}$ ) at 20  $^{\circ}\text{C}$  in phosphate buffer (10 mM, pH 7.4, triangles) and in urea (10 M, circles). (C) Circular dichroism spectra of a mixed solution of CC-19FCNA<sup>3.5</sup> and CC-20M<sub>se</sub>B<sub>N</sub><sup>3.5</sup> (1:1, 35  $\mu\text{M}$  total peptide concentration) at 20  $^{\circ}\text{C}$  in phosphate buffer (10 mM, pH 7.4, triangles) and in urea (10 M, circles).



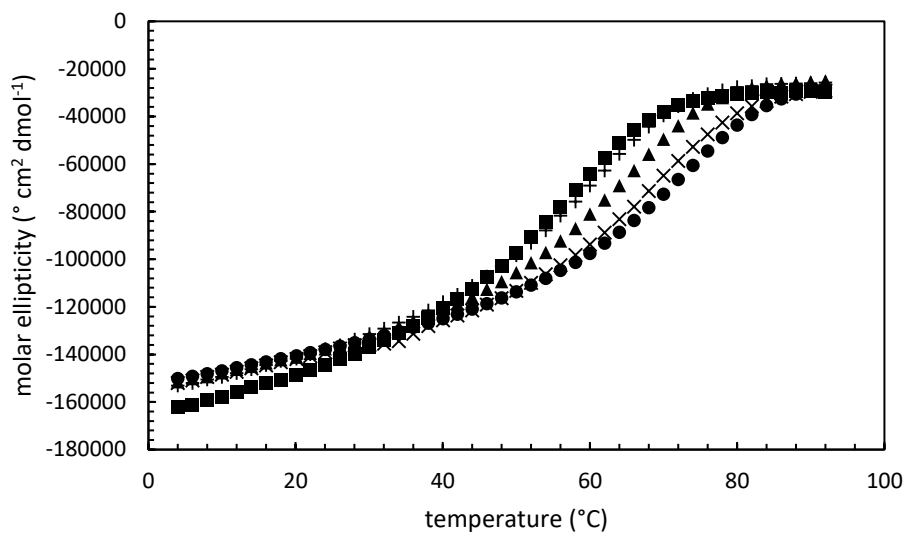
**Figure S2. (A)** Fluorescence emission spectra of CC-19F<sub>CNA</sub><sup>3.5</sup> (35 μM) at 20 °C in phosphate buffer (10 mM, pH 7.4, solid line) and in urea (10 M, dashed line). The excitation wavelength used was 240 nm. **(B)** Fluorescence emission spectra of a mixed solution of CC-19F<sub>CNA</sub><sup>3.5</sup> and CC-20M<sub>SeB</sub><sup>3.5</sup> (1:1, 70 μM total peptide concentration) at 20 °C in phosphate buffer (10 mM, pH 7.4, solid line) and in urea (10 M, dashed line). The excitation wavelength used was 240 nm.



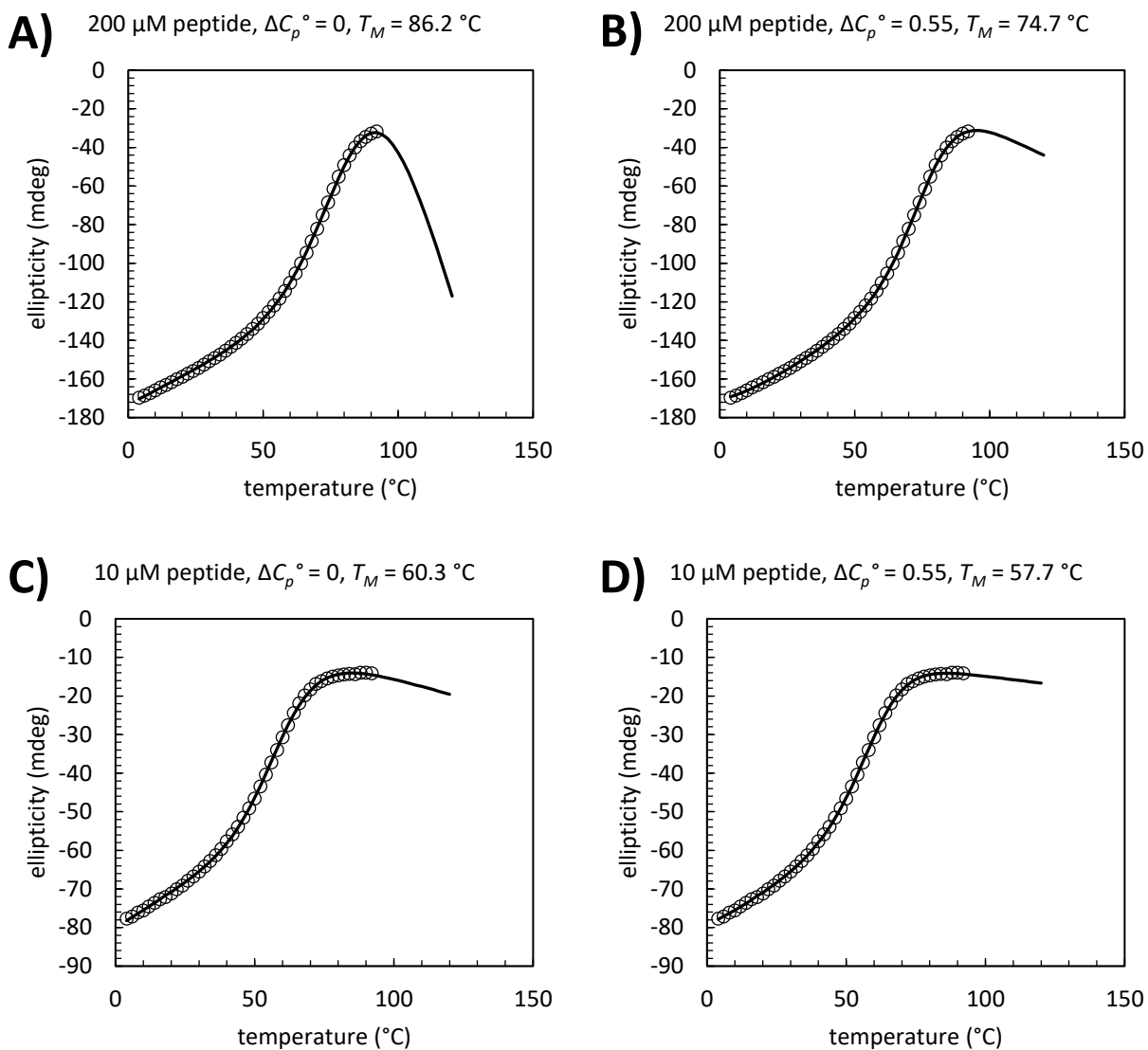
**Figure S3.** Circular dichroism spectra of CC-19WA<sub>N</sub><sup>3.5</sup> (triangles), CC-20M<sub>se</sub>B<sub>N</sub><sup>3.5</sup> (squares) and the dimer (circles) in phosphate buffer (10 mM, pH 7.4) at 20 °C with a 35 μM total peptide concentration.



**Figure S4.** (A) Circular dichroism spectra of CC-19W<sub>N</sub><sup>3.5</sup> (35 μM) at 20 °C in phosphate buffer (10 mM, pH 7.4, triangles) and in urea (10 M, circles). (B) Circular dichroism spectra of CC-20M<sub>Se</sub>B<sub>N</sub><sup>3.5</sup> (35 μM) at 20 °C in phosphate buffer (10 mM, pH 7.4, triangles) and in urea (10 M, circles). (C) Circular dichroism spectra of a mixed solution of CC-19W<sub>N</sub><sup>3.5</sup> and CC-20M<sub>Se</sub>B<sub>N</sub><sup>3.5</sup> (1:1, 35 μM total peptide concentration) at 20 °C in phosphate buffer (10 mM, pH 7.4, triangles) and in urea (10 M, circles).



**Figure S5.** Circular dichroism monitored thermal unfolding curves for the CC-19F<sub>CN</sub>A<sub>N</sub><sup>3.5</sup>/CC-20M<sub>Se</sub>B<sub>n</sub><sup>3.5</sup> dimer at 202 μM (circles), 101 μM (x), 49 μM (triangles), 19 μM (+) and 8 μM (squares). Experiments were performed in phosphate buffer (10 mM, pH 7.4) at 20 °C with a monomer ratio of 1:1. Curves were fit to equation (1) to determine  $T_M$ , the midpoint of the unfolding transition.



**Figure S6.** Dependence of apparent  $T_M$  values on the choice of  $\Delta C_p^\circ$  illustrated for  $[CC] = 200$   $\mu\text{M}$  and  $[CC] = 10$   $\mu\text{M}$ . **(A)** Fit to the 200  $\mu\text{M}$  data with  $\Delta C_p^\circ = 0$  kcal mol $^{-1}$  K $^{-1}$  yields a  $T_M$  of 86.2  $^\circ\text{C}$ , but also leads to a nonphysical post-transition baseline. **(B)** Fit to the 200  $\mu\text{M}$  data with  $\Delta C_p^\circ = 0.55$  kcal mol $^{-1}$  K $^{-1}$  yields  $T_M = 74.7$   $^\circ\text{C}$  with a more reasonable post-transition baseline. **(C)** and **(D)** display fits to melting data collected at  $[CC] = 10$   $\mu\text{M}$ . The choice of  $\Delta C_p^\circ$  has less dramatic effects.