

Structural Basis for Antisense and Antisense Duplexes with Modified Nucleotides

Ella Czarina Magat Juan^a, Takeshi Kurihara^a, Jiro Kondo^a, Takanori Ito^b, Yoshihito Ueno^c, Akira Matsuda^b and Akio Takénaka^a,
^a*Graduate School of Bioscience and Biotechnology, Tokyo Institute of Technology, Yokohama, Japan.* ^b*Graduate School of Pharmaceutical Sciences, Hokkaido University, Sapporo, Japan.* ^c*Faculty of Engineering, Gifu University, Gifu, Japan.* E-mail: ella@bio.titech.ac.jp

Oligonucleotides containing polyamines are currently being evaluated as potential antisense and antisense compounds. Those with 5-(*N*-aminohexyl)carbamoyl-2'-deoxyuridine (^NU) and its 2'-*O*-methyl derivative (^NU_m) exhibit improved nuclease resistance. Furthermore, these nucleotides stabilize duplex formation of the modified DNA and its target DNA or RNA strand. X-ray structures of these duplexes have shown good correlation between the conformational changes and the observed chemotherapeutic properties.

The amide groups of the modified uracil bases form six-membered rings through the intramolecular NH---O4 hydrogen bonds, so that the aminohexyl chains protrude into the major grooves. Some of the terminal ammonium groups are involved in intra-duplex interactions with phosphate oxygen anions, whereas the others interact with those of the adjacent duplex. Such interactions contribute to the stability of duplex formation. The 2'-*O*-methyl modification in ^NU_m shifts the ribose ring toward the C3'-*endo* conformation and influences duplex stability. Observed changes in the dimensions of the minor grooves and in the hydration structures are well correlated to nuclease resistance.

Keywords: antisense, antisense, crystal structure