Electronic supplementary information

Base recognition by L-nucleotides in heterochiral DNAs

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Note added after first publication: This Supplementary Information file replaces that originally published on 6 February 2012. The original version of Table S1 contained some incorrect nucleotide sequences in error. The correct sequences are provided in this updated version.

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1. General methods

Commercially available reagents were used without further purification. L-Nucleotides and their 5'-O-dimethoxytrityl-3'-(2-cyanoethyl)-*N*,*N*-diisopropylphosphoramidite derivatives were synthesized according to a previously reported procedure.¹ HPLC analyses were performed on a Shimadzu LC-10A system. A µBondasphere C18 5µm 100Å column (3.9×150 mm, Waters) was used with a linear gradient of acetonitrile in 50 mM triethylammonium acetate (TEAA, pH 7.0). Matrix-assisted laser desorption/ionization time-of-flight (MALDI TOF) mass spectra were acquired on a Voyager-DETM STR (Applied Biosystems) with 3-hydroxypicolinic acid as the matrix.

2. Oligonucleotide synthesis and characterization with MALDI-TOF mass spectrometry Oligodeoxyribonucleotides were synthesized on an Applied Biosystems model 392 automated DNA/RNA synthesizer. Reagents for the synthesis were purchased from Applied Biosystems Japan.

d(AAATCTGCG); m/z calcd for C₈₈H₁₁₃N₃₅O₅₁P₈ ([M+H]⁺), 2723.82, found: 2723.43; $d(CGCAGATTT); m/z calcd for C_{88}H_{113}N_{32}O_{53}P_8 ([M+H]^+), 2714.80, found: 2714.51;$ d(AALATCTGCG); m/z calcd for C₈₈H₁₁₃N₃₅O₅₁P₈ ([M+H]⁺), 2723.82, found: 2724.36; d(AAATCTLGCG); m/z calcd for C₈₈H₁₁₃N₃₅O₅₁P₈ ([M+H]⁺), 2723.82, found: 2723.53; d(AAATLCTGCG); m/z calcd for C₈₈H₁₁₃N₃₅O₅₁P₈ ([M+H]⁺), 2723.82, found: 2723.12; d(AAALTCTGCG); m/z calcd for C₈₈H₁₁₃N₃₅O₅₁P₈ ([M+H]⁺), 2723.82, found: 2722.57; d(CGCAGAATT); m/z calcd for C₈₈H₁₁₂N₃₅O₅₁P₈ ([M+H]⁺), 2723.82, found: 2723.77; d(CGCAGAGTT); m/z calcd for C₈₈H₁₁₂N₃₅O₅₂P₈ ([M+H]⁺), 2739.82, found: 2739.98; d(CGCAGACTT); m/z calcd for C₈₇H₁₁₂N₃₃O₅₂P₈ ([M+H]⁺), 2699.79, found: 2699.59; d(CGAAGATTT); m/z calcd for C₈₉H₁₁₃N₃₄O₅₂P₈ ([M+H]⁺), 2738.83, found: 2738.04; d(CGGAGATTT); m/z calcd for C₈₉H₁₁₃N₃₄O₅₃P₈ ([M+H]⁺), 2754.83, found: 2754.55; d(CGTAGATTT); m/z calcd for C₈₉H₁₁₄N₃₁O₅₄P₈ ([M+H]⁺), 2729.82, found: 2729.14; d(CGCAAATTT); m/z calcd for C₈₈H₁₁₃N₃₂O₅₂P₈ ([M+H]⁺), 2698.81, found: 2697.01; d(CGCACATTT); m/z calcd for C₈₇H₁₁₃N₃₀O₅₃P₈ ([M+H]⁺), 2674.78, found: 2673.62; $d(CGCATATTT); m/z calcd for C_{88}H_{114}N_{29}O_{54}P_8 ([M+H]^+), 2689.79, found: 2688.94;$ d(CGCAGGTTT); m/z calcd for C₈₈H₁₁₃N₃₂O₅₄P₈ ([M+H]⁺), 2730.80, found: 2729.11; d(CGCAGCTTT); m/z calcd for C₈₇H₁₁₃N₃₀O₅₄P₈ ([M+H]⁺), 2690.78, found: 2690.25; d(CGCAGTTTT); m/z calcd for C₈₇H₁₁₄N₂₉O₅₅P₈ ([M+H]⁺), 2705.79, found: 2704.63

3. Melting experiments

The concentrations of oligonucleotide solutions were calculated by using the equation and coefficients described by Bore.² The coefficients of the heterochiral oligomer were

assumed to be the same as those of the corresponding homochiral oligomer. Duplex solutions (6 mM) in 10 mM MgCl₂, 100 mM NaCl, and 70 mM MOPS (pH 7.1) were heated at 90 °C and cooled gradually to room temperature. Melting curves were measured at least twice at 270 nm on a JASCO V-560 spectrophotometer equipped with a programmable temperature control unit. The temperature was raised at a rate of 0.5°C/min and T_m values were obtained from the first-derivative plots of the melting curves.

complementary	strand	homo- and heterochiral strand				
		d(AADATCTGCG)		d(AALATCTGCG)		
d(CGCAGAXTT)	X = A	27.6 ± 0.0	(-14.5)	26.0 ± 0.4	(-7.6)	
	X = G	31.8 ± 0.4	(-10.3)	29.2 ± 0.2	(-4.4)	
	X = C	27.3 ± 0.3	(-14.8)	27.7 ± 0.5	(-5.9)	
	X = T	42.1 ± 0.3	F.M.	33.6 ± 0.2	F.M.	
		d(AAATCTDGCG)		d(AAATCTLGCG)		
d(CGXAGATTT)	X = A	22.1 ± 0.1	(-20.0)	19.3 ± 0.3	(-13.3)	
	X = G	22.5 ± 0.1	(-19.6)	31.6 ± 0.4	(-1.0)	
	X = C	42.1 ± 0.3	F.M.	32.6 ± 0.4	F.M.	
	X = T	25.0 ± 0.2	(-17.1)	22.4 ± 0.4	(-10.2)	
		d(AAAT <mark>DC</mark> TGCG)		d(AAAT <mark>LC</mark> TGCG)		
d(CGCAXATTT)	X = A	16.4 ± 0.0	(-25.7)	13.8 ± 0.2	-24.4	
	X = G	42.1 ± 0.3	F.M.	38.2 ± 0.2	F.M.	
	X = C	13.6 ± 0.2	(-28.5)	9.9 ± 0.1	(-28.3)	
	X = T	18.2 ± 0.2	(-23.9)	13.7 ± 0.1	(-24.5)	
		d(AAA <mark>DT</mark> CTGCG)		d(AAA <mark>LT</mark> CTGCG)		
d(CGCAGXTTT)	X = A	42.1 ± 0.3	F.M.	33.9 ± 0.1	F.M.	
	X = G	31.3 ± 0.1	(-10.8)	25.7 ± 0.5	(-8.2)	
	X = C	22.2 ± 0.2	(-19.9)	20.7 ± 0.5	(-13.2)	
	X = T	26.4 ± 0.2	(-15.7)	26.9 ± 0.1	(-7.0)	

4. Table S1. Melting temperature of mismatched homo- and heterochiral duplexes.^a

^{*a*}Samples contained 6 μ M duplex in 10 mM MgCl₂, 100 mM NaCl, and 70 mM MOPS (pH 7.1). Melting points are the average of at least two measurements ± standard deviation. Melting point differences from the fully matched (F. M.) duplex are shown in parenthesis.

5. References.

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