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Sulfonated Ni(II)porphyrin improves the detection of Z-DNA in condensed and non-condensed BZB DNA sequences

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Abstract

We report a very selective and sensitive spectroscopic detection of Z-DNA embedded in B-DNA in condensed as well as non-condensed DNA using anionic Ni(II) *meso*-tetrakis(4-sulphonatophenyl)porphyrin, **NiTPPS**. A combination of micromolar concentrations of Ni(II) and spermine⁴⁺ allowed us to prepare left-handed Z-DNA in short oligonucleotides without DNA condensation. A strong induced circular dichroism (ICD) signal was observed in the visible absorption region when **NiTPPS** was added to **BZ** DNA (Z-DNA fragment located at the end of a B-DNA tract with one B/Z DNA junction) and **BZB** DNA (Z-DNA sequence embedded in B-DNA having two B/Z DNA junctions). Almost no ICD signal was detected when **NiTPPS** was added to B-DNA. **NiTPPS** showed different binding modes with condensed and non-condensed Z-DNAs and allowed the distinction between condensed Z-DNA (positive bisignate CD couplet) and non-condensed Z-DNA (negative bisignate CD couplet).

Keywords

anionic nickel(II) porphyrin; left-handed Z-DNA; BZB sequences; induced circular dichroism; Z-DNA detection

Since the left handed form of DNA (Z-DNA) was first detected in 1972 [1], much work has been done to gain a scientific understanding of its biological role [2,3,4,5,6]. In vitro studies of Z-DNA have provided information regarding its structure and properties [7,8,9]. It is known that the transition of B- to Z-DNA in alternating pyrimidine-purine sequences can be induced by molar or millimolar or micromolar concentrations of cationic species (e.g. Ni²⁺, Co[NH₃]₆³⁺, spermine⁴⁺) [10,11]. Using circular dichroism (CD), it is possible to observe this transition experimentally. As shown in Fig. 1a, B-DNA exhibits a positive CD band at 280 nm and a negative CD band at 250 nm (blue curve) while Z-DNA has a negative CD band at 290 nm and a positive CD band at 260 nm [12,13,14,15].

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In real biological samples, however, the detection of Z-DNA is still an unsolved challenge because of the high B/Z DNA ratio and the spectroscopic interferences by proteins and other biological materials that also absorb in the UV region. In order to tackle this problem, we identified chiroptical probes that discriminate between B- and Z-DNA and do not absorb in the same region as DNA. The visible region above 300 nm is free from interferences and is ideally suited for induced circular dichroism (ICD) DNA recognition.

Metalloporphyrins have shown to be excellent chiroptical probes for detecting Z-DNA in alternating cytosine-guanine oligonucleotides (#b.p. <50) and polynucleotides (#b.p. ~1000) [16,17,18]. Recently we have reported that a cationic Zn(II)porphyrin (**ZnT4**, Fig. 1b) and an anionic Ni(II)porphyrin (NiTPPS, Fig. 1b) were able to spectroscopically detect the lefthanded Z-DNA under highly competitive conditions [19,20]. ZnT4 detected the Z-form embedded in B-DNA sequences with different B/Z ratios and nucleobase sequences via new ICD signals [19]. However, the sensing suffered from low specificity and intensity of the ICD signal. On the other hand, the anionic NiTPPS was used to sense the Ni(II)condensed Z-DNA in the presence of B-DNA (in this case two mixed oligonucleotides) via a strong ICD signal [20]. In order to improve the selectivity and sensitivity of detection of the Z-DNA fragment within oligonucleotide sequence, we have employed the anionic NiTPPS as a chiroptical probe. We have explored the detection of Z-DNA located at the end of a B-DNA tract with one B/Z DNA junction (BZ) as well as Z-DNA embedded in B-DNA having two B/Z DNA junctions (BZB(I)) in both condensed and non-condensed DNA samples (Fig. 2). Both DNAs have two 8-bromoguanines (depicted as X) incorporated in their sequences to promote the formation of Z-DNA fragment [21]. Natural DNA sequences **B** and **B**(I) with guanines instead of brominated guanines were used for comparison.

We started our study with a 33-mer **BZ** DNA. Addition of NiCl₂ (50 mM) to the **BZ** sequence (50 μ M) in Na-cacodylate buffer (1mM, pH = 7.0) promoted the expected B- to Z-DNA transition of the alternating cytosine-guanine part accompanied with Z-DNA condensation (Supplementary Material, Fig. S1 and S2). Addition of **NiTPPS** (5 μ M) gave rise to a strong bisignate CD signal with a positive CD band at 405 nm and negative CD band at 395 nm (Fig. 3). The absorption spectrum revealed hypochromicity (~34%) and a blue shift ($\Delta\lambda = 5.0$ nm) of the Soret band in the presence of **BZ**. Addition of **NiTPPS** to the non-brominated control sequence **B** under identical conditions did not give rise to ICD signals (Fig. 3a and S3).

Next, we explored the 42-mer **BZB(I)** sequence. The Z-form of the central portion of the **BZB(I)** was induced by NiCl₂ (50 mM) and confirmed by CD spectroscopy. The condensation of the **BZB(I)** was observed by a decrease of transmittance. The addition of **NiTPPS** (5 μ M) to a solution of **BZB(I)** (100 μ M + 50 mM NiCl₂) resulted in an intense bisignate CD signal in the Soret band region with a negative band at 390 nm and a positive band at 420 nm (Fig. 4a). The **NiTPPS** binding was accompanied by a hypochromic (H= ~ 50%) and hypsochromic ($\Delta\lambda$ = 5.0 nm) effect of the Soret band in UV-vis absorption spectrum (Fig. 5a). No ICD signal was observed upon addition of **NiTPPS** to the non-brominated control sequence **B(I)** from 0 to 10 μ M of **NiTPPS**, and a small bisignate signal is detected at 15 μ M (Fig. 4a and S6).

In order to explore the full potential of **NiTPPS**, we decided to test the spectroscopic sensing of Z-DNA fragments in the non-condensed DNA. We identified suitable experimental conditions that would allow the formation of Z-DNA fragments in short oligonucleotides at low ionic strength without DNA condensation. Spermine⁴⁺ failed to induce a Z-form in **BZ** and **BZB** [18]. However, the successive addition of micromolar concentrations of Ni(II) and spermine⁴⁺ allowed the B- to Z-DNA transition of the central GC tract without DNA aggregation. Spermine⁴⁺ is a ubiquitous cellular component and was

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added to shield the repulsion between phosphate backbone and anionic **NiTPPS**. The addition of **NiTPPS** yielded a negative exciton coupled CD signal (negative Cotton effect at 420 nm and positive Cotton effect at 400 nm) in the Soret band region (Fig. 5, red curve). A weak negative ICD signal was observed when **NiTPPS** was added to the control **B**(**I**) sequence (Fig. 5 and S7b). The absorption spectra did not show any significant changes of **NiTPPS** Soret band in presence of the **BZB(I**) sequence (see Fig. S8). The saturation of the negative bisignate ICD signal indicated that the ICD originated from the through space exciton coupling between the porphyrins' electronic dipoles when bound to DNA where limited number of suitable binding sites were available.

In conclusion, we report a very selective and sensitive spectroscopic detection of a Z-DNA fragment embedded in B-DNA in condensed as well as non-condensed DNA sequences using anionic Ni(II) porphyrin, **NiTPPS**. Anionic **NiTPPS** provided a more sensitive spectroscopic recognition of Z-DNA than previously reported cationic **ZnT4**. A combination of micromolar concentrations of Ni(II) and spermine⁴⁺ allowed us to prepare and detect the Z-DNA fragment in short non-condensed oligonucleotides with high sensitivity and selectivity. An intense ICD signal (~400 nm) was observed upon addition of **NiTPPS** to **BZ** and **BZB** DNA sequences. The ICD signal and its shape are dependent on the stage of Z-DNA condensation and allowed the distinction between condensed Z-DNA (positive bisignate CD couplet) and non-condensed Z-DNA (negative bisignate CD couplet).

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

Acknowledgments

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Appendix A. Supplementary data

Supplementary data associated with this article can be found online.



Fig. 1.

a) CD spectra of B-form (blue dashed curve) and Z-form (red curve) of poly(dG-dC)₂. b) Structure of cationic zinc(II) porphyrin (**ZnT4**, Zn(II)-*meso*-tetrakis(4-*N*-methylpyridyl)porphyrin) and anionic nickel(II) porphyrin (**NiTPPS**, Ni(II)-*meso*-tetrakis(4-sulphonatophenyl)porphyrin).

BZ	5'- <mark>(CG)₁₀CGCG</mark> ATAAACTCG-3' 3'- <mark>(GC)₁₀XCXC</mark> TATTTGAGC-5'
В	5'-(CG) ₁₀ CGCGATAAACTCG-3' 3'-(GC) ₁₀ GCGCTATTTGAGC-5'
BZB(I)	5'-GGTTTAT <mark>CXCX(CG)₁₀CGCG</mark> ATAAACC-3' 3'-CCAAATA <mark>GCGC(CG)₁₀XCXC</mark> TATTTGG-5'
B(I)	5'-GGTTTATCGCG(CG) ₁₀ CGCGATAAACC-3' 3'-CCAAATAGCGC(CG) ₁₀ GCGCTATTTGG-5'

Fig. 2. BZ, B, BZB(I), and B(I) sequences employed in current study (X depicts 8-bromoguanine, Z-DNA fragment is written in bold red).



0.0

200

300

400 500 600

 λ / nm

Fig. 3.

20

10

0

-10

-20

-30

200

300

CD / mdeg

a)

(a) CD and (b) UV-vis absorption spectra of **NiTPPS** (5 μ M) alone (black dotted curve) and in presence of **BZ** (100 μ M + 50 mM NiCl₂, red curve) and **B** (100 μ M + 50 mM MgCl₂, blue dashed curve) in Na- cacodylate buffer (1 mM, pH = 7.0). Inset: CD signal at 410 nm as a function of [**NiTPPS**] in presence of **BZ** (red circles) and **B** (blue triangles) sequences.

410 nm

500

2 4 6 8 [NiTPPS] / μΜ

400

 λ / nm



Fig. 4.

(a) CD and (b) UV-vis absorption spectra of **NiTPPS** (5 μ M) alone (black dotted curve) and in presence of **BZB(I)** (100 μ M + 50 mM NiCl₂, red curve) and **B(I)** (100 μ M + 50 mM MgCl₂, blue dashed curve). Inset shows the CD signal at 415 nm as a function of [**NiTPPS**] in presence of **BZB(I)** (red circles) and **B(I)** (blue triangles) sequences.



Fig. 5.

CD spectra of **NiTPPS** (5 μ M) in presence of **BZB(I)** (100 μ M + 100 μ M NiCl₂ + 6 μ M spermine, red curve) and in presence of **B(I)** (100 μ M + 100 μ M MgCl₂ + 6 μ M spermine, blue dashed curve). Inset: CD signal at 400 nm as a function of [**NiTPPS**] in presence of **BZB(I)** (red circles) and **B(I)** (blue triangles).