

DNAzymes

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A Light-Driven DNA Nanomachine for the Efficient Photoswitching of RNA Digestion**

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Recently, DNA has gained attention as one of the most promising molecules for use in bottom-up nanotechnology.^[1] In the last two decades, numerous DNA nanostructures with mechanical functions such as DNA tweezers, DNA walkers, and DNA gears have been constructed. [2,3] However, the practical use of DNA nanotechnology remains a great challenge. One of the problems limiting the application of DNA nanomachines is that oligo-DNAs or other small molecules have to be added as the "fuel" during each operation cycle, and "waste" molecules detrimentally accumulate in the system after several cycles.[3] Previously, we have tried to solve this problem by constructing a model photon-fueled nanomachine involving azobenzene moieties as photoswitches, based on the photoregulation of DNA hybridization.^[4,5] No waste was produced because only light, the cleanest source of energy, was used to drive the nanomachine, and the operation could be repeated for many cycles without loss of efficiency. [4a] Efforts should be made to construct nanomachines that work on the single-molecule level, which is highly favorable for nanotechnology applications. [4c] In the present study, we constructed a machinelike photoresponsive DNAzyme that can work at the singlemolecule level (intramolecular nanomachine). The change of its topological conformation can be regulated simply by light irradiation. For the first time, complete ON-OFF photoswitching of RNA digestion has been realized by regulating the higher order structure of a DNAzyme-RNA complex.

The 10–23 DNAzyme, captured from a DNA pool of random sequences by in vitro selection, was used here as the model system for constructing an RNA-cleaving nanomachine driven by photons. [6] As shown in Scheme 1 a, a photoresponsive machinelike DNAzyme (**Dz7X**) was con-

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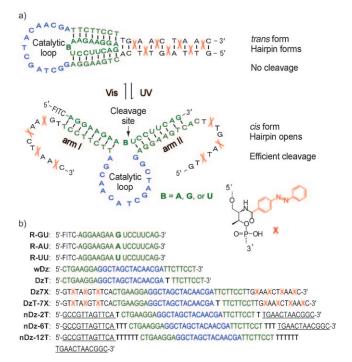
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structed by attaching complementary azobenzene-modified sequences to both ends of the 10–23 DNAzyme. As in our previous design, the two azobenzene-modified sequences were able to form an interstrand-wedged duplex after hybridization. [4c,d] When visible light is applied, the azobenzene residues (**X**) take the *trans* form, and a very stable duplex structure involving seven azobenzene units and nine base pairs is formed. [4c] The modified DNAzyme **Dz7X** can be regarded as a DNA hairpin structure with a big loop (Scheme 1a). In this case, the RNA-cleavage activity is expected to be suppressed because the topologically constrained higher order structure of the catalytic loop cannot form the correct conformation for cleavage, even when the RNA substrate hybridizes with both arms. On the other hand, when UV light is applied and azobenzene residues take the *cis*



Scheme 1. Design of the machinelike photoresponsive DNAzyme (a) and sequences of DNAzymes and RNA targets used in this study (b). The azobenzene-modified DNAzymes show high activity only when azobenzenes (in red; its structure (X) is also shown) take the cis form after irradiation with UV light. RNA substrate (in green) was labeled with the fluorophore FITC at the 5' end. The sequence of catalytic loop is shown in blue; two arms of each DNAzyme complementary to RNA substrate are shown in green. For the nDz series, a hairpin structure with a 12 bp long stem (underlined sequences) can form, and poly(deoxythymidine) spacers (bold letters) of various lengths are introduced between the loop and the stem.

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form, the steric hindrance of the nonplanar structures may cause the dissociation of part of the duplex so that the RNA is cleaved (Scheme 1 a). Notably, the opening and closing of the big hairpin can be carried out at the single-molecule level. Although several photoresponsive DNAzymes involving one azobenzene unit in the catalytic loop as the photoswitch have been reported, the strategies behind their function are completely different from that here. [7] Another merit of our design is that the modified DNAzyme should have greater exonuclease resistance than the wild-type one. [4e]

Figure 1 shows the results of the digestion of an RNA substrate by **Dz7X** at the GU site, which is one of the best substrate sites for the 10–23 DNAzyme.^[6] Here, single-

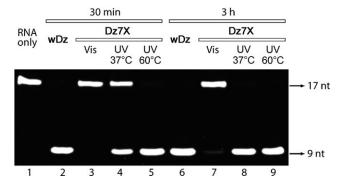


Figure 1. Cleavage of RNA target at the GU site using the photoresponsive DNAzyme Dz7X. Lane 1: control (RNA substrate); lanes 2 and 6: positive control of wild-type DNAzyme; lanes 3–5: 30 min of cleavage after irradiation with visible light at 37 °C, with UV light at 37 °C, and with UV light at 60 °C, respectively; lanes 6–9: 3 h of cleavage corresponding to the same samples in lanes 2–5. Conditions: 1.0 μM RNA, 5.0 μM DNAzyme, 1.0 M NaCl, 10 mM MgCl₂, pH 7.5 (Tris-HCl buffer), at 37 °C. The digestion products were analyzed using 20% denatured PAGE electrophoresis.

turnover conditions were used, and the concentration of Dz7X was five times higher than that of the RNA substrate. Either UV or visible light was applied before MgCl₂ was added to start the reaction. In the case of visible light, the digestion was very slow, and less than 1.8% of RNA was digested after 30 min of reaction at 37°C (lane 3, Figure 1). The melting temperature $(T_{\rm m})$ of the stem duplex part involving alternating azobenzene moieties and base pairs was measured to be 72°C, indicating that almost 100% of **Dz7X** was in the hairpin state (see Figure S1 in the Supporting Information). On the other hand, 51.4% of the RNA substrate was digested after 30 min when the DNAzyme had been irradiated with UV light at 37°C for 5 min (lane 4, Figure 1). The acceleration of RNA digestion by irradiation with UV light was as much as 29-fold. After 3 h reaction, only 11.2% of RNA was digested following visible light irradiation, and RNA was almost completely digested in the case of UV light irradiation (lanes 7 and 8, Figure 1). Thus, efficient photoregulation of RNA cleavage was realized by introducing multiple azobenzenes, as outlined in Scheme 1.

Compared with the results obtained with wild-type DNAzyme (**wDz**) (lane 2, Figure 1), the reaction of **Dz7X** after UV light irradiation (at 37°C) showed only about 40%

activity, indicating that not all hairpins were opened under these conditions. UV/Vis spectroscopic analysis showed that about 38% of azobenzene moieties were photoisomerized to the cis form when UV light was applied at 37 °C (see Figure S2 in the Supporting Information). When UV light irradiation was carried out at 60°C, at which point about 72% of azobenzene moieties had been photoisomerized to the cis form and almost all hairpins were open at 37°C, more than 90% activity was recovered compared with wDz. This result is reasonable because the RNA/DNAzyme complex in the open state has exactly the same conformation as that of wDz (Scheme 1a). The observed rate constants (k_{obs}) of trans-Dz7X, cis-Dz7X (UV at 37 °C), and wDz were measured to be 0.00071, 0.027, and $0.081 \, \mathrm{min^{-1}}$, respectively (see Figure S3 in the Supporting Information for the time-course of RNA cleavage). Thus, the difference in $k_{\rm obs}$ between the reactions with UV and Vis light irradiation was as large as 38-fold.

Similar results were also obtained for cleaving AU, UU, and GC sites (see Figure S4 in the Supporting Information for sequences of DNAzymes and RNA substrates with GC sites), although lower activity was observed (Figure 2). The differ-

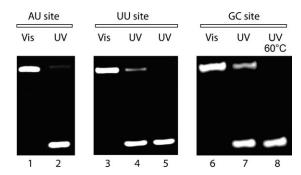


Figure 2. Photoregulation of RNA cleavage at AU, UU, and GC sites using azobenzene-modified DNAzymes. The digestion reaction was carried out for 5 h for the AU site, 18 h for the UU site, and 48 h for the GC site (see Figure S4 in the Supporting Information for sequences). For the UU site, besides Dz7X (lane 4), Dz7X-T with an additional dT was also used (lane 5). For lane 8, UV light irradiation was carried out at 60 °C. Both visible and UV light irradiation were carried out at 37 °C for all other experiments. Other conditions are the same as listed in Figure 1.

ence in activity of the DNAzymes between UV and visible light irradiation was 15- to 40-fold. Especially for the UU site, which was barely cleaved by wDz, we found that a modified DNAzyme (DzT) with an additional deoxythymidine unit (dT) at the junction position between arm II and the catalytic loop could greatly improve activity (lanes 4 and 5, Figure 2). The $k_{\rm obs}$ of ${\bf DzT}$ was 2.7-fold greater than that of ${\bf wDz}$ (data not shown). After UV light irradiation at 37°C a modified DNAzyme with an additional dT unit and seven azobenzene moieties (Dz7X-T) displayed an activity even comparable to that of wDz. Thus, with Dz7X-T, enhancement of activity and clear-cut photoregulation were realized simultaneously for cleaving the UU site. For all the RNA targets examined, the efficiency of the photoregulation was much higher than that achieved with photoresponsive DNAzymes reported previously.^[7] As most RNA-cleaving DNAzymes and ribozymes use a similar strategy for targeting RNA substrates with two arms, the strategy proposed here can be generalized for constructing other photoresponsive DNAzymes or ribozymes.[8]

Figure 3 shows real-time photoswitching of RNA digestion at the GU site at 37°C under multiple-turnover conditions. UV and visible light were applied alternatively

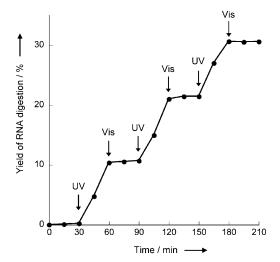


Figure 3. Photoswitching of RNA cleavage at the GU site by Dz7X. Visible light was applied at 0, 60, 120, and 180 min. UV light irradiation was carried out at 30, 90, and 150 min. The irradiation time for visible and UV light was 1 and 10 min, respectively.

at a fixed temperature (37°C). The digestion results clearly show that UV light turned on digestion and visible light completely turned it off (Figure 3). Photoregulation was completely reversible and almost no deviation was observed after three repetitions. As the introduced azobenzene moiety is very stable under physiological conditions, this ON-OFF regulation can in principle be repeated many times.^[9] Efficient suppression of RNA digestion by visible light is especially important for in vivo applications, where an RNA transcript can be digested at an exact point and time by control of UV light irradiation.

For evaluating the topological effect on RNA digestion, several DNAzymes with hairpin structures containing a stem 12 base pairs (bp) long and poly(dT) spacers of various lengths were designed (Scheme 1). As shown in Figure 4, when less than two dTs (only one dT in each side) were introduced, for which the topological constraint was much stronger, almost no digestion was observed (see Figure S5 in the Supporting Information for secondary structures). When six dTs were present, cleavage occurred (lanes 3 and 7, Figure 4). When the number of dTs increased to 12, the entire RNA substrate was digested after 4 h (lane 4, Figure 4). Interestingly, even for nDz-12T, only partial cleavage activity was recovered compared with that for wDz, indicating that the topological constraint influenced the digestion greatly.^[10] This supports our idea that photoregulation of RNA digestion by **Dz7X** occurs according to the strategy shown in Scheme 1. The change of topological conformation of the RNA/DNA-

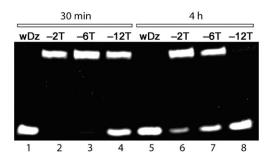


Figure 4. Effect of the topological constraint on RNA digestion by the 10-23 DNA enzyme (see sequences in Scheme 1 b).

zyme complex caused a large difference in cleavage activity between the trans and cis forms. Thus, we have succeeded in constructing a nanomachine that works on the single-molecule level for photoswitching RNA digestion.

In conclusion, clear-cut and reversible photoregulation of RNA digestion was attained by regulating the higher order structure of DNAzyme/RNA complexes. Our proposed strategy is applicable to other RNA-cleaving DNAzymes and ribozymes that target RNA substrates with two arms, allowing the construction of many photoresponsive DNAzymes or ribozymes. This newly constructed nanomachine is also promising for applications either in vivo for photoregulating gene expression with an antisense strategy, or in vitro for photoswitching the mechanical movement of a nanorobot.

Experimental Section

Oligonucleotides containing azobenzene residues were supplied by Nihon Techno Service Co., Ltd. (Tsukuba, Japan) and purified by either polyacrylamide gel electrophoresis or HPLC. DNA sequences containing only natural nucleotides were supplied by Integrated DNA Technologies, Inc. (Coralville, IA, USA). The concentration of all oligonucleotides was determined by UV/Vis spectroscopic analysis within 10% error.

Typical procedure for RNA digestion by a DNAzyme: First, a mixture of substrate RNA and the DNA enzyme was annealed in buffer (1m NaCl, 50 mm Tris-HCl, pH 7.5) by heating at 80 °C for 30 s followed by cooling to 37°C for 15 min. Then, either visible light (400-600 nm, 1 min, L-42 filter) or UV light (320-380 nm, 5-10 min, UV-D36C filter, 5.3 mW cm⁻²) irradiation was carried out at 60 or 37°C. The yield of photoisomerization was calculated from the absorbance change at 340 nm (V-530 UV/Vis spectrometer, Jasco, Tokyo, Japan). After this procedure, MgCl2 was added to start the cleavage reaction, and the reaction mixture was incubated at 37°C. For real-time photoswitching of RNA digestion, UV and visible light were applied alternatingly after MgCl2 had been added. During the reaction, a 4 µL aliquot of the reaction mixture was removed and added to 4 µL loading buffer (50 mm EDTA, 45 mm Tris-HCl, 45 mm borate, 7.0 m urea) to terminate the reaction. Finally, 8.0 µL of the resulting mixture was subjected to electrophoresis on a 20% polyacrylamide gel containing 7.0 m urea. Imaging and quantification of the digested RNA were carried out on a Fujifilm FLA-3000G fluorescent analyzer (Fuji Film, Tokyo, Japan).

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- [1] a) N. C. Seeman, A. M. Belcher, Proc. Natl. Acad. Sci. USA 2002, 99, 6451 – 6455; b) N. C. Seeman, P. S. Lukeman, Rep. Prog. Phys. 2005, 68, 237-270; c) M. K. Beissenhirtz, I. Willner, Org. Biomol. Chem. 2006, 4, 3392-3401; d) N. C. Seeman, Trends Biochem. Sci. 2005, 30, 119-125.
- [2] a) J. H. Chen, N. C. Seeman, *Nature* **1991**, *350*, 631 633; b) C. A. Mirkin, R. L. Letsinger, R. C. Mucic, J. J. Storhoff, Nature 1996, 382, 607-609; c) E. Winfree, F. Liu, L. A. Wenzler, N. C. Seeman, Nature 1998, 394, 539-544; d) W. M. Shih, J. D. Quispe, G. F. Joyce, Nature 2004, 427, 618-621; e) P. W. K. Rothemund, *Nature* **2006**, 440, 297-302; f) R. Chhabra, J. Sharma, Y. Liu, H. Yan, Nano Lett. 2006, 6, 978-983; g) R. P. Goodman, I. A. T. Schaap, C. F. Tardin, C. M. Erben, R. M. Berry, C. F. Schmidt, A. J. Turberfield, Science 2005, 310, 1661 -1665.
- [3] a) B. Yurke, A. J. Turberfield, A. P. Mills, F. C. Simmel, J. L. Neumann, Nature 2000, 406, 605 - 608; b) J. S. Shin, N. A. Pierce, J. Am. Chem. Soc. 2004, 126, 10834-10835; c) W. U. Dittmer, S. Kempter, J. O. Radler, F. C. Simmel, *Small* **2005**, *1*, 709–712; d) T. Liedl, T. L. Sobey, F. C. Simmel, *Nano Today* **2007**, *2*, 36– 41; e) Y. He, M. Su, C. Zhang, A. E. Ribbe, W. Jiang, C. Mao, Nature 2008, 452, 198-202.
- [4] a) X. G. Liang, H. Nishioka, N. Takenaka, H. Asanuma, ChemBioChem 2008, 9, 702-705; b) X. G. Liang, H. Nishioka, N. Takenaka, H. Asanuma, DNA Computing, Springer, New York, 2009, pp. 21-29 (Lecture Notes in Computer Science, Vol. 5347); c) X. G. Liang, T. Mochizuki, H. Asanuma, Small 2009, 5, 1761 – 1768; d) X. G. Liang, H. Nishioka, T. Mochizuki, H. Asanuma, J. Mater. Chem. 2010, 20, 575 – 581; e) X. G. Liang,

- M. Komiyama, H. Asanuma, Nucleosides Nucleotides Nucleic Acids 2008, 27, 332-350.
- [5] a) H. Asanuma, X. G. Liang, H. Nishioka, D. Matsunaga, M. Liu, M. Komiyama, Nat. Protoc. 2007, 2, 203-212; b) H. Asanuma, T. Ito, T. Yoshida, X. G. Liang, M. Komiyama, Angew. Chem. 1999, 111, 2547-2549; Angew. Chem. Int. Ed. 1999, 38, 2393-2395; c) X. G. Liang, H. Asanuma, H. Kashida, A. Takasu, T. Sakamoto, G. Kawai, M. Komiyama, J. Am. Chem. Soc. 2003, 125, 16408-16415.
- [6] a) S. W. Santoro, G. F. Joyce, Proc. Natl. Acad. Sci. USA 1997, 94, 4262-4266; b) J. Nowakowski, P. J. Shim, G. S. Prasad, C. D. Stout, G. F. Joyce, Nat. Struct. Biol. 1999, 6, 151-156.
- [7] a) Y. Liu, D. Sen, J. Mol. Biol. 2004, 341, 887-892; b) S. Keiper, J. S. Vyle, Angew. Chem. 2006, 118, 3384-3387; Chem. Int. Ed. 2006, 45, 3306 - 3309; c) H. Asanuma, H. Hayashi, J. Zhao, X. G. Liang, A. Yamazawa, T. Kuramochi, D. Matsunaga, Y. Aiba, H. Kashida, M. Komiyama, Chem. Commun. 2006, 5062-5064; d) D. Matsunaga, H. Asanuma, M. Komiyama, J. Am. Chem. Soc. 2004, 126, 11452-11453.
- [8] a) S. Schubert, J. Kurreck, Curr. Drug Targets 2004, 5, 667-681; b) G. F. Joyce, Annu. Rev. Biochem. 2004, 73, 791-836; c) K. E. Blount, O. C. Uhlenbeck, Annu. Rev. Biophys. Biomol. Struct. 2005, 34, 415-440; d) M. J. Fedor, J. Mol. Biol. 2000, 297, 269-291.
- [9] The cis-to-trans thermal isomerization does not influence the RNA digestion reaction at 37 °C. The half-life of cis-azobenzene used here is 45 h at 37 °C and 3.3 h at 60 °C. H. Nishioka, X. G. Liang, H. Kashida, H. Asanuma, Chem. Commun. 2007, 4354-
- [10] These results are also helpful for understanding the mechanism of RNA digestion by 10-23 DNAzyme, which is difficult to clarify as the detailed 3D structure of RNA/DNAzyme complex is unknown.

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