

*Supporting Information*

**Drastic enhancement of excess electron-transfer efficiency  
through DNA by inserting consecutive  
5-phenylethynyl-2'-deoxyuridines as a modulator**

Makiko Tanaka<sup>\*a</sup>, Kazuhiro Oguma<sup>a,b</sup>, Yoshio Saito<sup>b</sup> and Isao Saito<sup>\*a</sup>

<sup>a</sup>*NEWCAT Institute, School of Engineering, Nihon University, Koriyama, Fukushima  
963-8642, Japan*

<sup>b</sup>*Department of Chemical Biology and Applied Chemistry, School of Engineering,  
Nihon University, Koriyama, Fukushima 963-8642, Japan*

## Experimental

**Oligodeoxynucleotide synthesis and characterization.** Oligodeoxynucleotides (ODNs) were prepared using phosphoramidite chemistry on an ABI DNA Synthesizer. All phosphoramidites and reagents for DNA synthesis were purchased from Glen Research with the exception of the <sup>Ph</sup>U phosphoramidite. 5-Br-dU phosphoramidite for <sup>Br</sup>U and Pyrene-dU-CE Phosphoramidite for <sup>Py</sup>U were also purchased from Glen Research. 5-(Phenylethynyl)-2'-deoxyuridine (<sup>Ph</sup>U) was prepared according to the literature.<sup>1</sup> Calf intestinal alkaline phosphatase (Promega), Crotalus adamanteus venom phosphodiesterase I (USB), and Penicillium citrinum nuclease P1 (Yamasa Shoyu) were used for the enzymatic digestion of oligonucleotides. All aqueous solutions utilized purified water (Millipore, Milli-Q spUF). Reversed-phase HPLC was performed on CHEMCOBOND 5-ODS-H columns (10x150 mm, 4.6x150 mm) with a JASCO Chromatograph, Model UV-2075 plus at 260 nm. Mass spectra of ODNs purified by HPLC were determined with a MALDI-TOF mass spectrometer.

**UV-vis, and fluorescence measurements.** Absorption spectra were obtained using a Shimadzu UV-2550 spectrophotometer at room temperature using 1 cm path length cell. Fluorescence spectra were recorded on a Shimadzu spectrofluorophotometer (RF-5300PC).

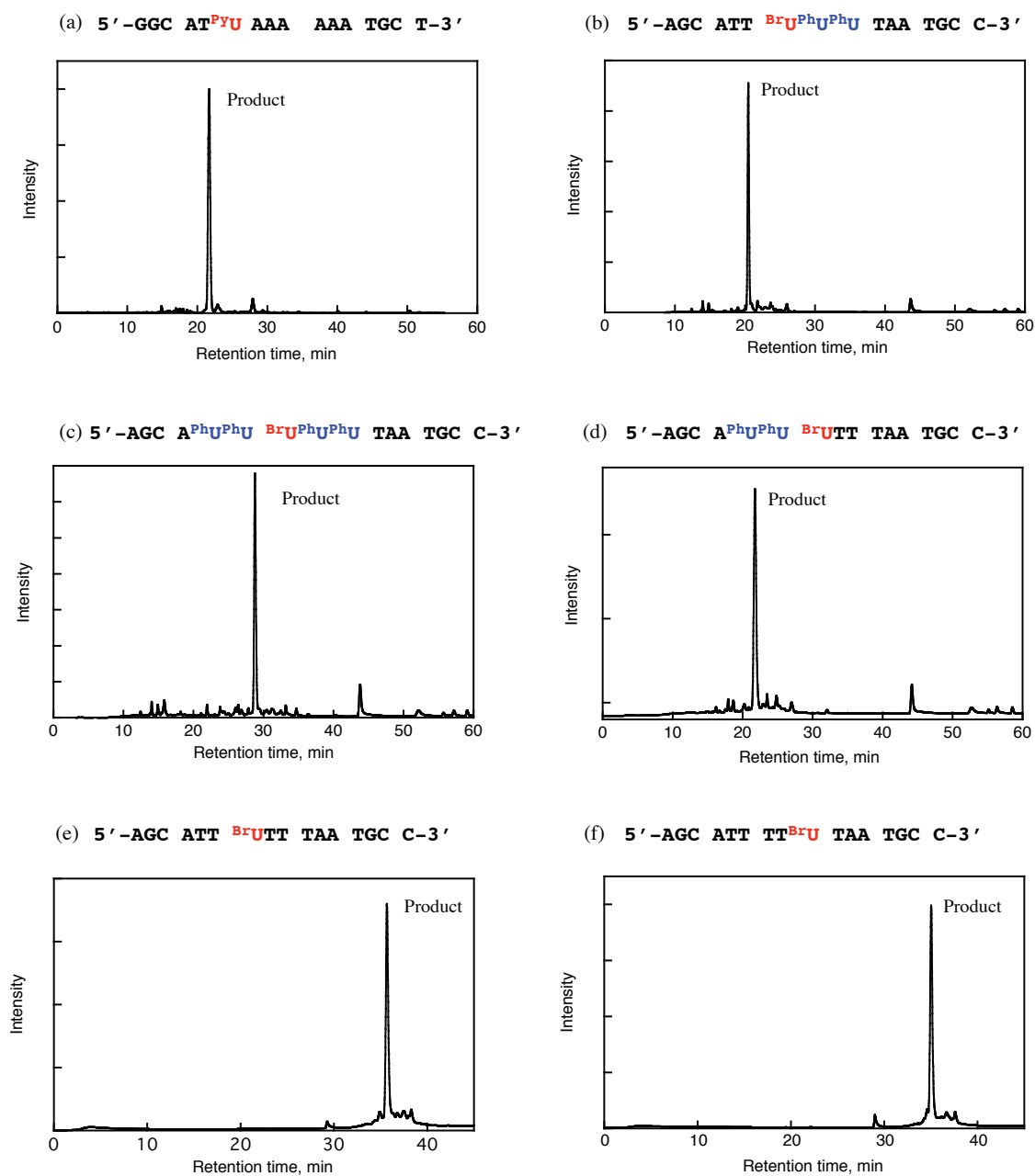
**Photoreduction Experiments.** Aliquots (10  $\mu$ M DNA, 100 mM NaCl, 50 mM sodium phosphate buffer, pH 7.0, total volume 30  $\mu$ L) for irradiation were prepared by annealing equimolar amounts of the desired DNA complements. Aliquots were then transferred to a lucent cell sealed with a rubber septum and deoxygenated with Argon for 20 min. Monochromatic light was irradiated with a Xe lamp (300 W; Asahi Spectra Co. Ltd.; MAX-303) equipped with high concentrating lens unit ( $\phi$  2.5), high quality band-pass filter (LX0400) for 400 nm, and ND filter (5-100%). The shutter timer is variable from 0.5 to 99999.9sec. After irradiation at 400 nm, duplex samples were digested by 37 °C incubation with Calf intestinal alkaline phosphatase, Crotalus adamanteus venom phosphodiesterase I, and Penicillium citrinum nuclease P1 for 12-14 h in order to yield the free nucleosides, and the samples were analyzed by reversed

phase HPLC CHEMCOBOND 5-ODS-H column (4.6x150 mm) at 290 nm. The percentage decomposition of  $^{Br}U$  was determined using dC as an internal standard for all HPLC traces. Irradiations were repeated at least three times and the results averaged.

**Electrochemical Measurements.** Measurements of the cyclic voltammetry (CV) were performed at 298 K, using a BAS 100W electrochemical analyzer in a deaerated DMF containing 0.10 M TBAPF<sub>6</sub> (tetra-*n*-butylammonium hexafluorophosphate) as a supporting electrolyte at 298 K. A conventional three-electrode cell was used with a platinum working electrode and a platinum wire as a counter electrode. The measured potentials were recorded with respect to the Ag/AgNO<sub>3</sub> ( $1.0 \times 10^{-2}$  M) reference electrode. The  $E_{red}$  values (vs Ag/AgNO<sub>3</sub>) are converted to those vs SCE by ferrocene (0.51V vs SCE in DMF)<sup>2</sup> as an internal reference. All electrochemical measurements were carried out under an atmospheric pressure of argon.

## References

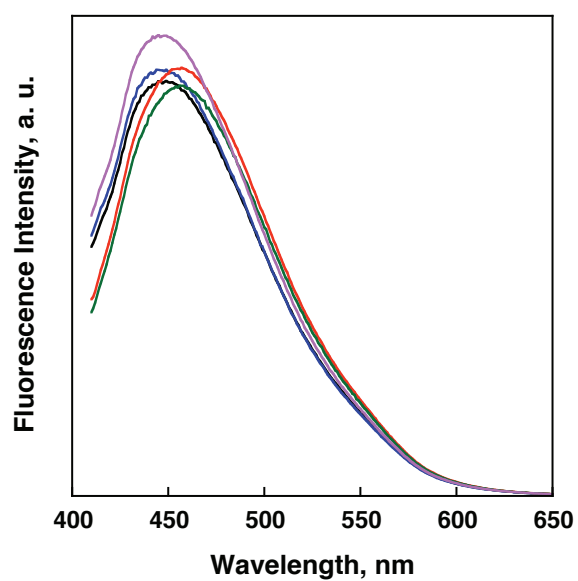
- 1 R. H. E. Hudson and A. Ghorbani-Choghamarani, *Org. Biomol. Chem.*, 2007, **5**, 1845-1848.
- 2 D. Dubois, G. Moninot, W. Kutner, M. T. Jones and K. M. Kadish, *J. Phys. Chem.*, 1992, **96**, 7137-7145.



**Fig. S1** HPLC profile of crude reaction mixture. HPLC analysis was carried out on a CHEMCOBOND 5-ODS-H column eluted with 0.05 M ammonium formate buffer containing acetonitrile. Gradient: 3-30 % over 60 min for (a), 3-40 % over 60 min for (b), (c) and (d), 3-20 % over 45 min for (e) and (f). The largest peaks were collected.

**Table S2. m/z values for modified ODNs determined by MALDI-TOF mass spectrometry.**

| Sequence   | Mass found | Mass calcd ([M+H] <sup>+</sup> ) |
|--|------------|----------------------------------|
| 5'-GGC AT <sup>PyU</sup> AAA AAA TGC T-3'  | 5123.4     | 5124.5                           |
| 5'-AGC ATT Br <sup>U</sup> Ph <sup>U</sup> Ph <sup>U</sup> TAA TGC C-3'                              | 5085.1     | 5084.3                           |
| 5'-AGC A <sup>PhU</sup> Ph <sup>U</sup> Br <sup>U</sup> Ph <sup>U</sup> Ph <sup>U</sup> TAA TGC C-3' | 5257.8     | 5256.5                           |
| 5'-AGC A <sup>PhU</sup> Ph <sup>U</sup> Br <sup>U</sup> TT TAA TGC C-3'                              | 5084.3     | 5084.3                           |
| 5'-AGC ATT Br <sup>U</sup> TT TAA TGC C-3'   | 4913.3     | 4912.1                           |
| 5'-AGC ATT TT <sup>BrU</sup> TAA TGC C-3'  | 4913.2     | 4912.1                           |



**Fig. S3** Fluorescence spectra of DNA I (blue line), II (green line), III (solid red line), IV (purple line), and V (black line); 2.0  $\mu$ M duplex, 50 mM phosphate buffer, pH 7.0, 100 mM NaCl. Excitation wavelength is 400 nm.

**Table S4. Sequence and melting temperatures for DNA duplexes containing a single <sup>Ph</sup>U base and a different counter base (A, T, G, C).**

| DNA sequence   | $T_m$ , °C <sup>a</sup> |
|--|-------------------------|
| 5' -CGC AAT <sup>Ph</sup> U <sup>TA</sup> ACG C-3'<br>3' -GCG TTA AAT TGC G-5'         | 50.6                    |
| 5' -CGC AAT <sup>Ph</sup> U <sup>TA</sup> ACG C-3'<br>3' -GCG TTA <b>T</b> AT TGC G-5' | 39.9                    |
| 5' -CGC AAT <sup>Ph</sup> U <sup>TA</sup> ACG C-3'<br>3' -GCG TTA <b>G</b> AT TGC G-5' | 40.4                    |
| 5' -CGC AAT <sup>Ph</sup> U <sup>TA</sup> ACG C-3'<br>3' -GCG TTA <b>C</b> AT TGC G-5' | 40.2                    |
| 5' -CGC AAC <sup>Ph</sup> U <sup>CA</sup> ACG C-3'<br>3' -GCG TTG AGT TGC G-5'         | 62.5                    |
| 5' -CGC AAC <sup>Ph</sup> U <sup>CA</sup> ACG C-3'<br>3' -GCG TTG <b>T</b> GT TGC G-5' | 50.6                    |
| 5' -CGC AAC <sup>Ph</sup> U <sup>CA</sup> ACG C-3'<br>3' -GCG TTG <b>G</b> GT TGC G-5' | 50.1                    |
| 5' -CGC AAC <sup>Ph</sup> U <sup>CA</sup> ACG C-3'<br>3' -GCG TTG <b>C</b> GT TGC G-5' | 48.2                    |

<sup>a</sup> Experimental conditions: [DNA duplex] = 2.5 μM in 50 mM sodium phosphate, pH7.0, 100mM NaCl.  $T_m$  is determined by monitoring the UV absorption at 260 nm.