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A Biomolecule-Compatible Visible Light-Induced Azide Reduction from a DNA-Encoded Reaction Discovery System

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Abstract

Using a system that accelerates the serendipitous discovery of new reactions by evaluating hundreds of DNA-encoded substrate combinations in a single experiment, we explored a broad range of reaction conditions for new bond-forming reactions. We discovered reactivity that led to a biomolecule-compatible, Ru(II)-catalyzed, visible light-induced azide reduction reaction. In contrast with current azide reduction methods, this reaction is highly chemoselective and is compatible with alcohols, phenols, acids, alkenes, alkynes, aldehydes, alkyl halides, alkyl mesylates, and disulfides. The remarkable functional group compatibility and mild conditions of this reaction enabled azide reduction to be performed on nucleic acid and oligosaccharide substrates without the detectable occurrence of side reactions. The reaction was also performed in the presence of a protein enzyme without loss of enzymatic activity, in contrast with two commonly used azide reduction methods. The visible light dependence of this reaction provides a means of photouncaging functional groups such as amines and carboxylates on biological macromolecules without using UV irradiation.

Reaction discovery provides new tools for chemical synthesis, illuminates new modes of reactivity, and can accelerate the discovery of molecules with desired properties in many fields of science. Traditional approaches to reaction discovery focus on one desired transformation at a time and typically search different reaction conditions for their ability to effect the desired transformation starting with one set of substrates. Recently, several research groups have taken a complementary approach to reaction discovery that searches many substrate combinations for new reactivity under various reaction conditions without focusing on any particular type of transformation.1-3 Our group previously developed a DNA-templated reaction discovery system using DNA hybridization to organize substrate pairs in a single solution such that bond formation transfers a biotin group to the DNA template that encodes the identities of the reactive substrates.1 *In vitro* selection, PCR amplification, and DNA microarray analysis subsequently revealed the identities of reactive

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substrate pairs. A second-generation DNA-encoded reaction discovery system expanded the set of compatible reaction conditions to include those that do not support DNA hybridization including organic solvents and elevated temperatures.3 Here we report results from the application of this second-generation system to explore regions of transition metal and reaction condition space, and describe the development of a resulting Ru(II)-catalyzed, visible light-induced azide reduction that is compatible with a wide variety of functional groups, including those present on biological macromolecules.

The reduction of azides to amines is an important transformation in organic synthesis. Readily prepared in regio- and stereoselective processes, azides are widely used as amine precursors.4 Current methods for the reduction of azides to amines include metal hydride reduction,4 catalytic hydrogenation,5 treatment with thiols,6 and the phosphine-based Staudinger reduction,7 as well as additional methods that require harsher conditions.4 While widely used, these methods have significant limitations. In general, azide reduction reactions involving metal hydrides or catalytic hydrogenation cannot be used on substrates containing acidic groups such as alcohols and acids, or unsaturated groups such as alkenes, alkynes, and carbonyls.5 Similarly, the Staudinger reduction is not compatible with disulfides and alkyl halides, 8,9 and can require elevated temperature and lengthy reaction times. 7 The limitations of existing azide reduction methods may explain the near absence of examples of azide reduction in biological macromolecules 10 despite the common use of both azides11-14 and amines15-17 as chemical handles in proteins, nucleic acids, and oligosaccharides. In addition to being highly tolerant of a variety of acidic and basic functional groups, a reaction compatible with biological macromolecules must also operate efficiently in neutral aqueous environments.18 The azide reduction discovered and developed in this work exhibits these key features of a biological macromolecule-compatible reaction.

Results and Discussion

Choice of Substrates and Reaction Conditions

The second-generation DNA-encoded reaction discovery system3 requires a pool of $n \times m$ DNA-linked substrate pairs that is assembled from two separate sub-pools of n and m substrates (Fig. 1a). We selected 30 substrates (two groups of 15) comprising 28 distinct functional groups based on two criteria: (i) substrates with high latent energy that also exhibit kinetic stability in unactivated form, and (ii) substrates anticipated to be compatible with DNA and pool assembly conditions (Fig. 1b). Collectively, these substrates represent $15 \times 15 = 225$ potentially reactive substrate pairs. We assembled the system using a modified version of our previously reported protocol starting with 5 nmol of DNA-linked substrates.3 Following enzymatic ligation and primer extension steps, we obtained ~1 nmol of the fully assembled pool of 15×15 pairwise substrate combinations. Because each reaction discovery experiment requires only 0.5 pmol of the substrate pool, this quantity of material is sufficient to assess the reactivity of these substrate combinations under ~2,000 different reaction conditions, collectively representing the evaluation of ~450,000 potential reactions. See the Supplementary Information for detailed procedures used to assemble and characterize the substrate pool.

We performed control experiments to validate the system using the known Cu(I)-catalyzed cycloaddition between alkynes and azides.19 We confirmed the ability of the system to detect alkyne-azide coupling under these conditions in the DNA-encoded library format and verified the formation of expected triazole products using DNA-linked substrates (Figs. 1b-c).

Next we determined regions of reaction condition space that are compatible with the reaction discovery system. Since the second-generation system does not rely on DNA hybridization to organize substrates into pairwise combinations, compatible reaction conditions include any that do not rapidly degrade nucleic acids. Reaction conditions that lead to the chemical modification or partial degradation of DNA are not necessarily excluded given the high sensitivity of PCR amplification and the tolerance of PCR to some types of chemical modifications; indeed, conditions observed to degrade DNA were shown to be compatible with our earlier reaction discovery system.1,3

Conditions that preserved the ability of the remaining DNA to be readily detected following 25 cycles of PCR were considered potentially compatible reaction conditions. These conditions included the presence of transition metals (Fe(II), Fe (III), Co(II), Ni(0), Ni(II), Cu(I), Cu(II), Ru(0), Ru(III), Rh(II), Rh(II), Pd(0), Pd(II), La(III), and Au(III)), common metal ligands (anions, phosphines, dienes, carbon monoxide, and nitrogen-based ligands), organic catalysts (amines and thiazolium salts), common oxidants (benzoquinones and peroxides), and single electron transfer reagents (hypervalent iodine, SmI₂, Mn(OAc)₃, and cerium ammonium nitrate (CAN)). A detailed list of potentially compatible reaction conditions is provided in Supplementary Fig. S2.1.

In Vitro Selection for Bond Formation and DNA Microarray Analysis

We treated 0.5 pmol of the substrate pool with a variety of compatible reaction conditions in 200 μ L of aqueous or organic solvent (Figs. 1b and 2; details are provided in the Methods section).3 After exposure to reaction conditions, the disulfide bonds in the substrate pool members were cleaved with tris(2-carboxyethyl)phosphine (TCEP), such that only DNA templates linked to (and encoding) substrates that have undergone bond formation remain covalently attached to biotin. These biotinylated DNA templates were captured with streptavidin-linked magnetic particles, washed, then eluted by heating in deionized water at 70 °C.20 The eluted DNA was amplified by PCR using fluorophore-linked primers (Cy3 for the samples subjected to reaction conditions and *in vitro* selections, and Cy5 for starting substrate pools), then hybridized in triplicate to DNA microarrays containing all 225 possible DNA sequences complementary to each pool member (Fig. 1a).

We observed a number of spots on the DNA microarrays with reproducibly high Cy3:Cy5 (green:red) ratios suggestive of potential bond-forming reactions between two substrates (Fig. 2). We grouped these initial positives into several categories. Some positives correspond to substrate combinations known to undergo bond formation under the reaction conditions tested (green circles in Fig. 2a). While these positives did not represent new reactions, they provided additional validation that the second-generation reaction discovery system can detect bond-forming reactivity. For example, we observed phenol A4 + malonic ester B11 and alkene A1 + malonic ester B11 when the system was treated with CAN in

methanol (Fig. 2a), consistent with previous reports that CAN mediates oxidative coupling reactions between malonic esters and electron-rich π systems (see Fig. 1b for the structures of substrates A1-A15 and B1-B15).21,22

A second group of positives represented reactions that are potentially new but consistent with known reaction mechanisms. For example, copper (II) chloride resulted in a positive signal corresponding to coupling between indole A3 and pyruvamide B4 (green circle in Fig. 2b). In the flask, treatment of indole 12 and ethyl pyruvate 13 in the presence of CuCl₂ or Cu(OTf)₂ led to the formation of a Friedel-Crafts-type coupling product (14) together with the bis-indole adduct (15) in aqueous solvents, while in organic solvents exclusive bis-indole formation was observed (Fig. 2b).23 We also observed substrate combinations that could be facilitated by a variety of Lewis acids or transition metals (orange circles in Figs. 2a-d). For instance, coupling between indole A3 and aldehyde B6, consistent with a Friedel-Crafts-type reaction (orange circles in Figs. 2b-d), and coupling between azide A8 and norbornene B3, consistent with a cycloaddition (orange circles in Figs. 2a, 2b and 2d), were observed under several tested reaction conditions involving transition metals including Fe(III), Co(II), Ni (II), Cu (II) and Ru(II).24,25

A third category of positives corresponded to potential coupling reactions that were reproducible in the DNA-encoded system but when tested in an intermolecular format did not result in the formation of a coupling product. For example, hypervalent iodine26 resulted in robust positive signals corresponding to the coupling of indole A3 with imidazole B9, as well as the coupling of benzyl alcohol A4 and imidazole B9 (green circles in Fig. 2c). When the corresponding reactions were attempted in flasks using small-molecule substrates, however, only the formation of oxidation byproducts 3-oxoindole 18 and benzaldehyde 20 were observed (Fig. 2c). We speculate that for this category of positives, bond-forming reactivity in the DNA-linked reaction discovery system is dependent in part on the highly dilute, intramolecular format of the system, or on interactions with groups in DNA that are absent when these reactions are explored in the flask.

A fourth category of positives correspond to bond-forming reactions that have not been previously reported, that are not easily predicted based on known reactivity, and that we observed to take place in an intermolecular format with either DNA-linked or non-DNA-linked substrates. For example, when the system was treated with Ru(bpy)₃Cl₂ in the presence of a visible light source (a 26 W compact fluorescent lamp (CFL) placed ~20 cm from the tube containing the substrate library), we observed positives that corresponded to coupling of aryl azide B12 and several functional groups (alkene A1, sulfone A14, phenol A4, and nitrile A13) (green circles in Fig. 2d). Although Ru(bpy)₃Cl₂ as a visible-light photoredox catalyst is well-studied in inorganic chemistry,27,28 its exploration in synthetic organic chemistry is more recent.29-32 Given the emerging importance of visible light-mediated organic transformations and the lack of previous reports of azide reactivity under these conditions, we chose to explore these reactions in greater detail.

Visible Light-Induced Azide Coupling and Reduction Reactions

We first studied the reaction between DNA-linked aryl azides and nitriles. Under conditions similar to those used in the DNA-encoded system (5 mM Ru(bpy)₃Cl₂ in 200 μ L of 1:9

water:acetonitrile with 30 mM Na₂CO₃ 20 cm from a 26 W CFL bulb), DNA-linked aryl azide **21** reacted with acetonitrile to generate a product consistent by LC/MS analysis with imidate adduct **22** (Fig. 2d). No significant product formation was observed in the absence of ruthenium complex or in the absence of a CFL light source.

Since bond formation between aryl azides and nitriles induced by visible light catalysis was unprecendented, we developed a mechanistic hypothesis to explain the above observations (Fig. 3a). Ground-state $[Ru(bpy)_3]^{2+}$ (23) is known to absorb visible light ($\lambda_{max} = 452$ nm) to generate the excited triplet $[Ru(bpy)_3]^{2+*}$ (24).27,28 When an electron donor is present, 24 can be reductively quenched to $[Ru(bpy)_3]^+$ (25).30,31,33 We speculate that 25 may reduce azide 26 through a one-electron transfer to generate azide radical anion 27. Following nitrogen gas extrusion and protonation, the resulting aminyl radical 28 may undergo a radical addition to the nitrile.34,35

With this mechanistic hypothesis in mind, we further studied these reactions in a non-DNA-linked format on a 0.1 mmol scale (Table 1). While the conditions that generated the coupling product of the DNA-linked azide and nitrile for unknown reasons did not lead to significant coupling product formation with azide **4** in the flask, the addition of the commonly used sacrificial electron donor diisopropylethylamine30,31 to these reaction conditions resulted in moderate yields (~60%) of the azide reduction product (**31**) after 96 h (Table 1, entry 1). The amine can form through formal hydrogen abstraction by the proposed aminyl radical intermediate (Fig. 3a),35,36 although future studies are needed to probe these and other mechanistic possibilities. We speculated that this reaction may offer significant advantages over existing azide reduction methods and sought to further develop the transformation.

Optimization of a Visible Light-Induced Azide Reduction Reaction

While the initial azide reduction reaction proceeded slowly and in moderate yields, the addition of formic acid or the Hantzsch ester (diethyl 1,4-dihydro-2,6-dimethyl-3,5-pyridinedicarboxylate) as hydrogen donors, supplementing or replacing tertiary amines as hydrogen donors,31 decreased the reaction time to 12 h and increased yields to ~90% (Table 1, entries 2 and 3). Without the ruthenium complex or the CFL light source, no azide reduction was observed under the above conditions (Table 1, entries 4 and 5). Further optimization revealed that the tertiary ammonium salt provided a neutral reaction media and when combined with the Hantzsch ester improved the reaction efficiency to 92-96% yields in 2 h in a variety of solvents (DMF, acetonitrile, or dichloromethane) (Table 1, entries 6-8). The use of acetate rather than formate as the ammonium counterion generated similar excellent product yields in 2 h (Table 1, entry 9). As little as 1 mol% Ru(bpy)₃Cl₂ was sufficient to afford the reduction product in 95% yield in 2 h (Table 1, entry 10).

Next we evaluated the functional group compatibility of the optimized azide reduction under the conditions shown in Table 1, entry 8 (5 mol% $Ru(bpy)_3Cl_2$, 10 equiv. $iPr_2NEt/HCOOH$, 1.5 equiv. Hantzsch ester in CH_2Cl_2). We observed that a variety of substrates containing protic functional groups including free indoles, acids, and alcohols are compatible with the reaction (Fig. 3b, substrates **32-35**). In addition, functional groups sensitive to hydrogenation including alkenes, alkynes, and aryl halides are also not affected by these reaction conditions

(Fig. 3b, substrates **36-39**). Likewise, functional groups sensitive to nucleophiles including alkyl halides, alkyl mesylates, and aldehydes also emerged intact under the reaction conditions (Fig. 3b, substrates **40-42**). An alkyl azide was also reduced in 72% yield in 24 h under modified conditions (Fig. 3b, substrate **43**).

Encouraged by the remarkable chemoselectivity of this reaction, we sought to develop an aqueous version to explore the possibility that this azide reduction might be applicable to biological molecules. Under the above optimized conditions (Table 1 entry 8) but in 1:1 CH₃CN:H₂O, the efficiency of the reaction was drastically reduced (Table 1, entry 11). Efficient product formation was restored when formic acid was removed, presumably increasing the availability of an unprotonated tertiary amine as a competent electron donor (Table 1, entry 12). The water-soluble reducing agent NADH could replace the Hantzsch ester with only a modest reduction in reaction efficiency (Table 1, entry 13). While effective, we speculated that the above basic conditions may compromise the reaction's compatibility with biological macromolecules. We discovered however that sodium ascorbate can act as both an electron donor and as a hydrogen donor37 without significantly lowering yields or lengthening reaction times (Table 1, entry 14). In the presence of sodium ascorbate the reduction proceeds efficiently in buffers of varying acidity (pH 4.0, pH 7.4, and pH 9.5; see Supplementary Fig. S3.1).

Compatibility of the Azide Reduction Reaction with DNA, Oligosaccharide, and Protein

Previous reports have demonstrated the use of $Ru(bpy)_3Cl_2$ to mediate reactions on protein substrates in crude cell extracts,38 and related complexes have been used as cell-permeable agents to image DNA in living eukaryotic and prokaryotic cells.39 These observations coupled with the excellent functional group tolerance and mild reaction conditions of the azide reduction raised the possibility that this reaction could be performed on functional group-rich biomolecules without inducing any side reactions. We tested this possibility by performing the reduction on, or in the presence of, nucleic acids, a protein, and an oligosaccharide and characterizing these macromolecules after each reaction.

We prepared DNA oligonucleotide **44** containing both an aryl azide and a disulfide and treated $0.5~\mu\text{M}$ of this substrate with 1 mM Ru(bpy)₃Cl₂ and 50 mM of sodium ascorbate in aqueous 200 mM Tris, pH 7.4 positioned 20 cm from a 26 W CFL bulb (standard aqueous reduction conditions). Azide reduction proceeded cleanly in 10 min without disulfide bond reduction (eq. 1 in Fig. 4a). In the absence of light, no azide reduction was observed after 15 h, and exposure to light after 15 h of darkness resulted in complete azide reduction in 10 min. In comparison, tris(2-carboxyethyl)phosphine (TCEP), a common Staudinger reaction agent, reduced both the azide and disulfide non-selectively (eq. 2 in Fig. 4a). Likewise, DNA oligonucleotide **47** containing an alkyl azide and a disulfide underwent selective azide reduction in 2 h when treated with 5 mM Ru(bpy)₃Cl₂ and 100 mM ascorbate (eq. 3 in Fig. 4a).

To test the compatibility of the azide reduction reaction with oligosaccharide substrates we prepared an azide-containing variant of naringin, a flavanone disaccharide (49) (Fig. 4b). When $100 \, \mu \text{M}$ of 49 was treated with standard aqueous reduction conditions, we observed

complete reduction in 30 min with no other transformations detected by HPLC, ¹H-NMR, or mass spectrometry (Fig. 4b).

Finally, we performed the azide reduction in the presence of a protein enzyme. We treated a solution containing $100 \,\mu\text{M}$ of naringin azide (49) and $10 \,\mu\text{M}$ of RNase A with standard aqueous reduction conditions. After 6 h, azide 49 was completely reduced by HPLC (Fig. 4c). We isolated the RNase A from the completed reaction by size-exclusion chromatography. No new species consistent with covalent modification of RNase A were observed by ESI mass spectrometry of the macromolecular fraction. Likewise, enzymatic assay of RNase A activity, which requires intact disulfide bonds,39 revealed 10% loss of catalytic activity from the azide reduction reaction developed in this work, while thiol-based reduction and the Staudinger azide reduction both resulted in the loss of 85-100% of enzyme activity (Fig. 4c).

Visible Light-Triggered Photouncaging of Carboxylic Acids

We hypothesized that the reaction described above may also enable azide-protected functional groups other than amines on functional group-rich biomolecules to be uncaged efficiently with visible light under mild conditions. Current photouncaging 40,41 reactions typically require UV light that can induce undesired side reactions including protein degradation, photobleaching, inhibition of photosynthesis, and nucleic acid damage.42

We prepared a DNA oligonucleotide-linked carboxylic acid caged as a 4-azidobenzyl ester 10,43 (52) and treated it with standard aqueous reduction conditions. Photouncaging of the free carboxylic acid through azide reduction and 1,6-elimination was complete within 10 min without any detected byproducts by LC/MS (Fig. 4d). As a deprotection strategy, this reaction proceeds rapidly under very mild conditions and is orthogonal to most currently used carboxyl deprotection strategies such as acid or base hydrolysis.44

Conclusion

Using a second-generation DNA-encoded reaction discovery system, we explored a broad range of transition metals and reaction conditions to identify combinations of substrates and reaction conditions that result in bond formation between substrates. From the reactivity revealed by resulting hits we developed a Ru(II)-catalyzed, visible light-induced azide reduction reaction that is efficient and can be conducted in organic or aqueous solvents, open to the air, at room temperature, and under neutral conditions. This reaction exhibits remarkable chemoselectivity in contrast with existing azide reduction methods and is compatible with alcohols, phenols, acids, alkenes, alkynes, aldehydes, alkyl halides, mesylates, and disulfides. The unusual functional group compatibility and mild required conditions of the reaction enabled azide reduction to be performed on oligonucleotide and oligosaccharide substrates, and in the presence of a protein enzyme, without compromising the structure or the enzymatic activity of the biomolecules. This reaction can also be used to photouncage groups such as amines and carboxylates on biomolecules without requiring UV light.

Methods

See the Supplementary Information for additional experimental details.

General Reaction Discovery Procedure

Each reaction discovery experiment was performed in 200 μ L total volume containing 0.5 pmol of the total substrate pool corresponding to 2.2 fmol of each unique substrate combination. Following exposure to the reaction conditions shown in Fig. 2, each solution was precipitated with ethanol to recover DNA-linked species. The pellet was dissolved in 30 μ L of water and 150 μ L of 0.1 M tris(2-carboxyethyl)phosphine hydrochloride (TCEP) in 1.0 M aqueous sodium phosphate, pH 8.0 was added to effect disulfide cleavage. After 30 min at 25 °C, this solution was combined with streptavidin-linked magnetic particles (14 μ L, corresponding to 20 pmol biotin-binding capacity, Roche Biosciences) suspended in 300 μ L of 10 mM Tris-Cl, 0.1 M NaCl, 1 mM EDTA, pH 7.5. After incubation 15 min at 25 °C, the supernatant was removed and the particles were captured by a magnetic separator, then rinsed once with 200 μ L 10 mM Tris-Cl, 1 M NaCl, 1 mM EDTA, pH 7.5, and once with 200 μ L H₂O. The particles were suspended in 40 μ L deionized water and incubated at 70 °C for 5 min to elute captured DNA.20 The supernatant was collected and the elution step was repeated once. The combined supernatants were used directly in PCR reactions for microarray analysis as previously described3 and detailed in Supplementary Information.

Representative Small-Molecule Azide Reductions

For aryl azides in organic solvent: to a solution of Ru(bpy) $_3$ Cl $_2$ (7.5 mg, 0.01 mmol, 0.05 equiv.), Hantzsch ester (76 mg, 0.3 mmol, 1.5 equiv.), iPr $_2$ NEt (350 μ L, 2.0 mmol, 10 equiv.) and HCOOH (75 μ L, 2.0 mmol, 10 equiv.) in 1 mL CH $_2$ Cl $_2$ was added azide 4 (43.6 mg, 0.2 mmol, 1.0 equiv.). The dark orange solution was stirred at 25 °C at a distance of 20 cm from a 26 W CFL bulb until TLC indicated the complete consumption of azide 4 (usually 2 h). The reaction mixture was directly subjected to silica gel chromatography. Isolated yields for other aryl azides are shown in Fig. 3b.

For aryl azides in mixed aqueous/organic solvent: to a solution of $Ru(bpy)_3Cl_2$ (3.8 mg, 0.005 mmol, 0.05 equiv.) and azide **4** (21.8 mg, 0.1 mmol, 1.0 equiv.) in 1 mL DMF/H₂O (1:1 ratio) was added sodium ascorbate (198 mg, 1.0 mmol, 10 equiv.) in two equal portions (the second portion was added after 24 h). The orange solution was stirred at 25 °C at a distance of 20 cm from a 26 W CFL bulb until TLC indicated the complete consumption of azide **4** (~48 h). The resulting reaction mixture was diluted with ethyl acetate, washed with water and brine, dried over Na_2SO_4 , concentrated in *vacuo* and subjected to silica gel chromatography.

For alkyl azides: To a solution of Ru(bpy)₃Cl₂ (15 mg, 0.02 mmol), Hantzsch ester (102 mg, 0.4 mmol) and iPr₂NEt (105 μ L, 0.6 mmol) in 2 mL DMF/MeOH (3:1 ratio) was added alkyl azide **43** (37.2 mg, 0.2 mmol). The dark orange suspension was stirred at 25 °C at a distance of 20 cm from a 26 W CFL bulb until TLC indicated the complete consumption of azide **43** (~24 h). To the reaction mixture was added di-*tert*-butyl dicarbonate (87.3 mg, 0.4 mmol) and triethylamine (56 μ L, 0.4 mmol) and stirred for 12 h. The resulting solution was

diluted with ethyl acetate, washed with water and brine, dried over Na₂SO₄, concentrated in *vacuo* and subjected to silica gel chromatography.

Representative DNA-Linked Azide Reduction

To a solution of 1 mM Ru(bpy) $_3$ Cl $_2$ and 50 mM sodium ascorbate in 200 μ L of aqueous 200 mM Tris-Cl, pH 7.4 was added DNA-linked azide to a final concentration of 0.5 μ M. After incubation at 25 °C at a distance of 20 cm from a 26 W CFL bulb for 10 min, the reaction was quenched by precipitation with ethanol. The pellet containing DNA-linked small molecules was dissolved in 20 μ L of 10 mM phosphate buffer, pH 8 and subjected to LC/MS analysis.

Naringin Azide Reduction in the Presence of RNase A

To a solution of 1 mM Ru(bpy) $_3$ Cl $_2$ and 50 mM sodium ascorbate in 200 μ L of aqueous 200 mM Tris-Cl, pH 7.4 was added naringin azide and RNase A to final concentrations of 100 μ M and 10 μ M, respectively. After incubation at 25 °C at a distance of 20 cm from a 26 W CFL bulb for 6 h, RNase A was collected through size-exclusion chromatography (Centri-Sep column, Princeton Separations) and characterized by mass spectrometry. After sample dilution 12-fold, RNase A enzyme activity was assayed by following cytidine 2′,3′-cyclic phosphate hydrolysis spectrophotometrically as previously described.45 The small-molecule fraction from polyethersulfone membrane ultrafiltration (Corning Life Sciences) was directly analyzed by reverse-phase HPLC to assess naringin azide reduction.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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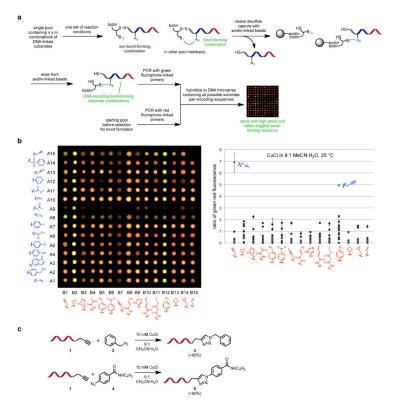


Figure 1.

DNA-encoded reaction discovery system and validation experiments. (a) Selection and analysis method for the detection of bond-forming reactions between DNA-linked small-molecule substrates. (b) Detection of the known Cu(I)-catalyzed cycloaddition19 between alkynes and azides. Conditions: 10 mM CuCl in CH₃CN, 25 °C, 10 min. The green:red ratio of each spot on the DNA microarray (left) is plotted on the right. Error bars represent the standard deviation of signals from three microarray replicates. (c) The reaction products of DNA-linked alkynes and non-DNA-linked azides are consistent with the anticipated triazole products. Conversion efficiencies estimated by LC/MS analysis are shown in parentheses.

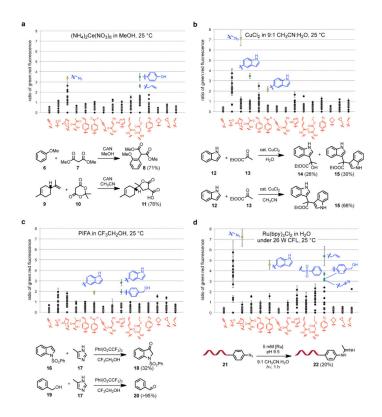


Figure 2. Selection results from four reaction conditions. The scatter plots show green:red fluorescence ratios of substrate combinations in the DNA-encoded library. The schemes show the outcomes of reactions in a flask. (a) In the presence of CAN coupling was observed between malonic esters and electron-rich π systems. 21,22 Conditions: 10 mM CAN in MeOH, 25 °C, 2 h. (b) CuCl₂ induced coupling of indoles and pyruvates, consistent with Friedel-Crafts-type coupling. Conditions: 1 mM CuCl₂ in H₂O, 25 °C, 16 h. (c) Phenyliodine bis(trifluoroacetate) (PIFA) generated positives consistent with imidazole-indole and imidazole-benzyl alcohol coupling, but in a non-DNA-linked format only substrate oxidation was observed. Conditions: 10 mM PIFA in CF₃CH₂OH, 25 °C, 2 h. (d) Ru(bpy)₃Cl₂ under visible light irradiation resulted in several positives including azide-nitrile coupling. Conditions: 10 mM Ru(bpy)₃Cl₂ in 100 mM aqueous sodium carbonate, pH 9.5, 20 cm from a 26 W CFL bulb, 25 °C, 1 h. For (a), (b), and (c), isolated yields are shown in parentheses; for (d), the conversion efficiency by LC/MS analysis is shown in parentheses.

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$$[Ru(bpy)_3]^{2+} = [Ru(bpy)_3]^{2+} = [Ru(bpy)$$

Figure 3. Development of the $Ru(bpy)_3Cl_2$ -mediated, visible light-induced azide reduction reaction. (a) Proposed mechanism for the azide-nitrile coupling reaction and the azide reduction reaction. (b) Substrate scope of the azide reduction reaction under the conditions shown in Table 1, entry 8 (except for substrate 43, which was reduced with iPr_2NEt and Hantzsch ester in 24 h using conditions described in the Methods section). Isolated yields are shown in parentheses.

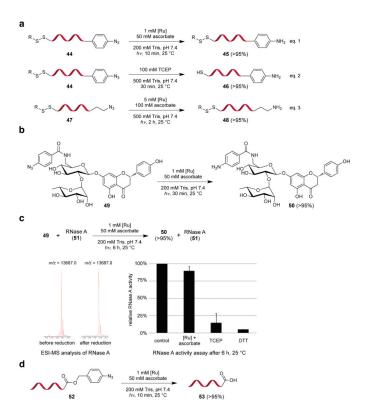


Figure 4.

Compatibility of the azide reduction reaction with biological molecules. Conversion efficiencies are shown in parentheses. (a) Azide- and disulfide-linked DNA oligonucleotides $(0.5 \,\mu\text{M}\text{ each})$ were reduced to the corresponding amines under the aqueous reaction conditions shown without inducing side reactions such as disulfide reduction. In contrast, TCEP reduced both the azide and disulfide groups. Reaction conversions were calculated by LC/MS analysis. (b) A functional group-rich azide analog of the disaccharide naringin (100 μM) was reduced cleanly to the corresponding amine. Conversion was calculated by HPLC analysis. (c) Ru(bpy)₃Cl₂-mediated, visible light-induced reduction of the naringin azide analog (100 μ M) in the presence of the protein enzyme RNase A (10 μ M in the reaction, $0.83 \,\mu\text{M}$ when assayed) did not alter the enzyme's covalent structure or activity. In contrast, thiol- or phosphine-induced azide reduction both resulted in loss of RNase A activity. Conditions are as follows. Control: 200 mM Tris, pH 7.4, 20 cm from 26 W CFL, 6 h, 25 °C. [Ru]+ascorbate: 1 mM [Ru], 50 mM ascorbate, 200 mM Tris, pH 7.4, 20 cm from 26 W CFL, 6 h, 25 °C. TCEP: 100 mM TCEP in 1 M phosphate, pH 8.0, 6 h, 25 °C. DTT: 500 mM DTT in 1 M phosphate, pH 8.0, 6 h, 25 °C. (d) Visible light-triggered photouncaging of carboxylic acid functional group in oligonucleotide substrates. Conversion was calculated by LC/MS analysis.

Table 1

Optimization of t

5% are isolated yields.	O NHC ₄ H ₉
the azide reduction reaction. Reported yields above 5% are isolated yields.	♠
the azide reduction reaction	NHC ₄ H ₉

entry	reaction conditions	%[Ru]	solvent	time	yield
1	2 equiv. iPr_2NEt , $h\nu$	2%	DMF	4 96 h	%09
2	10 equiv. iPr ₂ NEt/HCOOH, $h\nu$	2%	CH ₃ CN	12 h	%06
8	2 equiv. iPr ₂ NEt, 1.5 equiv. Hantzsch ester, hv	2%	CH ₃ CN	12 h	%68
4	10 equiv. iPr ₂ NEt/HCOOH, no light	2%	CH ₃ CN	48 h	<5%
S	10 equiv. iPr ₂ NE $t/HCOOH$, $\hbar \nu$	none	CH ₃ CN	70 d	<5%
9	10 equiv. iPr ₂ NEt/HCOOH, 1.5 equiv. Hantzsch ester, hv	2%	CH ₃ CN	2 h	92%
7	10 equiv. iPr ₂ NEt/HCOOH, 1.5 equiv. Hantzsch ester, hv	2%	DMF	2 h	92%
~	10 equiv. iPr ₂ NEt/HCOOH, 1.5 equiv. Hantzsch ester, hv	2%	$\mathrm{CH_2Cl_2}$	2 h	%96
6	10 equiv. iPr ₂ NEt/CH ₃ COOH, 1.5 equiv. Hantzsch ester, hv	2%	$\mathrm{CH_2Cl_2}$	2 h	%96
10	10 equiv. iPr ₂ NEt /HCOOH, 1.5 equiv. Hantzsch ester, $h\nu$	1%	$\mathrm{CH_2Cl_2}$	2 h	%56
111	10 equiv. iPr ₂ NEt/HCOOH, 1.5 equiv. Hantzsch ester, <i>hv</i>	2%	1:1 CH ₃ CN:H ₂ O	108 h	10%
12	2 equiv. iPr ₂ NEt, 1.5 equiv. Hantzsch ester, $h\nu$	2%	1:1 CH ₃ CN:H ₂ O	48 h	91%
13	2 equiv. iPr ₂ NEt, 1.5 equiv. NADH, $h\nu$	2%	1:1 CH ₃ CN:H ₂ O	4 09	64%
14	10 equiv. sodium ascorbate, hv	2%	1:1 DMF:H ₂ O	48 h	%06