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LNA units present in $[R_P\text{-PS}]\text{-(DNA\#LNA)}$ chimeras enhance the thermal stability of parallel duplexes and triplexes formed with (2'-OMe)-RNA strands†‡

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The results of CD measurements indicate that 2–4 LNA units distributed along 12 nt P-stereodefined phosphorothioate $[R_P\text{-PS}]\text{-(DNA\#LNA)}$ chimeras impose a C3'-endo conformation on the 2'-deoxyribonucleosides. Under neutral and slightly acidic conditions homopurine $[R_P\text{-PS}]\text{-(DNA\#LNA)}$ hybridizes with 9–12 nt Hoogsteen-paired (2'-OMe)-RNA strands to form parallel duplexes, which are thermally more stable than the reported earlier analogous complexes containing LNA-free $[R_P\text{-PS}]\text{-DNA}$ oligomers ($\Delta T_m = 7^\circ\text{C}$ per LNA unit at pH 5.4). Upon addition of the corresponding Watson–Crick paired (2'-OMe)-RNA strands, parallel triplexes are formed with further increased thermal stability.

Introduction

Nucleic acid triple-helical forms have been known for sixty years.¹ Their formation seems to play a role in sequence-specific recognition of a double helix,^{2–4} as well as in chromatin organization, DNA repair, transcriptional regulation and RNA processing.^{5,6} The third strand binds to an antiparallel duplex either in a parallel or an antiparallel orientation with respect to the purine strand, utilizing either a Hoogsteen or reverse Hoogsteen hydrogen bonding scheme, respectively.⁷ Hoogsteen G-C⁺ base pairing requires protonation of the N3 nitrogen atoms in cytosines⁸ (for CMP and dCMP, pK_a of 4.3 and 4.6 was found at 25 °C, respectively⁹). Triplexes may comprise RNA and/or DNA strands, with consequences for thermal stability.^{10,11,12} Usually, the Hoogsteen paired (Hp) parallel duplexes are thermally much less stable than the Watson–Crick paired (WCp) duplexes, but their stability increases when a Hp chain is linked to a pyrimidine chain (of inverted polarity) by a 3'-3' or 5'-5' linker.^{13–16} There are a few reports on parallel stretches in native DNA^{17–19} and in *Escherichia coli* mRNA.²⁰ Hp-DNA duplexes seem to be involved in regulation of cell processes and evolution of neurodegenerative diseases.^{21–23}

Since natural DNA molecules are easily degraded by phosphodiesterases, several modifications of the sugar-phosphate backbone have been proposed. Among them, phosphorothioate analogs of DNA (PS-DNA) were found to be very useful because their electronic and steric properties are remarkably close to those

of DNA.^{24–26} However, as short as 10–12 nt PS-DNA oligomers prepared by standard chemical methods (a phosphoramidite or an *H*-phosphonate approach) exist as mixtures of hundreds or even thousands of P-diastereomers.²⁷ Developed in this laboratory an oxathiaphospholane approach^{28,29} (utilizing the OTP monomers **1**, Chart 1) allows for preparation of P-stereodefined PS-DNA.

It was found that the antiparallel duplexes formed by $[R_P\text{-PS}]\text{-DNA}$ (rDNA) or the S_P -counterparts (sDNA) with DNA or RNA oligomers were thermally less stable than analogous complexes formed by unmodified oligomers.³⁰ However, we discovered that homopurine rDNA form thermodynamically highly stable parallel duplexes RNA&rDNA (e.g. **I**, “&” indicates the parallel orientation of strands) and even more stable parallel triplexes RNA&rDNA:RNA (e.g. **II**).^{31–33}

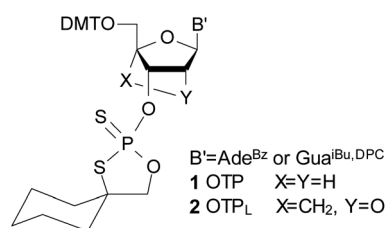
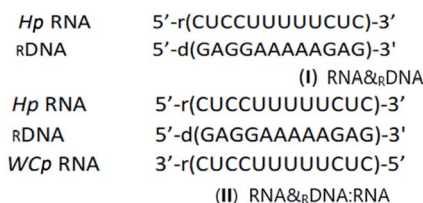


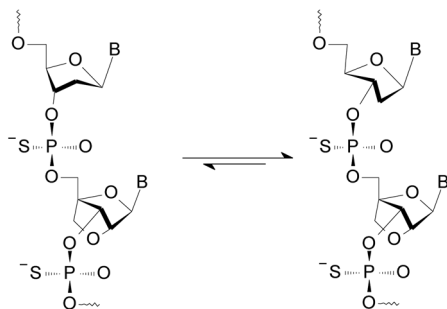
Chart 1 Structure of the 2-thio-1,3,2-oxathiaphospholane derivatives of nucleosides of the DNA (**1**) and LNA (**2**) type.

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‡ Dedicated to Professor Wojciech J. Stec on the occasion of his 80th birthday.





Scheme 1 An equilibrium leading to a C3'-endo conformation of the 2'-deoxyribonucleosides in an rDL strand (3).

Since only the homopurine [R_P -PS]-DNA oligomers form the complexes **I** and **II**, we hypothesized (based on IR measurements and molecular modeling) that the Hp-strand is anchored not only by the hydrogen bonding but also by water bridge(s) (via charge assisted hydrogen bonds³⁴) between the sulfur atoms of the R_P -phosphorothioate moieties and the O2 atoms in pyrimidine nucleobases. The overall A-like conformation of the RNA&rDNA and RNA&rDNA:RNA complexes, which is required for the interactions of that type, was imposed by the RNA strand(s), and was confirmed by CD measurements. The conformational factor is important because the complexes **I** and **II** are more stable when formed with the participation of (2'-OMe)-RNAs (m RNA), which are known for adopting more profound C3'-endo conformation than RNA molecules. To analyze further this phenomenon we employed LNA units³⁵ (often abbreviated as BNA, Bridged Nucleic Acids³⁶), in which the sugar rings adopt a profoundly rigid 3'-endo conformation with a very limited pseudorotational flexibility. Our experiments showed that 2-4 pyrimidine LNA units present in 9-12 nt Hp- m RNA strands gave rise to significantly enhanced thermal stability of the (m RNA#LNA)&rDNA complexes.³⁷ We wanted to check if a few LNA units (here denoted A_L or G_L) present in homopurine [R_P -PS]-(DNA#LNA) chimeric oligomers (rDL, **3**, Scheme 1) would impose a C3'-endo conformation on the 2'-deoxyribonucleosides in the phosphorothioate strand. If so, an energetic toll for the C2'-endo \rightarrow C3'-endo transition of the phosphorothioate central strand (Scheme 1) should be smaller giving rise to increased thermal stability of the parallel complexes. For that purpose we used recently developed P-diastereomerically pure OTP_L monomers **2** (Chart 1).³⁸

Results and discussion

Synthesized oligomers

In the codes shown in Table 1 and used throughout the text, the prefixes R/s indicate R_P -/ S_P -PS analogs, respectively, and the

§ The mechanism of stabilization (proposed by C. Hélène and also discussed by others^{12,42}), based on the hydrogen bonding between the 2'-OH group of ribose in the Hp-strand and the pro- R_P oxygen atom (or the R_P sulfur atom in our work) cannot operate because parallel triplexes and parallel duplexes formed with Hp- m RNAs are thermally more stable. Also hydrogen bonding pyrimidine-C2=O...H-S-P(O)(OR)₂, analogous to Cys-S-H...O=C in proteins,⁴³ is less likely because at neutral pH phosphorothioate diesters are fully ionized.

ending digit indicates the number of LNA units in a 12 nt oligomer. Accordingly, a code rA2 indicates an [R_P -PS]-(DNA#LNA) chimera bearing 10 2'-deoxyribonucleosides (dA, dG) and 2 A_L units, whereas sG4 stands for an [S_P -PS]-(DNA#LNA) analog bearing 4 G_L units. Four reference, non-palindromic, homopurine 12 nt oligonucleotides were synthesized (Table 1): (1) B0, a basal phosphate, LNA-free oligomer d(GGAGAAAGAGAG), (2) G4, an analog of B0 containing four G_L units, (3) rB0, an R_P -PS analog of B0; and (4) sG4, an S_P -PS analog of G4.

The [PS]-(DNA#LNA) chimeras rA2, rA4, rG2, rG4, and sG4 (all are isosequential to B0; Table 1) were manually synthesized on solid support using pure P-diastereomers of the oxathia-phospholane derivatives of DNA (**1**, OTP (ref. 29)) or LNA nucleosides (**2**, OTP_L (ref. 38)).

Homopyrimidine 12 nt WCp- and Hp- m RNAs (wR¹² and hR¹², respectively)¶ and their 9-nt congeners (wR⁹ and hR⁹), as well as the oligomers B0, G4 and 12-nt WCp-DNA (wD¹², d(CTCTCTTTCTCC), the fifth reference oligomer) were synthesized on an automated synthesizer using commercially available phosphoramidite monomers. The sequences of wR⁹ and hR⁹ are subsets of wR¹² and hR¹², truncated by three nucleotides either from the 3'-end (wR⁹) or from the 5'-end (hR⁹). The 9 nt segments located after the slashes in the sequences of homopurine oligomers (Table 1) are expected to interact with wR⁹ and hR⁹ in the mismatch-free manner, as shown in the complexes **III**, **IV**, and **V**. Other alignment of these strands is also possible but would lead to less stable, imperfect complexes, indicated as (im). As shown in Table 1, in the complexes formed by 9 nt m RNAs with rA2 or rG2, both LNA nucleotides hybridize within the 2- or 3-strand structures, whereas in rA4 and rG4 only three LNA units do this, as A_L -10 and G_L -11 (numbered from the 3'-end) stay outside. This partial involvement is denoted as rA4 {3} and rG4 {3}.

hR ⁹	5'- m (CUUUCUCUC)-3'
rDL	5'-d(GGAGAAAGAGAG)-3'
	(III)
rDL	5'-d(GGAGAAAGAGAG)-3'
wR ⁹	3'- m (CUUUCUCUC)-5'
	(IV)
hR ⁹	5'- m (CUUUCUCUC)-3'
rDL	5'-d(GGAGAAAGAGAG)-3'
wR ⁹	3'- m (CUUUCUCUC)-5'
	(V)

CD spectra for single-stranded chimeric rDL oligonucleotides

The impact of the LNA units on conformation of the rDL oligomers (single strands) was assessed by CD spectroscopy (Fig. 1). As a reference we used a spectrum recorded for wD¹², in which the 2'-deoxyribonucleosides exist in a 2'-endo form. It was found

¶ We assumed high melting temperatures of the investigated complexes, therefore, to avoid degradation of RNA oligomers at elevated temperatures we decided to use (2'-OMe)-RNA oligomers.



Table 1 Codes, sequences, and MALDI-TOF MS characteristics of the oligomers. Only 9-nt segments shown in the sequences after the slashes are involved in the interactions with wR⁹ and hR⁹

Code	Sequence (5' → 3' direction)	M. mass calc.	m/z found ^a
Reference	B0	[PO]-d(GGA/GAAAGAGAG)	3793
	G4	[PO]-d(GG _L A/G _L AAAG _L AG _L AG)	3905
	rB0	[R _p -PS]-d(GGA/GAAAGAGAG)	3969
	sG4	[S _p -PS]-d(GG _L A/G _L AAAG _L AG _L AG)	4081
	wD ¹²	[PO]-d(CTCTCTTCTCTCC)	3498
rDL	rA2	[R _p -PS]-d(GGA/GA _L AAGA _L GAG)	4025
	rA4	[R _p -PS]-d(GGA _L /GAA _L AGA _L GA _L G)	4081
	rG2	[R _p -PS]-d(GGA/G _L AAAG _L AGAG)	4025
	rG4	[R _p -PS]-d(GG _L A/G _L AAAG _L AG _L AG)	4081
			4075.5
^m RNA	wR ¹²	(2'-OMe)-CUCUCUUUCUCC	3774
	hR ¹²	(2'-OMe)-CCUCUUUCUCUC	3774
	wR ⁹	(2'-OMe)-CUCUCUUUC	2814
	hR ⁹	(2'-OMe)-CUUUCUCUC	2814

^a The relevant mass spectra for rDL and sG4 are given in Fig. S1–S5 (ESI).

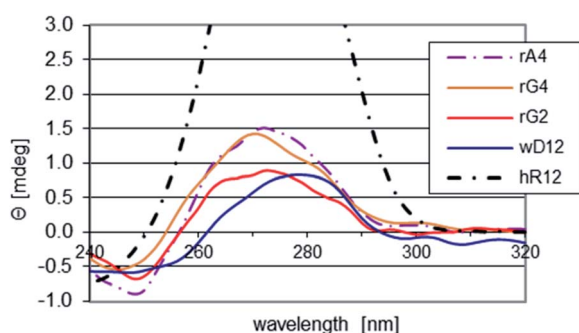


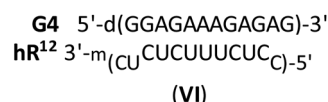
Fig. 1 CD spectra for the chimeric rDL and the reference hR¹² and wD¹² oligomers, recorded at pH 5.4 at room temperature.

that in the spectra recorded for rA4, rG2, and rG4 the isoelliptic points shifted from 263 nm to 255–257 nm, so the conformations changed towards that observed for the ^mRNA oligomer hR¹² (the isoelliptic point 250 nm, λ_{max} = 273 nm, Θ = 5.66 mdeg), in which the nucleosides adopt a 3'-endo form. That change is more developed for rA4 and rG4 as the intensities of the bands around 270 nm are *ca.* 50% higher than for rG2.

Melting and CD experiments in pH 7.2 buffer

For the annealing/melting and CD experiments, equimolar amounts of the homopurine oligomers (rDL, sG4, or the reference oligomers) and the 12 nt ^mRNAs (wR¹² and/or hR¹²) or the reference wD¹² were mixed. The temperatures of association (*T*_a, determined during annealing) and melting (*T*_m) were determined using the first order derivative method. The hysteresis values (*T*_m–*T*_a) for the two component mixtures did not exceed 3 °C, unless otherwise stated (Table 2).

The complex formed by the phosphate oligomer G4 with hR¹² was less stable than G4:wR¹² (*T*_m = 61 °C and 64 °C, respectively; Fig. S6, ESI†) and this indicates the formation of an imperfect duplex (VI).



The WCp duplex sG4:wR¹² was more stable than B0:wR¹² (Δ*T*_m = 11 °C) thus, the earlier identified stabilizing effect of the LNA units³⁹ overrode the commonly observed destabilizing effect of the phosphorothioate modification (*vide supra*³⁰). The *T*_m values found for the intended hR¹²&sG4 duplex (52 °C) and the hR¹²&sG4:wR¹² triplex (60 °C) were close to that for sG4:wR¹² (57 °C), indicating that the parallel complexes were not formed.

As mentioned earlier, rDNA or sDNA and WCp DNA or RNA matrices form the corresponding antiparallel duplexes of significantly lower thermal stability than natural DNA. Thus, taking into account *T*_m = 50 °C noted for the putative rB0:wR¹² complex, which was higher than for B0:wR¹² (*T*_m = 46 °C), and *T*_m = 62 °C noted for the mismatch-free hR¹²&rB0, one can conclude that the imperfect (im)wR¹²&rB0 complex (VII, 3 mismatches) was formed, rather than the rB0:wR¹². This phenomenon is supported by the relevant CD spectrum, in

Table 2 Melting temperatures [°C] for the complexes formed with the WCp and/or Hp 12 nt ^mRNA matrices. The samples were dissolved in 10 mM Tris–HCl, 100 mM NaCl, 10 mM MgCl₂ buffer (pH 7.2)

Homopurine oligomer	wR ¹²	hR ¹²	hR ¹² + wR ¹²
B0	51	47	53
G4	64	[57] ^b /61 (VI) ^a	[65]/69
rB0	50 (VII)	62 (I)	[67]/>80 (II)
rA2	[65]/73	[64]/79	[65]/>80 (II)
rG4	77 (VIII)	75 (I)	[80]/>80 (II)
sG4	57	52	60

^a Roman numerals in parentheses refer to the proposed structures of complexes. ^b Numbers in brackets indicate the temperature of association found during the annealing in the cases where the hysteresis value exceeded 3 °C.



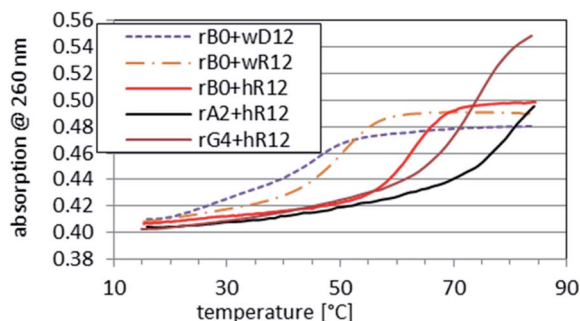
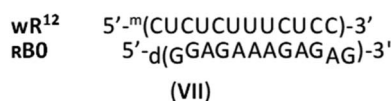


Fig. 2 Melting profiles recorded for rB0 ([R_P-PS]-DNA, no LNA units), rA2, and rG4 mixed with hR¹² at a 1 : 1 molar ratio in pH 7.2 buffer. The profiles for rB0 mixed with wD¹² or wR¹² are given as the references.

which an intense negative signal around 210 nm (similar to that for hR¹²&rB0) was noted (Fig. S7, ESI†) and indicates an enormous stabilizing effect of the [R_P-PS]-modification.



As expected, the parallel duplexes hR¹²&rA2 and hR¹²&rG4 were thermally substantially more stable ($T_m > 75^\circ\text{C}$, Fig. 2) than hR¹²&rB0 ($\Delta T_m = 17^\circ\text{C}$ and $\Delta T_m = 13^\circ\text{C}$, respectively). The latter effect, formally generated by 4 G_L units, was assumed to be more marked, but neutral pH apparently renders the C&G_L interactions less effective.

Interestingly, whereas (im)wR¹²&rB0 (VII) was less stable than hR¹²&rB0 ($\Delta T_m = -12^\circ\text{C}$, *vide supra*), a T_m value for rG4:wR¹² was higher than for hR¹²&rG4 (77°C vs. 75°C). This phenomenon might be explained by strong stabilizing effect of four G_L units at the WC interface, analogously to sG4:wR¹². However, the CD spectrum for rG4:wR¹² (Fig. 3, a green line) contained an intense negative signal around 210 nm and the isoelliptic point at 249 nm, and was similar to that for hR¹²&rG4:wR¹² (an orange line). Therefore, we suggest an additional favorable factor, where the G_L units effectively

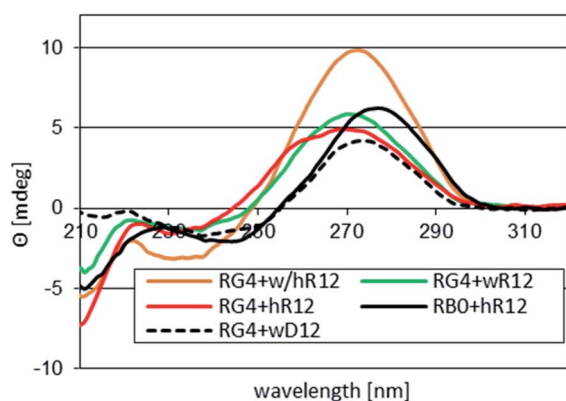
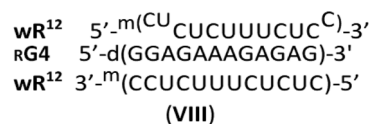


Fig. 3 CD spectra recorded for complexes formed by rG4 with 12 nt mRNAs in pH 7.2 buffer. Profiles for hR¹²&rB0 ([R_P-PS]-DNA, no LNA units) and for rG4:wD¹² are given as the references.

promote the C2'-endo \rightarrow C3'-endo conformational change of the central strand and allow formation of imperfect VIII.



This process seems to occur for rA2 mixed with wR¹² or hR¹² because the observed $T_m = 73^\circ\text{C}$ or 79°C , respectively, were close to those for rG4:wR¹² and hR¹²&rG4 ($T_m = 77^\circ\text{C}$ and 75°C , respectively). This suggestion is supported by an 8°C hysteresis, which for a bimolecular association should be very low (typically $<3^\circ\text{C}$), whereas it exceeds $13\text{--}15^\circ\text{C}$ for the trimolecular systems hR¹²&rB0:wR¹² and hR¹²&rA2:wR¹². Unlike commonly known triplexes containing a PO-DNA central strand, which dissociate in two discrete steps, those formed by rDNA undergo a single-step melting transition, followed by return of rDNA to the C2'-endo conformation. Thus, the reverse process is “more trimolecular” and in terms of entropy it is less favored than a bimolecular one, giving rise to the hysteresis. For wR¹²&rG4:wR¹² and hR¹²&rG4:hR¹² the observed hysteresis values were low because 4 G_L units substantially shift the conformation of the central PS-strand in the C3'-endo direction and make it “ready” for rapid association.

At pH 7.4, the triple-stranded complex hR¹²&rB1:wR¹² and its congeners containing rA2 or rG4 were thermally so stable that the inflection points could not be determined (the melting profiles not shown).

The above presented data indicate, that at neutral pH, the LNA units present in the rDL oligomers make the Hoogsteen interactions with mRNAs significantly stronger compared to the analogous rDNA. However, concomitantly, the rigid LNA nucleotides enhance hybridization at the WC interface strongly enough to promote the formation of the imperfect triplex VIII. This hybridization may result in the binding of non-target RNAs. This loss of specificity may be avoided using rDL oligomers with Watson-Crick base pairings being hampered due to the presence of N6-Me deoxyadenosine (m⁶dA) units. We reported recently that the m⁶dA units present in rDNA stabilize (by up to 4.5°C per modified unit) parallel duplexes formed by with Hp mRNAs compared to the analogous reference duplex containing only unmodified nucleobases, and prevent the formation of the corresponding parallel triplexes.⁴⁰

Melting and CD experiments performed using pH 5.4 buffer

Because the T_m values for complexes of the rDL oligomers with wR¹² or hR¹² formed in pH 5.4 buffer could not be determined, next melting experiments were done for the complexes with wR⁹ and/or hR⁹.

Compared to the reference rB0:wR⁹, the T_m values for complexes formed by rDL and sG4 with wR⁹ increased by $2\text{--}10^\circ\text{C}$ (Table 3 and Fig. 4) and this increase may be attributed to the stabilizing effect of the LNA nucleosides.

More interesting results were obtained for the complexes formed with hR⁹. The complex formed with sG4 (hR⁹&sG4{3}) or (im)sG4:hR⁹ was 25°C less stable than hR⁹&rG4{3}. Also, a T_m



Table 3 Melting temperatures [°C] for the complexes formed with WCp and/or Hp 9-nt ^mRNAs. Roman numerals in parentheses refer to the proposed structures of complexes. The samples were prepared in 10 mM Tris-HCl, 100 mM NaCl, 10 mM MgCl₂ buffer (pH 5.4)

	Homopurine oligomer	wR ⁹	hR ⁹	hR ⁹ + wR ⁹
1	G4	65	63 (IX)	68
2	RB0	47	54 (III)	77 (V)
3	RG2	54	67 (III or X)	>80 (V)
4	RA4{3}	50	75 (III or X)	>80 (V)
5	RG4{3}	49	75 (III or X)	>80 (V)
6	SG4	57	50 (IX)	57

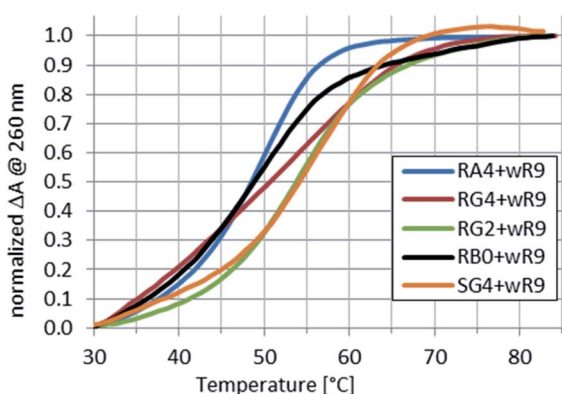


Fig. 4 Normalized melting profiles recorded for RA4, RG2, RG4, or SG4 mixed with wR⁹ at a 1 : 1 molar ratio in pH 5.4 buffer. A profile for RB0 ([R_P-PS]-DNA, no LNA units) is given as a reference.

value found for the putative hR⁹&SG4{3}:wR⁹ was equal to that for SG4:wR⁹, so the S_P internucleotide bonds present in SG4 did not promote the parallel hybridization, neither hR⁹ interacted with remarkably stable SG4{3}:wR⁹ to a measurable extent. The putative complexes hR⁹&G4{3} (no PS units) and hR⁹&SG4{3} (Fig. 5) were less stable (*T*_m = 63 °C and 50 °C, respectively) than the corresponding G4:wR⁹ (*T*_m = 65 °C) and SG4:wR⁹ (*T*_m = 57 °C) indicating the formation of the imperfect WCp duplexes (IX).

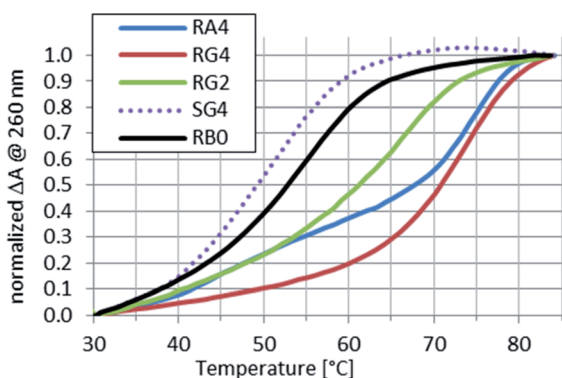
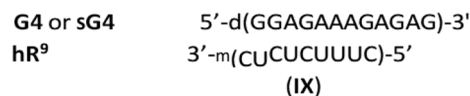
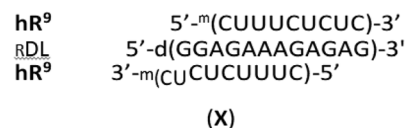


Fig. 5 Normalized melting profiles recorded for RA4, RG2, RG4, or SG4 mixed with hR⁹ at a 1 : 1 molar ratio in pH 5.4 buffer. A profile for RB0 ([R_P-PS]-DNA, no LNA units) is given as a reference.



The profiles shown in Fig. 5 indicate that hR⁹&RG2, hR⁹&RA4 {3}, and hR⁹&RG4{3} were remarkably more stable than hR⁹&RB0 (*ΔT*_m = 13 °C, 21 °C, and 21 °C respectively) with an average increase *ΔT*_m = 7 °C per the “active” LNA unit. But one cannot exclude partial formation of imperfect triplexes X, where a 7 nt part of hR⁹ acts as a WCp strand.



Of course, complexes analogous to X may be formed with participation of any fully or partially WCp RNA. This non-specific binding may be avoided using already mentioned rDL oligomers with the hampered Watson-Crick base pairings.

Despite very short 9 nt ^mRNAs used, the *T*_m values for the investigated triplexes hR⁹&rDL:wR⁹ exceeded 80 °C and could not be more precisely determined (data not shown). This phenomenon indicates that the C2'-endo → C3'-endo conformational shift caused by the LNA units is of great importance for the thermal stability of investigated parallel duplexes and triplexes, which may be beneficial in *in vitro* experiments with precisely selected RNA oligomers, but in cellular experiments may lead to unspecific hybridization with non-target oligomers present in the RNA pool.

Conclusions

The CD spectra recorded for single-stranded P-stereodefined phosphorothioate [R_P-PS]-(DNA#LNA) chimeras (rDL) indicate that 2-4 LNA units distributed along 12 nt oligomer changed the C2'-endo conformation (typical for PS-DNA) into the C3'-endo conformation. The results of melting experiments suggest that this change reduced an energetic toll for the C2'-endo → C3'-endo transition of the rDL strand giving rise to an increase in thermal stability of the parallel complexes with (2'-OMe)-RNAs compared to those formed by LNA-free [R_P-PS]-DNA oligomers (*ΔT*_m = 7 °C per the LNA unit at pH 5.4). Thermally more stable complexes are formed upon addition of the corresponding Watson-Crick-paired (2'-OMe)-RNA strands due to the formation of highly stable parallel triplexes. The LNA nucleotides enhance hybridization at the WC interface and may promote the formation of imperfect triplexes VIII. This hybridization may result in the binding of non-target RNAs. This loss of specificity may be avoided using rDL oligomers with the Watson-Crick base pairing hampered due to the presence of N6-Me-2'-deoxyadenosine (^m6dA) units.⁴⁰ Work on such modified oligonucleotides is in progress.

Experimental section

MALDI-TOF MS analyses of oligonucleotides were performed with the detection of negative ions, using a Voyager-Elite



instrument (PerSeptive Biosystems Inc., Framingham, MA) operating in the reflector mode, or a Shimadzu Biotech Axima Performance instrument operating in the linear mode.

Routine UV spectra were recorded on a CINTRA 10e spectrophotometer (GBC, Dandenong, Australia), using a quartz cuvette of 1 cm path length. UV monitored melting experiments were carried out in 1 cm path length cells, using a spectrophotometer CINTRA 4040 (GBC), equipped with a 6×1 Peltier thermocell.

For the UV-monitored (at 260 nm) thermal dissociation experiments the oligonucleotide samples were dissolved in pH 7.2 or pH 5.4 buffer containing 10 mM Tris-HCl, 100 mM NaCl, and 10 mM MgCl₂. The annealing was done from 85 °C to 15 °C with a temperature gradient of 1 °C min⁻¹. The melting profiles were recorded over a 15 → 85 °C range (0.5 °C min⁻¹). The temperatures of association (T_a) and melting (T_m) were calculated using the first order derivative method.

CD measurements were done on a Jasco J-815 dichrograph at room temperature, using a 0.5 cm path-length quartz cell (Hellma). The spectra were recorded over a 210–320 nm range with a 1.0 nm bandwidth, a scanning speed 50 nm min⁻¹, and a data pitch of 1 nm. After 3 spectra were accumulated, the baseline was subtracted and the resultant spectrum was smoothed with a Savitzky-Golay algorithm (the convolution width 7).

P-stereodefined oligonucleotides of the PS-DNA and PS-(DNA#LNA) series were synthesized manually at a 1 μmol scale, according to the previously published protocols.^{29,38} The first nucleoside unit was anchored to the solid support by a sarcosinyl linker.⁴¹ Routine coupling steps were performed using 20 mg of the OTP monomers. In the cycles where LNA-OTP_L monomers 2 were incorporated, double coupling was executed (20 mg + 20 mg) and in both steps the coupling time was extended to 20 minutes. All synthesized oligomers were routinely purified by two-step reverse phase HPLC (DMT-on and DMT-off). Relevant chromatograms for rDL and sG4 are shown in Fig. S8–S12, ESI†), and their identity was assessed by MALDI-TOF MS and purity by polyacrylamide gel electrophoresis (PAGE).

Unmodified DNA and RNA oligonucleotides were synthesized on a Gene-World synthesizer (K&A Laborgeraete GbR, Schaaheim, Germany).

Conflicts of interest

There are no conflicts to declare.

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Notes and references

- 1 G. Fensfeld, V. Davies and A. Rich, *J. Am. Chem. Soc.*, 1957, **79**, 2023.

- 2 J. R. Goñi, X. de la Cruz and M. Orozco, *Nucleic Acids Res.*, 2004, **32**, 354.
- 3 P. Jenjaroenpun, C. S. Chew, T. P. Yong, K. Choowongkamon, W. Thammasorn and V. A. Kuznetsov, *Nucleic Acids Res.*, 2015, **43**, D110–D116.
- 4 J. A. Brown, M. L. Valenstein, T. A. Yario, K. T. Tycowski and J. A. Steitz, *Proc. Natl. Acad. Sci. U. S. A.*, 2012, **109**, 19202.
- 5 F. A. Buske, J. S. Mattick and T. L. Bailey, *RNA Biol.*, 2011, **8**, 427.
- 6 A. Bacolla, G. Wang and K. M. Vasquez, *PLoS Genet.*, 2015, **11**(12), e1005696.
- 7 C. Hélène, in *Antisense Research and Application*, ed. S. T. Crook and B. Lebleu, and references cited therein, CRC Press Inc. Boca Raton, Ann Arbor, London, Tokyo, 1993, pp. 375–385.
- 8 G. Raghunathan, H. T. Miles and V. Sasisekharan, *Biopolymers*, 1994, **12**, 1573.
- 9 L. Lavelle and J. R. Fresco, *Nucleic Acids Res.*, 1995, **23**, 2692.
- 10 R. W. Roberts and D. M. Crothers, *Science*, 1992, **258**, 1463.
- 11 H. Han and P. B. Dervan, *Proc. Natl. Acad. Sci. U. S. A.*, 1993, **90**, 3806.
- 12 C. Escude, J.-C. Francois, J.-S. Sun, G. Ott, M. Sprinzl, T. Garestier and C. Hélène, *Nucleic Acids Res.*, 1993, **21**, 5547.
- 13 A. Aviñó, M. Frieden, J. C. Morales, B. G. de la Torre, R. Güimil-García, F. Azorín, J. L. Gelpí, M. Orozco, C. González and R. Eritja, *Nucleic Acids Res.*, 2002, **30**, 2609.
- 14 A. K. Shchvolkina, O. F. Borisova, M. A. Livshits, G. E. Pozmogova, B. K. Chernov, R. Klement and T. M. Jovin, *Biochemistry*, 2000, **39**, 10034.
- 15 E. R. Kandimalla and S. Agrawal, *Biochemistry*, 1996, **35**, 15332.
- 16 J. H. van de Sande, N. B. Ramsing, M. W. Germann, W. Elhorst, B. W. Kalisch, E. von Kitzing, R. T. Pon, R. C. Clegg and T. M. Jovin, *Science*, 1988, **241**, 551.
- 17 N. A. Tchurikov, B. K. Chernov, Y. B. Golova and Y. D. Nechipurenko, *FEBS Lett.*, 1989, **257**, 415.
- 18 N. A. Tchurikov, A. K. Shchvolkina, O. F. Borisova and B. K. Chernov, *FEBS Lett.*, 1992, **297**, 233.
- 19 O. F. Borisova, A. K. Shchvolkina, B. K. Chernov and N. A. Tchurikov, *FEBS Lett.*, 1993, **322**, 304.
- 20 N. A. Tchurikov, L. G. Chistyakova, G. B. Zavilgelsky, I. V. Manukhov, B. K. Chernov and Y. B. Golova, *J. Biol. Chem.*, 2000, **275**, 26523.
- 21 A. K. Jain and S. Bhattacharya, *Bioconjugate Chem.*, 2010, **21**, 1389.
- 22 A. K. Shchvolkina, O. F. Borisova, M. A. Livshits and T. M. Jovin, *Mol. Biol.*, 2003, **37**, 223.
- 23 M. W. Germann, C. N. Johnson and A. M. Spring, *Chimia*, 2009, **63**, 731.
- 24 S. Verma and F. Eckstein, *Annu. Rev. Biochem.*, 1998, **67**, 99.
- 25 P. Guga, *Curr. Topics in Med. Chem.*, 2007, **7**, 695.
- 26 P. Guga and M. Koziolkiewicz, *Chem. Biodiversity*, 2011, **8**, 1642.
- 27 W. J. Stec and A. Wilk, *Angew. Chem., Int. Ed. Engl.*, 1994, **33**, 709.

- 28 W. J. Stec, B. Karwowski, M. Boczkowska, P. Guga, M. Koziółkiewicz, M. Sochacki, M. Wieczorek and J. Błaszczyk, *J. Am. Chem. Soc.*, 1998, **120**, 7156.
- 29 P. Guga and W. J. Stec, in *Current Protocols in Nucleic Acid Chemistry*, ed. S. L. Beaucage, D. E. Bergstrom, G. D. Glick, and R. A. Jones, John Wiley & Sons, Hoboken, N.J., 2003, pp. 4.17.1.
- 30 M. Boczkowska, P. Guga and W. J. Stec, *Biochemistry*, 2002, **41**, 12483.
- 31 W. J. Stec, A. Grajkowski, B. Karwowski, A. Kobylańska, M. Koziółkiewicz, K. Misiura, A. Okruszek, A. Wilk, P. Guga and M. Boczkowska, *J. Am. Chem. Soc.*, 1995, **117**, 12019.
- 32 P. Guga, M. Boczkowska, M. Janicka, A. Maciaszek, S. Kuberski and W. J. Stec, *Biophys. J.*, 2007, **92**, 2507.
- 33 P. Guga, M. Janicka, A. Maciaszek, B. Rębowska and G. Nowak, *Biophys. J.*, 2007, **93**, 3567.
- 34 J. J. Novoa, I. Nobeli, F. Grepioni and D. Braga, *New J. Chem.*, 2000, **24**, 5.
- 35 A. A. Koshkin, S. K. Singh, P. Nielsen, V. K. Rajwanshi, R. Kumar, M. Meldgaard, C. E. Olsen and J. Wengel, *Tetrahedron*, 1998, **54**, 3607.
- 36 (a) S. Obika, D. Nanbu, Y. Hari, K. Morio, Y. In, T. Ishida and T. Imanishi, *Tetrahedron Lett.*, 1997, **38**, 8735; (b) S. Obika, D. Nanbu, Y. Hari, J. Andoh, K. Morio, T. Doi and T. Imanishi, *Tetrahedron Lett.*, 1998, **39**, 5401.
- 37 A. Maciaszek, A. Krakowiak, M. Janicka, A. Tomaszewska, M. Sobczak, B. Mikołajczyk and P. Guga, *Org. Biomol. Chem.*, 2015, **13**, 2375.
- 38 K. Jastrzębska, A. Maciaszek, R. Dolot, G. Bujacz and P. Guga, *Org. Biomol. Chem.*, 2015, **13**, 10032.
- 39 R. Kumar, S. K. Singh, A. A. Koshkin, V. K. Rajwanshi, M. Meldgaard and J. Wengel, *Bioorg. Med. Chem. Lett.*, 1998, **8**, 2219.
- 40 A. Maciaszek, K. Jastrzębska and P. Guga, *Org. Biomol. Chem.*, 2019, **17**, 4611.
- 41 T. Brown, C. E. Pritchard, G. Turner and S. A. Salisbury, *J. Chem. Soc., Chem. Commun.*, 1989, 891.
- 42 Y. Zhou, E. Kierzek, Z. P. Loo, M. Antonio, Y. H. Yau, Y. W. Chuah, S. Geifman-Shochat, R. Kierzek and G. Chen, *Nucleic Acids Res.*, 2013, **41**, 6664.
- 43 L. A. H. van Bergen, M. Alonso, A. Palló, L. Nilsson, F. De Proft and J. Messens, *Sci. Rep.*, 2016, **6**, 30369.

