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## Nucleosides, Nucleotides and Nucleic Acids

Publication details, including instructions for authors and subscription information:

<http://www.informaworld.com/smpp/title~content=t713597286>

### $\alpha$ -LNA ( $\alpha$ -D-Configured Locked Nucleic Acid)

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Online publication date: 09 August 2003

**To cite this Article** Christensen, Nanna K. , Petersen, Michael , Vester, Birte and Nielsen, Poul(2003) ' $\alpha$ -LNA ( $\alpha$ -D-Configured Locked Nucleic Acid)', *Nucleosides, Nucleotides and Nucleic Acids*, 22: 5, 1143 — 1145

**To link to this Article:** DOI: 10.1081/NCN-120022821

**URL:** <http://dx.doi.org/10.1081/NCN-120022821>

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## $\alpha$ -LNA ( $\alpha$ -D-Configured Locked Nucleic Acid)

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### ABSTRACT

Two pyrimidine  $\alpha$ -LNA nucleoside monomers have been synthesised and incorporated into  $\alpha$ -configured oligonucleotides. A fully modified mixed  $\alpha$ -LNA sequence displays unprecedented parallel stranded hybridisation with complementary RNA and a remarkable selectivity for RNA over DNA. Modelling shows  $\alpha$ -LNA:RNA to form an extended duplex with a very broad major groove.

*Key Words:* Locked nucleic acid; Parallel duplexes; Selective RNA-recognition.

LNA (Locked Nucleic Acid) is a nucleic acid analogue that displays unprecedented recognition of both single stranded DNA and RNA in all-modified LNA-oligomers as well as in LNA/DNA and LNA/RNA-mixmers.<sup>[1,2]</sup> The LNA monomers are bicyclo[2.2.1] nucleosides locked in an N-type conformation.<sup>[1,2]</sup>  $\alpha$ -DNA, i.e.,  $\alpha$ -D-configured oligodeoxynucleotides, forms stable parallel stranded duplexes with complementary DNA and RNA.<sup>[3]</sup> Conformationally restricted analogues of

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**Table 1.** Buffer: Na<sub>2</sub>HPO<sub>4</sub> (10 mM), NaCl (100 mM), EDTA (0.1 mM), pH 7.0; 1.5  $\mu$ M of each strand. All detected  $T_M$ 's were enhanced 3–6°C in a high salt buffer (NaCl, 750 mM). Boldface letters  $\sim\alpha$ -LNA.

$T_m$ ( $\Delta T_m$ per modification)/°C		Parallel DNA	ap DNA	Parallel RNA	ap RNA	Mismatch RNA (A to C)
$\alpha$ -DNA	5'- $\alpha$ - <sup>m</sup> CT <sup>m</sup> C <sup>m</sup> CTT <sup>m</sup> CTTT	43.0	< 10	32.0	< 10	
$\alpha$ -DNA/	5'- $\alpha$ - <sup>m</sup> CT <sup>m</sup> C <sup>m</sup> CTT <sup>m</sup> CTTT	30.5 (–12.5)	< 10	26.0 (–6.0)	< 10	
$\alpha$ -LNA	5'- $\alpha$ - <sup>m</sup> CT <sup>m</sup> C <sup>m</sup> CTT <sup>m</sup> CTTT	< 10	< 10	22.0 (–1.7)	< 10	
mixmers						
$\alpha$ -LNA	5'- $\alpha$ - <sup>m</sup> CT <sup>m</sup> C <sup>m</sup> CTT <sup>m</sup> CTTT	< 10	< 10	<b>61.5 (+2.9)</b>	< 10	47.0

$\alpha$ -DNA that contains bicyclic  $\alpha$ -configured nucleosides have been introduced.<sup>[4,5]</sup> However, these have been restricted towards S-type conformations<sup>[4,5]</sup> which are also the conformations preferred by  $\alpha$ -DNA. Subsequently,  $\alpha$ -LNA is the first  $\alpha$ -DNA analogue to be restricted in an N-type conformation due to the locked bicyclic carbohydrate skeleton.<sup>[6,7]</sup>

The successful synthesis of the  $\alpha$ -LNA thymine monomer was accomplished from diacetone- $\alpha$ -D-allose through standard steps ending with a Vorbrüggen nucleobase coupling reaction, a ring-closing reaction and, subsequently, separation of the  $\alpha$ - and ( $\beta$ -) LNA monomers.<sup>[6,7]</sup> The former was DMT-protected and converted by standard steps into two different phosphoramidites of either the thymine or the *N*-4-benzoyl-5-methylcytosine  $\alpha$ -LNA monomers.<sup>[7]</sup> These as well as the two corresponding  $\alpha$ -DNA phosphoramidites were used as building blocks for oligonucleotide sequences (Table 1).<sup>[7]</sup>

In  $\alpha$ -DNA/ $\alpha$ -LNA mixmers, the exchange of one or six  $\alpha$ -DNA monomers with  $\alpha$ -LNA monomers results in strongly decreased affinity against both DNA and RNA when compared to the unmodified  $\alpha$ -DNA sequence. On the other hand, a fully modified mixed  $\alpha$ -LNA sequence shows no recognition of DNA but a very strong recognition of RNA. This strong preference for RNA as well as for parallel over antiparallel duplex formation was confirmed by the use of a gel shift analysis. MD-simulations were performed assuming Watson-Crick base pairing (as no change in  $T_M$  was detected when lowering pH from 7 to 5) and suggested the  $\alpha$ -LNA/RNA duplex to be an extended duplex with a major groove width of 20 Å and the nucleobases of the  $\alpha$ -LNA monomers in *syn*-positions. Other duplex parameters were close to normal A- or B-type duplexes.

## ACKNOWLEDGMENTS

The Nucleic Acid Center is funded by the Danish National Research Foundation for the studies on nucleic acid chemical biology.

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