Supporting Information

Design and characterization of reversible head-to-tail boronate-linked macrocyclic nucleotides

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1. Experimental part

Except THF that was distilled on calcium with benzophenone, all the anhydrous solvents and reagents were commercial and used without further purification. Thin-layer chromatographies (TLC) were made on silica plate 60 F₂₅₄ Merck and revealed at 254 nm. Purifications on silica gel column chromatography were made with silica 40-63 µm Merck-Millipore. HRMS analysis were realised on Micromass Q-TOF spectrometer with an electrospray ionisation (ESI). NMR analysis were acquired at 25 °C on a Bruker Avance spectrometer operating at 400 MHz as well as on 600 MHz and 500 MHz Bruker Avance III spectrometers equipped with TCI and BBO cryo-probeheads respectively. Chemical shifts in organic solvents were reported in parts per million (ppm) referenced to the solvent residual peaks (CDCl₃ δ^{1} H = 7.26 and δ^{13} C = 77.16, CD₃OD δ^{1} H = 3.31 and δ^{13} C = 49.00, DMF-d₇ δ^{1} H = 2.75 and δ^{13} C = 29.76 for the high field methyl signal). Experiments in deuterated water were calibrated using DSS as reference for ¹H NMR. Indirect referencing was used for ¹³C NMR using the ¹³C/¹H ratio = 0.251449530.^[1] The pD of the solutions was measured with a Mettler-Toledo micro electrode and was adjusted using solutions of NaOD in D_2O and taking into account that $pD = pH_{reading} + 0.4$.^{[2] 1}H NMR data were reported as follows: chemical shift, multiplicity (s = singlet, d = doublet, t = triplet, q = quartet, dd = doublet of doublets, dt = doublet of triplets, m = multiplet, bs = broad singlet), coupling constant (J) in Hertz, integration and attribution. ¹¹B NMR data were acquired on the 400 MHz Bruker Avance spectrometer equipped with BBO probe head and the spectra were calibrated from BF₃.EtOEt as external reference. Diffusion Ordered Spectroscopy experiments (DOSY) were run with a sequence including longitudinal Eddy current delay with bipolar gradients and a double stimulated echo for convection compensation (sequence "dtebpgp3s" from the Bruker library). The magnetic field gradients were calibrated so that measuring the diffusion coefficient of water in D_2O at 25 °C yields a value of 1.902 x 10⁻⁹ m²s^{-1,[3]} The T_1 relaxation times were measured for the sample in DMF before to run the DOSY to properly set the relaxation delay (3 s). The DOSY spectra were acquired with optimized gradient lengths of 2.6 and 2.8 ms (δ), a diffusion delay of 50 ms (Δ) and with gradient values linearly ranging from 2 to 95 % of the maximal gradient (69.2 G/cm) in 32 steps. Translational diffusion coefficients (D) were extracted by non-linear curve fitting of the intensity decay with increasing gradient strength using the T_1/T_2 processing tool of the Topspin software. The analysis of the intensity attenuation curves was repeated fort at least 3 resonances for each compound in order to get reliable average D values.

5'-aldehyde-2',3'-isopropylideneuridine (5)



2',3'-O-isopropylideneuridine (2.3 g, 8.1 mmol) was dissolved in 100 mL of acetonitrile and IBX (3.4 g, 12.0 mmol) was added (insoluble). The mixture was stirred at 80°C for 5 hours. The progress of the reaction was monitored by TLC. After return to room temperature, the solution was filtered on celite and evaporated under reduced pressure to lead to a white solid corresponding to the compound **5** (2.0 g, 88 %) which did not required purification. **R**_f: 0.55 (DCM/MeOH : 95/5). ¹**H NMR** (400 MHz, CDCl₃) δ (ppm) : 1.35 (s, 3H, CH_{3 IP}), 1.52 (s, 3H, CH_{3 IP}), 4.55 (bs, 1H, H_{4'}), 5.10 (d, *J* = 6.4 Hz, 1H, H_{2'}), 5.20 (d, *J* = 6.0 Hz, 1H, H_{3'}), 5.51 (s, 1H, H_{1'}), 5.77 (d, *J* = 8.0 Hz, 1H, H₅), 7.29 (d, *J* = 8.0 Hz, 1H, H₆), 9.42 (s, 1H, CHO),

9.74 (s, 1H, H_N). ¹³**C NMR** (125 MHz, CDCl₃) δ (ppm) : 24.9 (CH_{3 IP}), 26.5 (CH_{3 IP}), 83.8 (C_{2'}), 84.9 (C_{3'}), 94.2 (C_{4'}), 100.2 (C₅), 102.9 (C_{1'}), 113.7 (C_{IV IP}), 144.5 (C₆), 150.7 (C₂), 163.8 (C₄), 199.6 (CHO). **HRMS (ESI⁺)** : *m*/z calculated for C₁₂H₁₄N₂O₆ [M+H]⁺ : 283.0930, found : 283.0931.

5'-deoxy-5'-vinyl-2',3'-isopropylideneuridine (6)



Chemical Formula: C₁₃H₁₆N₂O₅ Exact Mass: 280,11

The Nysted reagent (cyclo-dibromodi- μ -methylene[μ -(tetrahydrofuran)]trizinc, 16.2 mL, 7.08 mmol,) was dissolved in 20 mL of anhydrous THF and BF₃OEt₂ (1.75 mL, 14.0 mmol) was added dropwise at 0°C. The mixture was stirred at 0°C for 15 minutes and compound **5** (2.0 g, 7.08 mmol) dissolved in 5 mL of anhydrous THF was added dropwise. The mixture was stirred at room temperature for 4 hours. The progress of the reaction was followed by TLC. Sodium bicarbonate was added dropwise. 1-Butanol was then added and the organic phase was washed with sodium bicarbonate, dried on sodium sulphate, filtered and evaporated under reduced pressure. The crude material was purified by silica gel column chromatography (DCM/MeOH : 95/5) to lead to a white solid corresponding to the compound **6** (1.0 g, 50 %). **R**_f : 0.60 (EA/Cyclohexane 90/10). ¹**H NMR** (400 MHz, CD₃OD) δ (ppm) : 1.35 (s, 3H, CH_{3 IP}), 1.55 (s, 3H, CH_{3 IP}), 4.46 (m, 1H, H₄'), 4.74 (dd, *J* = 6.4 Hz, 4.4 Hz, 1H, H₃'), 5.04 (dd, *J* = 6.4 Hz, 2.1 Hz, 1H, H₂'), 5.22 (dt, *J* = 10.4 Hz, 1.4 Hz, 1H, H₆'cis), 5.33 (dt, *J* = 17.2 Hz, 1.4 Hz, 1H, H₆'trans), 5.68 (d, *J* = 8.0 Hz, 1H, H₅), 5.78 (d, *J* = 2.1 Hz, 1H, H₁'), 6.00 (ddd, *J* = 17.2 Hz, 10.4 Hz, 7.2 Hz, 1H, H₅'), 7.59 (d, *J* = 8.0 Hz, H₆). ¹³**C NMR** (125 MHz, CD₃OD) δ (ppm) : 25.6 (CH_{3 IP}), 27.5 (CH_{3 IP}), 85.5 (C₃'), 85.8 (C₂'), 89.7 (C₁'), 94.5 (C₄'), 102.8 (C₅), 115.5 (C_{IV IP}), 118.4 (C₆'), 136.9 (C₅'), 144.6 (C₆), 151.9 (C₂), 166.2 (C₄). **HRMS (ESI**⁺) : *m*/z calculated for C₁₃H₁₆N₂O₅ [M+H]⁺ : 281.1137, found : 281.1142.

5'-deoxy-5'-vinyl uridine (7)



Compound **6** (1.5 g, 5.35 mmol) was dissolved in 60 mL of milli-Q water and TCA (8.74 g, 54.0 mmol) was added. The mixture was stirred at room temperature for 4 hours. The progress of the reaction was monitored by TLC. Water was evaporated under reduced pressure and the crude material was purified by silica gel column chromatography (DCM/MeOH : 90/10) to lead to a white solid corresponding to the compound **7** (1.2 g, 93 %). **R**_f : 0.30 (DCM/MeOH : 90/10). ¹**H NMR** (400 MHz, CD₃OD) δ (ppm) : 3.96 (t, *J* = 5.8 Hz, 1H, H_{3'}), 4.20 (dd, *J* = 5.4 Hz, 3.9 Hz, 1H, H_{2'}), 4.34 (tt, *J* = 6.4 Hz, 1.4 Hz, 1H, H_{4'}), 5.28 (dt, *J* = 10.5 Hz, 1.3 Hz, 1H, H_{6'cis}), 5.40 (dt, *J* = 17.2 Hz, 1.4 Hz, 1H, H_{6'trans}), 5.74 (d, *J* = 8.4 Hz, 1H, H₅), 5.83 (d, *J* = 3.8 Hz, 1H, H_{1'}), 6.03 (ddd, *J* = 17.2 Hz, 10.5 Hz, 6.6 Hz, 1H, H_{5'}), 7.59 (d, *J* = 8.4 Hz, 1H, H₆).

¹³**C NMR** (125 MHz, CD₃OD) δ (ppm) : 75.0 (C_{3'} + C_{2'}), 85.6 (C_{4'}), 91.9 (C_{1'}), 102.9 (C₅), 118.4 (C_{6'}), 136.7 (C_{5'}), 142.4 (C₆), 152.2 (C₂), 166.1 (C₄). **HRMS (ESI⁻)** : m/z calculated for C₁₀H₁₂N₂O₅ [M-H]⁻ : 239.0668 found : 239.0669.

1-(5'-deoxyuridin-5'-yl)methylboronic acid (3)



To 5 mL of anhydrous diethylic ether were added dropwise borane dimethysulfide (3.0 mL, 15.0 mmol) and α -pinene (4.8 mL, 30.0 mmol). The mixture was stirred at 35°C for 4 hours. Compound **7** (450.0 mg, 1.9 mmol) dissolved in anhydrous THF was then added dropwise and the resulting mixture was stirred at room temperature for 15 hours. The progress of the reaction was monitored by TLC. The reaction was then quenched by addition of 1 mL of hydrochloric acid 0.1 M and diluted with ethyl acetate. The organic layer was washed with sodium chloride, dried on sodium sulphate, filtered and evaporated under reduced pressure. The crude material was purified by silica gel column chromatography (DCM/MeOH : 85/15) to lead to a white solid corresponding to the compound **3** (170.0 mg, 32 %). **R**_f : 0.25 (DCM/MeOH : 85/15). ¹**H NMR** (37 mM, 298 K, D₂O, pD 6.5, 400 MHz) δ (ppm) : 7.67 (d, *J* = 8.1 Hz, 1H, H₆), 5.89 (d, *J* = 8.1 Hz, 1H, H₅), 5.85 (d, *J* = 4.6 Hz, 1H, H_{1'}), 4.33 (t, *J* = 5.0 Hz, 1H, H_{2'}), 4.03 (t, *J* = 5.4 Hz, 2H, H_{6'/6''}). ¹³**C NMR** (37 mM, 298 K, D₂O, 125 MHz) δ (ppm) : 166.1 (Cq, C₄), 151.6 (Cq, C₂), 141.9 (CH, C₆), 102.4 (CH, C₅), 89.5 (CH, C_{1'}), 85.5 (CH, C_{4'}), 73.5 (CH, C_{2'}), 72.6 (CH, C_{3'}), 27.3 (CH₂, C_{5'}), 10.1 (CH₂, C_{6'}). ¹¹**B NMR** (37 mM, 298 K, D₂O, 128 MHz) δ (ppm) : 32.5 ppm. **HRMS (ESI⁺)** : *m*/z calculated for C₁₀H_{16s}N₂O₇ [M+H]⁺ : 287.10469, found : 287.10509.

Head-to-tail homodimer of 3 (htt₂)



¹**H NMR** (37 mM, 298 K, DMF-d₇, 600 MHz) δ (ppm) : 11.40 (bs, 1H, H_{N3}), 7.85 (d, J = 8.0 Hz, 1H, H₆), 5.71 (d, J = 8.0 Hz, 1H, H₅), 5.88 (d, J = 3.2 Hz, 1H, H₁'), 5.19 (m, 1H, H₂'), 4.81 (m, 1H, H₃'), 4.05 (q, J = 6.2 Hz, 1H, H₄'), 1.93 (m, 2H, H_{5'/5"}), 1.10-0.95 (m, 2H, H_{6'/6"}). ¹³**C NMR** (37 mM, 298 K, DMF-d₇, 125 MHz) δ (ppm) : 163.6 (Cq, C₄), 151.0 (Cq, C₂), 142.9 (CH, C₆), 102.6 (CH, C₅), 92.4 (CH, C₁'), 87.4 (CH, C₄'), 84.9 (CH, C₂'), 84.6 (CH, C₃'), 27.7 (CH₂, C₅'), 6.3 (CH₂, C₆'). ¹¹**B NMR** (37 mM, 298 K, DMF, 128 MHz) : very large peak around 34 ppm.

Head-to-tail homotrimer of 3 (htt₃)



¹**H NMR** (37 mM, 298 K, DMF-d₇, 600 MHz) δ (ppm) : 7.81 (d, *J* = 8.0 Hz, 1H, H₆), 5.72 (d, *J* = 8.0 Hz, 1H, H₅), 5.86 (d, *J* = 2.5 Hz, 1H, H_{1'}), 5.18 (m, 1H, H_{2'}), 4.80 (m, 1H, H_{3'}), 3.90 (q, *J* = 6.4 Hz, 1H, H_{4'}), 1.91 (m, 2H, H_{5'/5''}), 1.10-0.95 (m, 2H, H_{6'/6''}). ¹³**C NMR** (37 mM, 298 K, DMF-d₇, 125 MHz) δ (ppm) : 163.6 (Cq, C₄), 150.9 (Cq, C₂), 143.2 (CH, C₆), 102.6 (CH, C₅), 92.9 (CH, C_{1'}), 88.1 (CH, C_{4'}), 85.3 (CH, C_{2'}), 83.3 (CH, C_{3'}), 27.3 (CH₂, C_{5'}), 6.2 (CH₂, C_{6'}). ¹¹**B NMR** (37 mM, 298 K, DMF-d₇, 128 MHz) : very large peak around 34 ppm.

2. Oligonucleotides synthesis

All the natural oligonucleotides used were synthesized with an automatic DNA synthesizer Applied Biosystems 394 by classical phosphoramidite chemistry with commercial phosphoramidites and controlled-pore glass (Icaa-CPG) linked to 5'-O-DMTr-U or 5'-O-DMTr-dT through a 3'-O-succinyl linker. Analytical and semi-preparative high performance liquid chromatographies (HPLC) in reverse phase were made on a HPLC Dionex 600 system with UV detection at 260 nm and EC 75/4.6 Nucleodur 100-3 C18 (analytical) or VP 250/10 Nucleodur C18 HTec 5 μ m (semi-preparative) columns. Buffers used for the analytical HPLC were: 100 % TEAAc 0.05 M (buffer A) and 100 % ACN (buffer B) with a flow of 1 mL/min. Buffers used for the semi-preparative HPLC were : 1 % ACN in a solution of TEAAc 0.05 M (buffer A) and 80 % ACN in a solution of TEAAc 0.05 M (buffer B) with a flow of 4 mL/min. MALDI-TOF spectra were acquired on an Axima assurance spectrometer (Shimadzu Biotech) with a N₂ laser (337 nm). The matrix used was THAP or ATT in a solution of ACN/Ammonium citrate 0.1 M (1/1). The sample was mixed with the matrix with a ratio of 1/5. Oligonucleotides dosages were made on a Varian Cary 300 Bio UV spectrometer with a UV detection at 260 nm.

General procedure for linear oligonucleotides synthesis^[4,5]

RNA oligonucleotides U_2 and U_3 were prepared by automated synthesis on a 8 and 12 µmol scale respectively using commercial phosphoramidites with a 2'-O-PivOM protecting group and 3'phosphate CPG solid supports under the conditions shown in Table S1. A trichloroacetic acid solution in DCM was used as detritylation reagent, 5-benzylmercapotetrazole (BMT) as activator, a mixture of phenoxyacetic anhydride in THF/pyridine and *N*-methylimidazole in THF as capping solution and an iodine solution as oxidizing agent.

Step	Reaction	Reagent	Time (s)
1	Detritylation	3 % TCA in DCM	35
2	Coupling	0.1 M amidite in ACN + 0.3 M BMT in ACN	20
3	Capping	Ac ₂ O/THF/Pyridine + 10 % NMI in THF	8
4	Oxidation	0.1 M I ₂ /THF/H ₂ O/Pyridine	15

Table S1 : Standard conditions for oligonucleotides synthesis

General procedure for cyclic oligonucleotides synthesis

 $c(T_2)$ and $c(T_3)$ were synthesized according to the classical phosphoramidite chemistry with commercial phosphoramidites and 3'-phosphate CPG solid supports. The first cycle required a supplementary step of sulfurization with Beaucage reagent (3-phenyl-1,2,4-dithiazoline-5-one) after the first phosphoramidite incorporation (Table S2). Next cycles were executed according to the standard conditions. The column was then percolated with 1 mL of a solution of (PhO)₃PCH₃I (0.5 M in anhydrous DMF) at room temperature for 15 minutes, successively washed with 15 mL of DCM, 10 mL of ACN, 10 mL of DCM and dried with argon. The deprotection and cyclization were both achieved at the same time.

Step	Reaction	Reagent	Time (s)
1	Detritylation	3 % TCA in DCM	35
2	Coupling	0.1 M amidite in ACN + 0.3 M BMT in ACN	20
3	Capping	Ac ₂ O/THF/Pyridine + 10 % NMI in THF	8
4	Sulfurisation	0.05 M Beaucage reagent in ACN	60
5	Oxidation	0.1 M I ₂ /THF/H ₂ O/Pyridine	15

General procedure for deprotection, analysis and purification of oligonucleotides

DNA oligonucleotides were deprotected in a solution of ammonia 28 % at 55°C for 5 hours. RNA oligonucleotides were percolated for 3 minutes in a solution of DBU 1 M in ACN, dried with argon and then deprotected in a solution of ammonia 28 % at 37°C for 3 hours. Ammonia was then evaporated in SpeedVac and oligonucleotides were analysed by analytical HPLC and MALDI-TOF with a ATT matrix and ammonium citrate as co-matrix in negative mode. Oligonucleotides were then purified by semi-preparative HPLC (gradient 0-20-20 for linear U_2 and U_3 , 0-25-15 for cyclic $c(T_2)$ and $c(T_3)$).

Table 55. III/2 Obtained with MALDI-TOP analysis for pure Ois

Name	Yield (µmol)	<i>m/z</i> calcd. ^{<i>a</i>}	<i>m/z</i> found ^a
U2	2.79 (35 %) ^b	549.36	549.24
U ₃	2.46 (12 %) ^c	855.53	854.89
c(T ₂)	2.33 (29 %) ^b	623.45	623.45
c(T₃)	2.76 (35 %) ^b	927.64	927.57

^{*a*} *m/z* calculated for [M-H]⁻. ^{*b*} Isolated yields based on 8 μmol scale. ^{*c*} Isolated yields based on 12 μmol scale.

3. Estimation of the macrocycles size from diffusion coefficients

The translation diffusion coefficients can be connected to the molecular size through the Stoke-Einstein equation:

$$D = \frac{k_B T}{6\pi\eta f r_h} \tag{1}$$

where T is the sample temperature, k_B is the Boltzman constant, η is the solvent viscosity, r_H the hydrodynamic radius.

For small to medium-size molecules moreover, the equation denominator is overestimated and have to be correct using the f factor proposed by Gierer and Wirtz :^[6]

$$f_{GW} = (\frac{3r_s}{2r} + \frac{r}{r+r_s})^{-1}$$
(2)

where r and r_s and the solute and solvent hydrodynamic radii respectively.

It is then possible to relate the molecular weight of the solute i (MW_i) to its hydrodynamic radius according to the following expression:

$$r_i = \sqrt[3]{\frac{3 * MW_i}{4 \pi N_A \rho}}$$
 (3)

where N_A is Avogadro's number and ρ is the solute density.^[7]

Considering the polymeric nature of the different assemblies, one can estimate the molecular weight of each oligomer from the ratio of the diffusion coefficients. Assuming that both the monomer and the cyclic species do have spherical shape and relatively close densities and f factors (calculation of f according to equation (2) for the monomer, dimer and trimer respectively gave the following values for f: 0.63, 0.68 and 0.71), the dependence of D on the molecular weight for two molecules can be written :^[7]

$$\frac{D_1}{D_2} = \sqrt[3]{\frac{MW_2}{MW_1}}$$
 (4)

 Table S4: 1H-1H Vicinal coupling constants (in Hz).

	Uridine			3	
		D.0	Wet	Anhydrous	DMF-d ₇
	$D_2 0^{-1}$	D_2O	DMF-d ₇	Dimer ^b	Trimer ^c
³ J _{1'-2'}	4.4	4.7	5.3	3.2	2.5
³ J _{2'-3'}	5.3	5.0	5.3	7.8	nd
³ J _{3'-4'}	5.5	5.4	nd	5.1	5.6

^{*a*} from Blackburn B. J. et al.^[8] ^{*b*} measured on the diluted sample where the trimer **htt₃** was almost undetectable ^{*c*} measured on the concentrated sample (37 mM).

	NMR	QM ^b
	(single model) ^a	
φ _{1'2'}	123.0	117.9
Φ2'3'	6.2	21.5
φ _{3'4'}	-134.4	-161.2

 $\textbf{Table S5:} Exocyclic angles \ \varphi_{i \cdot j} \ (H_i - C_i - C_j - H_{j'}) \ in \ degrees \ calculated \ for \ the \ different \ models.$

^a for P = 52 and ϕ_m = 9°; ^b MP2/6-31G(d)

4. NMR spectra







Figure S2: ¹³C NMR spectrum of 5 in CDCl₃ (125 MHz)



Figure S3: ¹H NMR spectrum of 6 in CD₃OD (400 MHz)



Figure S4:¹³C NMR spectrum of 6 in CD₃OD (125 MHz)



Figure S5:¹H NMR spectrum of 7 in CD₃OD (400 MHz)



Figure S6: ¹³C NMR spectrum of 7 in CD₃OD (125 MHz)



Figure S8: ¹³C NMR spectrum of 3 in D₂O (pD 6.5, 125 MHz, 37 mM)



Figure S9: ¹¹B NMR spectrum of 3 in D₂O (pD 6.5, 128 MHz, 37 mM)



Figure S10 : ¹H NMR spectrum of the cyclic homopolymers of **3** (htt_2 and htt_3) in anhydrous DMF-d₇ (600 MHz, 37 mM)



Figure S11 : ¹³C NMR spectrum of the cyclic homopolymers of **3** (htt_2 and htt_3) in anhydrous DMF-d₇ (125 MHz, 37 mM)



Figure S12: ¹¹B NMR spectra of : (**A**) compound **3** in wet DMF-d₇ (4% H₂O, 128 MHz, 37 mM) and (**B**) the cyclic homopolymers of **3** (**htt**₂ and **htt**₃) in anhydrous DMF-d₇ (128 MHz, 37 mM). Data collected on the same machine and with the same acquisition parameters.



Figure S13: ¹H NMR spectra of the cyclic homopolymers of **3** in anhydrous DMF-d₇ at different concentrations: 37 mM (bottom), 1.4 mM (middle) and 0.14 mM (top). Blue and red arrows identify the signals of the major **htt**₂ dimer and minor **htt**₃ trimer respectively.



Figure S14 : Assignment of ¹H and ¹³C resonances for the two cyclic homopolymers of **3** in anhydrous DMF-d₇ on the ¹H (upper part) and ¹³C (lower part) NMR spectra . For clarity purpose, parts of the spectra free of signal were removed. Major htt_2 (•) and minor htt_3 (\blacktriangle) compounds are labelled. * This doublet can reasonably be assigned to the H₆ proton either from the free form or from a small linear assembly.



Figure S15 : Heteronuclear ¹H-¹³C NMR spectrum of **3** and cyclic homopolymers of **3** in D₂O (37 mM, pD 9.3). Peaks corresponding to bound methynes at position 2' and 3' are labelled.



Figure S16:¹ H NMR spectrum of cT₂ in anhydrous DMF-d₇.



Figure S18: ¹H NMR spectrum of U₂ in anhydrous DMF-d₇.



Figure S19: ¹H NMR spectrum of U_3 in anhydrous DMF-d₇.

5. Gaussian fits of the ¹H peaks intensity curve decay



Figure S20 : Gaussian fits of protons H_6 , $H_{1'}$ and $H_{4'}$ used to calculate the diffusion coefficients of **htt**₂ (upper part) and **htt**₃ (lower part) in anhydrous DMF-d₇ at 37 mM.



Figure S21 : Gaussian fits of protons H_6 , $H_{1'}$, H_5 and $H_{2'}$ used to calculate the diffusion coefficients of 3 in DMF-d₇ with 4 % of water (%v/v) at 34.3 mM.



Figure S22 : Gaussian fits to calculate the diffusion coefficients of $c(T_2)$ in anhydrous DMF-d₇.



Figure S23 : Gaussian fits to calculate the diffusion coefficients of $c(T_3)$ in anhydrous DMF-d₇.



Figure S24 : Gaussian fits to calculate the diffusion coefficients of U_2 in anhydrous DMF-d₇.



Figure S25 : Gaussian fits to calculate the diffusion coefficients of U_3 in anhydrous DMF-d₇.



Figure S26 : DOSY NMR spectrum of 3 in anhydrous DMF-d₇.

6. Conformational analysis of the sugar ring from the coupling constants

Coupling constants were calculated from the generalized Karplus equation (1):^[9]

$${}^{3}J_{HH} = P_{1}\cos^{2}\phi_{HH} + P_{2}\cos\phi_{HH} + P_{3} + \Sigma\Delta\chi_{i}\{P_{4} + P_{5}\cos^{2}(\zeta_{i}\phi_{HH} + P_{6}|\Delta\chi_{i}|)\}$$
(1)

using the empirical parameters $P_1 = 13.24$; $P_2 = -0.91$; $P_3 = 0$; $P_4 = 0.53$; $P_5 = -2.41$; $P_6 = 15.5$; $P_7 = 0.19$

and taking into account the electronegativity of the substituents according to the following equation:

$$\Delta \chi_{i} = \Delta \chi_{i}^{\alpha - \text{susbt}} - P_{7} \Sigma \Delta \chi_{i}^{\beta - \text{susbt}}$$
⁽²⁾

Following expressions were used to correlate between the parameters governing the conformation of the ribose:

$$\phi_{1'2'} = 123.3^{\circ} + 1.102 \phi_{m} \cos(P - 144^{\circ})$$
 (3)

$$\phi_{2'3'} = 0.2^{\circ} + 1.090 \phi_{\rm m} \cos P \tag{4}$$

$$\phi_{3'4'} = -124.9^{\circ} + 1.095 \phi_{\rm m} \cos(\mathsf{P} + 144^{\circ}) \tag{5}$$

7. Computational Method

As a starting point, structures of related c-di-GMP representing its "closed" and "open" conformers were extracted from the crystal structures 3I5A^[10] and 3HV8.^[11] The two guanines are on opposite faces/the same face in the open/closed conformer.^[12] Here, the boronate linkages bound to the C2' and C3' sugar carbons allowed to construct just a single conformer of the **htt**₂ dimer using the Molefacture module of the VMD software package.^[13] Finally, the structure of **htt**₂ was geometry optimized by means of the Gaussian 03 software package^[14] at the MP2/6-31G(d) level of theory.^[15]

8. Crystallization tests

Several methods were attempted to crystallize htt₂ and htt₃ :

- <u>Evaporation</u>: 10 mg of monomer **3** were dissolved in 1 mL of solvent (methanol, ethanol or water) in an opened test tube. The opened tube was let at room temperature or at 0°C for evaporation and crystallization.
- <u>Diffusion</u>: 10 mg of the monomer **3** were dissolved in 0.5 mL of methanol or DMF and 2 mL of another solvent (acetonitrile, diethyl ether, dichloromethane, dichloroethane, cyclohexane, pentane, toluene, dioxane or tetrahydrofuran) were slowly added. Test tubes were the placed at room temperature, at 0°C or at -20°C for several days.
- <u>Vapour diffusion</u>: 10 mg of the monomer **3** were dissolved in 0.5 mL of methanol or DMF. The test tube containing the mixture was placed into a closed falcon containing another solvent (acetonitrile, diethyl ether, dichloromethane, dichloroethane, cyclohexane or pentane). The falcon was placed at room temperature or 0°C for several days.
- <u>Dean Stark</u> : A suspension of the monomer **3** in a mixture of benzene/DMSO (9/1) or toluene/methanol (8/2) was heated under reflux with the use of a Dean Stark trap. The mixture was then slowly cooled to room temperature or 0°C. This method was also tested with 1 equivalent of triethylamine, 1,4-dipyridyl or 1,2-di(4-pirydil)ethylene.

Unfortunately, none of these methods allowed us to obtain any crystals.

9. Bibliography

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