**Supporting Information** 

Control of Electrostatic Interaction between Molecular Beacon Aptamer and Conjugated Polyelectrolyte for Detection Range-Tunable ATP Assay

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**Scheme S1.** Synthetic routes to CPEs. Reagents and conditions: (i)  $Pd(PPh_3)_4$ , aliquat, toluene, 2M  $K_2CO_3$ , reflux; (ii) trimethylamine in tetrahydrofuran/methanol, room temperature, 48 h.

Polymers	M <sub>n</sub> (g∕mol)ª	M <sub>w</sub> (g/mol)⁵	PDI
NP2	21,400	42,100	1.97
NP4	15,300	38,300	2.50
NP6	19,100	34,400	1.80

 Table S1. Molecular weights of neutral precursor polymers.

<sup>a</sup>Number-average molecular weight. <sup>b</sup>Weight-average molecular weight.

Table S2. Summary of	optical properties of	f CPEs in Tris-HCl buffer.
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MDc	) (pm)a	) (pm)b	${\cal D}_{PL}{}^{c}$	$\boldsymbol{\varepsilon}_{max}^{d}$
IVIE S	N <sub>abs</sub> (IIIII)"	Λ <sub>Ρ</sub> ι (ΠΠΙ) <sup>2</sup>	(%)	(M <sup>-1</sup> cm <sup>-1</sup> )
MP2	379	420	35	$4.5 \times 10^{4}$
MP4	378	421	40	$5.4 \times 10^{4}$
MP6	374	418	70	$5.6 \times 10^{4}$

<sup>a</sup>Maximum absorption wavelength. <sup>b</sup>Maximum PL wavelength. <sup>c</sup>PL quantum efficiency. <sup>d</sup>Molar absorption coefficient at  $\lambda_{abs}$ .

S3



**Fig S1.** Absorption (solid) and PL (dashed) spectra of MPs and 6-FAM in 20 mM of Tris-HCl buffer (pH 7.4). [MPs] =  $3 \mu$ M.



**Fig S2.** PL Quantum efficiency of MPs depending on [NaCl] in 20 mM of Tris-HCl buffer (pH 7.4). [NaCl] = 0~100 mM. [MPs] =  $2.5 \mu$ M. Excitation at 380 nm.

Table S3. Equilibrium constants for conformational transformation of MBAs



	MBA sequence (5' - 3')ª	ΔG <sup>0</sup>	Equilibrium constant (K <sub>open-hairpin</sub> )			
MBA1	GCGC GCGG GGAG TATT GCGG AGGA GCGC GC	-5.8 kcal/mol	$1.2 \times 10^{4}$			
MBA2	GCGC GCGC GGGG AGTA TTGC GGAG GAGC GCGC	-9.7 kcal/mol	$6.4 \times 10^{6}$			
MBA3	GCGC GCGC GC <mark>GG GGAG TATT GCGG AGGA</mark> GCGC GCGC GC	-13.5 kcal/mol	3.5 × 10 <sup>9</sup>			
All MPA word labeled with [6 EAN] at E' and [DAPCYL] at 2' termini						

aAll MBA were labeled with [6-FAM] at 5' and [DABCYL] at 3' termini

## \* Calculation of Gibbs free energy

http://mfold.rna.albany.edu

The equilibrium constants for conformational change of MBAs were obtained at the above website where the standard Gibbs free energy change,  $\Delta G^{\circ}$ , was calculated for folding of nucleic acids with particular base sequence.

(37 °C,  $[Na^+]$  = 100 mM and  $[Mg^{2+}]$  = 0 M)

## \* Equilibrium Constant (K)

 $\Delta G = \Delta G^{\circ} + RTInK$ 

In equilibrium state,  $\Delta G = 0$ 

∴∆G° = -RTInK

 $lnK = \Delta G^{\circ}/-RT$ 

 $K = e^{(\Delta G^{\circ} / -RT)}$ 

Where R is gas constant (1.9863 cal/k·mol), T is temperature (310.15 K) and G is Gibbs free energy.



**Fig S3.** Comparison of fluorescence enhancement of (a) MBA1 and (b) MBA3 with increasing [MPs] =  $0 \sim 0.9 \mu$ M. [MBA1] = [MBA3] = 20 nM. Excitation at 490 nm.