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Analysis of the Interaction between the Cocaine-Binding Aptamer and its Ligands using Fluorescence Spectroscopy

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13 Abstract

14 We used fluorescence spectroscopy to measure the binding affinity and provide new
15 insights into the binding mechanism of cocaine and quinine with the cocaine-binding DNA
16 aptamer. Using the intrinsic fluorescence of quinine and cocaine, we have observed
17 quenching of ligand fluorescence upon binding of aptamer. Quantification of this quenching
18 provides an easy method to measure the binding constant using small amounts of sample.
19 The observed quenching coupled with a red shift of the Stokes shift in the emission
20 spectrum indicates that quinine and cocaine interact with the aptamer through stacking
21 interactions.

22

23 Key words

24 Fluorescence spectroscopy, fluorescence quenching, aptamers, small molecule-DNA
25 interactions

26

27 Introduction

28 The cocaine-binding aptamer has become a widely employed model system for the
29 development of aptamer-based biosensors. The different sensors, utilizing this aptamer,
30 report ligand binding using a variety of methods including color change, electric and
31 fluorescent outputs amongst numerous other examples.¹⁻¹⁵ The secondary structure of the
32 cocaine-binding aptamer is comprised of a three-way junction with a tandem AG mismatch
33 and a dinucleotide bulge located near the junction (Figure 1).¹⁶ When stem 1 of the aptamer
34 is six base pairs long (MN4, Figure 1) the aptamer is folded both in the free state and the
35 bound state. However, if stem 1 is three base pairs long (MN19, Figure 1) the aptamer is
36 loosely structured in the free state and tightens up or becomes structured upon ligand
37 binding.¹⁶⁻¹⁸ It is this ligand-dependant structural change that is exploited in most of the
38 biosensor applications of this aptamer.

39
40 One unusual feature of the cocaine-binding aptamer is that it binds quinine tighter than
41 cocaine, the ligand for which it was originally selected.¹⁹⁻²² Quinine originated from the
42 bark of the *cinchona* tree and has been widely used for centuries to treat malaria. Quinine
43 was also one of the first fluorophores identified and is a standard for calibrating
44 fluorescence spectrometers.²³ The standard emission spectra of quinine in 1 N H₂SO₄ show
45 that quinine has two excited states at ~250 nm and ~350 nm with one emission state at
46 ~450 nm.²⁴ The standard fluorescence lifetime (τ) and quantum efficiency (ϕ) of quinine in
47 1 N H₂SO₄ are (19.2 ± 0.1) ns and 0.545 ± 0.003, respectively.²⁵ These standard values can
48 be converted and employed in any experimental condition using a comparative conversion
49 method.^{26,27} Cocaine is also fluorescent and absorbs the UV light at 232 nm and 274 nm

50 while it emits the fluorescent light at 315 nm at room temperature and neutral pH
51 range.^{24,28}

52
53 In this study, we use the intrinsic fluorescence of both quinine and cocaine to study the
54 binding of quinine and cocaine to two sequence variants of the cocaine-binding aptamer.
55 Fluorescent methods have long been used in aptamer studies but they typically employ a
56 fluorescent tag on the aptamer.²⁹ To the best of our knowledge this is the first study using
57 the intrinsic fluorescence of a ligand to study aptamer-small molecule interactions. On the
58 basis of our observed fluorescence quenching, as aptamer is added to ligand, the K_d of the
59 aptamer-ligand interaction is measured. From a comparison of the change in fluorescence
60 intensity and Stokes shift observed with ethidium bromide, cocaine and quinine binding we
61 conclude that cocaine and quinine form stacking interactions when binding the aptamer
62 and do not bind via an intercalation mechanism.

63

64 **Materials and Methods**

65 ***Sample preparation.***

66 Aptamer samples were obtained from Integrated DNA Technologies (IDT) and had their
67 measured mass confirmed by IDT to be the same as expected using ESI mass spectrometry.
68 The DNA aptamer samples were dissolved in autoclaved distilled deionized H₂O (ddH₂O)
69 and then exchanged three times using a 3-kDa molecular weight cut-off Amicon
70 concentrator with sterilized 2 M NaCl followed by four exchanges into ddH₂O. Quinine
71 hemisulfate monohydrate and cocaine hydrochloride stock powder samples were obtained
72 from Sigma-Aldrich (catalog numbers 145912 and C5776, respectively). Except when it is

73 specified, all aptamer and ligand samples were dissolved in 20 mM sodium phosphate
74 (pH 7.4), 140 mM NaCl before use. These buffer conditions are similar to what was used in
75 our previous calorimetry-based experiments.^{16,20-22} The aptamer and ligand concentrations
76 were determined by a Cary 100 ultraviolet (UV) spectrophotometer using the extinction
77 coefficients supplied by the manufacturers. To avoid the effect of any undesirable quencher
78 species and molecular oxygen all working samples were prepared under sterile conditions,
79 filtered through a 0.2- μm microfilter and degassed with a MicroCal Thermo Vac unit for 5
80 minutes at 4 °C. To induce the intramolecular folding of MN4, MN19, SS1, MS3 and ATP3
81 aptamers, DNA samples were incubated at 95 °C for 3 minutes and immediately immersed
82 in ice-water for 5 minutes before they were mixed with the ligand. The sequence of the MS3
83 aptamer is as published in Neves et al.³⁰ and the sequence of ATP3 is the same as the 27-
84 mer DNA aptamer published by Lin and Patel.³¹

86 ***Fluorescence quenching experiments.***

87 Steady-state fluorescence scans were performed employing a Cary Eclipse
88 spectrofluorometer and 10-mm fused quartz cuvettes. Each experiment was performed at
89 15 °C and 23 °C. The temperature was maintained constant throughout each experiment
90 using a Cary Peltier controller. Next, the spectrofluorometer was optimized for the limit of
91 detection to maintain constant photomultiplier tube (PMT) voltage, signal-to-noise ratio
92 (SNR) and spectral bandwidth (SBW) parameters. For each ligand-aptamer titration, the
93 total ligand concentration was kept constant and at least half of the expected K_d value. The
94 observed fluorescence intensities from 3-6 replicates were corrected for the inner-filter
95 effect to compensate the loss of the incident intensity by:

$$F = F_{obs} \times 10^{\frac{(A_{ex} + A_{em})\ell}{2}}$$
(1)

where F is the corrected fluorescence, F_{obs} is the observed intensity in the absence of the inner-filter effect, A_{ex} and A_{em} are the absorbance values of the aptamer at the excitation and emission wavelengths of the ligand, and (ℓ) is the light path^{32,33}. For the simplicity of the parameters referred to in this study, all of the observed fluorescence intensities were corrected and denoted as fluorescence intensity (F). The obtained fluorescence intensities were averaged and normalized as relative fraction units (RFU) of F_0 .³⁴

For the analysis of 1:1 ligand-aptamer complex, the most valid calculation to quantify the dissociation constant (K_d) is made by a quadratic function:

$$\frac{F_0 - F}{F_0 - F_b} = \frac{[L]_t + [A]_a + K_d - \sqrt{([L]_t + [A]_a + K_d)^2 - 4[L]_t[A]_a}}{2[L]_t}$$
(2)

where F and F_0 are the fluorescence intensities of the ligand in the presence and absence of the aptamer respectively, F_b is the fluorescence of a fully bound ligand-aptamer, $[L]_t$ is the total concentration of ligand, and $[A]_a$ is the concentration of added aptamer.³³ To quantify the binding affinities, each binding isotherm was plotted as a function of bound to free ligand (F/F_0) versus the total aptamer concentration in the solution. Then, the isotherms were fitted to the non-linear regression function:

$$\frac{F}{F_0} = F_1 + (F_2 - F_1) \frac{K_d^n}{K_d^n + x^n}$$
(3)

where n denotes the number of binding sites; F_2 and F_1 are the vertical and horizontal asymptotes respectively. The K_d in Eq. 3 is derived from the quadratic binding function (Eq.

117 2).³³ The fitting model (Eq. 3) was defined and developed applying OriginPro 2016 C
118 scripts.

119
120 In the quinine-aptamer binding assays, quinine was excited at 234 nm. Then, emission
121 scans were performed from 270 nm to 450 nm to exclude the interference of Raman and
122 Rayleigh scattering peaks, and to detect the maximum fluorescence intensity of quinine at
123 ~383 nm. For the cocaine-aptamer binding assays, cocaine was excited at 232 nm. The
124 emission scans were carried out from 270 nm to 450 nm, and the maximum fluorescence
125 intensity of cocaine was detected at ~315 nm. To confirm the quenching results were
126 specific for a functional cocaine-binding aptamer, both ligands were titrated against the
127 non-binding SS1 aptamer at 15 °C and under the same conditions performed for MN4 and
128 MN19.

129
130 In the dynamic quenching analyses, the Stern-Volmer isotherms of the acquired maximum
131 fluorescence intensities were plotted as a function of free to bound ligand (F_0/F) versus the
132 total aptamer concentration. The Stern-Volmer plots were fitted to the mixed static-
133 dynamic quenching models:

$$134 \quad \frac{F_0}{F} = (1 + K_{sv}[Q])(1 + K_a[Q]) \quad (4)$$

$$135 \quad \frac{F_0}{F} = (1 + K_{sv}[Q])e^{V[Q]} \quad (5)$$

136 where the association constant (K_a) and the Stern-Volmer constant (K_{sv}) become mutually
137 dependent constants, and (V) stands for the volume per mole of the ligand-aptamer
138 complex within the static interaction proximity.³⁵⁻³⁷

139

140 To compare the effect of intercalation versus base-stacking interactions, the MN4 and
141 MN19 aptamers were titrated into 1 μ M ethidium bromide and excited at 230 nm, 286 nm
142 and 486 nm separately. Each emission scan was acquired from 550 to 700 nm to detect the
143 fluorescence of ethidium bromide at \sim 613 nm at 23 $^{\circ}$ C. Similar to binding affinity analyses
144 of quinine and cocaine, the maximum emitted intensities of ethidium bromide were
145 averaged, corrected and analyzed as ratios to F_0 . Furthermore, wavelengths of the emission
146 maxima were recorded to determine the Stokes shift of each fluorescence scan. The
147 difference in Stokes shifts between free and bound states ($\Delta\lambda_b^f$) of the ligands (ethidium
148 bromide, quinine and cocaine) with the MN4 and MN19 aptamers were noted for
149 comparison analyses.

150

151 To quantify the bimolecular quenching rate constant (k_q) of cocaine, the fluorescence
152 lifetime (τ) of cocaine in the absence of aptamers were measured using the time-resolved
153 mode of the Cary Eclipse spectrofluorometer at 15 $^{\circ}$ C and 23 $^{\circ}$ C. The rate of the
154 fluorescence intensity as a function of time (t) was fitted to

$$155 \quad F(t) = I_0 e^{-t/\tau} \quad (6)$$

156 where I_0 denotes the incident light intensity²³. For the calculation of k_q in quinine-aptamer
157 binding experiments, we used the standard τ_0 values available in the literature (18.5 ns and
158 17.5 ns) at 15 $^{\circ}$ C and 23 $^{\circ}$ C respectively.²⁵ The k_q was computed using:

$$159 \quad \frac{F_0}{F} = 1 + K_{sv}[Q] = 1 + K_a[Q] = 1 + k_q\tau_0[Q] \quad (7)$$

160 where the Stern-Volmer constant (K_{SV}) represents a dynamic binding constant in a
161 collisional interaction. In an exclusively static quenching, the K_{SV} is replaced with the K_a .^{38,39}

162

163 **Results**

164 *Effect of aptamer binding on ligand fluorescence.*

165 Upon addition of aptamer to quinine or cocaine, the fluorescence of the ligand was
166 quenched (Figure 2). We utilized this quenching to quantify the binding affinity and
167 dynamics of MN4 and MN19 to both quinine and cocaine. The observed fluorescence
168 emission maxima were corrected for the inner-filter effect using Eq. 1 for each aptamer-
169 ligand pair accounting for the absorbance of DNA at the excitation wavelength used for the
170 ligand (Supporting Figure 1). We found that the titrations of MN4 and MN19 into a constant
171 concentration of quinine, while irradiated at 234 nm, quenched the maximum fluorescence
172 emission at ~383 nm (Figure 2a). Similarly, the titrations of MN4 and MN19 aptamers in
173 cocaine, while excited at 232 nm, quenched the maximum fluorescence emission at
174 ~315 nm (Figure 2b). The nonlinear fitting analyses of the acquired binding isotherms
175 using Eq. 3 (Figure 3) yielded the K_d values reported in Table 1. The K_d values of all four
176 aptamer-ligand combinations decreased as the temperature was raised from 15 °C to 23 °C
177 (Table 1).

178

179 In order to confirm that the fluorescence quenching we observe is a result from specific
180 binding, we analysed the change in fluorescence of quinine and cocaine upon addition of
181 the SS1 cocaine-binding aptamer. This aptamer has the same sequence as MN4 except that
182 both AG bases (A21/G29, A7/G30) are switched to be GA base pairs (G21/A29, G7/A30)

183 (Figure 4a). These two changes result in an aptamer that does not bind to quinine as
184 assessed by ITC methods (Figure 4b). When SS1 was titrated into cocaine or quinine, the
185 observed fluorescence reduced in intensity; however, when corrected for the inner filter
186 effect no reduction in binding was observed (Figure 4c, d). As a comparison, the observed
187 and corrected data for MN4-quinine is also shown in Figure 4e. As a further control we
188 tested the MS3 and ATP3 DNA sequences as additional negative controls for binding and
189 also observed no change in the observed fluorescence upon addition of MN4 (Supporting
190 Figure 2).

191
192 The shift of the emission maximum of quinine was measured in the free and ligand-bound
193 states for both MN19 and MN4 aptamers. When quinine was bound by MN4, the emission
194 maxima of quinine shifted (3.05 ± 0.02) nm toward the infrared region (Figure 5). A slightly
195 shorter red shift of (2.49 ± 0.01) nm occurred when quinine was bound by MN19. With
196 cocaine binding, the emission maximum of MN4•cocaine shifted (1.07 ± 0.03) nm toward
197 the infrared region, and with MN19 red-shifted by (0.46 ± 0.01) nm (Figure 5). These
198 differences in Stokes shift are statistically different as confirmed by a t-test with a p-value
199 less than 0.0001.

200
201 To provide a comparison for quinine and cocaine binding, we titrated MN4 and MN19 into
202 ethidium bromide (EtBr) and monitored the change in fluorescence of EtBr. When bound
203 by both MN4 and MN19, the fluorescence intensity of EtBr increased (Figure 5b).
204 Additionally, the MN4•EtBr and MN19•EtBr complexes resulted a blue shift in the EtBr
205 emission spectrum of (10.9 ± 0.02) nm and (11.5 ± 0.1) nm, respectively (Figure 5).

206
207 As a control ligand binding we also analysed the interaction of benzoylecgonine with MN4.
208 Benzoylecgonine is a metabolite of cocaine and the cocaine-binding aptamer is typically
209 described as only weakly binding or not interacting with this molecule.^{1,21,40,41} Using
210 differential scanning calorimetry (DSC) methods Harkness and coworkers have determined
211 an affinity of MN4 for benzoylecgonine of 604 μM at 30 $^{\circ}\text{C}$.²² Benzoylecgonine has the same
212 fluorescent properties as cocaine and as shown in Supporting Figure 3 its fluorescence is
213 quenched upon addition of MN4 with a resulting K_d value of (91 ± 52) μM at 15 $^{\circ}\text{C}$. This
214 binding affinity agrees reasonably well with the expected affinity of 220 μM that is
215 calculated using the thermodynamic parameters previously reported.²² We will also note
216 that the lower K_d value measured here at a lower temperature is consistent with our
217 binding measurements that show that the affinity of the cocaine-binding aptamer increases
218 as the temperature is decreased (Table 1).

219

220 ***Analysis of fluorescence quenching mechanisms.***

221 We analyzed the mechanism of the fluorescence quenching of quinine and cocaine with
222 MN4 and MN19 binding by Stern-Volmer analysis. The Stern-Volmer isotherm of MN4-
223 quinine at 15 $^{\circ}\text{C}$ produced a non-linear plot with K_{SV} constant of (7.2 ± 0.3) μM^{-1} , using the
224 linear portion of the binding curve. The isotherm saturated with excess MN4 indicating that
225 the ligand is fully bound (Figure 6a). In contrast, at 23 $^{\circ}\text{C}$ the MN4-quinine titration
226 resulted in a linear Stern-Volmer plot with K_{SV} constant of (1.31 ± 0.03) μM^{-1} (Figure 6a). As
227 both of these plots have a linear region, when the temperature is increased, the K_{SV} value
228 decreases. Therefore, we conclude that the MN4-quinine quenching follows a static

229 mechanism.^{35,38,39,42}

230

231 The titrations of MN4-cocaine at 15 °C and 23 °C showed non-linear Stern-Volmer plots
232 with K_{SV} constants of $(0.24 \pm 0.02) \mu\text{M}^{-1}$ and $(0.11 \pm 0.00) \mu\text{M}^{-1}$, respectively (Figure 6b).

233 The titrations of MN19-quinine at 15 °C and 23 °C also showed non-linear Stern-Volmer
234 plots with K_{SV} constants of $(2.71 \pm 0.06) \mu\text{M}^{-1}$ and $(0.81 \pm 0.01) \mu\text{M}^{-1}$, respectively (Figure
235 6c). For both of these aptamer-ligand pairs, the curve shows that quenching occurs through
236 a mixed static-dynamic process.^{35,38,39,42}

237

238 For the titration of MN19 with cocaine at 15 °C, we observe a linear Stern-Volmer plot with
239 a K_{SV} constant of $(3.4 \pm 0.06) \times 10^{-3} \mu\text{M}^{-1}$. We obtain non-linear plot with K_{SV} constant of
240 $(4.75 \pm 0.06) \times 10^{-3} \mu\text{M}^{-1}$ at 23 °C (Figure 6d). This switch from a linear to a quadratic plot
241 and an increase in K_{SV} value with temperature indicates that the quenching mechanism
242 changes from a mostly static to a mostly dynamic mechanism.

243

244 The fluorescence lifetime (τ) of free cocaine was measured using time-resolved
245 fluorescence spectroscopy. For free cocaine, τ was measured to be $(2.56 \pm 0.73) \mu\text{s}$ and
246 $(2.23 \pm 0.83) \mu\text{s}$ at 15 °C and 23 °C, respectively (Supporting Figure 4). Using the τ value
247 the bimolecular quenching rate constants (k_q) for cocaine was determined using eq. 7 and
248 reported in Table 1. For quinine, the τ value is too short to measure using our
249 instrumentation. Instead, we used the τ values of quinine available in the literature²⁵ to
250 quantify the k_q values of quinine interacting with MN19 and MN4 (Table 1).

251

252 Discussion

253 We have used the observed fluorescence quenching of quinine and cocaine with aptamer
254 binding to measure the affinity of these ligands to both the MN4 and MN19 cocaine-binding
255 aptamer constructs (Table 1). The values measured here agree within experimental error
256 with our previously reported values using ITC values.^{16,21,30} The benefits of using this
257 fluorescence technique to measure binding affinity are the significantly (over 40 fold)
258 lower amounts of material needed for fluorescence methods compared to ITC methods, and
259 the faster time it takes perform the titration in the fluorescence experiment than in the ITC
260 run (though the ITC experiment is automated). The experiment performed in this study,
261 where the intrinsic change in fluorescence intensity upon binding is used to measure
262 affinity, is not new but has rarely been used to study aptamer-ligand interactions. This
263 method should be easily implemented to other aptamer-ligand pairs as long as the ligand
264 for the aptamer has fluorescence properties.

265
266 The fluorescence studies performed here also provide new insights into the binding
267 interaction of cocaine and quinine with the cocaine-binding aptamer. As we previously
268 noted,²⁰ the values of the thermodynamic binding parameters (ΔH and $T\Delta S$) place cocaine
269 and quinine into the intercalating-type of DNA ligands as classified by Chaires⁴³. However,
270 quinine and cocaine are not known to be intercalating molecules, nor do they seem likely to
271 be intercalators as their structures possess only one or two fused aromatic rings. Instead,
272 we have thought that these two ligands interact with the cocaine-binding aptamer in a
273 stacking manner where one face of the aromatic ring of the ligand interacts in a π - π

274 stacking manner with a base or multiple bases in the aptamer. It is likely that stacking
275 interaction contribute significantly to binding as 6-methoxyquinoline, the aromatic portion
276 of quinine, is bound by the cocaine-binding aptamer ten-fold tighter than cocaine.²¹

277 In support of this stacking mechanism, we measured the fluorescence binding properties of
278 a known intercalator, ethidium bromide, and compared them with those of cocaine and
279 quinine. The fluorescence intensity of ethidium bromide increases when bound by the
280 aptamer and we observe a blue shift of 11-12 nm with MN4 and MN19 binding (Figure 5).
281 These values are consistent with previously reported changes in fluorescence for ethidium
282 bromide intercalating into DNA.^{44,45} In contrast, cocaine and quinine exhibit fluorescence
283 quenching and a red shift when binding MN19 or MN4 (Figure 5). These differences
284 indicate that in the bound state, quinine and cocaine are still at least partially solvent
285 accessible as would be expected in a stacking arrangement. Jagtep *et al.* demonstrated that
286 overlap of conjugated π -systems in stacking interactions results in a distinctive emission
287 transition to the low-energy range, and this red shift increases with greater π -system
288 overlap.⁴⁶ Our results show that quinine binding by MN19 and MN4 yields a greater red
289 shift than when the same aptamers bind cocaine. Detecting a smaller red shift in cocaine-
290 aptamer emission spectra corresponds to cocaine having one aromatic ring as opposed to
291 two fused aromatic rings in quinine. This is consistent with stacking interactions.

292

293 To conclude, the change in the fluorescence spectrum of quinine and cocaine ligands as a
294 function of cocaine-binding aptamer concentration is a powerful and sensitive tool to

295 quantify the binding affinities of cocaine-binding aptamers as well as providing insights
296 into their binding mechanisms.

297

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303

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371 **Tables**
372

Table 1. Dissociation constants and binding parameters of quinine and cocaine with the MN4 and MN19 aptamers constructs.¹

Ligand	Aptamer	T (°C)	K_d (μM)	K_{SV} (μM^{-1})	k_q ($\mu\text{M}^{-1}\text{s}^{-1}$)
Quinine	MN4	15.0	0.094 ± 0.003	7.2 ± 0.3	3.91×10^8
		23.0	0.745 ± 0.004	1.31 ± 0.03	7.08×10^7
	MN19	15.0	0.47 ± 0.01	2.71 ± 0.06	1.46×10^8
		23.0	1.09 ± 0.02	0.81 ± 0.01	4.39×10^7
Cocaine	MN4	15.0	3.92 ± 0.07	0.24 ± 0.02	1.55×10^1
		23.0	7.7 ± 0.1	0.11 ± 0.00	7.10×10^0
	MN19	15.0	21.1 ± 0.6	$(3.4 \pm 0.9) \times 10^{-3}$	2.19×10^{-1}
		23.0	28.8 ± 0.3	$(4.75 \pm 0.06) \times 10^{-3}$	3.10×10^{-1}

¹ Assays carried out in 20 mM sodium phosphate buffer (pH 7.4), and 140 mM NaCl. The error range stated here is the standard deviation after fitting to a mean of three to six replicates.

373
374

375 **Figure Captions**

376

377 **Figure 1.** Structures of the ligands and the DNA aptamers used in this study. Dashed lines
378 between nucleotides indicate Watson-Crick base-pairing in the secondary structure
379 whereas diamonds show the AG base pairs. Solid lines display the phosphodiester bonds in
380 the backbone of the aptamers.

381

382 **Figure 2.** The raw fluorescence emission spectra of (a) quinine and (b) cocaine titrated
383 with the MN4 aptamer. Fluorescence scans were carried out in 20 mM sodium phosphate
384 buffer (pH 7.4), and 140 mM NaCl at 23 °C. Aptamer aliquots quenched fluorescence, and
385 aptamer was added until the fluorescence of the ligand remained unchanged between
386 additions. (a) Fluorescence emission spectra of 0.06 μM quinine hemisulfate excited at 234
387 nm. (b) Fluorescence emission spectra of 4.8 μM cocaine hydrochloride excited at 232 nm.

388

389 **Figure 3.** Steady-State fluorescence quenching analysis of quinine and cocaine ligands
390 binding MN4 and MN19 aptamers in 20 mM sodium phosphate, (pH 7.4) 140 mM NaCl at
391 15 °C (blue triangle) and 23 °C (red square). Displayed here are the titrations of (a) MN4-
392 quinine; (b) MN4-cocaine; (c) MN19-quinine and (d) MN19-cocaine. The corrected and
393 normalized relative fraction fluorescence (RFU) for each ligand is expressed on the y-axis,
394 where F_0 and F are inner-filter corrected emission maxima in the absence and presence of
395 the corresponding aptamer. Each data point represents an average of 3-6 experiments with
396 the error bar representing one standard deviation.

397

398 **Figure 4.** Putative secondary structure of the SS1 aptamer construct (**a**). This aptamer has
399 the tandem AG base pairs as typically seen in the functional cocaine-binding aptamer
400 changed to be GA base pairs. Isothermal titration calorimetry (ITC) experiment (**b**) where
401 the SS1 aptamer is titrated with quinine shows no ligand binding occurs with this aptamer.
402 ITC data was acquired in 20 mM sodium phosphate (pH 7.4) 140 mM NaCl at 20 °C.
403 Negative control titration of the non-binding aptamer SS1 with quinine and cocaine at
404 15 °C. Shown is the emission of cocaine (**c**) and quinine (**d**) ligands versus SS1 aptamer
405 concentration in 20 mM sodium phosphate (pH 7.4) and 140 mM NaCl. The blue diamonds
406 show the observed non-specific quenching of fluorescence due to the inner-filter effect of
407 the aptamer. The red circles are the corrected fluorescence using Eq. 3. Isotherms in (**e**)
408 display a comparison between the observed and corrected fluorescence quenching in MN4-
409 quinine.

411 **Figure 5.** The fluorescence emission spectra of quinine and ethidium bromide titrated with
412 MN4. Fluorescence scans were carried out in 20 mM sodium phosphate buffer (pH 7.4), and
413 140 mM NaCl at 23 °C. Aptamer aliquots were added until the fluorescence of the ligand
414 remained unchanged. (**a**) Fluorescence emission spectra of 0.06 μM quinine hemisulfate
415 excited at 234 nm. (**b**) Fluorescence emission spectra of 1 μM ethidium bromide excited at
416 230 nm. (**c**) Bar graph of the change in Stokes shift for the indicated combinations of ligand
417 and aptamer.

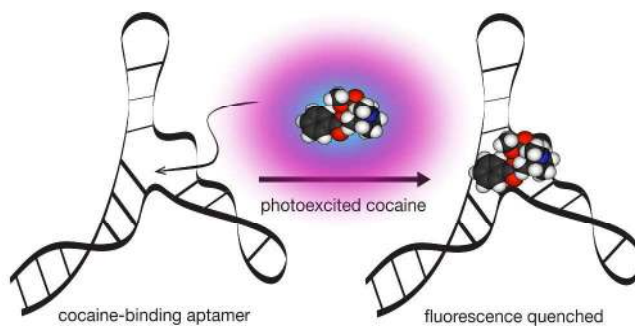
419 **Figure 6.** Dynamic and static fluorescence quenching analysis of quinine and cocaine
420 ligands binding MN4 and MN19 aptamers at 15 °C (blue triangles) and 23 °C (red squares).

421 Shown here are the Stern-Volmer plots of (a) MN4-quinine; (b) MN4-cocaine; (c) MN19-
422 quinine and (d) MN19-cocaine. The corrected and normalized relative fraction
423 fluorescence (RFU) for each ligand is expressed on the y-axis, where F_0 and F are inner-
424 filter corrected emission maxima in the absence and presence of the corresponding
425 aptamer. Each data point represents an average of 3-6 experiments with the error bar
426 representing one standard deviation.

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428 Graphical Abstract



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430 **Supporting Information**

431 **Figure S1.** The spectral overlap of unbound ligands and aptamers examined in this study.
432 Spectra on left are the normalized UV absorbance of (a) MN19, and (c) MN4 aptamers in
433 arbitrary units (a.u.). Spectra on right are the normalized emission fluorescence of (b) free
434 cocaine, and (d) free quinine. All data acquired in 20 mM sodium phosphate (pH 7.4)
435 140 mM NaCl at 23 °C. Quinine emission spectrum does not overlap with the absorbance
436 spectra of aptamers whereas cocaine emission spectrum overlaps with the absorbance
437 spectra of aptamers.

438
439 **Figure S2.** Negative control titrations of the non-binding MS3 (black squares) and ATP3
440 (red circles) aptamers with quinine at 15 °C. Shown is the emission of quinine versus
441 aptamer concentration in 20 mM sodium phosphate (pH 7.4) and 140 mM NaCl. The data
442 shown is the corrected fluorescence using Eq. 3.

443
444 **Figure S3.** Steady-State fluorescence quenching analysis of benzoylecgonine binding MN4
445 in 20 mM sodium phosphate, (pH 7.4) 140 mM NaCl at 15 °C. The corrected and normalized
446 relative fraction fluorescence (RFU) for the ligand is shown on the y-axis, where F_0 and F
447 are inner-filter corrected emission maxima in the absence and presence of the
448 corresponding aptamer. Each data point represents an average of 3 experiments with the
449 error bar representing one standard deviation.

450
451 **Figure S4.** Fluorescence time-resolved analysis of unbound cocaine. Shown are the
452 fluorescence lifetime decay of cocaine in 20 mM sodium phosphate (pH 7.4) 140 mM NaCl

453 at 15 °C (blue) and 23 °C (red). The lifetime measured $(2.56 \pm 0.73) \mu\text{s}$ and $(2.23 \pm 0.83) \mu\text{s}$
454 at 15 °C and 23 °C respectively.

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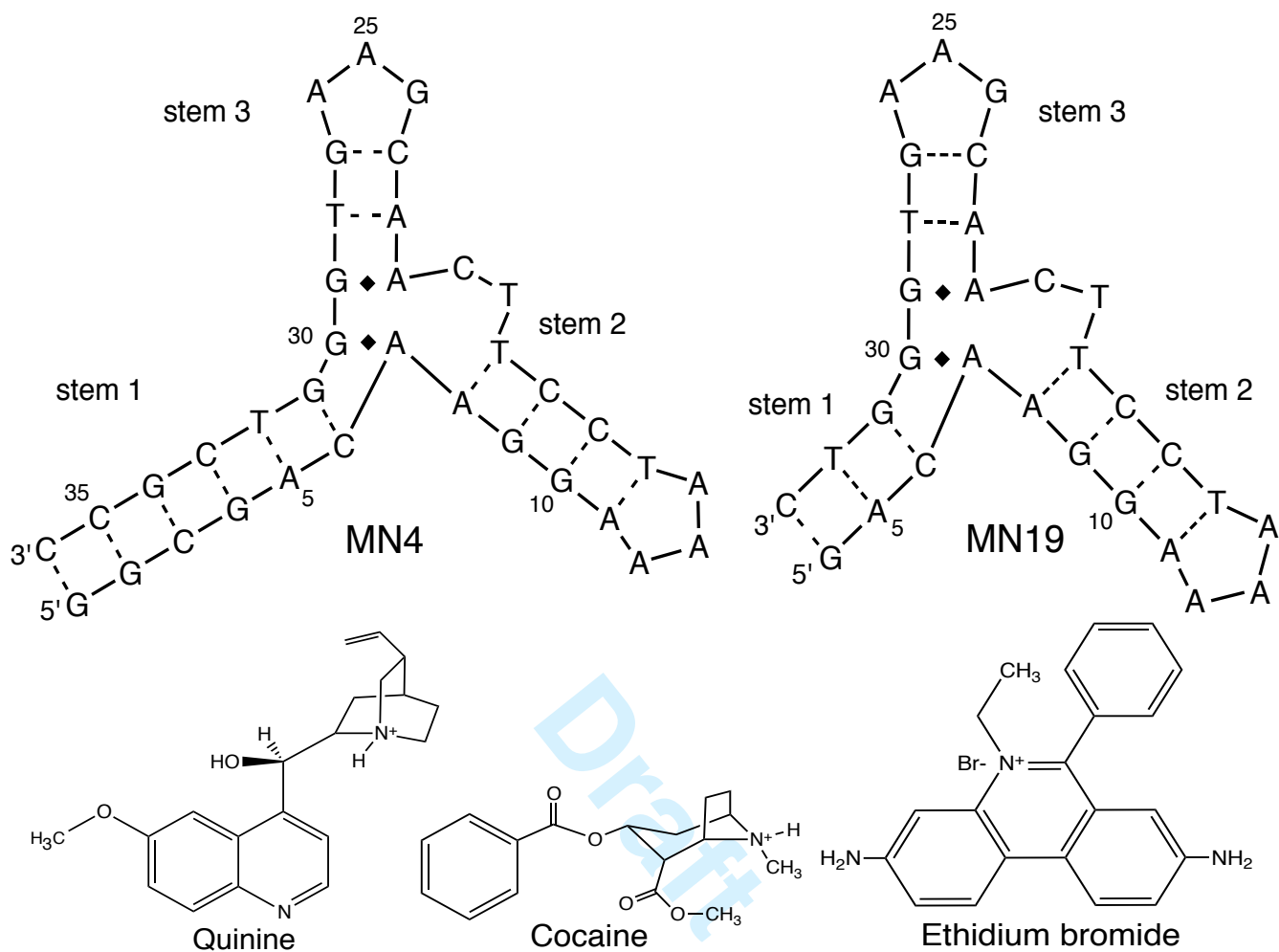


Figure 1

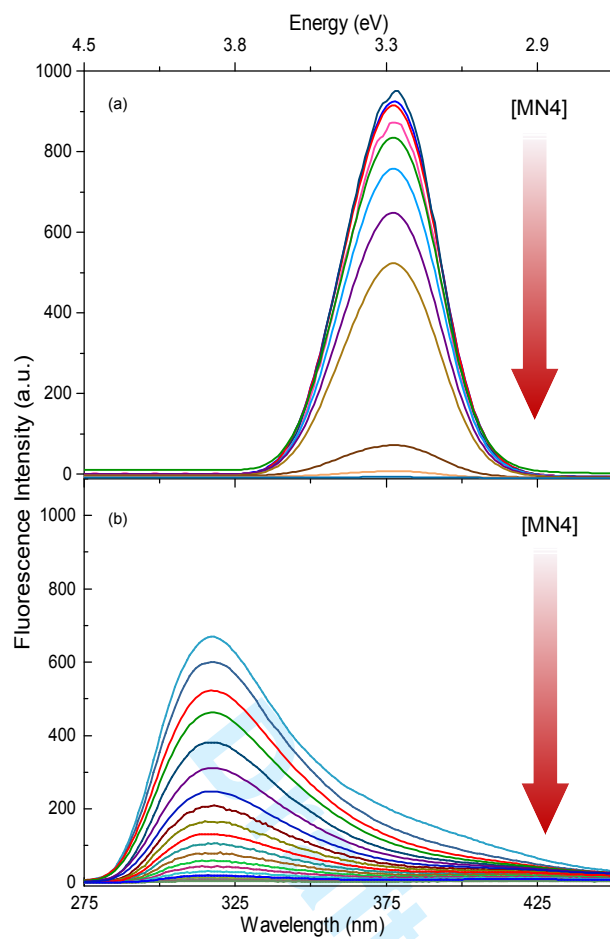


Figure 2

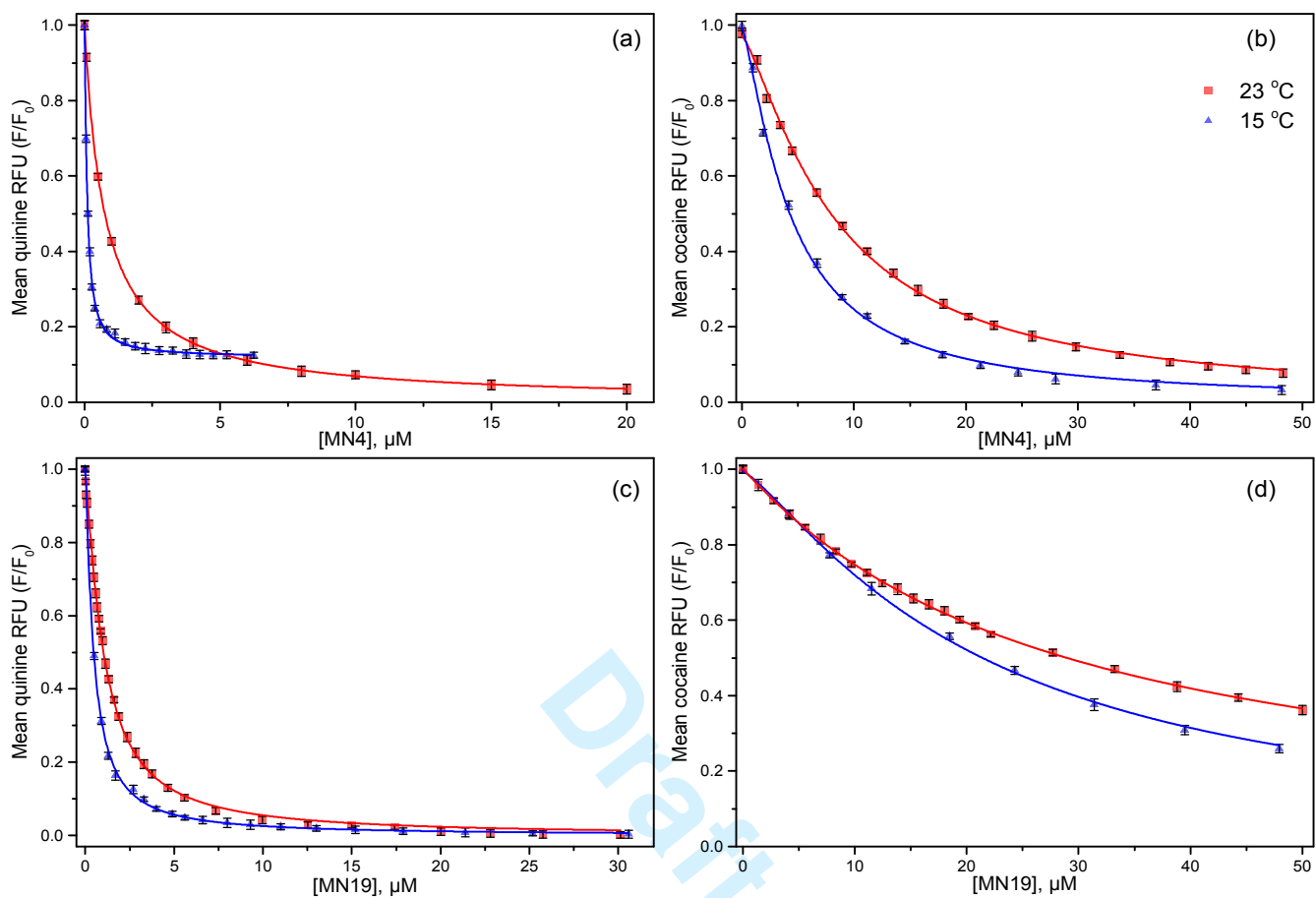


Figure 3

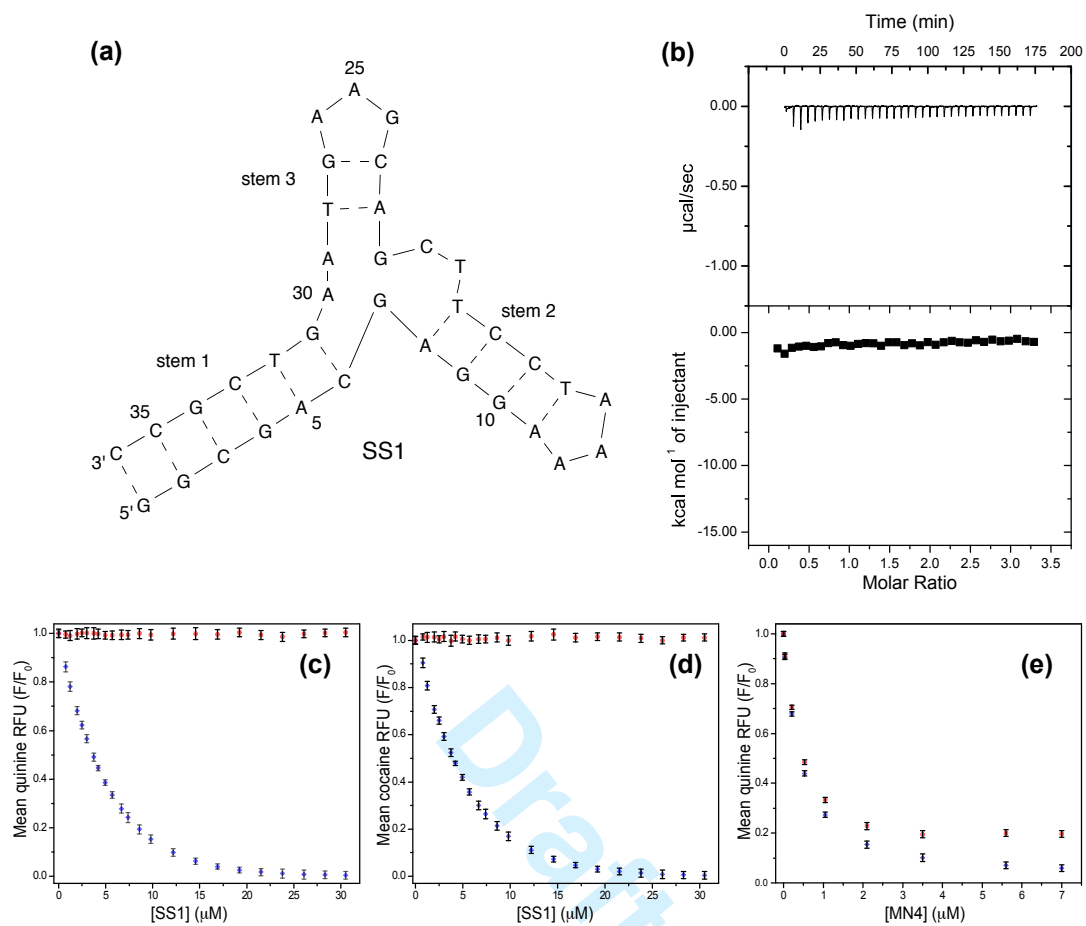


Figure 4

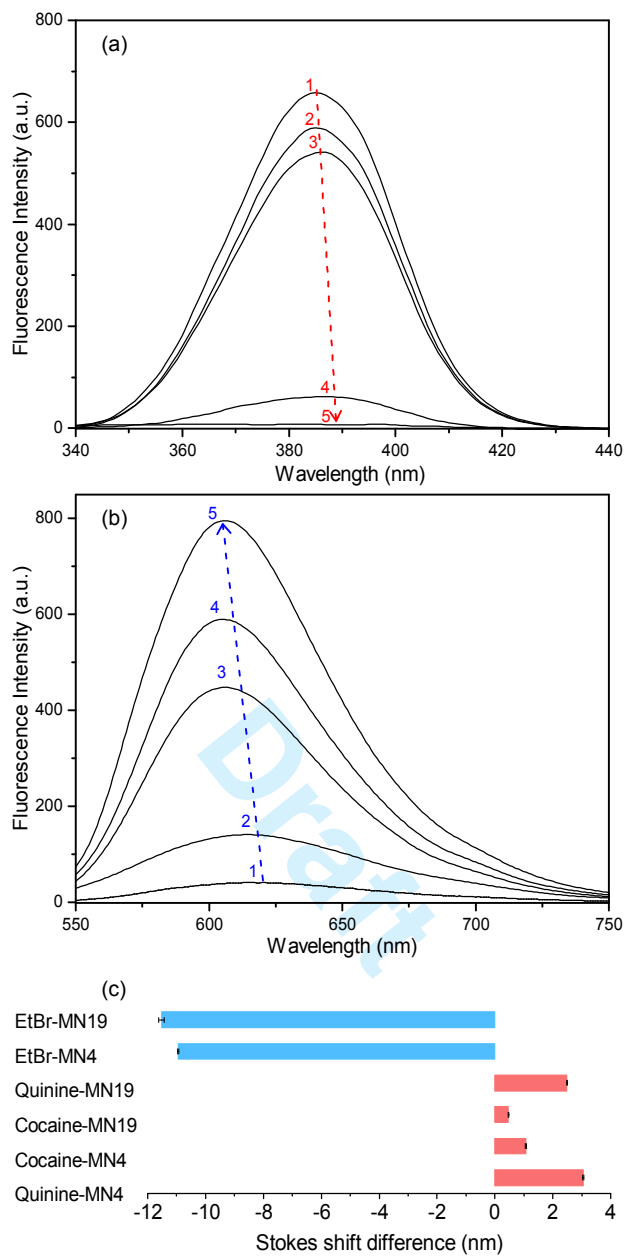


Figure 5

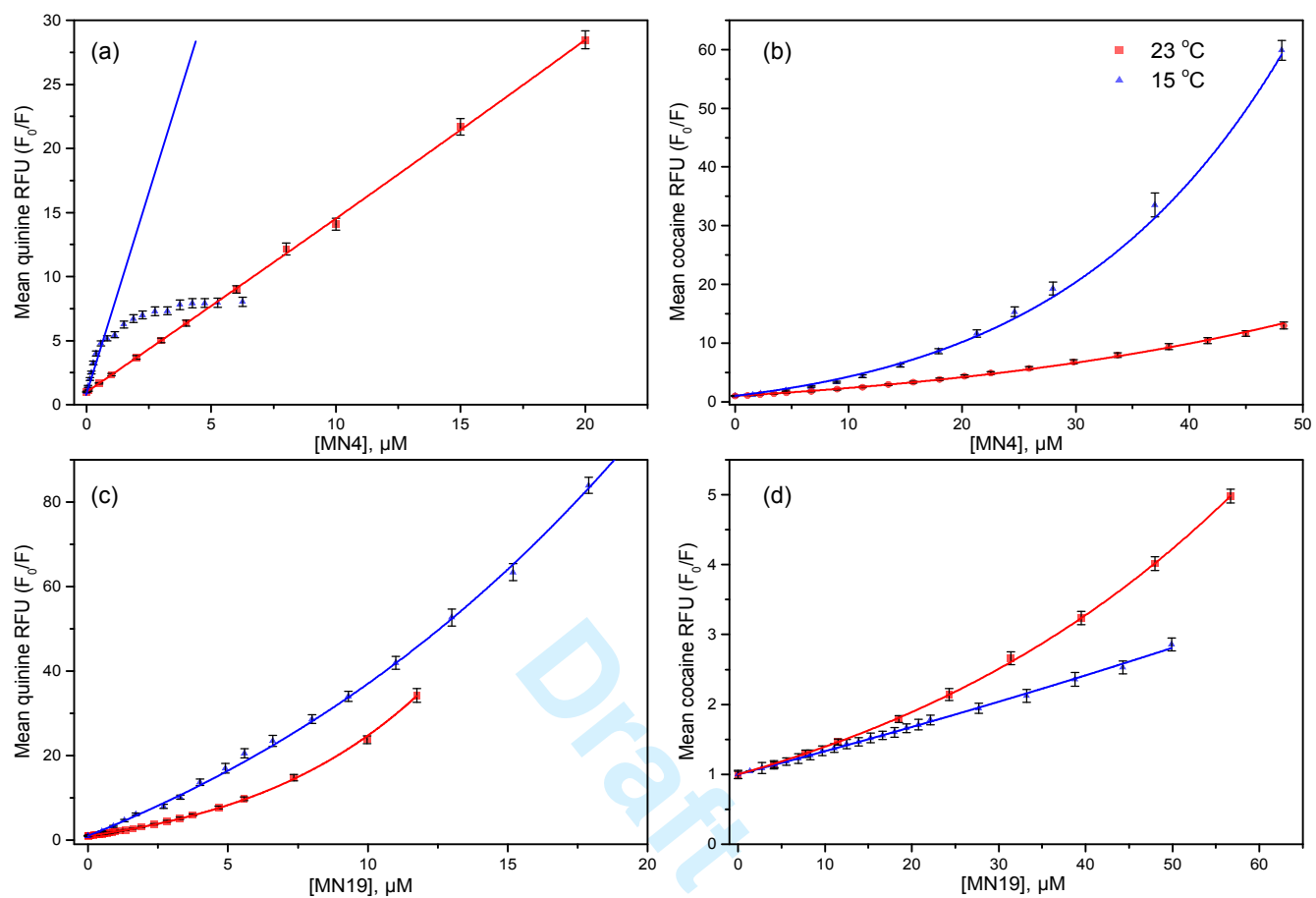


Figure 6