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Reactivity of Parallel-Stranded DNA to Chemical Modification Reagents

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ABSTRACT: Four 25-nt long oligonucleotides containing dA and dT (D1, D2, D3, and D4) which are capable of forming parallel-stranded (ps) or antiparallel-stranded (aps) duplexes have been synthesized [Rippe, K., Ramsing, N. B., & Jovin, T. M. (1989) *Biochemistry* 28, 9536-9541]. In the present study, the OsO₄-pyridine complex (Os,py), diethyl pyrocarbonate (DEPC), KMnO₄, and the 1,10-phenanthroline-cuprous complex [(OP)₂Cu⁺] were used to investigate the conformation-dependent reactivity of ps, aps, and single-stranded (ss) oligonucleotides. The products were analyzed by polyacrylamide gel electrophoresis with single-nucleotide resolution. The results confirm the duplex nature of the ps combinations of oligonucleotides and reveal structural differences in comparison with the aps molecules. Under conditions in which ss-DNA is substantially sensitive to Os,py, both the ps and aps duplexes are very unreactive. A similar result was observed with KMnO₄ and DEPC, although with the latter reagent the modification pattern of the labeled strands D1* and D4* was slightly different for the parallel than for the antiparallel duplex. The (OP)₂Cu⁺ complex efficiently cleaves the aps but not the ps duplex and shows a preference for TAT steps. We also tested the effect of monovalent and divalent cation concentrations on the chemical reactivity of the ps, aps, and ss species. Elevated NaCl concentration leads to a dramatic increase in the Os,py and KMnO₄ modification of ss molecules and the ps, but not the aps, duplex. We attribute the apparent reaction with ps-DNA to a destabilization of this conformation under the conditions of reaction. In contrast, all reactions with DEPC are somewhat depressed at high salt concentration. The effects of MgCl₂ and temperature on the chemical reactivity with Os,py were also determined. The helix-coil transition of both the ps and the aps duplexes can be monitored by chemical modification with the OsO₄-pyridine reagent.

The antiparallel orientation of DNA strands is a characteristic feature of all right-handed B-type and left-handed Z-type helices. In 1986, N. Pattabiraman presented force field calculations indicating the possibility that a stable duplex composed of d(A)₆ and d(T)₆ could form with both strands oriented in the same 5'-3' direction and associated via *reverse* Watson-Crick base pairing with the glycosidic bonds in a trans

orientation (Pattabiraman, 1986). The calculated stability was similar to that of conventional antiparallel DNA, encouraging numerous experimental approaches for examining the existence and properties of parallel-stranded hairpins (van de Sande et al., 1988; Ramsing et al., 1989; Germann et al., 1989; Shchylorkina et al., 1989; Tchurikov et al., 1989) and of parallel-stranded duplexes (Ramsing & Jovin, 1988; Germann et al., 1988; Ramsing et al., 1989; Rippe et al., 1989, 1990; Rippe & Jovin, 1989; Jovin et al., 1990). On the basis of the results available to date, it is clear that DNA can adopt a parallel-stranded (ps)¹ conformation that is only moderately

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less stable than antiparallel B-DNA. Parallel-stranded DNA (ps-DNA) differs from B-DNA with respect to numerous thermodynamic, spectroscopic, and drug binding characteristics and as a substrate for DNA-specific enzymes.

The present investigation exploited chemical probes developed recently for studying DNA conformations in solution and their transitions in sequences of interest. The objectives were to (i) specify further the structural differences between ps-DNA and aps-DNA, (ii) establish means for identifying the ps conformation in natural sequences, and (iii) demonstrate the utility of the chemical probes in studies of oligonucleotides, a surprisingly undeveloped area of application. Since the synthetic oligonucleotides used in this study were composed exclusively of dA and dT nucleotides, the structure-sensitive chemical probes that allow detection of the modification of dA and dT residues or cleavage products with single-nucleotide resolution were chosen. Several chemicals, including Os₂py [Galazka et al., 1986; Johnston & Rich, 1985; for a review, see Paleček et al. (1990)], DEPC (Johnston & Rich, 1985; Herr, 1985), KMnO₄ (Borowiec et al., 1987; Lyamichev et al., 1989), and the (OP)₂Cu⁺ complex (Kuwabara et al., 1986; Sigman, 1986; Sigman & Spassky, 1989), can be used to detect structure-dependent hypersensitivity of a given set of bases. Os₂py, which adds to the C5–C6 of the pyrimidine rings, site-specifically (Lukášová et al., 1982) modifies single-stranded and distorted regions in DNA, such as the loops of cruciform structures (Lilley & Paleček, 1984; Klysik et al., 1988), B–Z junctions (Nejedlý et al., 1985; Galazka et al., 1986; Johnston & Rich, 1985; Klysik et al., 1988), intramolecular triplexes (Vojtišková & Paleček, 1987; Hanvey et al., 1988; Voloshin et al., 1988; Vojtišková et al., 1988; Parniewski et al., 1989), and looped-out bases and bubbles (Bhattacharya & Lilley, 1989). It reacts primarily with thymines, forming osmate ester derivatives, and more slowly with cytosine and guanine (Klysik et al., 1988). DEPC carbethoxylates the N-7 position of purines in a duplex conformation dependent way (Johnston & Rich, 1985; Herr, 1985; Klysik et al., 1988). It is reactive with dA and dG residues of DNA in the left-handed state and extensively modifies bases within the cruciform loops (Furlong & Lilley, 1986; Sholten & Nordheim, 1986). B-DNA is insensitive to modification with either Os₂py or DEPC. KMnO₄ has also been shown to be useful in studies of DNA conformation (Borowiec et al., 1987; Lyamichev et al., 1989) and is also more reactive with ss than with double-stranded species. Detection of Os₂py-, DEPC-, or KMnO₄-modified bases is accomplished by piperidine cleavage of the labeled strand and electrophoresis of the resulting products on a sequencing gel next to the Maxam and Gilbert sequencing reactions of the same fragment (Johnston & Rich, 1985; Herr, 1985; Klysik et al., 1988; Furlong & Lilley, 1986; Sholten & Nordheim, 1986). The copper–phenanthroline [(OP)₂Cu⁺] complex has been used to distinguish between the conformational details of B-DNA (Kuwabara et al., 1986; Sigman, 1986; Sigman & Spassky, 1989). A remarkable feature of the (OP)₂Cu⁺-mediated reaction is its mechanism of scission. The minor groove is required for the binding of the complex to DNA and subsequent cleavage, rendering the reagent highly specific for double-stranded B-type helices.

In this report, we present results on chemical modification of ps-, aps-, and ss-DNAs. The set of chemical probes serves

to distinguish ps duplexes from their aps counterparts and ss-DNA on the basis of differences in the extents of modification and their dependence on monovalent and divalent cation concentrations and temperature.

MATERIALS AND METHODS

Oligonucleotide Synthesis and End Labeling. Oligonucleotides D1, D2, D3, and D4 were synthesized and end-labeled with ³²P as described previously (Rippe et al., 1989). After the kinase reaction, oligonucleotides were purified by preparative, denaturing, 15% polyacrylamide gel electrophoresis followed by extraction and ethanol precipitation (Maniatis et al., 1982).

Duplex Formation. Unless otherwise specified, the DNA strand combinations were mixed in a buffer containing 50 mM NaCl, 50 mM Tris-HCl, pH 8.0, and 10 mM MgCl₂. The concentration of the unlabeled strand was usually 20 times higher than that of the end-labeled strand (0.8 μM). The melting temperature under these conditions was estimated from values determined previously in optical measurements by applying a correction according to the expression

$$T_m^{-1} = T_{m,\text{ref}}^{-1} - (R/\Delta H_{\text{vH}}) \ln(4C_i/C_t) \quad (1)$$

where $T_{m,\text{ref}}$ is the reference T_m of an equimolar mixture with a strand concentration C_t under the same salt conditions as the chemical modification experiment carried out in the presence of an excess concentration, C_i , of one of the strands. Equation 1 is derived from eq 5 and 14 in Ramsing et al. (1989) and the following expression for the dissociation constant K in the case of one strand being in excess:

$$K \approx C_i\alpha/(1 - \alpha) \quad (2)$$

where α is the fractional degree of transition to the coil form. The values of the van't Hoff enthalpy difference ΔH_{vH} and $T_{m,\text{ref}}$ for the helix-coil transition in 10 mM MgCl₂ were those determined previously from measurements of absorption hyperchromicity (Rippe et al., 1989). The samples were heated to 70 °C for 10 min and cooled slowly to room temperature before starting the reaction, the volume of which was usually 50 μL.

OsO₄-Pyridine Modification. Modification with Os₂py was performed according to Galazka et al. (1986). The 25 mM OsO₄ (Sigma) stock solution was kept frozen (–20 °C). DNA solutions in the appropriate buffer were supplemented with pyridine (2% v/v), and the reaction was started by adding OsO₄ stock solution to a final concentration of 2 mM. After 2 min, the reaction was stopped by adding 4 volumes of ethanol. Then the DNA was pelleted, washed with 75% ethanol, dissolved in 1 M piperidine, and heated at 90 °C for 30 min. The samples were dried in a Savant vacuum centrifuge, dissolved in formamide, and loaded on a 15% polyacrylamide [19:1 acrylamide/bis(acrylamide)] sequencing gel. Exposure times of the Kodak XAR-5 X-ray film were adjusted according to the number of counts per lane loaded on each individual gel. In reference to the reactions with Os₂py and the other reagents, the terms overmodification and overreaction are used to indicate that the extent of cleavage greatly exceeded the limit of a single site per molecule, thereby biasing the distribution.

DEPC Modification. Modifications with DEPC were performed as described previously (Klysik et al., 1988). Solutions of the synthetic oligonucleotides in appropriate buffers were made as in the case of Os₂py reactions except that OsO₄ and pyridine were omitted. Reactions were started at room temperature by adding one-tenth of the sample volume of

¹ Abbreviations: ps, parallel-stranded; ps-DNA, parallel-stranded DNA; aps, antiparallel-stranded; aps-DNA, antiparallel-stranded DNA; ss, single-stranded; Os₂py, OsO₄-pyridine complex; DEPC, diethyl pyrocarbonate; (OP)₂Cu⁺, 1,10-phenanthroline-cuprous complex; nt, nucleotide(s).

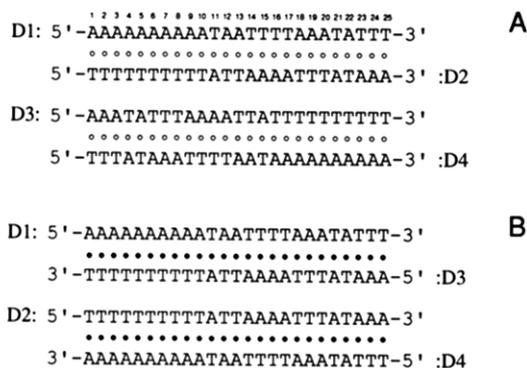


FIGURE 1: Sequences of oligonucleotides and their duplexes. The Watson-Crick base pairs of the antiparallel duplex *aps*-D1-D3 and *aps*-D2-D4 are shown by the symbol (●) and the *reverse* Watson-Crick (trans Crick-Watson) base pairs of the parallel-stranded duplexes (*ps*-D1-D2, *ps*-D3-D4) by the symbol (○). (A) Parallel-stranded duplexes *ps*-D1-D2 and *ps*-D3-D4. (B) Antiparallel-stranded duplexes *aps*-D1-D3 and *aps*-D2-D4.

DEPC and vortexing. Modification was terminated by precipitation of DNA with 4 volumes of ethanol. Further workup was as in the case of *Os,py* modification.

KMnO₄ Reaction. Reactions with KMnO₄ were performed according to Yeung et al. (1988). DNA was exposed to KMnO₄ (80 μg mL⁻¹) for 15 min at room temperature, and the reaction was terminated by adding 10 μL of allyl alcohol. After precipitation with ethanol, the DNA was treated with piperidine, heated to 90 °C for 30 min, dried, and loaded on the sequencing gel after dissolving in formamide.

(OP)₂Cu⁺ Reaction. DNA cleavage by phenanthroline-copper complex was done in 50 mM Tris-HCl, pH 7.2, 2 mM MgCl₂, 4.8 mM 3-mercaptopropionic acid, 340 μM 1,10-phenanthroline (Sigma), and 80 μM CuSO₄. The reaction was stopped by adding 2,9-dimethyl-1,10-phenanthroline (Sigma) to a final concentration of 2 mM. After lyophilization, the samples were dissolved in formamide and loaded on a sequencing gel.

Densitometric Tracings of the Autoradiograms. The X-ray films were transilluminated and the transmitted images acquired with a Series 200 slow-scan CCD camera (Photometrics, Tucson, AZ) with a dynamic range of 14 bits and a spatial resolution of 4.2 pixels mm⁻¹. The images were processed on an Apple Macintosh IIcx computer using combinations of the programs TCL-Image (Delft Center for Image Processing, distributed by Multihouse, Amsterdam; the Macintosh version was kindly supplied by Dr. I. T. Young) and Image (W. Rasband, National Institutes of Health, Bethesda, MD). Optical densities (OD) were computed after flat-field correction, and the central 10 lines of each lane were averaged (for the scan plots) or depicted in a 3-D representation. Good linearity was obtained up to an optical density of 1.5. Plots were generated with the commercial programs Passage and DeltaGraph; superposed scans were displaced vertically for enhanced clarity.

RESULTS AND DISCUSSION

The physicochemical and biochemical characteristics of the 25-nt long oligonucleotides D1, D2, D3, and D4 and their duplexes have been reported previously (Rippe et al., 1989; Ramsing et al., 1989; Rippe & Jovin, 1989; Jovin et al., 1990). Figure 1 depicts the possible combinations of oligonucleotides that lead to the formation of *ps* or *aps* duplexes. Thus, oligonucleotides D1 + D2 and D3 + D4 mixed in stoichiometric amounts form *ps* duplexes (Figure 1A), whereas D1 + D3 and D2 + D4 form conventional *aps* helices (Figure 1B). The other

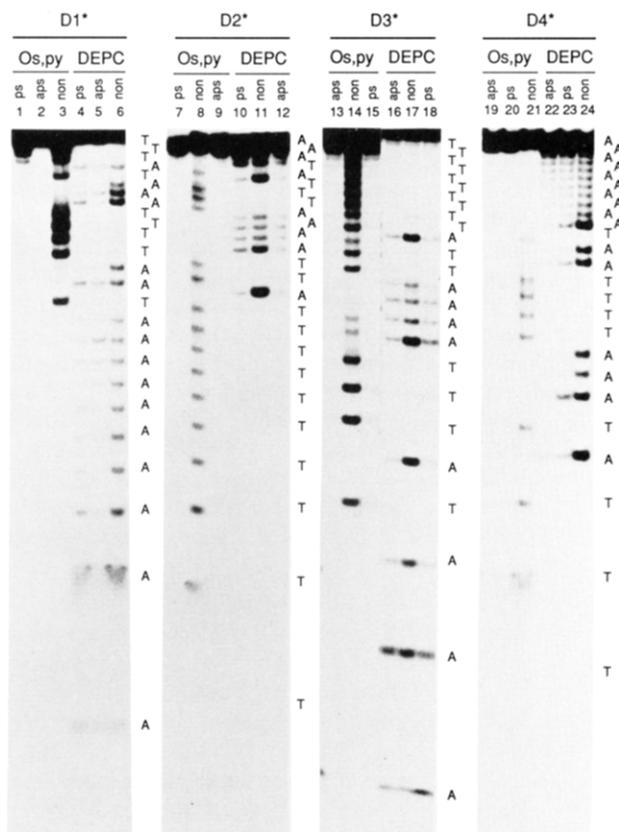


FIGURE 2: Autoradiograph from sequencing gel of *Os,py*- and DEPC-modified duplexes (*aps* and *ps*) and nonpairing (*non*) species. The modification reactions contained mixtures of the given labeled strand (indicated by asterisks) and about 20-fold excess of the unlabeled strand necessary to constitute the *ps*, *aps*, or nonpairing combination (cf. Figure 1). For example, in the first panel, D1* is labeled, and the lanes denoted with *ps* (lane 1), *aps* (lane 2), and *non* (lane 3) correspond to the combinations *ps*-D1*-D2, *aps*-D1*-D3, and D1* + D4. Modifications were performed in 50 mM NaCl, 50 mM Tris-HCl, pH 8.0, and 10 mM MgCl₂ at room temperature. After modification and piperidine cleavage, the samples were run on a 15% sequencing gel before autoradiography. Each lane contained the same number of counts loaded. Modification patterns obtained for single-stranded D1*, D2*, D3*, and D4* oligonucleotides (data not shown) were identical with those resulting from reactions of nonpairing mixtures D1* + D4, D2* + D3, D2 + D3*, and D1 + D4*.

combinations, D1 + D4 or D2 + D3, are not complementary in either orientation; neither parallel nor antiparallel duplexes have been detected (Rippe et al., 1989).

In this study, the duplexes preformed *in vitro* were subjected to chemical modification and chemical cleavage. The collected data were compared with results obtained by similar treatment of nonpairing combinations of oligonucleotides or of one strand employed as single-stranded references.

Modifications of *ps*-, *aps*-, and *ss*-DNA with *Os,py* and DEPC. Upon mixing labeled D1 fragment (D1*) with unlabeled D2 or D3, the *ps* or *aps* duplexes *ps*-D1*-D2 and *aps*-D1*-D3 are formed under the buffer conditions used (10 mM MgCl₂, 50 mM NaCl, and 50 mM Tris-HCl, pH 8.0) (Rippe et al., 1989). D1* mixed with D4 remains single-stranded (*ss*); that is, duplex formation is not detected. In a similar way, the end-labeled D2 (D2*) can be used to generate *ps*-D1-D2* and *aps*-D2*-D4, or when mixed with D3, to establish a single-stranded control. The same combinations can be produced with labeled D3* and D4*.

The reactivities of the given combinations with *Os,py* and DEPC are shown in Figure 2. As expected, both chemicals react extensively with the unpaired mixture of the oligonucleotides, or the single-stranded oligonucleotides (data not

shown). Os₄py overreacts with dT, whereas DEPC overmodifies dA residues. The aps or ps duplexes, however, are much less sensitive to these chemicals, regardless of which strand is labeled and analyzed. In particular, Os₄py shows virtually no reaction with aps DNA, but a slight reactivity can be detected for the ps counterparts on the original autoradiographs. The differences between modification intensities of the given bands for aps or ps duplexes and ss or nonpaired species obtained with DEPC are also clear. Most reactive is the nonpairing combination of oligonucleotides; the aps and ps duplexes are less sensitive to this chemical. Some minor but reproducible differences between the intensities of individual bands derived from D1* and D4* strands forming aps or ps duplexes are also apparent (cf. Figure 1, lanes 4, 5 and 22, 23). These results are consistent with previous studies (Rippe et al., 1989; Ramsing et al., 1989), and we conclude that the appropriate combinations of synthetic oligonucleotides form aps or ps duplexes that are less accessible to chemicals such as Os₄py or DEPC than DNA in a single-stranded conformation.

Effect of NaCl on Os₄py and DEPC Reactivity. The stability of double-stranded ps or aps DNA depends on a number of physicochemical factors including the counterion concentration in the DNA solution. Salt-induced changes in DNA conformation such as in the B-Z transition are very well documented (Rich et al., 1984; Jovin et al., 1987). Thus, it was of interest to compare the change in the Os₄py and DEPC reactivity to aps and ps duplexes and ss-DNA as a function of the salt concentration. An increase in NaCl concentration up to 4 or 5 M results in a slight increase in the extent of Os₄py modification of the aps-D1·D3* duplex [Figure 3 (top), lanes 13–16; and Figure 3 (bottom), panel D]. Interestingly, a substantial increase in reactivity of 1 dT residue (closest to the 5' end, T₄; see Figure 1A) out of the 16 T residues present on this strand is evident. The behavior of the ps duplex is substantially different. At 0.5 or 1.5 M NaCl, weak Os₄py modifications are observed [Figure 3 (top), lanes 21 and 22; Figure 3 (bottom), panel F]. Upon further increases in salt concentration, the ps duplex is overmodified, as reflected by the diminution of the radioactive material at the position of the intact fragment (top of the gel) [Figure 3 (top), lanes 23 and 24; Figure 3 (bottom), panel F]. The nonpairing combination of the oligonucleotides D2 + D3* [Figure 3 (top), lanes 17–20; Figure 3 (bottom), panel E] as well as D3* alone (data not shown) is also sensitive to the salt increase and become overmodified at the highest NaCl concentration used. We conclude that elevated salt concentrations increase the intrinsic reactivity of the Os₄py-pyridine reagent. Modification of aps-D1·D3* by Os₄py changes only slightly as a function of salt concentration; thus, the different behavior of the ps-D3*·D4 duplex must reflect distinctive features of the parallel-stranded conformation. In fact, measurements of absorption hyperchromicity show that high salt concentrations reduce the thermodynamic stability of the ps helix. For example, the *T_m* of ps-D1·D2 and ps-D3·D4 in 1.6 M NaCl is 46 °C (Rippe et al. 1989) but decreases to 32 °C in 4.9 M NaCl and 10 mM MgCl₂. In addition, the helix-coil transition is less cooperative; ΔH_{VH} decreases by ~30%. These changes do not reflect a conformational transition in the native state as judged from the invariant wavelength-dependent hyperchromicity pattern (Ramsing et al., 1989; Rippe et al., 1989) and from circular dichroism spectra. (Experimental and theoretical studies of these high salt effects on ps-DNA will be reported elsewhere.) Due to the reduced cooperativity of the helix-coil transition of ps-DNA, we calculate a value of

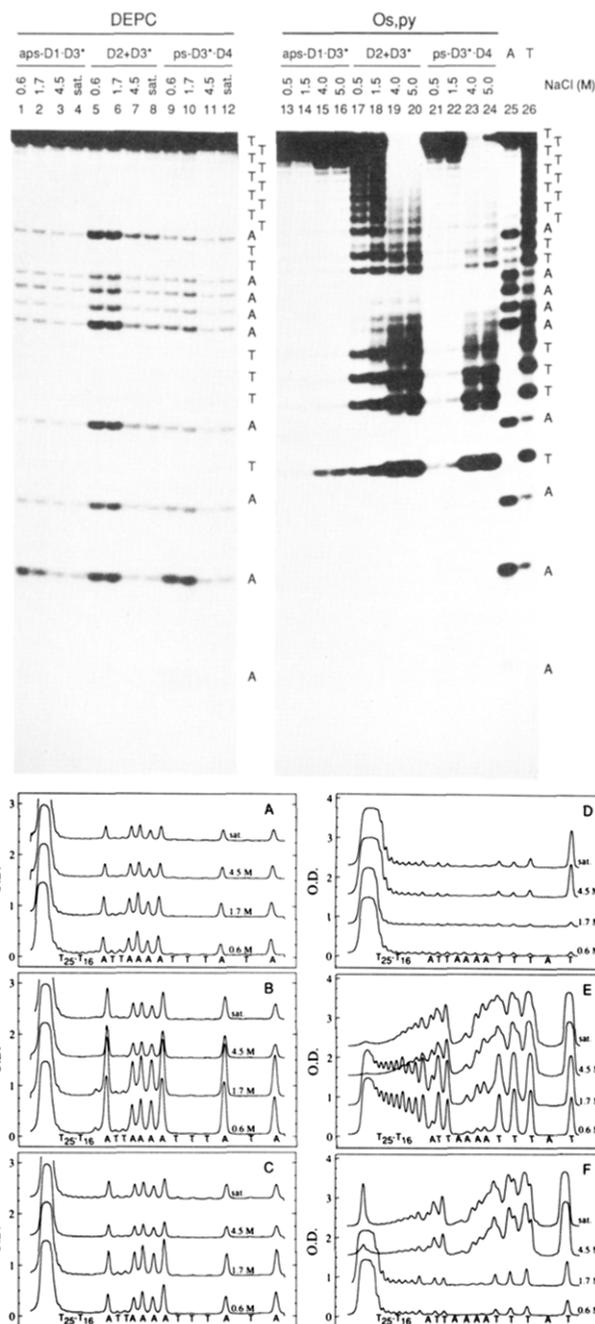


FIGURE 3: DEPC and Os₄py modification of aps-D1·D3*, ps-D3*·D4, and nonpairing (non) combination of oligonucleotides D2 + D3* as a function of increasing NaCl concentration. (Top) Autoradiograph from sequencing gel of reaction products. Reactions were performed at room temperature in the buffer containing 50 mM NaCl, 50 mM Tris-HCl, pH 8.0, and 10 mM MgCl₂ supplemented with the indicated concentration of NaCl. The abbreviation "sat." refers to the solution saturated with NaCl and A and T to the products of the Maxam-Gilbert sequencing reaction for adenines and thymines, respectively. (Bottom) Densitometric scans of the autoradiograph presented in the top part of the figure. A-C show the results for the DEPC modification: (A) aps-D1·D3*; (B) D2 + D3*; (C) ps-D3*·D4. Panels D-F show the results for the modification with the Os₄py-pyridine reagent. (D) aps-D1·D3*; (E) D2 + D3*; (F) ps-D3*·D4.

α at 25 °C (7 °C below the optically determined *T_m*) of ~0.2, indicating that a considerable fraction is in the ss coil state and thereby susceptible to the Os₄py reagent. The apparent *T_m* of the ps duplex would be depressed still further due to the increased reactivity of the unlabeled single strand in excess to Os₄py at high NaCl concentrations. That is, a continuous diminution of *C_i* during the reaction would have the effect of shifting the transition curve to lower temperatures (eq 1 and

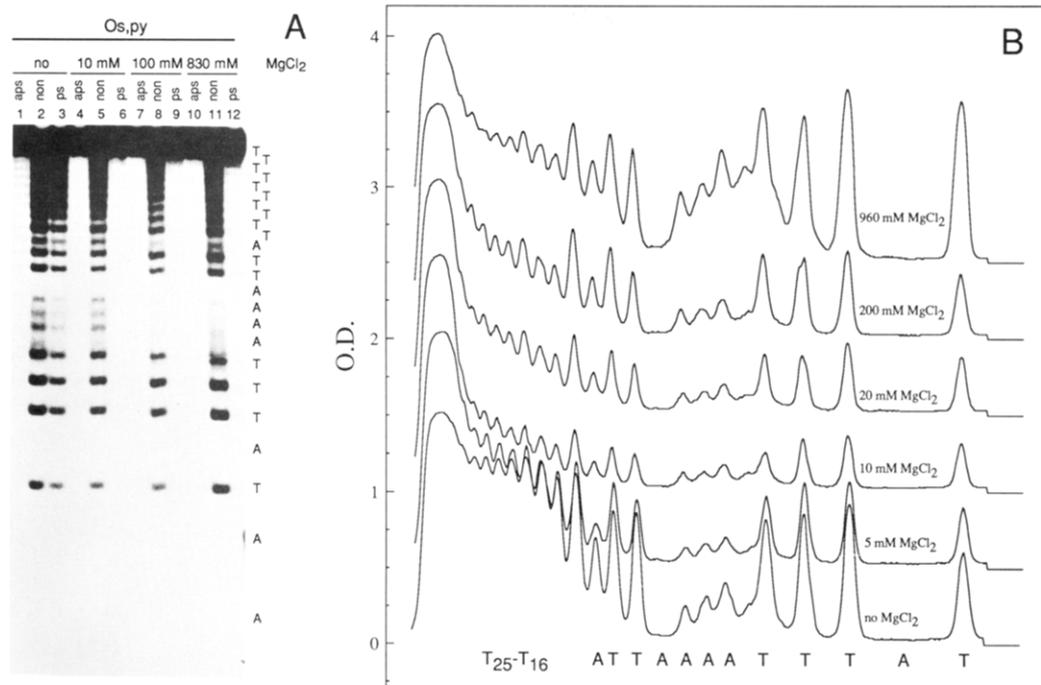


FIGURE 4: Effect of MgCl₂ on Os.py reactivity. (A) Autoradiograph of products from reaction with aps-D1·D3* (aps), ps-D3*·D4 (ps), and D2 + D3* (non) separated on a sequencing gel. Modifications were conducted in a buffer containing 50 mM Tris-HCl, pH 8.0, and 50 mM NaCl at room temperature after supplementing the mixture with the indicated concentration of MgCl₂. (B) Effect of MgCl₂ on Os.py reactivity on single-stranded D3*. The experiment was performed essentially as in (A). The plot derives from the quantitation of the autoradiograph.

2). In contrast, the T_m of the aps duplexes is much less affected by the high salt condition, decreasing from 60 °C at 1.6 M NaCl (Rippe et al., 1989) to 58 °C in 4.9 M NaCl/10 mM MgCl₂ (data not shown), a value well above the reaction temperature of 25 °C. In addition to the effect of high salt concentration, the OsO₄-pyridine reagent itself may destabilize the ps helix (see Helix-Coil Transition of ps and aps Duplexes Monitored by OsO₄-Pyridine Modification). In summary, we interpret the apparent hyperreactivity of the ps-DNA with Os.py at high salt concentration as the result of an increase in the Os.py reactivity of ss-DNA, and a destabilization of the ps duplex with a concomitant reduction of the helix-coil transition cooperativity.

Substantially different effects of increasing NaCl concentration can be demonstrated with DEPC (Figure 3). Little change in sensitivity is detected with the ps duplex [Figure 3 (top), lanes 9–12; Figure 3 (bottom), panel C], the aps duplex [Figure 3 (top), lanes 1–4; Figure 3 (bottom), panel A], and the nonpairing combination D2 + D3* [Figure 3 (top), lanes 5–8; Figure 3 (bottom), panel B], all of which show a slight decrease in reactivity. The reaction of DEPC with unpaired dA is much less than that of Os.py with unpaired dT residues [Figure 2; Figure 9 in Bhattacharya and Lilley (1989)]. Thus, the differences in modification of the helix and coil forms, already relatively minor at low salt concentration, are reduced even further under high salt conditions. We conclude that DEPC is an effective probe of DNA conformation only at moderate ionic strength.

Effect of MgCl₂ on Reactivity of ps-, aps-, and ss-DNA with Os.py. We tested the effect of Mg²⁺ ions on the Os.py reactivity of ss DNA (D3* or D2 + D3*) and the ps-D3*·D4 and aps-D1·D3* duplexes (Figure 4A). From the densitometric trace of the autoradiogram of the sequencing gel shown in Figure 4B, it is clear that the D3* oligonucleotide at first shows a slightly lower reactivity as the MgCl₂ concentration increases. Still higher MgCl₂ concentrations cause a slight elevation in Os.py modification, but the changes in the intensities of the bands are relatively small.

In the absence of Mg²⁺, the ps duplex displays a modification rate that is only slightly lower than that obtained for the nonpairing combination of oligonucleotides, that is, substantially higher than in the case of the aps complex (Figure 4A, lanes 1–3). This result can be interpreted in terms of the thermodynamic stability of the ps duplex as compared with the aps reference, and correlates well with previous findings (Rippe et al., 1989). The salt conditions used are high enough to allow aps but not ps duplex formation at room temperature (22 °C). The experiments were performed in the buffer containing 50 mM NaCl and 50 mM Tris-HCl, pH 8, buffer. The T_m for aps-D1·D3 under these conditions was previously determined to be around 40 °C, whereas the T_m for the ps-D3·D4 duplex is about 22 °C (Rippe et al., 1989).

Due to the increased stability of both duplex structures, the aps- and ps-DNAs are less sensitive to Os.py modification than the nonpairing combination of oligonucleotides at all Mg²⁺ concentrations used (Figure 4A, lanes 4–12). In contrast to the findings obtained with the NaCl experiment (Figure 3), no dramatic increase in sensitivity is observed over the range of MgCl₂ concentrations.

Helix-Coil Transition of ps and aps Duplexes Monitored by OsO₄-Pyridine Modification. As demonstrated in the previous sections, the Os.py reactivity toward the ps duplex and ss-DNA is greatly influenced by NaCl (Figure 3). We expected that the modification kinetics might be relatively invariant over a substantial range of temperature. In fact, the Os.py reagent was used in earlier studies of temperature-dependent cruciform and triple-stranded DNA formation (Klysik et al., 1988; Hanvey et al., 1989). We addressed the question of whether one can monitor the helix-coil transitions of ps and aps duplexes by detecting the change in sensitivity of DNA to Os.py modification. Results of such an experiment are shown in Figure 5. It is apparent that little change in the elevated modification rate can be detected with the nonpairing combination of D2 + D3* oligonucleotides over the temperature range 22–62 °C [Figure 5 (top), lanes 9–16]. Substantial increases in Os.py sensitivity occur, however, for ps

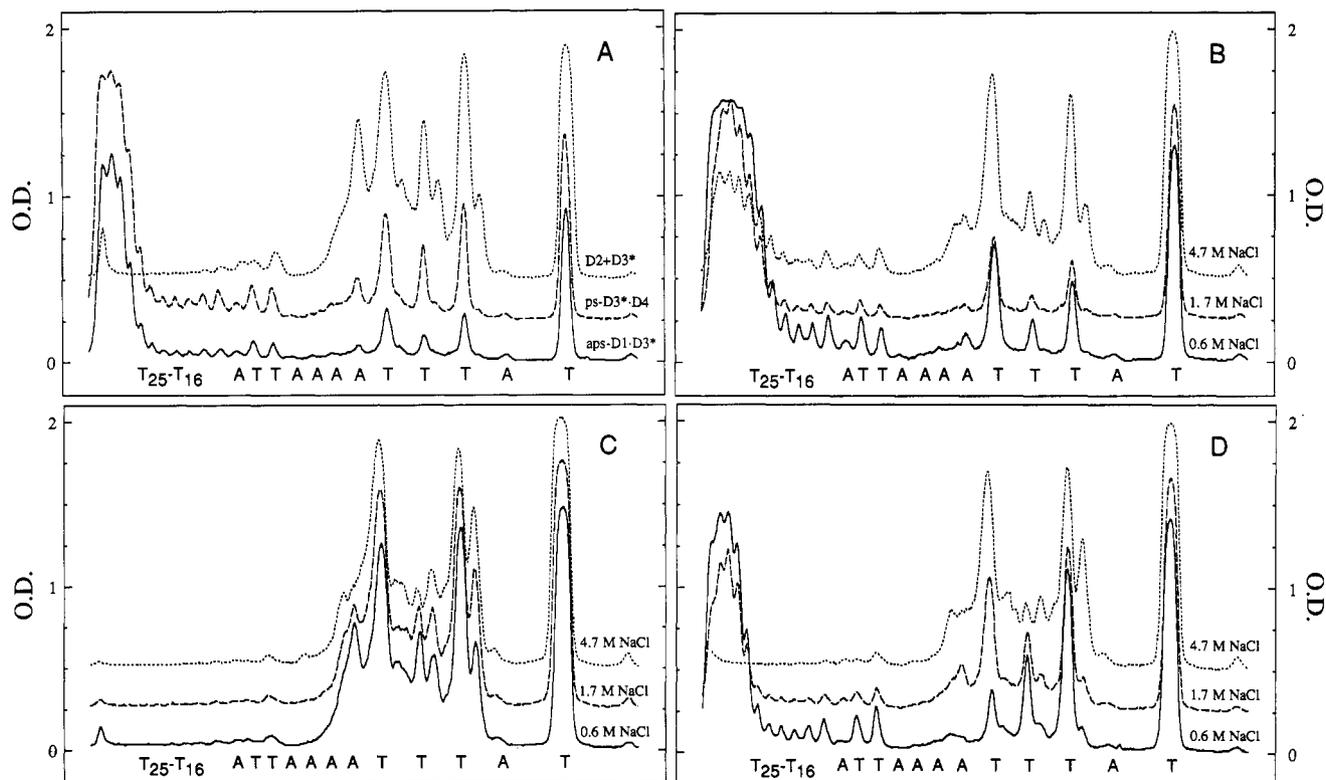


FIGURE 6: Reactivity of ps, aps, and nonpairing combinations of oligonucleotides to KMnO_4 as revealed by densitometric tracings. Reactions with all combinations of oligonucleotides were conducted at room temperature in 50 mM Tris-HCl, pH 8.0, and 10 mM MgCl_2 , in addition to the given NaCl concentration. (A) All combinations in 50 mM NaCl. (B) aps-D1-D3* in 0.6, 1.7, and 4.7 M NaCl. (C) D2 + D3* in 0.6, 1.7, and 4.7 M NaCl. (D) ps-D3*-D4 in 0.6, 1.7, and 4.7 M NaCl.

attach to the deoxyribose moiety, leading to the scission of the phosphodiester bonds. This efficient chemical nuclease has been shown to cleave B-DNA and A-DNA at a reduced rate, but not Z-DNA and single-stranded species (Pope & Sigman, 1984; Sigman et al., 1985).

It was of interest to determine with single-nucleotide resolution whether $(\text{OP})_2\text{Cu}^+$ complex could actually discriminate between the aps and ps duplexes. The aps and ps duplexes as well as ss oligonucleotides and the mixture of nonpairing oligonucleotides were cleaved with this chemical, and the resulting products were subjected to sequencing gel electrophoresis (Figure 7A,B). As expected, the most efficient cleavage occurs in the case of aps duplexes. The predominant cleavage sites are located at the two TAT steps on the aps-D1-D3* (Figure 7A) or at the TAT and at one out of two TAA steps on the aps-D2-D4* duplex (Figure 7B). The TAT steps on B-DNA helix are known to be the most sensitive sites for $(\text{OP})_2\text{Cu}^+$ cleavage (Veal & Rill, 1988). Little or no degradation of the D3* or D4* oligonucleotides occurs when ps DNA is formed with unlabeled D4 and D3, respectively. The cleavage pattern obtained for ss-D3* and ss-D4* or for the nonpairing combinations (D2 + D3* and D1 + D4*) is still different from that obtained for their aps and ps counterparts. For example, ss-D3* or D2 + D3* are slightly sensitive to $(\text{OP})_2\text{Cu}^+$ scission at the TAT step closest to the 3' end of D3*. The second TAT step closer to the 5' end remains insensitive. (In aps-D1-D3*, both TAT sites are sensitive.) The ss-D4* and D1 + D4* cleavage patterns are also identical and differ from the pattern obtained for the aps-D2-D4* duplex. The TAT and 5'-TAA steps efficiently cleaved in the aps duplex formed with D4* seem to be much less sensitive in the case of ss-D4*, on which new sites appear at the second TAA sequence that is closer to the 3' end (Figure 7B). Previous work has demonstrated that the $(\text{OP})_2\text{Cu}^+$ reagent does not cleave ss-DNA (Pope & Sigman, 1984; Sigman et al., 1985).

Table I: Reactivity of Chemical Reagents with Different Combinations of Oligonucleotides^a

reagent	[MgCl_2]	[NaCl]	aps	ps	ss ^b
Os ₂ py	low	low	-	- ^c	++
	low	high	+	++++ ^d	+++
	high	low	-	-	++
DEPC	low	low	+ ^e	+ ^e	++
	low	high	+ ^f	+ ^f	+
KMnO_4	low	low	+	+	+++
	low	high	++	+++	+++
$(\text{OP})_2\text{Cu}^+$	low	low	+++ ^g	-	+

^aRelative activities at 22 °C: +++, very high; ++, high; +, moderate; -, none or very low. ^bSingle strands D1, D2, D3, and D4 or nonpairing combinations D2 + D3 and D1 and D4. ^cBoth the ps and aps duplexes display a very low sensitivity to Os₂py under these conditions, but slight reactivity can be detected for the ps duplex on the original autoradiographs. ^dHigh activity attributable to reaction with single strands formed upon dissociation of the duplex. ^eLow reactivity, but the modification patterns of the same strand involved in the parallel or antiparallel duplex formation are different. ^fSlightly decreased reactivity at higher salt concentration. ^gPreferred cutting sites are the TAT steps.

Our results obtained with ss oligonucleotides most likely reflect formation of transient and partially paired duplexes in solution. Antiparallel species with 14 or less paired bases can be drawn for the D series of oligonucleotides (Rippe et al., 1989).

CONCLUSIONS

The spectroscopic, thermodynamic, and substrate properties of ps duplexes formed from synthetic oligonucleotides, including those used in this study (D1, D2, D3, and D4), have been described in previous reports (Ramsing & Jovin, 1988; Rippe et al., 1989; Ramsing et al., 1989; Rippe & Jovin, 1989; Jovin et al., 1990). It has also been shown that parallel-stranded DNA can be formed by using oligonucleotides with incorporated dG and dC nucleotides (Rippe et al., 1990). Next to B-DNA, ps-DNA seems to be the most stable right-handed

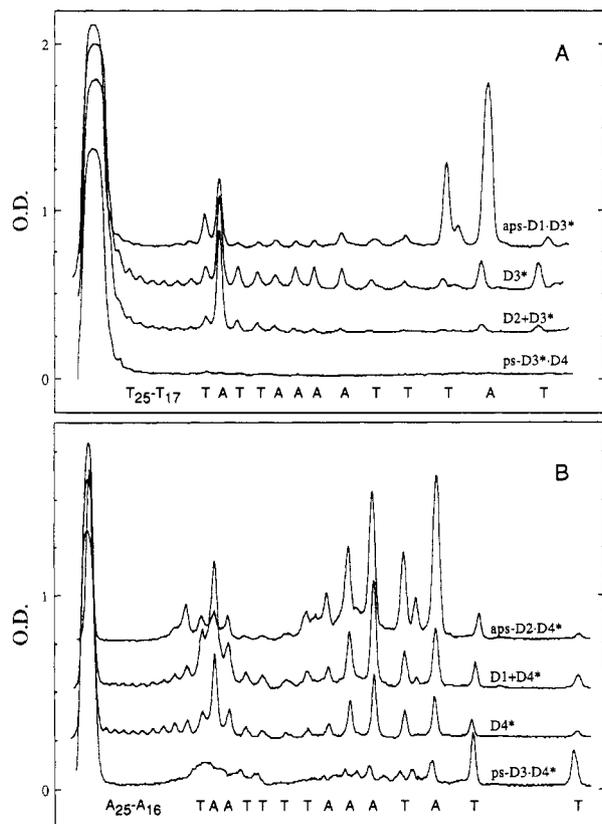


FIGURE 7: Copper-phenanthroline $[(OP)_2Cu^+]$ cleavage pattern of ps, aps, and nonpairing combinations of labeled D3* and D4*. The figure shows densitometric scans of an autoradiograph of the DNA sequencing gel on which the reaction products were separated. The assignment of the peaks was derived from the migration of the Maxam-Gilbert sequencing products. (A) Strand D3* labeled. $(OP)_2Cu^+$ cleavage pattern of ps-D3*.D4, aps-D1.D3*, D2 + D3* (nonpairing combination), and D3*. (B) Strand D4* labeled. $(OP)_2Cu^+$ cleavage pattern of ps-D3.D4*, aps-D2.D4*, D1 + D4* (nonpairing combination), and D4*.

helical conformation known to date which can be readily achieved under physiological conditions. The dramatic structural differences of ps duplexes as compared with B-DNA are consistent with the chemical modification properties monitored with single-nucleotide resolution as described in this report. All of the chemicals employed can be used to ascertain if a given strand is single-stranded or is involved in aps or ps duplex formation, as summarized in Table I. Os.py has a slightly higher reactivity for the ps than the aps duplex at low salt concentration. Under high-salt conditions, the same reagent overreacts with the ps but not the aps duplex, an effect we attribute to a destabilization of the helix and a parallel increase in single-strand reactivity. DEPC distinguishes between the ss and duplex states of DNA and exhibits different patterns of modification within a given sequence in the ps and aps conformations. $KMnO_4$ exhibits a selectivity similar to that of Os.py. Finally, $(OP)_2Cu^+$ reveals specific cleavage patterns in the case of aps-DNA but does not attack ps-DNA, presumably because the latter conformation lacks the specific minor groove stereochemistry of B-DNA.

The synthetic oligonucleotides of the D series (D1, D2, D3, and D4) are particularly suitable for studies of structural properties reflected at the level of single residues. We have demonstrated that the modification property of a given sequence can be determined in the single-stranded state and compared to its chemical susceptibility either in the aps or in the ps duplex. Thus, this work further evaluates and extends the methodological arsenal that can be applied to distinguish

between aps-, ps-, and ss-DNAs in solution, particularly in the case of defined oligonucleotides. It is worth noting that the present approach may be useful in a more general way since it reduces substantially the amount of material required for distinguishing between conformations below that employed with spectroscopic techniques. In addition, it may serve to derive certain thermodynamic parameters such as T_m values in cases where a spectroscopic determination is not feasible because of the presence of highly ultraviolet-absorbing substances (e.g., solvents and drugs) in the sample.

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Registry No. D1 + D2, 123411-22-3; D3 + D4, 123411-23-4; D1 + D3, 123411-24-5; D2 + D4, 123411-25-6; D1, 123411-18-7; D2, 123411-19-8; D3, 123411-20-1; D4, 123411-21-2; DEPC, 1609-47-8; $(OP)_2Cu^+$, 17378-82-4; $KMnO_4$, 7722-64-7; NaCl, 7647-14-5; $MgCl_2$, 7786-30-3.

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Metabolism of the Carbocyclic Nucleoside Analogue Carbovir, an Inhibitor of Human Immunodeficiency Virus, in Human Lymphoid Cells[†]

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ABSTRACT: Carbovir (CBV) is a highly selective carbocyclic nucleoside inhibitor of HIV replication in human lymphocytes and is potentially useful in the treatment of AIDS [Vince et al. (1988) *Biochem. Biophys. Res. Commun.* 156, 1046-1053]. Using human lymphoid cells severely deficient in nucleoside kinases, we were able to identify the route of activation of CBV metabolism. The present studies have demonstrated that CBV is anabolized to the mono-, di-, and triphosphates and to guanosine 5'-triphosphate in CCRF-CEM cells. Conversion to GTP amounted to 15-20% of the total analogue nucleotides formed in the cells and may arise from CBV through depurination and salvage via HGPRT. Evidence was obtained that neither deoxycytidine kinase, adenosine kinase, or mitochondrial deoxyguanosine kinase is primarily involved in the initial step of phosphorylation of CBV in CCRF-CEM cells. In contrast, earlier studies [Johnson & Fridland (1989) *Mol. Pharmacol.* 36, 291-295] showed that a cytosolic 5'-nucleotidase catalyzes the activation of CBV to the monophosphate. Other biochemical effects examined showed that the nucleobases hypoxanthine and adenine, but not guanine, their respective nucleosides, and the dideoxynucleosides 2',3'-dideoxyinosine, 2',3'-dideoxyguanosine, and 3'-azido-3'-deoxythymidine produced significant increased accumulation of CBV nucleotides in CEM cells. The exact mechanism for this potentiation of CBV phosphorylation has not been elucidated but may be due to a modulating effect of intracellular nucleotides on 5'-nucleotidase activity.

The carbocyclic nucleoside analogue carbovir (carbocyclic 2',3'-didehydro-2',3'-dideoxyguanosine, NSC-614846, CBV)¹ is a novel guanosine derivative which has shown potent in vitro

activity against the human immunodeficiency virus, the etiological agent of AIDS. CBV inhibits HIV replication and

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¹ Abbreviations: CBV, carbovir; HIV, human immunodeficiency virus; AIDS, autoimmune deficiency syndrome; AZT, 3'-azido-3'-deoxythymidine; ddC, 2',3'-dideoxycytidine; TP, 5'-triphosphate; ddNTP, 2',3'-dideoxynucleoside 5'-triphosphate; HPLC, high-pressure liquid chromatography; HGPRT, hypoxanthine-guanine phosphoribosyltransferase; ddG, 2',3'-dideoxyguanosine; dGuo, 2'-deoxyguanosine; PNP, purine nucleoside phosphorylase.