Nucleic Acid Template-Directed Assembly of Metallosalen-DNA Conjugates

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Watson-Crick base pairing organizes DNA duplex formation necessary for genetic information storage in biological systems. DNA and RNA templates also direct the specific binding of nucleotide substrates during diverse enzyme-catalyzed reactions in replication, transcription, and DNA repair pathways. Recently, nucleic acid recognition properties have been extended to nonbiological systems, where DNA base pairing has been used to drive the template-directed chemical ligation of oligonucleotides,¹ and the assembly of nanostructures and novel materials.²

We have been interested in expanding the versatility of nucleic acid base pairing for the addressable synthesis of bioconjugates in aqueous solution. Metallosalen-DNA (4, Scheme 1) represents an ideal system to demonstrate the concept of nucleic acid template-directed molecular synthesis. Salens, which are constructed from two salicylaldehydes and a diamine, serve as ligands for a broad range of metal ions. Many metallosalens are compatible with aqueous conditions³ and have demonstrated utility as DNA cleavage reagents⁴ and versatile catalysts for enantioselective transformations.^{3a,5} Template-directed synthesis of metallosalen-DNA conjugates offers a unique approach to a new class of metal-DNA hybrids. Metal-DNA conjugates previously have been employed as probes of DNA structure and electron transfer,⁶ "chemical nucleases" for targeted nucleic acid cleavage,⁷ and

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Scheme 1. Template-Directed Metallosalen–DNA Assembly^a



^{*a*} **3a** (DNA), SPACER = TT or **3b** (RNA), SPACER = UU.

Scheme 2^a



^a Key: (a) HO(CH₂)₃OH, acidic alumina, PhCH₃, reflux; (b) NaOH, BzCl, THF; (c) [(i-Pr)₂N]₂POCH₂CH₂CN, (i-Pr)₂NH₂•CHN₄, CH₂Cl₂.

scaffolds for metal-mediated base pairing motifs.8 Thus, metallosalen-DNA may offer a new bioconjugate platform for DNAorganized materials, nucleic acid cleavage and detection strategies, and in vitro evolution of novel ribozymes and deoxyribozymes.9

Our approach to template-directed synthesis of metallosalen-DNA is illustrated in Scheme 1. The DNA-metallosalen building blocks consist of two DNA oligonucleotides modified, at either the 3' or 5' end, with salicylaldehyde moieties (1 and 2). The modified strands are aligned on a complementary nucleic acid template (3), bringing the salicylaldehyde groups into proximity in a duplex. The metallosalen conjugate then is assembled by addition of an appropriate metal and diamine. Herein we report the efficient DNA and RNA template-directed synthesis and characterization of purified metallosalen-DNA conjugates.

A salicylaldehyde phosphoramidite (8, Scheme 2) was synthesized as a precursor to salicylaldehyde-DNA conjugates 1 and 2, necessary for metallosalen-DNA assembly. The protecting groups for 8, including a benzoate ester for the phenol and a 1,3dioxane for the aldehyde, were chosen for their compatibility with DNA synthesis and postsynthetic deprotection. Starting from known salicylaldehyde derivative 5,¹⁰ dioxane 6 was prepared by alumina-catalyzed acetalization. Direct benzoylation of 6 with benzovl chloride afforded 7, which was converted to phosphoramidite 8 by standard methods.¹¹

Oligonucleotide 2 was synthesized by DNA phosphoramidite chemistry (3'-to-5'), using 8 in the final coupling step. Oligonucleotide 1, bearing a 3'-terminal salicylaldehyde, was produced by 5'-to-3' DNA synthesis using commercial nucleoside 5'phosphoramidites. Standard DNA deprotection with concentrated ammonia removed the phenolic benzoate group, and subsequent

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Figure 1. Gel electrophoresis assay of metallosalen–DNA assembly. (Lane 1) 10 bp marker. (Lanes 2–6) Mn assembly reactions at pH 8.0 for 1 h. (Lane 2) All components: $Mn(OAc)_2$, **1**, **2**, and **3b**, EN. (Lane 3) –EN. (Lane 4) – $Mn(OAc)_2$. (Lane 5) –**2**. (Lane 6) –**3b**. (Lane 7) Ni(II) assembly reaction with all components at pH 6.5 for 24 h. (Lane 8). Labeled **1**. All reactions included 10 mM buffer (HEPES for Mn and MES for Ni) and 150 mM NaCl.

incubation in 15 mM NaOAc/HOAc buffer (pH 4.0) for 2 h at 37 °C cleaved the dioxane group to the aldehyde. Oligonucleotides 1 and 2 were purified by denaturing polyacrylamide gel electrophoresis (PAGE) or reverse-phase HPLC (RP-HPLC), and their identities were confirmed by matrix-assisted laser desorption time-of-flight (MALDI-TOF) mass spectrometry (for 1: [M]⁻ calcd, 4860.17; found, 4860.86; for 2: [M]⁻ calcd, 4835.15; found, 4835.37).

The DNA template-directed assembly of a Mn-metallosalen-DNA conjugate (Mn-4) is demonstrated in Figure 1. Assembly reactions were monitored by gel electrophoresis using radiolabeled 1 strand as a tracer; metallosalen–DNA formation was reflected in a shift of the labeled strand to lower gel mobility upon conjugation. When DNA 1 and 2 were annealed to template 3a and incubated for 1 h at 37 °C in the presence of 100 μ M ethylenediamine (EN) and 400 µM Mn(OAc)₂, a new DNA complex, corresponding to Mn-4, was produced in \sim 65% yield (Lane 2). Removal of assembly components EN (lane 3) or strand 2 (lane 5) prevented complex formation. In the absence of manganese (lane 4), ~4% maximum yield of unmetalated 4 was observed. Most importantly, DNA template 3a (lane 6) was required for the assembly reaction. While the templated reaction was complete in 1 h, untemplated complex formation was detectable only after 8 h. Taken together, the results in Figure 1 demonstrate that metallosalen-DNA formation is DNA templateand metal ion-dependent.12 Two additional contributors to Mn-4 assembly efficiency were the identity of the template spacer units and the pH of the reaction. Two nucleotide residues (TT) provided an optimal spacer for Mn-4 conjugate formation, consistent with our molecular modeling of DNA-metallosalen duplexes. Optimal yields of Mn-4 assembly were observed at pH 8 but decreased with increasing pH as predicted by metal speciation data and the limited solubility of manganese ions with increasing pH.¹³

Purified metallosalen–DNA conjugate **Mn-4** was characterized by two complementary methods: MALDI-TOF MS and nucleoside composition analysis. Preparative synthesis of metallosalen– DNA (~4 nmol) was performed using a complementary RNA



Figure 2. RP-HPLC analysis of **Ni-4** nucleoside/Ni-salen enzymatic digestion products. dU, deoxyuridine (internal standard). dI, deoxyinosine. *, peak present in the digestion blank.

template, **3b**.¹⁴ Following **Mn-4** assembly with **3b**, selective RNase H digestion of the RNA template strand of the DNA/RNA hybrid facilitated the purification of **Mn-4**. After isolation, **Mn-4** was stable in buffered aqueous solutions for several days at 25 °C. MALDI-TOF characterization of purified **Mn-4** provided the mass expected for metalated **Mn-4** ([M]⁻ calcd, 9773.35; found, 9772.83). To verify the base and salen composition, **Mn-4** was subjected to nuclease digestion and quantitative RP-HPLC analysis.¹⁵ The component nucleosides and salen ligand were present in the ratios expected for **Mn-4**.¹⁶

Metallosalen–DNA conjugates containing Ni(II) (Ni-4) were assembled by an analogous template-directed strategy. Ni-4 was synthesized using 2 μ M each of 1, 2, and 3a (or 3b), 300 μ M Ni(OAc)₂, and 150 μ M EN at pH 6.5. Ni-4 assembly proceeded in 74% yield at 37 °C in 24 h (Figure 1, lane 7).¹⁷ MALDI-TOF MS characterization ([M]⁻ of Ni-4 calcd, 9777.06; found, 9777.64) verified the identity of the metal–DNA conjugate. Ni-4 was further characterized by enzymatic digestion to deoxynucleosides and Ni–salen, which were present in the correct ratios for 4 as shown by quantitative RP-HPLC analysis in Figure 2.

We have demonstrated the efficient nucleic acid templatedirected synthesis and characterization of a new class of metal– DNA conjugates, metallosalen–DNA. The chemical diversity of metallosalen–DNA assembly processes and the applications of these metal–DNA hybrids currently are under investigation. Nucleic acid template-directed molecular synthesis offers a powerful approach for the addressable synthesis of new DNA bioconjugates, which may offer significant potential for applications including targeted nucleic acid cleavage, biosensors, and catalysis.

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Supporting Information Available: Experimental procedures, spectroscopic and analytical data for compounds 1-8, HPLC and MALDI-TOF data for compounds 1, 2, and 4, base composition data for 4 (PDF). This material is available free of charge via the Internet at http://pubs.acs.org.

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