Evaluation of Metal-Mediated DNA Binding of Benzoxazole Ligands by Electrospray Ionization Mass Spectrometry

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The binding of a series of benzoxazole analogs with different amide- and ester-linked side chains to duplex DNA in the absence and presence of divalent metal cations is examined. All ligands were found to form complexes with Ni2+, Cu2+, and Zn2+, with 2:1 ligand/metal cation binding stoichiometries dominating for ligands containing shorter side chains (2, 6, 7, and 8), while 1:1 complexes were the most abundant for ligands with long side chains (9, 10, and 11). Ligand binding with duplex DNA in the absence of metal cations was assessed, and the long side-chain ligands were found to form low abundance complexes with 1:1 ligand/DNA binding stoichiometries. The ligands with the shorter side chains only formed DNA complexes in the presence of metal cations, most notably for 7 and 8 binding to DNA in the presence of Cu2+. The binding of long side-chain ligands was enhanced by Cu2+ and to a lesser degree by Ni2+ and Zn2+. The cytotoxities of all of the ligands against the A549 lung cancer and MCF7 breast cancer cell lines were also examined. The ligands exhibiting the most dramatic metal-enhanced DNA binding also demonstrated the greatest cytotoxic activity. Both 7 and 8 were found to be the most cytotoxic against the A549 lung cancer cell line and 8 demonstrated moderate cytotoxicity against MCF7 breast cancer cells. Metal ions also enhanced the DNA binding of the ligands with the long side chains, especially for 9, which also exhibited the highest level of cytotoxicity of the long side-chain compounds. (J Am Soc Mass Spectrom 2008, 19, 209–218) © 2008 American Society for Mass Spectrometry

The discovery of the selective cytotoxic activity of UK-1 [1–3], a bis(benzoxazole) isolated as a secondary metabolite from Streptomyces, has stimulated the development of other benzoxazole and benzimidazole compounds with similar anticancer activities [4–6]. As a topoisomerase II inhibitor, one of the unique properties of UK-1 is its ability to bind biologically important divalent metal cations [3] and its metal-mediated DNA binding [3, 7]. While UK-1 has demonstrated cytotoxicity against a number of cell lines, it does not inhibit the growth of bacteria, yeast, or fungi [1, 8], making the mechanism of this selective cytotoxic activity and the metal binding properties of UK-1 and new analogs of great interest.

In a recent study, we used electrospray ionization mass spectrometry (ESI-MS) in conjunction with cytotoxicity assays to examine several simple analogs of UK-1 to explore the metal ion binding requirements of these compounds, assess metal-mediated DNA binding and evaluate anticancer and antibacterial activity [6]. Interestingly, the only ligand that exhibited anticancer activity, 2 (Figure 1), was also the only metal-mediated DNA binder with a preference for Ni2+ as determined by ESI-MS experiments. Two other compounds, 4 and 5, formed complexes with DNA in a nonspecific, nonmetal-mediated manner, and showed antibacterial but not cytotoxic behavior. These results suggested a correlation between metal-mediated binding and anticancer activity of the compounds. To improve upon the solubility of the benzoxazoles in aqueous solutions and to allow further examination of the metal-mediated DNA binding and anticancer activity, a series of analogs of 2 have been synthesized with different ester and amide-linked side chains (Figure 1) [9].

In the present study, ESI-MS was used to screen the binding of the new ligands to a series of divalent metals including Mg2+, Ni2+, Cu2+, and Zn2+. Ligand binding to duplex DNA in the presence and absence of metal ions was also assessed. ESI-MS has been shown to be a useful tool for the analysis of noncovalent ligand/DNA

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complexes due to its sensitivity, low sample consumption, and fast analysis times that make it amenable to high throughput screening [10-12]. Early studies focused on examining well-studied, commercially available duplex DNA-binding compounds including both minor groove binders [13-16] and intercalators [17-19] to establish that the binding stoichiometries, selectivities, and specificities observed by ESI-MS correlate with known solution behavior. More recent studies have extended the use of the technique to novel compounds, including studies by our group examining the metal-dependent binding of UK-1 and related benzoxazoles and benzimidazole analogs [6, 7]. Our previous studies have demonstrated that one of the main advantages of a mass spectrometry-based analysis technique for metal-mediated DNA binding ligands over traditional spectroscopic techniques is that information about ligand/metal and ligand/metal/DNA binding stoichiometries is obtained. Moreover, typical solution methods to study the interaction of small molecules with DNA include absorption and fluorescence spectroscopy, yet the application of these methods to the types of metal ion ligands described here is made difficult by the complex equilibria involved and the effects of metal ions. The changes in UV/Vis absorbance spectra for UK-1 and analogs upon DNA binding are subtle compared to those due to metal ion binding. While UK-1 and related 2-hydroxyphenyl benzoxazoles are fluorescent, this fluorescence is quenched upon metal ion binding, which prohibits the use of the intrinsic fluorescence of these compounds in studies of their metal-mediated DNA binding. Displacement-based DNA binding assays employing DNA-binding dyes such as ethidium bromide are also precluded in the study of the metal-mediated DNA binding of UK-1 and analogs due to the quenching of the dye’s fluorescence by various transition metals of interest. Thus, a general and rapid alternative method, such as ESI-MS, for studying the metal-mediated DNA binding interactions of these compounds is needed.

For the present study, anticancer cytotoxicity assays were also done so that the activities of these analogs could be correlated to the metal binding behavior elucidated in the mass spectrometry studies and compared with other previously reported UK-1 analogs. Previous studies had identified 2 as a simplified analog of UK-1 that retains the selective cytotoxicity of the natural product [4]. The analogs of 2 examined here were designed to increase the potential for metal ion complexation and metal-mediated DNA binding by modifying the ester side chain of 2, especially through the incorporation of additional sites for metal ion coordination.

**Experimental**

The syntheses of 2 and 12 have been previously reported [4, 20]. The synthesis of the benzoxazole analogs 6, 7, 8, 9, 10, and 11 will be reported separately [9]. The oligodeoxynucleotides (ODNs) used in this study, d(GCGGGATGGGCG), d(CGCCCCATCCCCGC),
d(GCGGGAATTGGCG), d(CGCCCAATTCCGC), d(GCGGAAATTTCG), and d(CGCCAAAATTCG) were purchased from IDT Technologies (Coralville, IA) as ammonium salts. Duplex DNA was annealed by preparing equimolar (1 mM) concentrations of the non-self-complementary ODNs in 250 mM ammonium acetate. Concentrations were verified spectroscopically using Beer’s law and the extinction coefficients for the DNA strands provided by the manufacturer. The solutions were heated to 90 °C and allowed to cool to room-temperature over 7 h. Analytical solutions containing a ligand and metal salt, a ligand and DNA duplex, or a ligand, metal salt, and DNA duplex were prepared at equimolar (10 μM) concentrations unless noted otherwise, in 50 mM ammonium acetate solutions with 50% methanol.

The samples were directly infused at 3 μL/min into a Thermo Electron (San Jose, CA) LCQ mass spectrometer. For the DNA binding experiments, the instrument was operated in the negative ion mode with an electrospray voltage of 3.5 kV and a heated capillary temperature of 90 to 110 °C with sheath and auxiliary gas flows of 40 and 10 arbitrary units, respectively. Ligand/metabolion ion solutions were examined in the positive ion mode using an electrospray voltage of 4.5 kV and the same heated capillary and gas flow rates used for the solutions containing DNA.

Cytotoxicity was determined using the AlamarBlue cell viability assay as described previously [4]. Briefly, aliquots of 100 μL cell suspension (1–3 × 10^3 cells) were placed in microtiter plates in an atmosphere of 5% CO₂ at 37 °C. After 24 h, 100 μL of culture media and 2 μL of the compound in DMSO were added to each well in duplicate, and the plates incubated an additional 72 h at 37 °C. Compounds, along with MitoXyn-C as a positive control, were evaluated at final concentrations ranging from 0.001 to 50 μM. Cell viability was determined by removing the culture media from each well, and adding 200 μL of fresh media and 20 μL of AlamarBlue reagent (Biosource, Camarillo, CA), followed by an additional 6 h incubation before fluorescence measurement. Fluorescence was measured by a Beckman Coulter DTX880 plate reader with excitation at 530 nm and emission at 590 nm. The percent growth was calculated from the fluorescence data using the eq:

\[ \text{% Growth} = 100 \times \frac{F_t - F_0}{F_c - F_0} \]  

(1)

where \( F_t \) is the averaged measured fluorescent intensities of AlamarBlue reagent at the time just before the exposure of the cells to the test substance, \( F_c \) is the averaged measured fluorescent intensities of AlamarBlue reagent after 72 h exposure of the cells to the test substance at a particular concentration, and \( F_0 \) is the averaged measured fluorescent intensities of AlamarBlue reagent after 72 h exposure of the cells to the vehicle without the test substance.

The concentration of compound required to inhibit growth by 50% (IC₅₀) was determined by nonlinear regression fitting the %growth data to the eq:

\[ y = \text{Min} + \text{Max} - \text{Min})/(1+10^{[x-\log_{10}(\text{IC}_{50})]} \times H) \]  

(2)

where \( x \) is the compound concentration, \( y \) is the % growth; \( \text{Min} = \) the minimum response plateau (0% Growth); \( \text{Max} = \) the maximum response plateau (100% growth); \( H \) is the Hill slope co-efficient.

Cytotoxicity assays were carried out versus controls in which DMSO but no benzoxazole ligands were added to the cells. The concentration of DMSO employed in these assays does not affect the growth of the cells.

Results and Discussion

Ligand Binding to Metals

The binding of the benzoxazole ligands with metals including Mg²⁺, Ca²⁺, Ni²⁺, Cu²⁺, and Zn²⁺ was screened by ESI-MS to determine the binding preferences of the analogs and identify the metal ions that are most promising for enhancing ligand binding. Solutions containing a metal ion and a ligand were prepared at equimolar concentrations in the same ammonium acetate buffer used for subsequent DNA binding experiments and analyzed in the positive ion mode (Figure 2). No complexes were formed between any of the ligands and either Mg²⁺ and Ca²⁺. Figure 2 shows mass spectra for solutions containing Mg²⁺ with 8 (Figure 2a) and 9 (Figure 2b), demonstrating the absence of ligand/metal complexes. To ensure that the binding between the ligands and these metals was not concentration-dependant, the metal ion concentration was increased 10-fold to 100 μM; however, no complexation was observed.

These results appear at first to contradict the results of previous studies in which the analogs of UK-1, including 2, were found to form complexes with Mg²⁺ [4]. However, the prior binding studies were undertaken in a solvent containing 100% methanol due to the low solubility of the early analogs in aqueous buffers [4]. When methanol was used as a solvent in the current ESI-MS experiments, 2, 6, 7, 8, 9, 10, and 11 formed abundant complexes with Mg²⁺ (spectra not shown). These results demonstrate the importance of the solvent in the metal ion binding of the benzoxazoles, presumably due to the differences in metal ion solvation in different solvents. To maintain consistency with the buffer used for the DNA binding experiments, the 50 mM ammonium acetate buffer with 50% methanol (vol/vol) was used for the remaining metal ion binding experiments.

Ligands 2, 6, 7, 8, 9, 10, and 11 formed complexes with the divalent transition metals Ni²⁺, Cu²⁺, and to a lesser degree, Zn²⁺. Typical ligand/metal ion binding stoichiometries ranged from 1:1 to 2:2, with the 2:2 complexes most likely being dimers of the 1:1 complexes. Ligands with the shorter side chains, 6, 7, and 8
formed abundant 2:1 complexes, as shown in Figure 2c for a solution containing 8 with Cu²⁺. The short side-chain ligands were also capable of forming 1:1 complexes, however a water adduct was always bound to the complex, conceivably to fill the coordination shell of the metal. The long side-chain compounds, 9, 10, and 11, were able to form abundant 1:1 complexes without a solvent adduct as demonstrated by the spectrum of 9 with Cu²⁺ shown in Figure 2d. Likely, the long side chain is able to wrap partially around the metal ion upon binding, thereby filling the coordination shell without addition of solvent molecules.

The only benzoxazole ligand that did not form complexes with Ni²⁺, Cu²⁺, and Zn²⁺ was 12 (spectra not shown). This compound is the only analog examined in this study that contains a methyl ester group rather than a hydroxyl group on the phenyl moiety of the compound. The 2-(2’-hydroxyphenyl)benzoxazole moiety is also present in synthetic metal ion chelators [21, 22], and is believed to play a key role in the metal ion chelation of the compounds [3]. The lack of metal ion binding observed by 12 in the ESI-MS experiments further implicates the role of the hydroxyl phenyl group in the metal ion binding of the benzoxazoles and makes 12 a good negative control compound for ligand/metal binding examined by ESI-MS.

The collisionally activated dissociation (CAD) mass spectra of the ligand/metal complexes were also examined by ESI-MS. Upon collisional activation, the 2:1 complexes containing 6, 7, and 8 dissociate via the ejection of a neutral ligand with the rapid addition of a water molecule (always present in the trap in trace amounts) to the resulting 1:1 complex (spectra not shown). This is the same fragmentation pattern reported for complexes containing the anti-cancer analog 2 with Cu²⁺ and Ni²⁺ [6]. When the resulting 1:1 complexes were subjected to a second stage of CAD (MS²), the complexes did not undergo further observable fragmentation. Likely, the attached water molecule is dislodged during collisional activation, but then rapidly reattaches before ion detection. This type of solvent adduction process has been commonly observed for transition-metal complexes in a quadrupole ion trap [23–27]. This was the same fragmentation pattern observed for the dissociation of the watersolvated 1:1 complexes containing 6, 7, and 8 and a transition-metal.

Collisional activation of the 2:1 benzoxazole:metal complexes that contained the ligands with longer side chains, 9, 10, and 11, showed dissociation via the loss of one ligand, leaving 1:1 complexes (spectra not shown). Upon MS², these 1:1 complexes produced different fragmentation pathways that were dependent on the ligand. Complexes containing 7 and 9, the ligands with a polyethylene glycol side chain, dissociated via the loss of small portions of the side chain, such as C₂H₄O,
while the metal ion remained bound to the remainder of the ligand. These losses are not observed for 10, which contains a six carbon alkyl chain with a terminal hydroxyl group. Instead, the 1:1 complexes containing 10 dissociate via the loss of the metal ion with the spontaneous addition of methanol or water. In general, these initial ESI-MS experiments not only revealed that Ni$^{2+}$, Cu$^{2+}$, and Zn$^{2+}$ are apparently the favored metals for ligand binding, but also showed differences in the preferred binding stoichiometries and fragmentation patterns for complexes containing ligands with long versus short side chains.

**DNA Binding of the Ligands Without Metals**

In previous ESI-MS studies, benzoxazole and benzimidazole ligands that exhibited antibacterial activity were found to bind to DNA in the absence of metal cations, while compounds exhibiting anticancer activity, notably UK-1 and 2, only formed complexes with duplex DNA in a metal-mediated manner. Therefore, the DNA binding of the new analogs of 2 (Figure 1) in the absence of metal cations was of great interest and was examined by ESI-MS in the present study. Three non-selfcomplementary DNA duplexes, d(GCGGGATGGGCG/CGCCCCATCCC), d(GCGGAAATTTGGCG/CGCCAAATTTCGC), and d(GCGGAAATTTGGCG/CGCCAAATTTCGC), were used for the initial screening study. The duplex sequences were selected to have different degrees of G/C and A/T base pair composition to investigate possible sequence preferences of the ligands. Solutions containing one ligand and one duplex at equimolar (10 μM) concentration were prepared in 50 mM ammonium acetate with 50% methanol. The high methanol composition was necessary to ensure the ligands remained soluble in the analytical solutions. To allow for comparison of the DNA binding of the new analogs with 2, experiments involving 2 were included in the present study to maintain consistency with the earlier study [6].

As demonstrated by the series of spectra shown in Figure 3 for solutions containing one of six ligands with d(GCGGAAATTTGGCG/CGCCAAATTTCGC), three of the ligands, 9 (Figure 3d), 10 (Figure 3e), and 11 (Figure 3f) were found to form low abundance complexes with 1:1 ligand/DNA binding stoichiometries, while 6 (Figure 3a), 7 (Figure 3b) and 8 (Figure 3c) did not form any complexes detectable by ESI-MS. As expected, 2 likewise did not bind to DNA in the absence of metal ions (spectra not shown). The binding results of the ligands with the other two duplexes, d(GCGGGATGGGCG/CGCCCCATCCC) and d(GCGGAAATTTGGCG/CGCCAAATTTCGC), were simi-

**Figure 3.** ESI-mass spectra of solutions containing duplex 3/4 and (a) 6, (b) 7, (c) 8, (d) 11, (e) 9, and (f) 10. Solutions contained a ligand, metal salt, and DNA duplex at equimolar (10 μM) concentrations in 50 mM ammonium acetate solutions with 50% methanol.
lar to those shown in Figure 3, suggesting that these ligands do not have significant sequence selectivities. The ligands that do bind to the duplex all contain ester- or amide-linked side chains that are longer than those compounds that did not bind to duplex DNA, suggesting the side chains could play a role in promoting nonmetal-mediated DNA binding of the benzoazoxoles.

The nonmetal-mediated binding of 9, 10, and 11 differs from that of the antibacterial ligands 4 and 5 examined in our previous study. The complexes formed by 4 and 5 had greater relative abundances, and the binding stoichiometries ranged from 1:1 to 3:1 and were highly dependent on ligand concentration [6]. The binding behavior of ligands 4 and 5 was similar to that of commercial intercalators examined in previous ESI-MS studies [17, 28]. While 9, 10, and 11 formed complexes with duplexes without metals, the binding stoichiometries never exceed 1:1, and their relative abundances are low.

DNA Binding of the Ligands with Metals

Metal cations are documented to play a key role in the DNA binding of UK-1 and 2; the 1:1 binding of divalent metal ions by these ligands can result in cationic complexes with increased electrostatic attraction towards DNA. In analogy with the magnesium-dependent DNA binding by quinobenzoazoxines [29], the metal ion-mediated binding of UK-1 and analogs with DNA may involve shared coordination of the ligand and phosphate groups on the DNA with the metal ion. This metal-mediated binding behavior is thought to be related to the anti-cancer activity of these ligands over other nonmetal-mediated benzoazole and benzimidazole compounds. Previous studies of the solution metal ion binding ability of UK-1 and analogs had indicated that Mg$^{2+}$ and Zn$^{2+}$ were implicated in the metal-mediated DNA binding by these compounds [4], while in ESI-MS studies Ni$^{2+}$, Co$^{2+}$, and Zn$^{2+}$ promoted the greatest duplex binding for UK-1 [7], and the duplex binding of 2 was found to be mediated by Ni$^{2+}$ [6].

To determine if the binding of the new benzoazoxole ligands is also metal-mediated, ESI-MS experiments were undertaken for solutions containing equimolar (10 μM) concentrations of a benzoazole ligand, duplex d(GGGGAAATTGGCC/GGCCCAATTCCGCG), and a metal salt. Based on the first section of results described above, only the metals that were found to form complexes with the ligands, Ni$^{2+}$, Cu$^{2+}$, and Zn$^{2+}$, were used in this phase of the study. The most significant enhancement in duplex/ligand binding was observed with Cu$^{2+}$ as demonstrated by the spectra shown in Figure 4, which correspond to solutions containing Cu$^{2+}$, the duplex and either 6 (Figure 4a), 7 (Figure 4b), or 8 (Figure 4c). While none of these three ligands bind DNA in the absence of metal ions (i.e., Figure 3a, b, and c), complexation is observed for all three in the presence of copper, with the most significant degree of binding observed for 7 and 8. Complexes containing ligand/metal/DNA binding stoichiometries of 1:1:1 and 2:2:1 were observed for 7 and 8, while only low abundance 1:1:1 were formed with 6. Similar results were obtained for 2 (spectra not shown). The ions that are not labeled in Figure 4 correspond to sodium adducts of the DNA and ligand/Cu$^{2+}$/DNA complexes, with the source of sodium contamination likely being the benzoazoxole solutions.

While 9, 10, and 11 were found to bind to duplex DNA even in the absence of metal cations, their binding was enhanced in the presence of Cu$^{2+}$, most significantly for 9 and 10 (spectra not shown). To compare the changes in the degree of ligand binding upon addition of the metal salt, the fraction of bound DNA values were calculated by expressing the sum of the abundances of all ions attributed to DNA/ligand complexes as a fraction of the total abundances of all DNA-containing ions (both free DNA and DNA/ligand complexes) as has been previously reported for other ligand/DNA complexes [30]. The abundances for all of the sodium adducts associated with each complex were included in the calculation, and only ions in the 5-charge state were used since this was the dominant charge state in the mass spectra. Inclusion of the abundances for the 6-charge state does not change the results. The increase in the extent of binding by 9, 10 and 11 is reflected in the fraction of bound DNA values summarized in Table 1. The results here and summarized in Table 1 demonstrate that Cu$^{2+}$ has the greatest impact on ligand binding, enhancing the binding of 9, 10, and 11, and promoting binding by the other ligands with the most dramatic metal-mediated behavior seen for 7 and 8, and to a lesser degree, 6.

In general, Ni$^{2+}$ had a small impact on ligand binding as demonstrated by the spectra shown in Figure 5. Extensive binding of Ni$^{2+}$ to the duplex DNA is observed in Figure 5, but this does not translate into an enhancement of ligand binding. Ligand 8 was the only new analog that exhibited binding in the presence of Ni$^{2+}$ yet did not bind to DNA without metal ions, as shown in Figure 5c. A low abundance 1:1:1 complex containing 8 is formed (Figure 5c), but with lower abundance relative to the complexes formed by 8 in the presence of Cu$^{2+}$ (Figure 4c). The duplex binding of 9 was also enhanced in the presence of Ni$^{2+}$ (spectra not shown). For 9, the fraction of bound DNA increased from 0.17 in the absence of metal ions to 0.34 in the presence of Ni$^{2+}$ (Table 1). Experiments with 2 confirmed the results of our previous study that found the binding of the ligand to be Ni$^{2+}$-mediated. The fraction of bound DNA values based on the 2/Ni$^{2+}$/DNA complexes was 0.18, which is similar to that of the 8/Ni$^{2+}$/DNA complexes, 0.14.

Figure 5a and b show that neither 6 nor 7 bound to the DNA with Ni$^{2+}$, nor did Ni$^{2+}$ enhance the binding of 10 or 11 (spectra not shown). As summarized in Table 1, the fraction of bound DNA values for 10 and 11 in the absence of metal ions were 0.09 and 0.20, respectively.
While both ligands formed 1:1 ligand/Ni$^{2+}$/DNA complexes, the relative abundances of these complexes were either the same as or lower than the abundances of the complexes formed in the absence of metal ions. This is reflected in the fraction of bound DNA values for solutions containing the ligands with Ni$^{2+}$ and DNA which were calculated to be 0.13 for 10 and 0.08 for 11.

Zn$^{2+}$ did not have a significant impact on the benzoxazole ligand binding to DNA. Zn$^{2+}$ did not promote DNA binding by 6, 7, 8, or 2, the ligands that likewise did not bind to DNA in the absence of metals (spectra not shown). Ligand 10 formed 1:1 ligand/Zn$^{2+}$/DNA complexes, however the fraction of bound DNA value for ligand binding in the presence of zinc was 0.17, which is not significantly different than it was in the absence of metal, 0.20 (spectra not shown). The DNA binding of 9 and 11 was enhanced by Zn$^{2+}$, as indicated by an increase in their fraction of bound DNA values (spectra not shown). For 11, the fraction of bound DNA without metals was 0.09 and in the presence of Zn$^{2+}$ it increased to 0.17, while the values increased from 0.17 to 0.27 for 9.

Aside from 2, the ligands displaying the most pronounced metal-mediated DNA binding based on the ESI-MS results are 7 and 8. The binding of 6 was also mediated by Cu$^{2+}$, however the complexes formed by the ligand are lower in relative abundance than those formed by 7 and 8. This result is somewhat unsurprising due to the similarities in the structures of 2, 7, and 8. The ester linkage of 2 is changed to an amide-linkage

Table 1. Fraction of bound DNA values$^a$ for the benzoxazole ligands and d(GCGGAATTGGCG/CGCCAACCTCCGC) with Ni$^{2+}$, Cu$^{2+}$, or Zn$^{2+}$ and in the absence of metals. Solutions contained equimolar (10 μM) concentration of ligand, DNA, and, where appropriate, metal ion$^b$

<table>
<thead>
<tr>
<th>Ligand</th>
<th>No metal</th>
<th>Ni$^{2+}$</th>
<th>Cu$^{2+}$</th>
<th>Zn$^{2+}$</th>
</tr>
</thead>
<tbody>
<tr>
<td>2</td>
<td>0</td>
<td>0.18</td>
<td>0.17</td>
<td>0</td>
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<tr>
<td>6</td>
<td>0</td>
<td>0</td>
<td>0.14</td>
<td>0</td>
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<td>0.40</td>
<td>0</td>
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<tr>
<td>8</td>
<td>0</td>
<td>0.14</td>
<td>0.27</td>
<td>0</td>
</tr>
<tr>
<td>9</td>
<td>0.17</td>
<td>0.34</td>
<td>0.27</td>
<td>0.27</td>
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<tr>
<td>10</td>
<td>0.20</td>
<td>0.13</td>
<td>0.32</td>
<td>0.17</td>
</tr>
<tr>
<td>11</td>
<td>0.09</td>
<td>0.08</td>
<td>0.15</td>
<td>0.17</td>
</tr>
</tbody>
</table>

$^a$All values ±0.05. This error represents the largest standard deviation for experiments that were repeated on three separate days.

$^b$All solutions were prepared in a solvent composed of 50 mM ammonium acetate with 50% methanol (vol/vol).
in 8. While both 8 and 2 exhibited metal-mediated binding, copper had a more positive impact on 8 binding than it did on 2. Likewise, Ni\(^{2+}\) promoted greater binding for 2 than it did for 8. This difference may result from the preferential coordination of Cu(II) to nitrogen atoms over oxygen atoms [31]. Similar differences can be seen for the binding of 7, which also has an amide-linked side chain that is extended by two carbon atoms compared to 8 and 2. Compared with 2, 7 demonstrates greater DNA binding with copper and no binding with Ni\(^{2+}\).

While the longer side chains of 9, 10, and 11 enhance nonmetal-mediated DNA binding, they tend to preclude the dramatic metal-mediated DNA binding behavior observed for the ligands with the shorter side chains. The longer side chains of the ligands are believed to enhance the metal binding of the ligands by wrapping around and coordinating to the metals. The ester-linked polyethylene glycol side chain of 9 produced the best enhancement of metal-mediated binding, as all three metal ions, Ni\(^{2+}\), Cu\(^{2+}\), and Zn\(^{2+}\), increased the binding of the ligand to DNA relative to the binding in the absence of any metal. The results for 11, with the shorter, amide-linked polyethylene glycol side chain were not as favorable. While the binding was enhanced by Cu\(^{2+}\) and Zn\(^{2+}\), the overall fraction of bound DNA values were generally lower than the values for the other ligands in the presence of metals. With an amide-linked side chain that did not contain ethylene glycol groups, the binding of 10 was only enhanced by Cu\(^{2+}\). The binding results of the ligands with the longer side chains suggests that the longer, polyethylene glycol side chain of 9 is the most favorable for overall metal-mediated binding.

**Cytotoxicity Assays**

There are some interesting correlations between the degree of metal-mediated binding behavior of the benzoxazole ligands determined by ESI-MS and their cytotoxicity to two cancer cell lines. As summarized by the IC\(_{50}\) values shown in Table 2, the new ligands with the greatest cytotoxicity against the A549 lung cancer cell line were 7, 8, and 9, all having IC\(_{50}\) values that were similar to 2. Both 7 and 8 exhibited the most dramatic degree of metal-mediated DNA binding as their fraction of bound DNA values increased from 0 in the absence of metal ions to 0.40 and 0.27, respectively, in the presence of Cu\(^{2+}\). Ligand 8 was also found to form complexes in the presence of Ni\(^{2+}\). While 9 was not a
truly metal-mediated DNA binding ligand, it exhibit the most consistent and dramatic enhancement in DNA binding in the presence of metal ions compared to all the ligands that bound to DNA in the absence of metal ions.

Ligands 8 and 9 also showed cytotoxicity against the MCF7 breast cancer cell lines with IC50 values of 10 ± 8 μM and 13 ± 2 μM, respectively. Both of these values are on par with the IC50 value of 2 (4 ± 2 μM). Interestingly, 7, which showed activity against A549 cells comparable to that of 2 was relatively inactive against MCF7 cells (IC50 = 30 ± 10 μM), and 6, which was relatively inactive against A549 cells, retained some activity against MCF7 cells (IC50 = 15 ± 5 μM). At this point the reasons for these differences are unclear, but we note that of the two analogs of 2 that incorporate the most conservative structural changes, 6 and 8, only 8 demonstrates enhanced metal-mediated DNA binding relative to 2, and this analog also displays the most comparable cytotoxicity against both cancer cell lines relative to 2. Ligand 9 also demonstrated good cytotoxic activity, and while it was not a truly metal-mediated DNA binder, its DNA complexation was consistently enhanced by the presence of Cu2+, Ni2+, and Zn2+. In contrast, compound 12, which does not bind metal ions, was the only compound in this series that did not display cytotoxicity towards either cell line.

Conclusions

The correlation between significant metal-mediated or metal-enhanced binding determined by ESI-MS and anticancer activity of the benzoxazoles ligands assessed by cytotoxicity assays is demonstrated in this study. ESI-MS experiments reveal that Cu2+ and Ni2+ form the most abundant complexes with 6, 7, 8, 9, 10, and 11, while less abundant complexes are formed with Zn2+. For the complexes containing short side chains, 6, 7, and 8, 2:1 ligand/metal ion binding stoichiometries were predominant, whereas the compounds with longer side chains, 9, 10, and 11, formed abundant 1:1 complexes. DNA binding experiments reveal that the analogs with longer side chains formed complexes with duplex DNA in the absence of metal ions, while those with shorter side chains did not.

Of the ligands that did not bind to duplex DNA in the absence of metal ions, the DNA binding by 7 and 8 was enhanced most dramatically by Cu2+, Ni2+ influenced duplex binding for 8 and 2, and enhanced the binding by 9. The metal ion with the least substantial effect was Zn2+, which only enhanced the binding of 9 and 11, two ligands that formed complexes with DNA regardless of the presence of metals.

Of the compounds examined in this study, both 7 and 8 were also found to be the most cytotoxic against the A549 lung cancer cell line and 8 demonstrated moderate cytotoxicity against MCF7 breast cancer cells. Metal ions also enhanced the DNA binding of the ligands with the long side chains, most notably for 9, which also exhibited the highest level of cytotoxicity of the long side-chain compounds.

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