Repeated and Folded DNA Sequences and Their Modular Ag₁₀⁶⁺ Cluster

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Figure S1. *A minimal DNA length develops* Ag_{10}^{6+} : Absorption (A) and circular dichroism (B) spectra of (C₂A)₃ (dashed), (C₂A)₁₀ (dotted), and (C₂A)₁₂ (solid) oligoncleotides with silver clusters. (C) Mass spectra of (C₂A)_{3,10,12} oligoncleotides with silver clusters. The spectra for the oligonucleotides (C₂A)₁₀ and (C₂A)₁₂ are similar to those with (C₂A)₆ (see Figure 1).



Figure S2. A minimal DNA length develops Ag_{10}^{6+} : (A) Absorption (solid), excitation (dashed), and emssion (dotted) spectra of the cluster with $(C_2A)_7$. A 6 nm shift the absorption and excitation maxima may indicate a mixture of species, and we are currently investigating these spectra. (B) Emission spectra acquired with excitation wavelvenths of 300- 600 nm. A consistent emission band indicates that a single emissive species forms. (C and D) Excitation (dashed) and emission (dotted) spectra for the clusters with $(C_2A)_4$ and $(C_2A)_5$, respectively. Similar spectra develop with the longer $(C_2A)_x$ sequences and thus suggest that the Ag_{10}^{6+} cluster has a smaller subuit that is emissive.



Figure S3: A minimal DNA length develops Ag_{10}^{6+} : (A) Size exclusion chromatographs of 30 μ M solution of (C₂A)₆ (Solid) and (C₂A)₆-Ag_{10}^{6+} (dashed) in 10 mM citrate buffer with 40 mM NaClO₄. For (C₂A)₆-Ag₁₀^{6+}, the two peaks are the unlabeled strand and the strand with the cluster. The dotted line was acquired at 440 nm, which corresponds to the absorption of the cluster. The peaks due to the DNA and cluster lie within 0.02 min, which supports a single type of cluster conjugate. (B) A calibration curve relates the retention times to the hydrodynamic radii using single-stranded thymine oligonucleotides dT₁₀, dT₁₅, dT₂₀, and dT₃₀.}



Figure S4. A minimal DNA length develops Ag_{10}^{6+} : (A) Absorption spectra of $(C_2A)_6$ (solid), $(CAC)_6$ (dashed), and $(C(CAC)_6$ (dotted) oligoncleotides with silver clusters. (B) Emission spectra of $(C_2A)_6$ (solid) and $(CAC)_6$ (dashed) oligonucleotides. Permuting the sequence alters the favored environment for the silver clusters.



Figure S5: A minimal DNA length develops Ag_{10}^{6+} : Expanded views of the -4 to -7 charge states for $(C_2A)_6/Ag_{10}^{6+}$ complexes. The isotopologue masses were predicted using the specific formulas, and the blue tick marks represent the predicted masses from these formulas (see each graph). The fully protonated strand has the formula $C_{168}H_{217}O_{100}N_{66}P_{17}$.



Figure S6. A minimal DNA length develops Ag_{10}^{6+} : Absorption (A) and circular dichroism (B) spectra of $(C_2T)_3$ (dashed), $(C_2T)_{10}$ (dotted), and $(C_2T)_{12}$ (solid) oligoncleotides with silver clusters. (C) Mass spectra of $(C_2T)_{3,10,12}$ oligoncleotides with silver clusters. The spectra for the oligonucleotides $(C_2T)_{10}$ and $(C_2T)_{12}$ are similar to those with $(C_2T)_6$ (see Figure 2).



Figure S7: *Multiple nucleobases coordinate* Ag_{10}^{6+} : (A) Fluorescence spectra of (C₂T)₆ at different pH values. The maximum change occurs from pH = 9.2 to 10.2, which brackets the pK_a ~ 9.7 for the N3 of thymine. (B) Fluorescence intensity is strong at high pH, quenches at low pH, and recovers at high pH. Collectively, these spectra suggest that the cluster and H⁺ compete for the N3 binding site in thymine.



Figure S8: *Multiple nucleobases coordinate* Ag_{10}^{6+} : Expanded views of the -6 and -7 charge states for $(C_2A)_8$ - Ag_{10}^{6+} , $-Ag_{11}^{7+}$, $-Ag_{12}^{8+}$, and $-Ag_{13}^{9+}$ complexes. The isotopologue masses were predicted using the specific formulas for these complexes and are indicated by the blue tick marks. The fully protonated strand has the formula $C_{224}H_{289}O_{134}N_{88}P_{23}$.



Figure S9: Longer strands add Ag^+ : (A) Absorption and Circular Dichroism (B) spectra of the $(C_2A)_6$ - Ag_{10}^{6+} complex before and after adding 4 equivalents of Ag^+ followed by dialysis.



Figure S10: Longer strands add Ag^+ : (A) Mass spectra of the -6 charge state of the $(C_2T)_6$ - Ag_{10}^{6+} cluster without and with 4 and 8 added equivalents of Ag^+ : $(C_2T)_{10}$ followed by 100-fold dilution and dialysis. The mass spectra show higher order complexes $(C_2T)_{10}$ - Ag_{10}^{6+} , $-Ag_{11}^{7+}$, $-Ag_{12}^{8+}$, and $-Ag_{13}^{9+}$.



Figure S11: Longer strands accumulate Ag^+ : (A) Continuous variation analysis for Ag^+ binding to native $(C_2A)_6$. The concentration of oligonucleotide is based on the C_2A repeats, and the total concentration of Ag^+ and C_2A is 36 μ M. The ellipticity was measured at 260 nm and the χ is the mole fraction with respect to C_2A . Linear extrapolations from the two limiting concentrations intersect at $\chi \sim 0.46$.