

STRUCTURES OF POLY(dA-dT,ip⁵dU) CONTAINING VARIOUS SMALL AMOUNTS
OF THE ANTIHERPETIC 5-ISOPROPYL-2'-DEOXYURIDINE

János Sági, Jitka Štokrová¹, Michaela Vorlíčková², Alena Spánová²,
Jaroslav Kypr², Éva Ruff and László Ötvös

Central Research Institute for Chemistry, Hungarian Academy of
Sciences, H/1525 Budapest, PO Box 17, Hungary

¹Institute of Molecular Genetics, Czechoslovak Academy of
Sciences, CS/16637 Prague, Czechoslovakia

²Institute of Biophysics, Czechoslovak Academy of Sciences,
CS/61265 Brno, Czechoslovakia

Received February 10, 1992

Three different concentrations of the antiherpetic agent 5-isopropyl-2'-deoxyuridine (ip⁵dU) were introduced into the synthetic DNA poly(dA-dT) to analyze resulting copolymers by electron microscopy, UV absorption and CD spectroscopy. The poly(dA-dT,ip⁵dU) containing 1.3 and 4.3% ip⁵dU did not much differ from the parent poly(dA-dT) but poly(dA-dT,ip⁵dU) with 7.1% ip⁵dU behaved in an unusual way. Results are explained by the notion that if bulky isopropyls occur sufficiently close to each other then stable hairpins protruding from the double helix are formed, presumably to accommodate the ip⁵dU-s into the loops.

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The thymidine analogue 5-isopropyl-2'-deoxyuridine (ip⁵dU) is a commercially available drug (HEVIZOS) (1) for the local treatment of patients with herpes labialis, zoster or progenitalis (2). Antiherpes action of 5-substituted dU-s at the molecular level is assumed to originate from i/ the inhibition of viral DNA polymerase by the modified triphosphate formed in the virus-infected cells by the herpes-encoded thymidine kinase, and ii/ from the altered biochemical properties of the newly synthesized herpesvirus DNA containing incorporated 5-substituted dUMP

ABBREVIATIONS

5-isopropyl-2'-deoxyuridine is abbreviated as ip⁵dU. The ip⁵ is also used in ip⁵dUMP, ip⁵dUTP and the copolymer poly(dA-dT,ip⁵dU). UV, CD and EM stand for ultraviolet, circular dichroism and electron microscopy, respectively.

molecules (3-6). This modified viral DNA was modelled earlier by poly(dA-dT) where dTMP could partially be replaced by ip⁵dUMP (7). Structure of poly(dA-dT,ip⁵dU) is characterized in the present study by EM, UV absorption and CD spectroscopies in order to elucidate contribution of conformation to the altered biochemical properties observed earlier (7). For this purpose poly(dA-dT,ip⁵dU)-s containing 0, 1.3, 4.3 and 7.1 % ip⁵dU were synthesized by *E. coli* Klenow DNA polymerase enzyme.

EXPERIMENTAL

Synthesis

Poly(dA-dT) and poly(dA-dT,ip⁵dU)-s were prepared by methods applied earlier (7). Reaction mixtures contained 95%-5%, 75%-25% and 25%-75% of dTTP : ip⁵dUTP to obtain poly(dA-dT,ip⁵dU)-s with about 1%, 4% and the maximum amount (7-8%) of ip⁵dUMP, based on the results of analytical reactions (7). In the 25 ml mixtures 80-120 OD_{max} units of high-molecular weight copolymers, excluded on Bio-Gel A-5m (Bio-Rad Labs) column, were formed.

Nucleoside compositions were determined from the enzymic hydrolysates of the polymers by HPLC (ISCO, Supersil 3 μ m Si 4x100 mm column, EtAc-MeOH {98%-2%} eluent, 0.6 ml/min). Poly(dA-dT,ip⁵dU)-s contained 1.3%, 4.3% and 7.1% ip⁵dU, that is 2.6, 8.6 and 14.2% of dT were replaced by ip⁵dU.

Heterogeneity of the copolymers synthesized was determined by neutral and alkaline agarose (0.5 %) gel electrophoresis with 18 marker DNAs ranging from 468 to 48502 base pairs. Neutral distribution of lengths and the main band in brackets were₅ as follows for poly(dA-dT) with 0, 1.3, 4.3 and 7.1% ip⁵dU: >100,000 - 810 (10,500), 5,200 - 300 (1800), 9,300 - 290 (3,200) and 3,000 - 270 (740), respectively.

Methods

UV absorption-temperature melting profiles were determined essentially as described earlier (8), with a heating rate of 0.5 °C/min. Spectra were recorded on a Hewlett-Packard 8452A diode array spectrophotometer.

CD measurements were carried out using a Jobin-Yvon dichrograph Mark IV calibrated with isoandrosterone. Concentrations of the polymers, calculated by using $\epsilon(P) = 6640 \text{ M}^{-1}\text{cm}^{-1}$ for poly(dA-dT) and the poly(dA-dT,ip⁵dU)-s, ranged within 30-50 μ g/ml.

The samples for EM analysis were taken from stock solutions used for CD measurements. Each polydeoxynucleotide was analysed by three techniques, namely modified aqueous, formamide and BAC, as described previously (9-11). Electron micrographs were recorded at magnification of 10,000 in a JEOL JEM 1200EX microscope operating at 60 kV, enlarged fivefold.

RESULTS

Poly(dA-dT) analogues were first studied by the absorbance versus temperature melting profiles. Figure 1 shows that only thermal transition of poly(dA-dT,ip⁵dU) of 7.1% ip⁵dU was unusual,

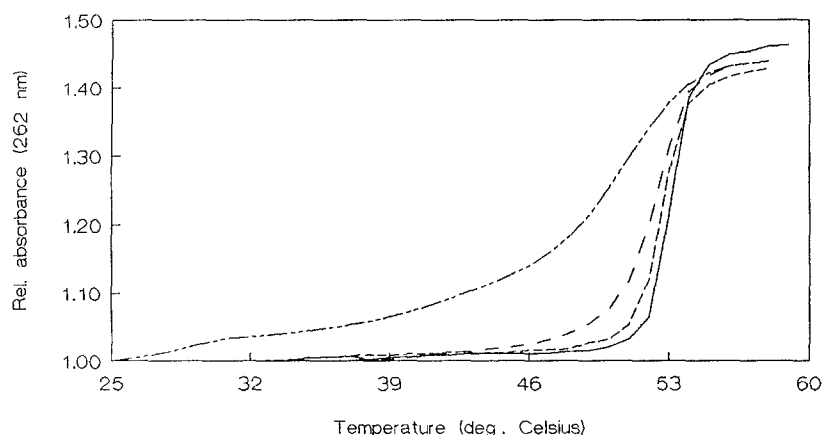


Figure 1. UV melting profiles of poly(dA-dT,ip⁵dU)-s with 0%, —; 1.3%, ----; 4.3%, — · —; and 7.1% ip⁵dU, ····. Melting profiles were determined in 25 mM NaCl, 10 mM Tris.HCl (pH 7.2), 1 mM EDTA solution. Based on the first derivative of the transition curve, T_m values of 53.0, 52.9, 52.5 and 50.9°C were obtained for poly(dA-dT,ip⁵dU)-s with 0, 1.3, 4.3 and 7.1% ip⁵dU, respectively. T_m 's were reproducible within $\pm 0.2^\circ\text{C}$. Thermal hyperchromicities at 262 nm were between 43.1 and 45.8%. Widths of transitions, defined by the temperature range between 25 and 75% of the thermal hyperchromicity, were 1.2, 1.5, 2.3 and 7.6°C for polydeoxynucleotides of the above sequence.

its UV absorption at 260 nm increased gradually far before the main cooperative part of melting of the duplex. Substitution caused only small decrease in the stability of the duplexes.

CD spectrum of poly(dA-dT) exhibits temperature-induced changes prior to melting. In addition, denaturation is accompanied by a positive band red-shift (12). We observed similar changes with poly(dA-dT,ip⁵dU) molecules containing 0, 1.3 or 4.3% ip⁵dU (Figure 2, right panel). However, the positive band of poly(dA-dT,ip⁵dU) of 7.1% ip⁵dU-content showed a different temperature-dependence and the melting itself exhibited only a low degree of cooperativity. The red-shift of the positive band maximum was cooperative with poly(dA-dT)-s up to 4.3% ip⁵dU but it turned into a gradual dependence where ip⁵dU was 7.1% (Figure 2, panel right, bottom).

Further insight into the perturbations was obtained by EM, though the polydeoxynucleotides were inhomogeneous in length (see Experimental) that makes detailed statistical analysis of the EM data impossible. Three different EM techniques were used: the "aqueous" (9), the "formamide" (9,11) and the "BAC" spreading technique (10,11). The "aqueous" technique (9) is presented only

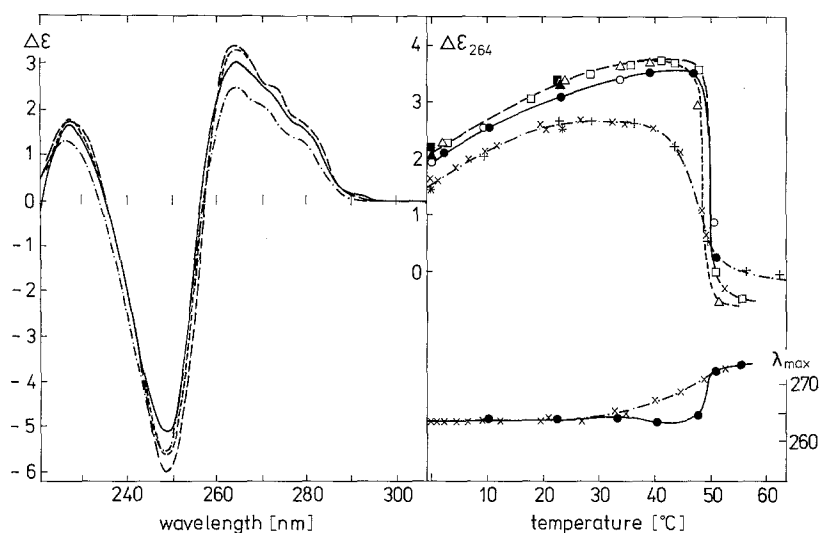


Figure 2.

Left panel: CD spectra of poly(dA-dT), —; and poly(dA-dT,ip⁵dU)'s with ip⁵dU-content of 1.3%, — — —; 4.3%, - - -; and 7.1%, - · - · -; determined in 25 mM NaCl, 10 mM Tris.HCl (pH 7.3), 1 mM EDTA at 23.2°C.

Right panel: Temperature-induced changes in the CD spectra monitored by ellipticity at 264 nm (top of right panel) and by the shift of the positive band maximum (bottom). Poly(dA-dT), —; and analogues with ip⁵dU-content of 1.3%, — — —; 4.3%, - - -; and 7.1%, - · - · -, respectively. Different symbols on the same curves, like o and ●, □ and ■, Δ and ▲, or x, + and * represent repeated determinations.

here which visualized poly(dA-dT) as filaments of variable lengths containing long branches (Figure 3A). Each technique used showed that the introduction of ip⁵dU into poly(dA-dT) made the macromolecules shorter and that the number of non-branched molecules decreased whereas the number of branches per molecule increased with the increasing ip⁵dU content (Figures 3B-3D).

DISCUSSION

The crystal structure of ip⁵dU shows that one of the two isopropyl methyls protrudes significantly (1.59 Å) from the attached base plane, making the formation of regular B-DNA impossible (13). However, the thermal melting profiles of poly(dA-dT,ip⁵dU) with 4.3% or less ip⁵dU were normal (Figures 1 and 2). In other words, if on average every 24th nucleotide in poly(dA-dT) is replaced by ip⁵dUMP, poly(dA-dT) can buffer the perturbations introduced by the bulky isopropyl groups in the double helix major groove. Poly(dA-dT) is either different enough

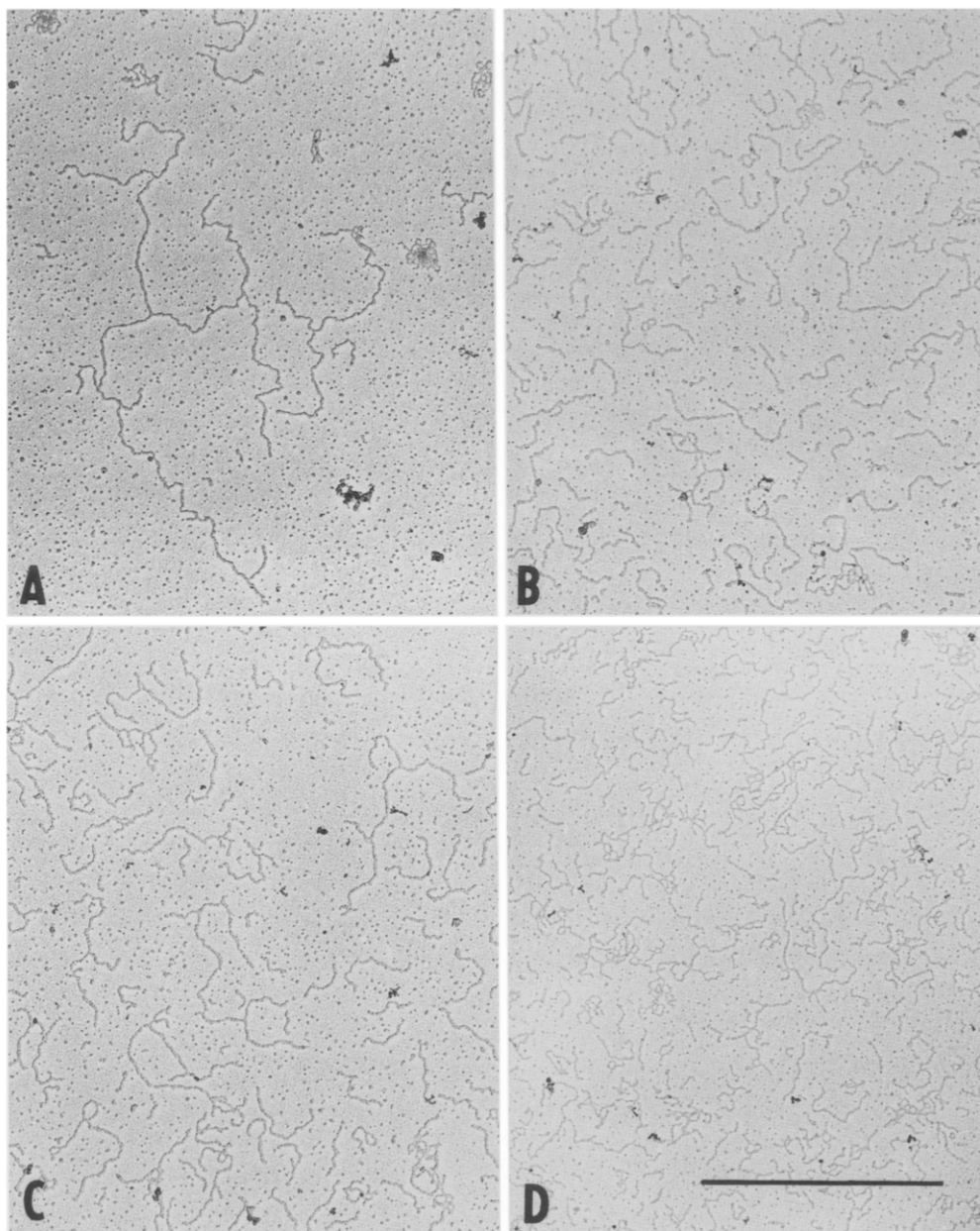


Figure 3.
Electron microscopic pictures of poly(dA-dT,ip⁵dU)'s containing 0%, A; 1.3%, B; 4.3%, C; and 7.1% ip⁵dU, D obtained by the "aqueous" technique. The bar is 1 μ m.

from regular B-DNA (14,15) or flexible enough (16) to accommodate solitary ip⁵dU-s.

If the replacement occurs with a higher frequency, i.e. each 14th nucleotide is replaced by ip⁵dUMP (7.1% ip⁵dU), the melting

profiles become qualitatively different. This can be explained by independent melting out of small molecule parts before the melting of the whole molecule takes place. Incorporation of ip^5dU is probably not uniform. The DNA strands where several $\text{dA-ip}^5\text{dU}$ dimers follow each other, even if DNA polymerase could synthesize it, the duplex can no longer tolerate their presence. Sequential irregularities rather tend to bulge out and form hairpins where ip^5dU -s are assumed to be in the loop. These hairpins seem to be stable as compared to hairpin structures in $\text{poly}(\text{dA-dT,ip}^5\text{dU})$ -s with 4.3% and lower ip^5dU -content. The latter stem-loops, visualized only by EM, are metastable in solution and thus are not shown by the T_m -curves. So far, hairpins have been found to be stable in response to negative supercoiling in circular DNA molecules (17,18), in artificial linear DNA constructs (12) and in oligodeoxynucleotide structures (e.g. 19) or in RNAs (20).

$\text{Poly}(\text{dA-dT,ip}^5\text{dU})$ molecules become shorter as the ip^5dU -content increases. Presumably the polymerase enzyme dissociates more frequently from the distorted replication complex than from $\text{poly}(\text{dA-dT})$. Shortening is shown by EM (Figure 3) and agarose gel electrophoresis (Experimental). EM also clearly shows that the number of branches increases in $\text{poly}(\text{dA-dT})$ with the amount of ip^5dU . Thus, incorporated ip^5dU induces branching. Branches in $\text{poly}(\text{dA-dT})$ are long-stem stem-loops structures. Up to 4.3% ip^5dU -content stem-loops are metastable in $\text{poly}(\text{dA-dT,ip}^5\text{dU})$ and can be seen only on the EM pictures. Spectroscopy shows only the more stable hairpin-loop structures induced by 7.1% ip^5dU .

The presence of stable hairpins in $\text{poly}(\text{dA-dT,ip}^5\text{dU})$ containing 7.1% ip^5dU is well compatible with its strongly enhanced sensitivity to the single-strand-specific nuclease S1, as compared to $\text{poly}(\text{dA-dT})$ (7). The increased sensitivity can be explained by the presence of real single-stranded regions in $\text{poly}(\text{dA-dT,ip}^5\text{dU})$, occurring in the hairpin loops, similarly as observed with oligodeoxynucleotide loop structures (19). In this way, herpes-virus DNA synthesized *in vivo* in the presence of the antiherpetic ip^5dU may be destabilized against some cellular endonucleases if a sufficient level of substitution by ip^5dUMP for dTMP is reached.

ACKNOWLEDGMENTS

Authors (J.S.) wish to thank Dr. K. Ganzler for the HPLC analyses. This work was supported in part by grants No. 1015 and 1754 from the Hungarian Research Fund (OTKA).

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