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Synthesis and hybridizing properties of P-stereodefined chimeric [PS]-{DNA:RNA} and [PS]-{DNA:(2'-OMe)-RNA} oligomers†

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Oxathiaphospholane derivatives of 2'-OMe-ribonucleosides and 2'-O-TBDMS-ribonucleosides (M N-OTP and T N-OTP, respectively; nucleobase protected) were synthesized and separated into pure P-diastereomers. X-ray analysis showed the R_P absolute configuration of the phosphorus atom in the fast-eluting diastereomer of T A-OTP. The fast- and slow-eluting P-diastereomers of M N-OTP and T N-OTP were used in the solid-phase synthesis of phosphorothioate dinucleotides (M N_{PS}T and N_{PS}T, respectively), which were subsequently hydrolyzed with R_P -selective phosphodiesterase svPDE and S_P -selective nuclease P1 to determine the absolute configuration of the phosphorus atoms. P-Stereodefined phosphorothioate ([PS]) 10-mer chimeric oligomers [PS]-{DNA:(2'-OMe)-RNA} and isosequential [PS]-{DNA:RNA} containing two M N_{PS} or N_{PS} units were synthesized. Melting experiments performed for their complexes with Watson-Crick paired DNA matrix showed that M N_{PS} or N_{PS} units decrease the thermal stability of the duplexes ($\Delta T_{\rm m} = -0.5 \div -5.5$ °C per modification) regardless of the absolute configuration of the P-atoms. When the (2'-OMe)-RNA matrix was used an increase in $T_{\rm m}$ was noted in all cases ($\Delta T_{\rm m} = +1 \div +7$ °C per modification). The changes in thermal stability of the duplexes formed by [PS]-chimeras with DNA and (2'-OMe)-RNA matrices do not correlate with the absolute configuration of the phosphorus atoms.

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Introduction

In the last five decades, many DNA analogs have been chemically synthesized, mainly to improve their stability against nucleases and in this way to ensure long-lasting recognition of the target nucleic acid. This increased stability was expected to produce new probes useful for example for controlling gene expression or biochemical manipulations. To some extent, this goal has been achieved by the introduction of phosphorothioate analogs of DNA ([PS]-DNA),¹ although nonspecific binding to proteins² and immunological side effects³,⁴ initially hindered their application *in vivo*. [PS]-RNA oligomers were obtained shortly after using phosphoramidite or *H*-phosphonate monomers with sulfurization. Note: interestingly, it has been shown that [PS]-DNA probes in the absence of transfecting agents enter mammalian cells more efficiently than DNA.⁵ An explanation

based on the dynamic covalent exchange of phosphorothioates with cellular thiols and disulfides has recently been published.⁶ Some drugs containing [PS]-DNA and [PS]-RNA-derived units have been approved by the FDA and other agencies, and several others are in advanced clinical trials.⁷

Noteworthy, in 2007 phosphorothioation of DNA was discovered in bacteria,⁸ and in 2019, a DNA phosphorothioation-based antiviral system was discovered in archaea.⁹ The identified chimeric phosphate:phosphorothioate compounds ([PO]:[PS]) contained the phosphorothioate phosphorus atoms only of R_P absolute configuration.¹⁰ In 2020, the R_P phosphorothioate modification was detected in RNA isolated from prokaryotes and eukaryotes.¹¹

Although the phosphorothioate internucleotide bond is isoelectronic with the natural phosphate moiety, the properties of [PS]-DNA differ from natural DNA ([PO]-DNA) in several aspects, mainly due to the different steric requirements of the sulfur atom, the altered affinity for metal ions ("soft" sulfur vs. "hard" oxygen), and unsymmetrical distribution of the negative charge in the diester phosphorothioate anion. 12 Depending on the intended application, these altered properties may be considered either disadvantageous or beneficial, but in the vast majority of biology-related experiments, one should consider the stereochemistry of the internucleotide phosphorothioate moieties. Since the substitution of sulfur for a nonbridging oxygen atom creates a stereogenic center, for example, a [PS]-

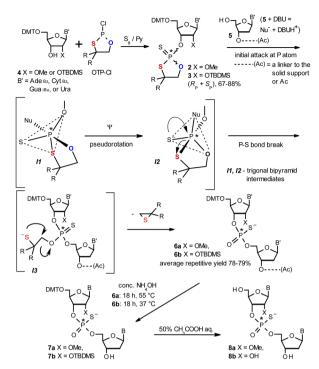
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 \dagger Electronic supplementary information (ESI) available: HPLC profiles for the separation of P-diastereomers of 2 and 3; HR MS, $^1\text{H},\,^{13}\text{C}$ and ^{31}P NMR spectra for P-diastereomers of 2 and 3; the crystal data and data collection and refinement parameters for the fast-eluting detritylated 3a (B' = Ade^{B^2}); MALDI-TOF mass spectra recorded for chimeric PS-oligonucleotides. CCDC 2063388. For ESI and crystallographic data in CIF or other electronic format see https://doi.org/10.1039/d2ra04855h

Chart 1 Classes of the OTP derivatives of 2'-deoxyribo- and ribonucleosides. $B' = Ade^{Bz}$, Cyt^{Bz} , Gua^{iBu} , or Ura. For those with R = H or Me, see ref. 13 and 16, respectively.

oligonucleotide decamer synthesized by a non-stereocontrolled method exists as a mixture of $2^9 = 512$ P-diastereomers, and the content of a single diastereomer is less than 0.2%. Since each Pdiastereomer can interact with a biological target (usually with fixed chirality) in slightly different ways, the mixtures of hundreds of P-diastereomers may be useless for detecting the interactions of interest. To overcome this problem, stereocontrolled synthetic methods have been developed. 13-15 Among them, the approach utilizing so called oxathiaphospholane monomers (OTP; developed in this laboratory)13,16 has yielded the most P-stereodefined oligomers so far, and some of them showed interesting P-stereodependent properties in biochemical studies. 17-21 So far, the OTP method has mainly used separated P-diastereomers of 3'-O-(2-thio-4,4-pentamethylene-1,3,2oxathiaphospholane) derivatives of 2'-deoxyribonucleosides (1, X = H, Y = S, R, R = -(CH₂)₅-, Chart 1), although 2-selenamonomers (1, X = H, Y = Se, R,R = $-(CH_2)_5$ -) have been used in the synthesis of P-stereodefined phosphoroselenoate analogs of DNA.²² In 1996, a set of four monomers 3 (X = O-TBDMS, R = H) was synthesized and chromatographically separated into Pdiastereomers.23 They were used in the solid-phase synthesis of dinucleoside phosphorothioates NPSN, which were obtained in 66-98% yields. However, the separation of fast- and slow-eluting monomers was very difficult and inefficient. In other attempts to synthesize P-stereodefined [PS]-RNA, the synthesis of up to 10-mers was achieved using a method based on nucleoside-3'-Ooxazaphospholidine monomers with the 2'-O-TBDMS protecting group.24 Good coupling yields (97-99%) were observed, but stereoselectivity was not perfect (≥96 : 4). A diastereoselectivity ≥98:2 was observed when oxazaphospholidine monomers with 2'-O-(2-cyanoethoxymethyl) group were used.25

This discovery prompted our attempt to develop a fully stereospecific method for the synthesis of [PS]-(2'-OMe)-RNA ([PS]-MRNA) and [PS]-RNA oligonucleotides with P-stereodefined phosphorothioate internucleotide linkages. (For simplicity the (2'-OMe)-, (2'-O-TBDMS)-, 2'-OH- and 2'-deoxyribonucleosides will be referred to as MN, TN, N, and dN, respectively). In a next step we will attempt to develop a method for the synthesis of {[PO]-DNA:[PS]-MRNA},‡ {[PO]-DNA:[PS]-RNA}, and {[PO]-RNA:[PS]-RNA} chimeras that could be used as model compounds in biochemical studies.



Scheme 1 Synthesis of OTP monomers 2 and 3 (R,R = $-(CH_2)_5-$) and the mechanistic principle of the coupling step.

In this report, we present the synthesis of OTP monomers 2 (X = OMe, MN-OTP, Chart 1) and 3 (X = O-TBDMS, TN-OTP) and their separation into pure P-diastereomers, as well as the result of the successful crystallographic analysis of detritylated 3a (B' = Ade^{Bz}). The P-diastereomerically pure monomers 2 and 3 were used for the synthesis of chimeric dinucleotides 8a and 8b, respectively (Scheme 1), and the absolute configuration of the P atoms was determined enzymatically. Melting curves were recorded to determine the thermal stability of duplexes formed by P-stereodefined chimeric [PS]-{DNA:MRNA} and [PS]-{DNA:RNA} oligonucleotides (10 nt in length) with Watson-Crick paired DNA and MRNA matrices.

Results and discussion

The mechanistic principle of the OTP method

The synthesis of OTP monomers 2 and 3 and the mechanistic principle of the coupling step are shown in Scheme 1. It has been documented²⁶ that the condensation does not follow a stereoinvertive S_N2P mechanism but is a stereoretentive process. The initial attack of a nucleoside 5 occurs from the side opposite to the most electronegative atom bonded to the phosphorus center (the oxygen atom in the oxathiaphospholane ring of 2 or 3, marked in blue; Scheme 1) to form a trigonal bipyramidal intermediate *I1*. Subsequently, a pseudorotation process furnishes the intermediate *I2*, in which the thioalkyl leaving group occupies the axial position, followed by cleavage of the P–S bond to form the triester intermediate *I3*. The condensation process concludes with the elimination of episulfide. Finally, deprotection of the resulting intermediates 6 and

[‡] A colon in structures such as {[PO]-RNA:[PS]-RNA} or [PS]-{DNA:^MRNA} indicates that the nucleotides of both types are contained in the same oligonucleotide strand.

7 and purification steps yield [PS]-oligomers 8. The range of Poligonucleotides phosphorothioate stereodefined extended by the synthesis of [PS]-LNA27 and [PS]-GNA28 oligomers as well as oligomers containing 3'-amino-2',3'-dideoxy nucleosides ([NPS]-DNA).29 Unfortunately, oligomers of the latter two series were obtained in low yields due to unexpected side reactions and unfavorable conformational factors.

Preparation of P-diastereomerically pure MN-OTP and TN-OTP monomers

Protected ribonucleosides 4 (Scheme 1, $B' = Ade^{Bz}$, Cyt^{Bz} Gua^{iBu} , or Ura; X = OMe or O-TBDMS), were phosphitylated with 2-chloro-4,4-pentamethylene-1,3,2-oxathiaphospholane (OTP-Cl, 1.2 equiv.) and then sulfurized with elemental sulfur.30 The resultant monomers 2 and 3 were isolated in 67-88% yields. The P-diastereomers (formed in nearly equimolar amounts) were separated by preparative HPLC on a silica gel column (Fig. S1a and b, ESI†), and their structures were confirmed by HR MS (Fig. S2a-h and S3a-h, ESI†). (For ¹H, ¹³C and ³¹P NMR spectra see ESI, Fig. S4-S6,† respectively). Interestingly, ³¹P NMR analysis (Tables 1 and 2) showed that all fast-eluting isomers of 3 had lower chemical shifts (δ) than their sloweluting counterparts (as observed for dN-OTP monomers¹⁶ and

LNA-OTP²⁷) whereas this relationship was compounds 2.

X-ray crystallography analysis of the detritylated oxathiaphospholane monomer 3a ($B' = Ade^{Bz}$)

Previously published results on the hydrolysis of dinucleotides d(N_{PS}N) with snake venom phosphodiesterase (svPDE) and nuclease P1 (nP1) (R_{P} - and S_{P} -specific phosphodiesterase, respectively) showed that the compounds obtained from fastand slow-eluting monomers dN-OTP (1, X = H, Y = S, R = Me or $R_1R = -(CH_2)_5$ -) contain the internucleotide bonds with the P atoms of the absolute configuration R_P and S_P, respectively.¹⁶ The same correlation was found for the analogous OTP derivatives of LNA nucleosides.27 X-ray analysis of LNA-thymidine-3'-O-(2-thio-4,4-pentamethylene-1,3,2-oxathiaphospholane) confirmed the stereoretentive course of the condensation step. The "locked" structure naturally resulted in the C3'-endo

conformation of the sugar ring.

In the present studies, a crystal suitable for X-ray analysis was obtained from the 5'-OH deprotected fast-eluting monomer 3a ($B' = Ade^{Bz}$). Detritylation was carried out in anhydrous acetonitrile using a sodium hydrogen sulfate suspension deposited on silica gel.31 The crystal structure was resolved (Fig. 1) with the refinement parameter R = 0.0563 and ESI.† The

Table 1 Characteristics of the MN-OTP monomers 2a-d

| | ^M A-OTP (2a) | ^M C-OTP (2b) | ^M G-OTP (2c) | $^{\mathrm{M}}$ U-OTP (2 d) |
|--|-------------------------|----------------------------------|-------------------------|-------------------------------------|
| Yield ^a (%) | 78 | 88 | 69 | 83 |
| Rf^b (TLC) | 0.72 | 0.83 | 0.70 | 0.81 |
| Eluent for TLC/HPLC ^c | 80:20 | 70:30 | 100 | 40:60 |
| MM calc. (Da) | 893 | 869 | 875 | 766 |
| For fast-2/slow-2 | | | | |
| Rt ^d (min) | 16/24 | 13/18 | 12/17 | 19/26 |
| $HR MS^e (m/z)$ | 892.2630/892.2625 | 868.2510/868.2510 | 874.2706/874.2715 | 765.2079/765.2082 |
| δ^{31} P NMR ^f (ppm) | 106.555/106.132 | 106.690/106.579 | 107.042/106.688 | 106.719/106.571 |

^a Yield of the isolated mixture of P-diastereomers. ^b Eluent CHCl₃: MeOH (9:1, v/v). ^c Separation of P-diastereomers at an AcOEt: hexane ratio (v/ v). ^d A Phenomenex Luna 5u Silica gel column (100 Å; 250 × 10 mm; flow rate 5 mL min⁻¹, isocratically). ^e Recorded with a SYNAPT G2-Si High Definition Mass Spectrometer. f In anhydrous CDCl₃.

Table 2 Yield, chromatographic and spectroscopic data for the ^TN-OTP monomers **3a-d**

| | ^T A-OTP (3a) | ^T C-OTP (3b) | ^T G-OTP (3c) | ^T U-OTP (3d) |
|--|---|---|---|---|
| Yield ^a (%) Rf ^b (TLC) Eluent for TLC/HPLC ^c MM calc. (Da) | 70 0.64 50:50 993 | 72 0.74 70:30 969 | 67 0.62 100 975 | 73 0.71 40:60 866 |
| For fast-3/slow-3 Rt^d (min) $HR MS^e$ (m/z) δ ³¹ P NMR^f (ppm) | 15/17 992.3309/992.3311 105.880/107.519 | 35/40 968.3198/968.3192 106.966/107.022 | 10/15 974.3427/974.3418 106.244/107.342 | 25/35 865.2776/865.2775 106.500/107.369 |

^a Yield of the isolated mixture of P-diastereomers. ^b Eluent CHCl₃: MeOH (9:1, v/v). ^c Separation of P-diastereomers at an AcOEt: hexane ratio (v/ v). ^d A Phenomenex Luna 5u Silica gel column (100 Å; 250 × 10 mm; flow rate 5 mL min⁻¹, isocratically). ^e Recorded with a SYNAPT G2-Si High Definition Mass Spectrometer. f In anhydrous CDCl₃.

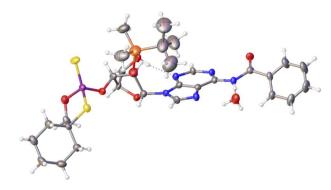


Fig. 1 X-ray ORTEP diagram of the detritylated *fast*-eluting P-diastereomer of 3a showing the $R_{\rm P}$ absolute configuration. The displacement ellipsoids are shown at the 50% probability level. ³² Colors: red – oxygen, blue – nitrogen, pink – phosphorus, yellow – sulfur, orange – silicon, black – carbon, white – hydrogen.

crystal data and the data acquisition and refinement parameters are listed in Table S1 (ESI†). It should be noted that despite the presence of a 2'-oxygen atom, the (normally strong) anomeric effect is not decisive in this case and the sugar ring exists in the C2'-endo conformation (Fig. 2), perhaps due to the steric requirements of the bulky TBDMS group.

Crystallization experiments with all other DMT-labeled as well as detritylated OTP derivatives were unsuccessful.

Synthesis and enzymatic analysis of ${}^MN_{PS}T$ and $N_{PS}T$ dinucleotides

Since X-ray analysis was successful only for the derivative of *fast*-eluting **3a**, enzymatic analysis was performed to complete the stereochemical assignment in the remaining **2** and **3**. For this purpose, 4 pairs of P-diastereomeric diesters $^{M}N_{PS}T$ (**6a**, X = OMe) and 4 pairs of P-diastereomeric diesters $N_{PS}T$ (**6b**, X = OH) were prepared from 3'-O-acetyl-thymidine (**5**, B' = Thy, Scheme **1**) and diastereomerically pure (*fast*- or *slow*-eluting) ^{M}N -OTP **2a–d** or ^{T}N -OTP **3a–d**, respectively. The base-labile protecting groups in intermediates **6** were removed with concentrated ammonia. Subsequently, products **7a** (X = OMe; derived from **2**) were routinely detritylated with 50% AcOH_{aq}. The DMT and TBDMS moieties in the freeze-dried products **7b** (X = O-TBDMS;

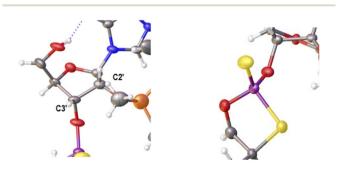


Fig. 2 Parts of the X-ray ORTEP diagram show: left image—a DNA-like C2'-endo conformation of detritylated fast-eluting 3a; right image—spatial orientation of substituents on the phosphorus atom. Colors as in Fig. 1.

Table 3 MALDI TOF MS data for $^{M}N_{PS}T$ 8a (X = OMe) and $N_{PS}T$ (8b, X = OH) synthesized from P-diastereomerically pure ^{M}N -OTP 2a-d and ^{T}N -OTP 3a-d, respectively

| $^{\mathrm{M}}$ N-OTP (2a-d) or $^{\mathrm{T}}$ N-OTP (3a-d) | | MALDI TOF MS m/z [$m/z_{\rm calc}$] | | | |
|--|--------|---|---------------------------|--|--|
| B' | Isomer | $^{M}N_{PS}T$ (8a, X = OMe) | $N_{PS}T$ (8b, $X = OH$) | | |
| Ade ^{Bz} | Fast | 600.1283 [601] | 586.1066 [587] | | |
| | Slow | 600.1283 [601] | 586.1064 [587] | | |
| Cyt^{Bz} | Fast | 576.1171 [577] | 562.1013 [563] | | |
| • | Slow | 576.1174 [577] | 562.1011 [563] | | |
| Gua ^{iBu} | Fast | 616.1229 [616] | 602.1076 [602] | | |
| | Slow | 616.1232 [616] | 602.1074 [602] | | |
| Ura | Fast | 577.1006 [578] | 563.0858 [564] | | |
| | Slow | 577.1007 [578] | 563.0848 [564] | | |
| | | | | | |

derived from 3) were removed by treatment with $3HF \times NEt_3$. The MALDI TOF MS data for the resulting dinucleotides 8a and 8b are given in Table 3.

Small amounts (2–3 $\rm OD_{254}$ units) of all 16 P-diastereomeric dinucleotides **8** were isolated using RP-HPLC and further treated with the svPDE and nP1 enzymes, which are $R_{\rm P}$ - and $S_{\rm P}$ -stereoselective, respectively. Subsequent RP-HPLC analysis revealed that all dinucleotides **8** obtained from the *fast*-eluting **2a–d** and **3a–d** were hydrolyzed with svPDE and not with nP1. Their counterparts obtained from the *slow*-eluting monomers were hydrolyzed with nP1 and not with svPDE. These results, together with the X-ray data collected for **3a**, confirm the stereoretentive course of the condensation step.

Relative reactivity of ^MN-OTP and ^TN-OTP compared with dN-OTP

It has been reported previously that when monomers 1 were used, 92-94% yields were observed with single coupling, allowing 18-23 nt long PS-DNA oligomers to be assembled.18 Previous experiments with the OTP derivatives of LNA nucleosides showed that double coupling was necessary to achieve an acceptable yield of the condensation step.27 Using this modified protocol, chimeric [PS]-{DNA:LNA} oligomers were obtained with LNA units in every third position.§ This lower reactivity was tentatively attributed to steric hindrance due to the C2'-O-CH₂-C4' bridge near the reacting phosphorus center. However, unsatisfactory attempts to synthesize oligomers with 3'-amino-2',3'-dideoxy nucleosides ([NPS]-DNA)29 and subsequent molecular modeling experiments indicated the importance of the deoxyribose ring conformation in the OTP monomers. Importantly, only the OTP derivative of 3'-amino-3'-deoxythymidine allowed the synthesis of [NPS]-DNA 8-mer, while none of the monomers with the other three nucleobases allowed more than four coupling steps. Therefore, competition experiments were performed to determine the relative reactivity of selected purine and pyrimidine monomers 2 and 3 compared

§ Uniformly modified [PS]-LNA oligomers were not synthesized because of expected too strong rigidity of the resultant strand.

Table 4 The composition (mol%, by ^{31}P NMR) of the mixtures of ^{M}N -OTP/dN-OTP and ^{T}N -OTP/dN-OTP used in the reaction with thymidine attached to the solid support, and the composition (mol%, by HPLC) of the resultant mixtures of dinucleotides $^{M}N_{PS}T$ /dN_{PS}T and $^{T}N_{PS}T$ /dN_{PS}T

| | N = A | | N = U | |
|--|-------|-------------|-------|-------|
| OTP substrates, dinucleotide products ^{ab} | Fast | Slow | Fast | Slow |
| ^M N-OTP (2): dN-OTP (1) | 49:51 | 44:56 | 53:47 | 46:54 |
| $^{\mathrm{M}}\mathrm{N}_{\mathrm{PS}}\mathrm{T}:\mathrm{d}\mathrm{N}_{\mathrm{PS}}\mathrm{T}$ | 26:74 | 42:58 | 31:69 | 32:68 |
| Selectivity ratio ^c | 0.36 | <u>0.91</u> | 0.40 | 0.55 |
| $^{\mathrm{T}}$ N-OTP (3): dN-OTP (1) | 42:58 | 44:56 | 53:47 | 55:45 |
| $^{\mathrm{T}}\mathrm{N}_{\mathrm{PS}}\mathrm{T}^{d}:\mathrm{d}\mathrm{N}_{\mathrm{PS}}\mathrm{T}$ | 22:78 | 6:94 | 19:81 | 15:85 |
| Selectivity ratio ^c | 0.39 | 0.08 | 0.21 | 0.14 |

 $[^]a$ The dinucleotide products were detritylated before cleavage from the solid support. b In the adenosine containing products, the benzoyl protecting group was removed with concentrated ammonia. c The selectivity ratios are calculated by dividing % composition of products by % composition of substrates. d The TBDMS group in $^TN_{PS}T$ dinucleotides was not removed.

to the corresponding dN-OTP. For this purpose, the approximately equimolar mixtures 1/2 and 1/3 (measured by ³¹P NMR) were reacted with thymidine bound to a solid support (10 mg of each OTP monomer was used, i.e., a 14-18-fold excess). The resultant MN_{PS}T/dN_{PS}T and TN_{PS}T/dN_{PS}T mixtures were detritylated, detached from the solid support, deprotected with concentrated ammonia (the TBDMS group in the TNPST dinucleotides was not removed), and analyzed using RP-HPLC. The required standards were prepared independently. The results (Table 4) generally show that the MN-OTP monomers studied were two to three times less reactive compared with the corresponding dN-OTP, as the selectivity ratio values (calculated by dividing the percent composition of the dinucleotide products by the percent composition of the OTP substrates) ranged from 0.36 to 0.55. Only for the slow-eluting MA-OTP the difference was almost negligible (Table 4, selectivity ratio 0.91, double underlined). The reactivity of ^TN-OTP was 5-12 times lower, with the lowest selectivity ratio value of 0.08 (Table 4, underlined). Reactivity comparable to MN-OTP was found only for the fasteluting ^TA-OTP (selectivity ratio 0.39).

The generally low reactivity observed for 2 and 3 suggested that the synthesis protocol optimized for [PS]-DNA³⁴ and later modified for the synthesis of [PS]-LNA oligomers²⁷ would be inadequate. In addition, the widely varying selectivity ratios indicated that several factors might influence reactivity, so that optimization of the protocol for all 16 diastereomers of ^MN-OTP

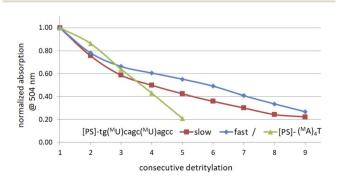


Fig. 3 Decay of absorption (at 504 nm) of the DMT-cation released in consecutive detritylation steps.

and ^TN-OTP would be a lengthy process with uncertain prospects for success. An attempt to synthesize uniformly modified [PS]-(^MA)₄T, in which the most reactive *slow*-eluting ^MA-OTP (2a, see Table 4) was double-coupled (thymidine was bound to the support), yielded only a small amount of product. The results of the DMT cation assay, shown in Fig. 3 (green line), clearly indicate that although the first coupling was carried out with 86% yield, the next three coupling steps were much less efficient with 74, 67 and 49% yield, respectively. The synthesis of [PS]-(^MA)₄T showed that the synthesis of [PS]-^MRNA and [PS]-RNA oligomers with mixed sequences and longer than penta- or hexamers will be very difficult.

Synthesis and properties of chimeric [PS]-{DNA:^MRNA} and [PS]-{DNA:RNA} oligonucleotides

Because of the relatively low repetitive yield of coupling steps in syntheses with monomers 2 or 3, we focused on the synthesis of P-stereodefined chimeric [PS]-{DNA:MRNA} and [PS]-{DNA:RNA} oligonucleotide decamers (collectively referred to as [PS]-chimeras or PSCh) with base sequence 5'-tgtcagctag-3', where lowercase letters represent DNA units. In the sequences of analogous chimeric oligomers, the introduced N_{PS} are indicated in uppercase letters (*e.g.* A or U), whereas the MN_{PS} units are indicated in uppercase letters with a superscript M prefix (*e.g.* MA or MU).

In the synthesis of each oligomer, the dN-OTP monomer was replaced in two selected cycles by ^{M}N -OTP or ^{T}N -OTP carrying the same nucleobase, except for the T-OTP monomer, which was replaced by ^{M}U -OTP or ^{T}U -OTP congeners. The codes, specific sequences, and HR MS data for two synthesized reference [PS]-DNA and sixteen chimeric [PS]-oligomers are listed in Table 5. [Note: in codes such as **UR** or M **US**, the capital letters **R** or **S** denote [PS]-oligomers with phosphorus atoms with either R_{P} or S_{P} absolute configuration, respectively].

The DMT cation decay assay was performed for each synthesis, and calculations for the steps using MN-OTP or TN-OTP (double coupling was performed) gave the average coupling yields of 78 and 79%, respectively (Table S2, ESI†). After cleavage from the support and removal of the base-labile protecting groups, the DMT-labeled oligomers were purified

Table 5 Codes, sequences, and HR MS characteristics of the [PS]-DNA and chimeric [PS]-{DNA:MRNA} and [PS]-{DNA:RNA} oligomers under study

| | | MS ES (m/z) | | | | |
|-----------------|---|---------------|-----------|--|--|--|
| Code | [PS]-oligomer ^a sequence $(5' \rightarrow 3')$ | Calc. | Found | | | |
| [PS]-DNA | [PS]-DNA from 1a-d | | | | | |
| BR | tgtcagctag | 3185.34 | 3185.3401 | | | |
| BS | tgtcagctag | 3185.34 | 3185.3301 | | | |
| [PS]-{DN | A: ^M RNA} from 1a–d and 2a–d | | | | | |
| MAR | tgtc ^M Agct ^M Ag | 3245.36 | 3245.3501 | | | |
| MAS | | | 3245.3501 | | | |
| ^M CR | tgt ^M Cag ^M Ctag | 3245.36 | 3245.3401 | | | |
| ^M CS | | | 3245.3500 | | | |
| ^M GR | t ^M Gtca ^M Gctag | 3245.36 | 3245.4001 | | | |
| ^M GS | | | 3245.3401 | | | |
| ^M UR | tg ^M Ucagc ^M Uag | 3217.33 | 3217.3000 | | | |
| ^M US | | | 3217.3000 | | | |
| [PS]-{DN | A:RNA} from 1a–d and 3a–d | | | | | |
| AR | tgtcAgctAg | 3217.33 | 3217.3181 | | | |
| AS | | | 3217.3101 | | | |
| CR | tgtCagCtag | 3217.33 | 3217.3101 | | | |
| CS | | | 3217.3000 | | | |
| GR | tGtcaGctag | 3217.33 | 3217.3101 | | | |
| GS | | | 3217.3101 | | | |
| UR | tgUcagcUag | 3189.30 | 3189.3000 | | | |
| US | | | 3189.3000 | | | |

 $[^]a$ In the sequences, the units containing 2'-deoxyribo- and ribonucleosides are indicated in lowercase and uppercase letters, respectively. The $^MN_{PS}$ units are marked with a superscript M prefix (^MN) .

by RP-HPLC, and the collected samples were detritylated and isolated by RP-HPLC. Fig. 4 shows typical chromatograms recorded during the DMT-ON and DMT-OFF purification steps; here those for ${}^{\mathbf{M}}\mathbf{GS}$ ($S_{\mathbf{P}}$). MALDI TOF MS spectra for PSCh are shown in ESI.†

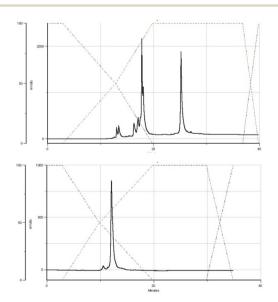


Fig. 4 HPLC profiles recorded for ^MGS during DMT-ON (the upper panel) and DMT-OFF (the lower panel) purification steps.

Melting experiments for canonical and hybrid duplexes

For the melting experiments, the oligomers were mixed with the equimolar amount of Watson–Crick-paired DNA matrix (**Md**) or mRNA matrix (**Mm**) (2.0 μ M concentration each) in pH 7.2 buffer. The observed changes in UV absorbance followed the two-state mechanism and the recorded curves had the expected S-shape. An example set of melting curves is shown in Fig. 5. Melting temperatures (compiled in Table 6) were calculated using the numerical fitting method provided by the UV spectrophotometer manufacturer. The $\Delta T_{\rm m}$ values were calculated by subtracting the control $T_{\rm m}$ from the values recorded for the duplexes tested.

Melting experiments for duplexes formed by [PO]-chimeras

When analyzing the results of melting experiments with duplexes formed by [PO]-{DNA: MRNA} chimeras (MA, MC, MG, and MU), an oligonucleotide with the sequence [PO]-5'-tgtcagctag-3' (Basal DNA oligomer, Bd) was used as a reference. Compared to the reference Bd/Md duplex ($T_{\rm m}=41\,^{\circ}{\rm C}$, Table 6), the [PO]-{DNA: MRNA}/Md duplexes were $1 \div 9\,^{\circ}{\rm C}$ less thermally stable ($T_{\rm m}=32\,\div\,40\,^{\circ}{\rm C}$, Fig. 6, green bars), with the pyrimidine units (MCPO, MUPO) destabilizing the duplexes more than the purine units.

Melting experiments with the Mm matrix showed that the purine units (${}^{\rm M}{\rm A}_{\rm PO}$, ${}^{\rm M}{\rm G}_{\rm PO}$) slightly stabilized the duplexes ($\Delta T_{\rm m}$ $= 1 \div 4$ °C, Fig. 6, black bars), compared to the reference **Bd**/ **Mm** duplex ($T_{\rm m} = 51$ °C). The duplexes containing the ${}^{\rm M}{\rm C}_{\rm PO}$ or ^MU_{PO} nucleotides were less stable than **Bd/Mm** and the destabilizing effect was strongest for ${}^{\rm M}$ U ($\Delta T_{\rm m} = -5$ °C). These results differ from those of Wengel and coworkers, who showed that one to three LNA nucleotides introduced into the DNA strand at either the adenine (5'-AGCACCAG) or thymine (5'-TGCTCCTG) residues increased the thermal stability of the corresponding {DNA:LNA}/RNA heteroduplexes in 110 mM Na⁺ buffer.³³ For two introduced A-LNA or T-LNA units, $\Delta T_{\rm m} = 10$ and 16 °C, respectively, were found. It can be argued that RNA matrices were used in these experiments, but Dolot and coworkers showed that crystals of DNA/RNA and DNA/MRNA duplexes have very similar geometries.34

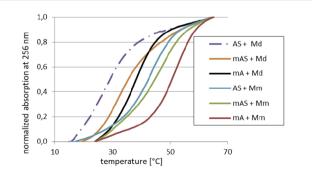


Fig. 5 Normalized melting curves for selected duplexes (see legend) dissolved in pH 7.2 buffer (10 mM Tris–HCl, 100 mM NaCl, and 10 mM MgCl $_2$). The curves designations mA and mAS (in the legend) correspond to $^{\rm M}$ A and $^{\rm M}$ AS, respectively.

Table 6 Melting temperatures derived from the melting profiles recorded at pH 7.2 for [PO]- or [PS]-oligomers mixed with the Watson–Crick paired DNA (Md) or (2'-OMe)-RNA (Mm) matrices at a 1:1 molar ratio

| | Sequence $(5' \rightarrow 3')^a$ | Md | | Mm | |
|---------------------------|--|---|---|---|---|
| Code | | $T_{\mathrm{m}}\left[^{\circ}\mathrm{C}\right]$ | $\Delta T_{\mathrm{m}} \left[{}^{\circ}\mathrm{C} \right]$ | $T_{\mathrm{m}}\left[^{\circ}\mathrm{C}\right]$ | $\Delta T_{ m m} \left[^{\circ} m C \right]$ |
| [PO]-D | NA reference oligomer | | | | |
| Bd | tgtcagctag | 41 | _ | 51 | _ |
| [PO]-{I | DNA: ^M RNA} chimeras ^b | | | | |
| M _A | tgtc ^M Agct ^M Ag | 38 | -3 | 52 | +1 |
| $^{\mathbf{M}}\mathbf{C}$ | tgt ^M Cag ^M Ctag | 36 | -5 | 48 | -3 |
| ^M G | t ^M Gtca ^M Gctag | 40 | -1 | 55 | +4 |
| ^M U | tg ^M Ucagc ^M Uag | 32 | -9 | 46 | -5 |
| [PS]-D | NA reference oligomers | | | | |
| BR | tgtcagctag | 35 | _ | 32 | _ |
| BS | 0 0 0 | 35 | _ | 35 | _ |
| [PS]-{[| ONA:RNA} chimeras ^c | | | | |
| AR | tgtcAgctAg | 30 | -5 | 40 | +8 |
| AS | | 27 | -8 | 43 | +8 |
| CR | tgtCagCtag | 30 | -5 | 38 | +6 |
| CS | | 34 | -1 | 39 | +4 |
| GR | tGtcaGctag | 31 | -4 | 39 | +7 |
| GS | | 30 | -5 | 37 | +2 |
| UR | tgUcagcUag | 27 | -8 | 40 | +8 |
| US | | 31 | -4 | 40 | +5 |
| [PS]-{E | ONA: MRNA} chimeras ^c | | | | |
| MAR | tgtc ^M Agct ^M Ag | 30 | -5 | 44 | +12 |
| MAS | | 32 | -2 | 47 | +12 |
| MCR | tgt ^M Cag ^M Ctag | 31 | -4 | 41 | +9 |
| ^M CS | | 32 | -4 | 39 | +4 |
| ^M GR | t ^M Gtca ^M Gctag | 30 | -5 | 46 | +14 |
| ^M GS | Ü | 31 | -4 | 48 | +13 |
| ^M UR | tg ^M Ucagc ^M Uag | 29 | -6 | 40 | +8 |
| ^M US | | 24 | -11 | 41 | +6 |

 $[^]a$ In the sequences, the DNA and RNA units are written in lowercase and uppercase letters, respectively. The $^{\rm M}{\rm N}_{\rm PS}$ units are marked with a superscript M prefix ($^{\rm M}{\rm N}$). b The $\Delta T_{\rm m}$ values were calculated using the $T_{\rm m}$ values for the corresponding duplexes **Bd/Md** or **Bd/Mm**. c The $\Delta T_{\rm m}$ values were calculated using the $T_{\rm m}$ values for the corresponding duplexes **B(R,S)/Md** or **B(R,S)/Mm**.

Melting experiments for the PSCh/Md duplexes. The basal complexes BR/Md and BS/Md (P-stereodefined [PS]-DNA/DNA; collectively referred to as B(R,S)/Md) were equally stable ($T_{\rm m}=35~^{\circ}{\rm C}$). Compared with the phosphate B/Md reference complex, they were 6 $^{\circ}{\rm C}$ less thermally stable (Table 6). Compared with B(R,S)/Md, in all cases, regardless of the absolute configuration of the P atoms, ribonucleotide units ($N_{\rm PS}$) as well as (2'-OMe)-ribonucleotide units ($N_{\rm PS}$) present in the P-stereodefined [PS]-chimeras further reduced the thermal stability of the duplexes (Fig. 6, blue bars), as the $\Delta T_{\rm m}$ values ranged from $-1~^{\circ}{\rm C}$ for the most stable duplex CS/Md ($T_{\rm m}=34~^{\circ}{\rm C}$) to $-11~^{\circ}{\rm C}$ for $^{\rm M}{\rm US/Md}$ ($T_{\rm m}=24~^{\circ}{\rm C}$). Within groups of [PS]-oligonucleotides carrying $^{\rm M}{\rm N_{PS}}$ or $N_{\rm PS}$ with the same nucleobase (e.g., $^{\rm M}{\rm AR}$, $^{\rm M}{\rm AS}$, AR, and AS) the patterns of changes were different.

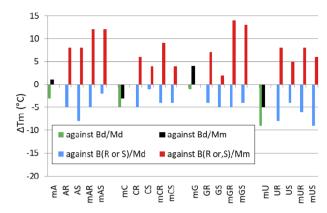


Fig. 6 Changes in melting temperature of duplexes ($\Delta T_{\rm m}$; °C) formed by [PO]-{DNA: MRNA} chimeras with Md (green bars) or Mm (black bars), and PSCh oligomers with Md (blue bars) and Mm (red bars). The $\Delta T_{\rm m}$ values were calculated using the $T_{\rm m}$ values for the corresponding duplexes Bd/Md, Bd/Mm, B(R,S)/Md and B(R,S)/Mm. Designations along the horizontal axis such as mAR or mGS correspond to MAR and MGS, respectively.

Melting experiments for the PSCh/Mm duplexes. The BR/ Mm and BS/Mm complexes (collectively referred to as B(R,S)/ **Mm**) had $T_{\rm m}=32$ and 35 °C, respectively (Table 6), and were similarly stable to B(R,S)/Md. Compared with the phosphate B/ Mm reference complex, they were less stable by 19 and 16 °C, respectively. In contrast to the PSCh/Md duplexes, the use of the (2'-OMe)-RNA matrix (Mm) resulted in an increase in the thermal stability of the duplexes. The most stable ${}^{\mathbf{M}}\mathbf{GS}/\mathbf{Mm}$ had $T_{\rm m} = 48 \,^{\circ}\text{C}$, but compared to **B(R,S)/Mm** the largest $\Delta T_{\rm m} = 14 \,^{\circ}\text{C}$ was observed for MGR/Mm (Fig. 6, red bars), while the smallest effect ($\Delta T_{\rm m}=2~^{\circ}{\rm C}$) was observed for GS/Mm. The complexes formed by Mm with PSCh containing MN_{PS} units (especially those containing the purine derivatives) were more stable than those with PSCh containing the NPS nucleotides. Within groups of [PS]-oligonucleotides carrying MNPS or NPS with the same nucleobase the patterns of changes were different.

Summary of melting experiments. The analysis of the $\Delta T_{\rm m}$ values obtained from the melting experiments (they are shown in Fig. 6) leads to three conclusions. First, compared to B(R,S)/Md and B(R,S)/Mm, the ${}^MN_{PS}$ and N_{PS} units present in PSCh decreased the thermal stability of duplexes formed with Md and increased the stability of those formed with Mm. Second, the strongest stabilizing effects ($\Delta T_{\rm m}$ > 10 °C) were caused by purine ^MN_{PS} residues in oligonucleotides interacting with Mm. Third, the changes in thermal stability of duplexes formed by PSCh with Md and Mm did not correlate with the absolute configuration of phosphorus atoms. For example, $\Delta T_{\rm m}$ for **GR/Mm** is 5 °C larger than for GS/Mm, whereas the $\Delta T_{\rm m}$ values for AR/Mm and AS/Mm and for MAR/Mm and MAS/Mm are the same (8 and 12 °C, respectively). However, the $\Delta T_{\rm m}$ for the duplexes formed by PSCh with the S_P absolute configuration of phosphorus atoms were in no case larger than for the duplexes with their R_P counterparts.

Conclusions

The 2-thio-1,3,2-oxathiaphospholane monomers of 2'-OMe and 2'-O-TBDMS series (MN-OTP and TN-OTP, 2 and 3, respectively)

were synthesized in good yield and effectively separated into pure P-diastereomers. The X-ray crystallographic analysis revealed the R_P absolute configuration of the phosphorus atom in the fast-eluting diastereomer 3a (B' = Ade^{Bz}). Using the R_P selective phosphodiesterase svPDE and the S_P -selective Nuclease P1 it was found that in both monomer series the fastand slow-eluting P-diastereomers are precursors of dinucleotides with $R_{\rm P}$ and $S_{\rm P}$ absolute configuration of the phosphorus atoms, respectively. Importantly, no traces of P-epimerization were observed during coupling steps. P-Stereodefined chimeric [PS]-{DNA:MRNA} and isosequential [PS]-{DNA:RNA} 10-mer oligomers with two MN_{PS} or N_{PS} units were synthesized. Melting experiments using complexes with Watson-Crick paired DNA matrix showed that MNPS or NPS units decrease the thermal stability of the duplexes ($\Delta T_{\rm m} = -0.5 \div -5.5$ °C per modification), regardless of the absolute configuration of the Patoms. When the (2'-OMe)-RNA matrix was used an increase in $T_{\rm m}$ was observed in all cases ($\Delta T_{\rm m} = +1 \div +7$ °C per modification). The changes in thermal stability of duplexes formed by [PS]-chimeras with DNA and (2'-OMe)-RNA matrices did not correlate with the absolute configuration of phosphorus atoms.

Despite of double-coupling, the average repetitive yields of the condensation steps with either 2 or 3 were rather moderate (78–79%). Nevertheless, even these moderate yields should allow the synthesis of {[PO]-DNA:[PS]-MRNA} and {[PO]-RNA:[PS]-RNA} chimeras that could be used as model compounds in biochemical studies. To achieve this goal, the conditions for an effective combination of phosphoramidite and OTP synthesis methods need to be worked out. This work is in progress.

Experimental section

 1 H NMR and 31 P NMR spectra were acquired using Bruker instruments (AV-200, DRX-500, or Avance Neo 400; operating for 1 H at 200, 500, and 400 MHz, respectively). Chemical shift (δ) values are in ppm, referenced to internal tetramethylsilane (TMS) or residual solvent protons for 1 H NMR and external 85% $\rm H_{3}PO_{4}$ for 31 P NMR.

High-resolution mass spectra (HR MS) were acquired using a Synapt G2 Si mass spectrometer (Waters Corporation, Milford, MA) equipped with an ESI source and a quadrupole time-of-flight mass analyzer. Measurements were performed in negative ion mode with the capillary and sampling cone voltages set at 2.7 kV and 20 V, respectively. The temperature of the source was 110 °C. To ensure satisfactory accuracy, data were acquired in a centroid mode and readings were corrected during acquisition using leucine-enkephalin as an external reference (Lock-SprayTM) generating reference ions at m/z 554.2615 Da ([M–H] $^-$) in negative ESI mode. Data sets were processed using MassLynx 4.1 software (Waters).

MALDI-TOF MS analyses of oligonucleotides were performed with negative ion detection using a Voyager Elite instrument (PerSeptive Biosystems Inc, Framingham, MA) operating in reflector mode or an Axima Performance instrument (Shimadzu Biotech Corp., Japan) operating in linear mode.

TLC silica gel 60 plates F254 developed in $CHCl_3/MeOH$, 9: 1 (v/v), were used for routine analyses. HP TLC Silica gel 60 plates with a UV-F₂₅₄ indicator were used to evaluate the separability of the P-diastereomers of 2 and 3.

A Varian binary HPLC system (two PrepStar 210 pumps, 25 mL pump heads, one ProStar 320 UV/vis detector at 275 nm) was used for HPLC separation of the P-diastereomers of monomers 2 and 3. A Phenomenex Luna 5 μm silica column (100 Å; 250 \times 10 mm; flow rate 5 mL min $^{-1}$) was used in search of the appropriate conditions. Preparative separation was performed with a Pursuit XRs silica gel column (10 μm , 250 \times 21.2 mm) eluted at a flow rate of 25 mL min $^{-1}$. Gradient-grade HPLC solvents from Sigma-Aldrich, Baker, or ChemPur were used.

Crystallographic data were collected using an XtaLAB Synergy, Dualflex, HyPix diffractometer at T = 100.00(10) K.

Routine UV spectra were recorded with a CINTRA 4040 spectrophotometer (GBC, Dandenong, Australia). UV-monitored melting experiments were performed at 260 nm in cuvettes with 1 cm path length using a UV-vis NIR spectrophotometer V-770 (Jasco, Japan) equipped with a 6 \times 1 Peltier thermal cell.

For thermal dissociation experiments, oligonucleotide samples were dissolved in pH 7.2 buffer containing 10 mM Tris–HCl, 100 mM NaCl, and 10 mM MgCl₂. Annealing was performed from 85 °C to 15 °C with a temperature gradient of 1 °C min⁻¹. Melting profiles were recorded over a range of 15 \rightarrow 85 °C (0.5 °C min⁻¹). Melting temperatures ($T_{\rm m}$) were calculated using the software provided by Jasco.

P-Stereodefined PSCh oligonucleotides were synthesized manually at a 1 µmol scale, using the slightly modified previously published protocols.16,35 The first nucleoside unit was solid support by a DBU-resistant anchored to the [-C(O)CH2CH2C(O)sarcosinyl-succinoyl linker N(CH₃)CH₂C(O)-LCAA-CPG].³⁶ Routine coupling steps were performed using 20 mg of the dN-OTP monomers 1. In the cycles where the monomers 2 or 3 were incorporated, double coupling was executed (20 mg + 20 mg) and in both steps the coupling time was extended to 20 minutes. The assembled DMT-labeled oligomers were cleaved from the support (25% NH₄OH, r.t., 3 h) and the protecting groups from nucleobases were removed with 25% NH₄OH_{aq} over 18 h ([PS]-{DNA: MRNA}, 55 °C; [PS]-{DNA: TRNA}, 37 °C). The samples were concentrated under reduced pressure and the DMT-labeled oligonucleotides were isolated using RP-HPLC. The collected fractions were concentrated under reduced pressure. The [PS]-{DNA:MRNA} oligomers were detritylated with 50% AcOHaq for 20 minutes and the volatile components were evaporated. The TBDMS and DMT groups in [PS]-{DNA: TRNA} oligomers were simultaneously removed by treatment with 3HF × NEt₃ at room temperature for 20 hours. The reagent was quenched with sterile water and all chimeric [PS]-oligomers were isolated by means of RP-HPLC. The identity of oligonucleotides was confirmed by MALDI-TOF MS (for the relevant spectra see ESI†).

Unmodified DNA and (2'-OMe)-RNA oligonucleotide matrices (**Md** and **Mm**, respectively) were synthesized on a Gene-World synthesizer (K&A Laborgeraete GbR, Schaafheim,

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Germany) and routinely isolated by a two-step RP-HPLC method.

Synthesis of oxathiaphospholane monomers 2 and 3 – a general procedure

Oxathiaphospholane monomers 2 and 3 were synthesized analogously to a general procedure published for the synthesis of standard OTP monomers.34 Briefly, a suitable nucleoside 4 (2 mmol, N-protected if necessary; Scheme 1) was dried overnight in a 25 mL round bottom flask with two necks (with a magnetic stirrer inside) at high vacuum (0.01 mm Hg). To the flask purged with dry argon, 10 mL of anhydrous pyridine was added using a gas-tight syringe. To the stirred solution, 0.36 mL 2-chloro-4,4-pentamethylene-1,3,2-oxathiaphospholane (OTP-Cl, 0.51 g, 2.4 mmol) was added at room temperature for 5 minutes using a gas-tight syringe. After approximately 5 minutes, 0.2 g of dry elemental sulfur (approximately 6 mmol) was added, and stirring was continued at room temperature for 12 hours. Excess sulfur was filtered off, the solvent evaporated, and the residue dissolved in 2 mL of acid-free chloroform (distilled with pyridine). The crude product was applied to a silica gel column (glass tube 15×3 cm, height of gel suspension layer - 5 cm, 230-400 mesh) and the column was eluted with acid-free chloroform. Appropriate fractions (8-10 mL each, analyzed by TLC on silica gel 60 plates, Rf's are given in Table 1) were combined, and the solvent was evaporated under reduced pressure (15-20 mm Hg) at a water bath temperature not exceeding 30 °C. Anhydrous toluene (5-6 mL) was then added and the solution was evaporated to dryness excluding moisture. Finally, the samples were dried under high vacuum (oil pump) for 12 hours and stored at -15 °C. The yields of the isolated mixtures of P-diastereomers are given in Tables 1 and 2.

Detritylation of the fast-eluting diastereomer of 3a

To a solution of 120 mg (0.12 mmol) of *fast*-eluting isomer of 5'-O-DMT-2'-O-TBDMS- N^6 -Bz-adenosine-3'-O-(2-thio-4,4-pentamethylene-1,3,2-oxathiaphospholane) in 2 mL of anhydrous acetonitrile a sodium hydrogen sulfate/silica gel reagent (50 mg) was added and the suspension was stirred at room temperature. The reaction progress was monitored by TLC using a 9:1 (v/v) chloroform: methanol mixture as an eluent (Rf of the substrate 0.57; Rf of the product 0.31). After 26 hours the reaction went to complete. The silica gel was filtered off and the solvent was evaporated *in vacuo*. The product was isolated by semipreparative HPLC (a silica gel column eluted with ethyl acetate) and evaporation at reduced pressure furnished white solid material (52 mg, 62%).

Details of X-ray data collection and reduction

Single colorless transparent plate-shaped crystals of the detritylated *fast*-eluting P-diastereomer of **3a** were obtained by recrystallization from a mixture of ethyl acetate and methanol (4:1 v/v). A suitable crystal of $0.11 \times 0.08 \times 0.02$ mm dimension was selected and mounted on a suitable support. During data collection the crystal was kept at a steady temperature T=

100.00(10) K. The structure was solved with the *XT* structure solution program³⁷ using an intrinsic phasing solution method and *Olex2* (ref. 38) as a graphical interface. The model was refined with *ShelXL* (version 2018/3; ref. 39), using the least squares minimization.

In solution synthesis of dinucleoside 3',5'-phosphorothioates ^MN_{PS}T (8a) and N_{PS}T (8b)

To a sample of a P-diastereomerically pure (fast- or slow-eluting) oxathiaphospholane monomer 2 or 3 (5 mg, ca. 5 µmol) and 3'-O-Ac-thymidine (3 mg, ca. 11 μmol), 600 μL of dry acetonitrile and 5 µL of 1,8-diazabicyclo-[5.4.0]undec-7-ene (DBU, 1.1 molar equivalent over the OTP monomers) were added under dry argon. After 2 h the reaction was complete. The acetonitrile was removed under reduced pressure and the products 6 were treated (in a tightly closed vessel) with concentrated ammonia solution (1 mL) for 18 h either at 55 °C (6a) or at 37 °C (6b), except for B = Ura (2 h at room temperature). The products 7awere routinely detritylated using 50% AcOH_{aq}. In case of 7b, approximately 10% of the ammoniacal solution was evaporated to dryness and the DMT and TBDMS moieties were removed with 3HFxNEt₃ (150 μL, 17 h, room temperature) followed by evaporation under reduced pressure. The dimers 8 were isolated using RP-HPLC. Conditions: a C18 column, 250 imes 4.6 mm, 5 μ m; elution at 1.0 mL min⁻¹ with a gradient of 0.1 M TEAB, pH 7.3 to 40% CH₃CN in 0.1 M TEAB over 20 minutes.

Conflicts of interest

There are no conflicts to declare.

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Notes and references

- F. Eckstein, Angew. Chem., Int. Ed. Engl., 1983, 22, 423;
 W. J. Stec, G. Zon, W. Egan and B. Stec, J. Am. C. Soc., 1984, 106, 6077;
 F. Eckstein, Annu. Rev. Biochem., 1985, 54, 367;
 F. Eckstein, Antisense Nucleic Acid Drug Dev., 2000, 10, 117.
- 2 C. A. Stein, Biochim. Biophys. Acta, 1999, 1489, 45.
- 3 A. M. Krieg, Antisense Nucleic Acid Drug Dev., 2001, 11, 181.
- 4 A. M. Krieg, P. Guga and W. J. Stec, *Oligonucleotides*, 2003, **13**, 491.
- 5 S. T. Crooke, S. Wang, T. A. Vickers, W. Shen and X. Liang, *Nat. Biotechnol.*, 2017, 35, 230.
- 6 Q. Laurent, R. Martinent, D. Moreau, N. Winssinger, N. Sakai and S. Matile, *Angew. Chem., Int. Ed.*, 2021, **60**, 19102.

- 7 S. T. Crooke, P. P. Seth, T. A. Vickers and X. Liang, J. Am. Chem. Soc., 2020, 142, 14754.
- 8 L. Wang, S. Chen S, T. Xu, K. Taghizadeh, J. S. Wishnok, X. Zhou, D. You, Z. Deng and P. C. Dedon, *Nat. Chem. Biol.*, 2007, 3, 709.
- 9 L. Xiong, S. Liu, S. Chen, Y. Xiao, B. Zhu, Y. Gao, Y. Zhang, B. Chen, J. Luo, Z. Deng, X. Chen, L. Wang and S. Chen, *Nat. Commun.*, 2019, **10**, 1.
- 10 P. Guga and M. Koziołkiewicz, *Chem. Biodiversity*, 2011, 8, 1642.
- 11 Y. Wu, Y. Tang, X. Dong, Y. Y. Zheng, P. Haruehanroengra, S. Mao, Q. Lin and J. Sheng, ACS Chem. Biol., 2020, 15, 1301.
- 12 P. A. Frey and R. D. Sammons, Science, 1985, 228, 541.
- 13 W. J. Stec, A. Grajkowski, M. Koziołkiewicz and B. Uznański, *Nucleic Acids Res.*, 1991, **19**, 5883.
- 14 A. Wilk, A. Grajkowski, L. R. Phillips and S. L. Beaucage, *J. Am. Chem. Soc.*, 2000, **122**, 2149.
- 15 N. Oka, M. Yamamoto, T. Sato and T. Wada, *J. Am. Chem. Soc.*, 2008, **130**, 16031.
- 16 W. J. Stec, B. Karwowski, M. Boczkowska, P. Guga, M. Koziołkiewicz, M. Sochacki, M. Wieczorek and J. Błaszczyk, J. Am. Chem. Soc., 1998, 120, 7156.
- 17 M. Boczkowska, P. Guga and W. J. Stec, *J. Biomol. Struct. Dyn.*, 1999, **16**, 1291.
- 18 T. Inagawa, H. Nakashima, B. Karwowski, P. Guga, W. J. Stec, H. Takeuchi and H. Takaku, *FEBS Lett.*, 2002, **528**, 48.
- 19 P. Guga, Curr. Top. Med. Chem., 2007, 7, 695.
- 20 P. Guga, M. Boczkowska, M. Janicka, A. Maciaszek, S. Kuberski and W. J. Stec, *Biophys. J.*, 2007, 92, 2507.
- 21 P. Guga, M. Janicka, A. Maciaszek, B. Rębowska and G. Nowak, *Biophys. J.*, 2007, **93**, 3567.
- 22 P. Guga, A. Maciaszek and W. J. Stec, Org. Lett., 2005, 7, 3901.
- 23 A. Sierzchała, A. Okruszek and W. J. Stec, J. Org. Chem., 1996, 61, 6713.
- 24 N. Oka, T. Kondo, S. Fujiwara, Y. Maizuru and T. Wada, *Org. Lett.*, 2009, 11, 967.

- 25 Y. Nukaga, K. Yamada, T. Ogata, N. Oka and T. Wada, *J. Org. Chem.*, 2012, 77, 7913.
- 26 W. J. Stec, A. Grajkowski, A. Kobylańska, B. Karwowski, M. Koziołkiewicz, K. Misiura, A. Okruszek, A. Wilk, P. Guga and M. Boczkowska, *J. Am. Chem. Soc.*, 1995, **117**, 12019.
- 27 K. Jastrzębska, A. Maciaszek, R. Dolot, G. Bujacz and P. Guga, *Org. Biomol. Chem.*, 2015, **13**, 10032.
- 28 A. Tomaszewska-Antczak, K. Jastrzębska, A. Maciaszek, B. Mikołajczyk and P. Guga, *RSC Adv.*, 2018, **8**, 24942.
- 29 E. Radzikowska, R. Kaczmarek, D. Korczyński, A. Krakowiak, B. Mikołajczyk, J. Baraniak, P. Guga, K. A. Wheeler, T. Pawlak and B. Nawrot, RSC Adv., 2020, 10, 35185.
- 30 R. Wallin, M. Kalek, A. Bartoszewicz, M. Thelin and J. Stawinski, *Phosphorus*, *Sulfur*, and *Silicon*, 2009, **184**, 908; N. Wang, P. Saidhareddy and X. Jiang, *Nat. Prod. Rep.*, 2020, 37, 246.
- 31 R. K. Kannasani, V. V. S. Peruri and S. R. Battula, *Chem. Cent. J.*, 2012, **6**, 136.
- 32 L. J. Farrugia, J. Appl. Crystallogr., 2012, 45, 849.
- 33 H. Kaur, J. Wengel and S. Maiti, *Biochemistry*, 2008, 47, 1218– 1227.
- 34 R. Dolot, A. Maciaszek, B. Mikołajczyk and B. Nawrot, *Crystals*, 2022, **12**, 760.
- 35 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 and Sons, Hoboken, N. J., 2003, p. 4.17.1.
- 36 T. Brown, C. E. Pritchard, G. Turner and S. A. Salisbury, *J. Chem. Soc., Chem. Commun.*, 1989, 891.
- 37 G. M. Sheldrick, Acta Crystallogr., Sect. A: Found. Adv., 2015, 71. 3.
- 38 L. J. Bourhis, O. V. Dolomanov, R. J. Gildea, J. A. K. Howard and H. Puschmann, *Acta Crystallogr., Sect. A: Found. Adv.*, 2015, 71, 59.
- 39 G. M. Sheldrick, Acta Crystallogr., Sect. C: Struct. Chem., 2015, 71, 3.