

Supplementary Materials

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S1: HPLC Profile of ODN3 and ODN5 Formation.

HPLC Conditions:

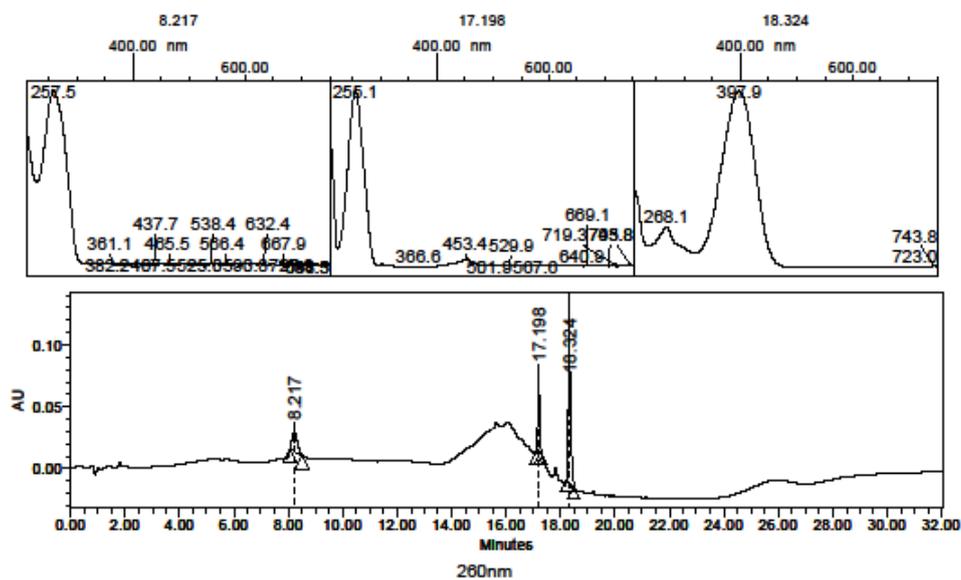
Column: X-Bridge OST C18 2.5 μ m, 4.6x50 mm

Gradient:

Mobile phase: A:TEEA 50 mM, pH 7 B: CH₃CN

t (min)	A(TEEA)	B(CH ₃ CN)
0	95	5
10	90	10
15	20	80
20	20	80
22	95	5

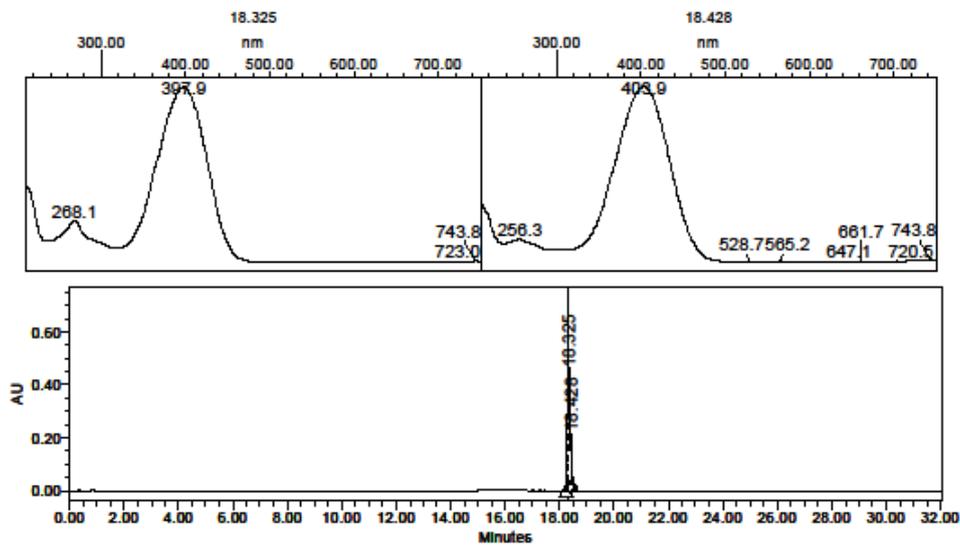
ODN3 t = 0 h, 25 °C, (detector λ = 260 nm)



Pureté et comparaison avec la librairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	8.217	191385	15.81		
2	17.198	293519	24.25		
3	18.324	725268	59.93		

ODN3 t = 0 h, 25 °C, (detector λ = 390 nm)

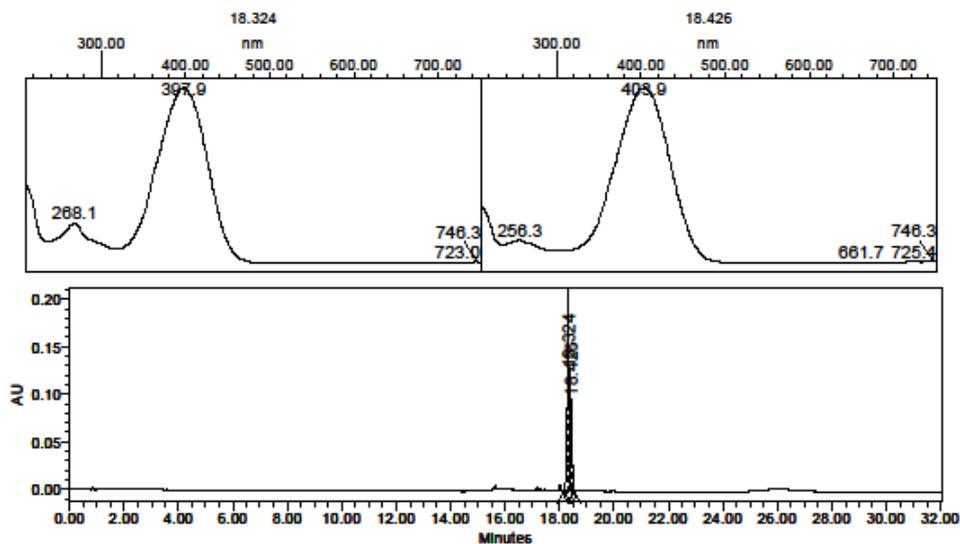


390nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	18.325	3369582	93.98		
2	18.428	215978	6.02		

ODN3 t = 0 h, 25 °C, (detector λ = 440 nm)

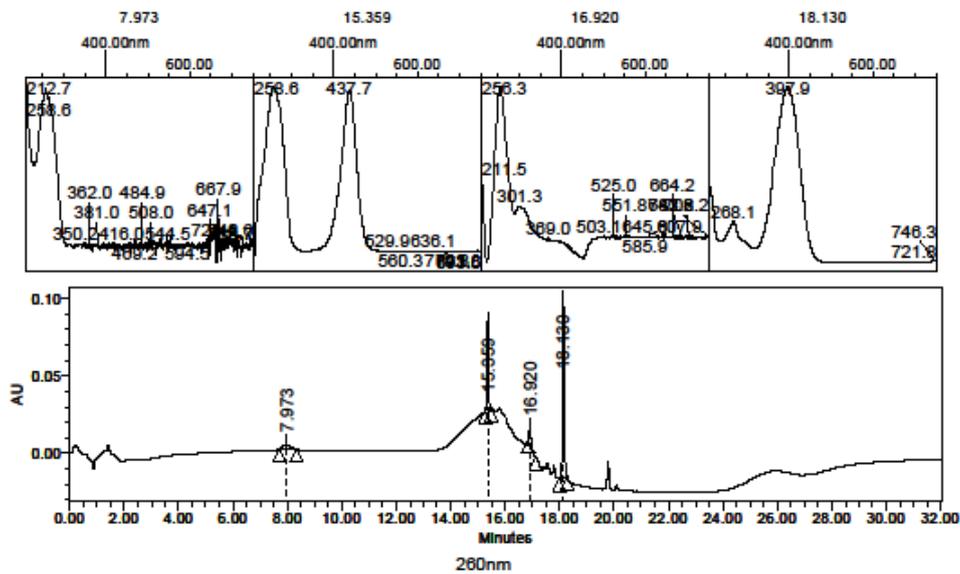


440nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	18.324	1007584	74.28		
2	18.426	348845	25.72		

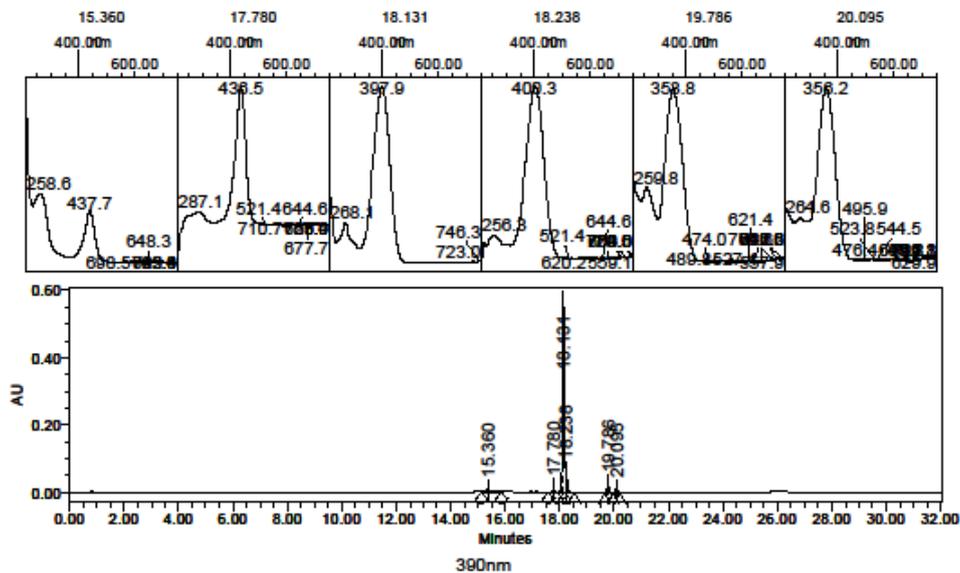
ODN3 t = 1 h, 25 °C, (detector λ = 260 nm)



Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	7.973	51269	5.87		
2	15.359	216215	24.77		
3	16.920	54280	6.22		
4	18.130	551261	63.14		

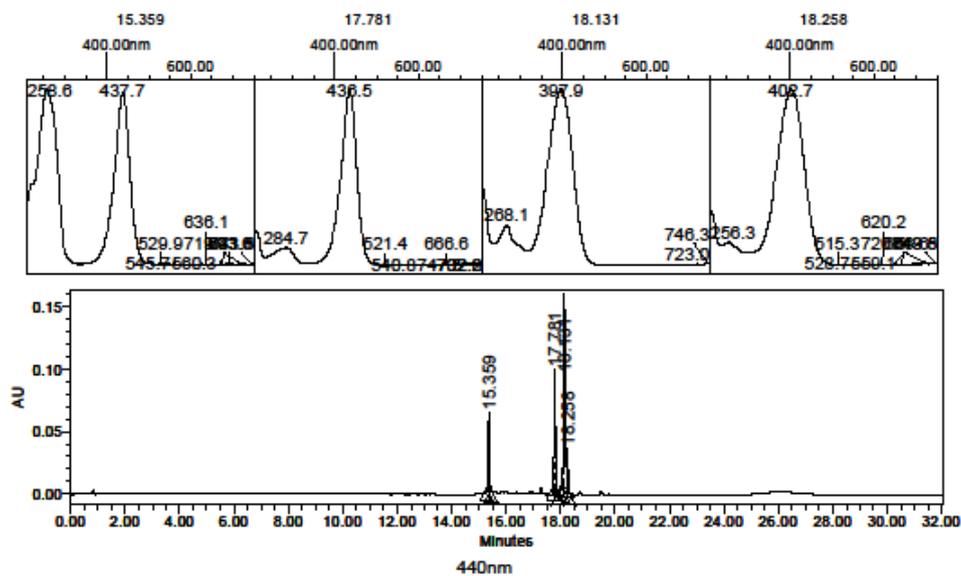
ODN3 t = 1 h, 25 °C, (detector λ = 390 nm)



Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	15.360	154199	4.64		
2	17.780	48580	1.46		
3	18.131	2726216	82.06		
4	18.238	246110	7.41		
5	19.786	104656	3.15		
6	20.095	42283	1.27		

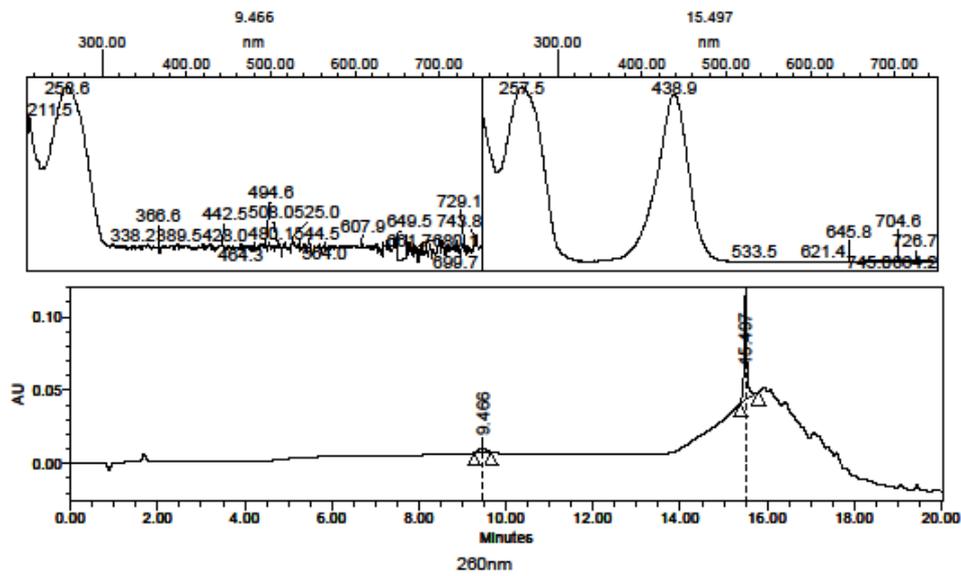
ODN3 t = 1 h, 25 °C, (detector λ = 440 nm)



440nm
Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	15.359	198531	14.78		
2	17.781	360985	26.88		
3	18.131	736310	54.83		
4	18.258	47145	3.51		

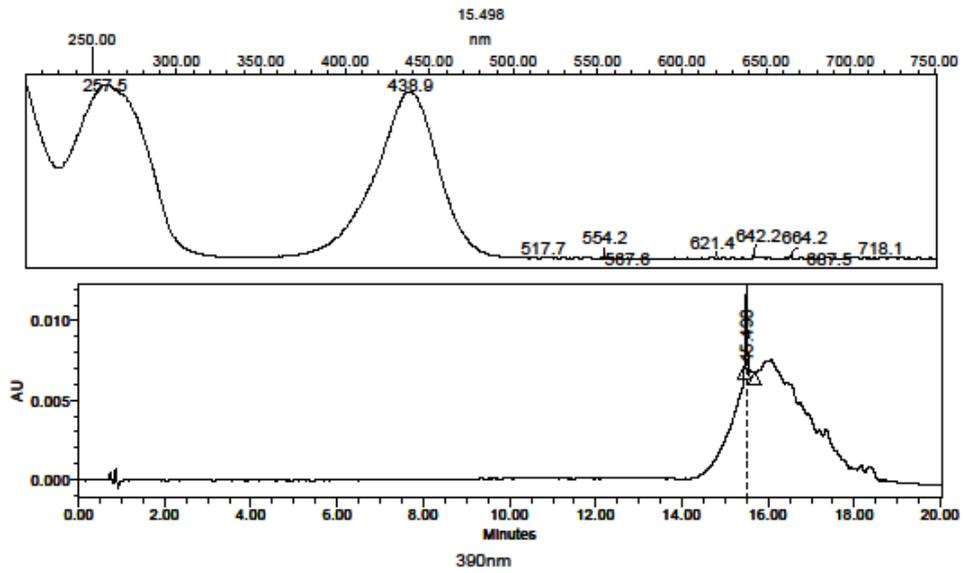
purified ODN3 purified (detector λ = 260 nm)



260nm
Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	9.466	35413	9.56		
2	15.497	334981	90.44		

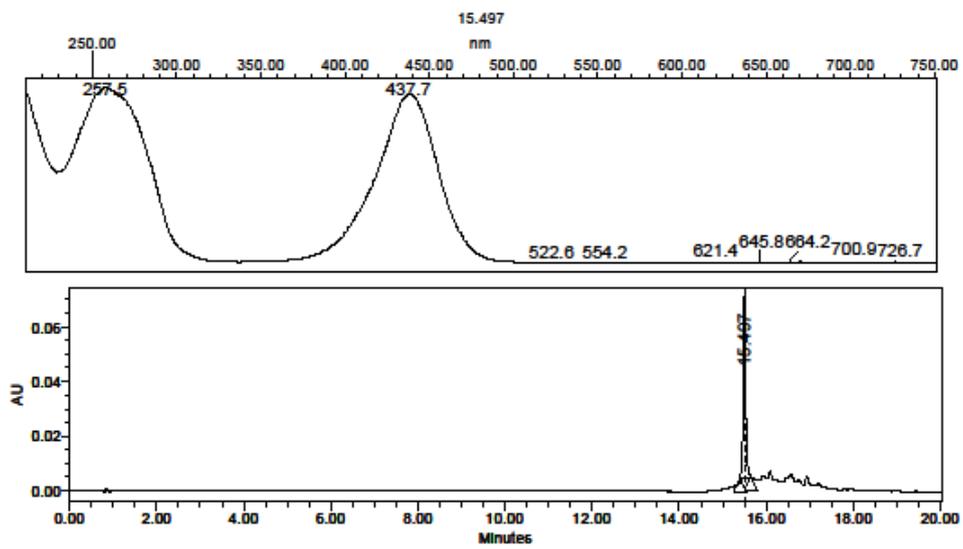
purified ODN3 purified (detector $\lambda = 390$ nm)



Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	15.498	15920	100.00		

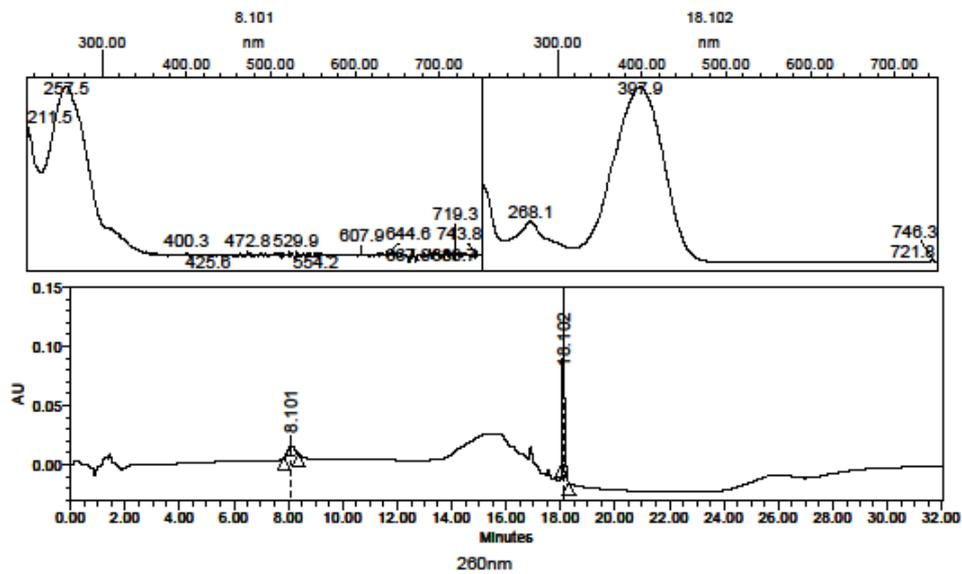
purified ODN3 purified (detector $\lambda = 440$ nm)



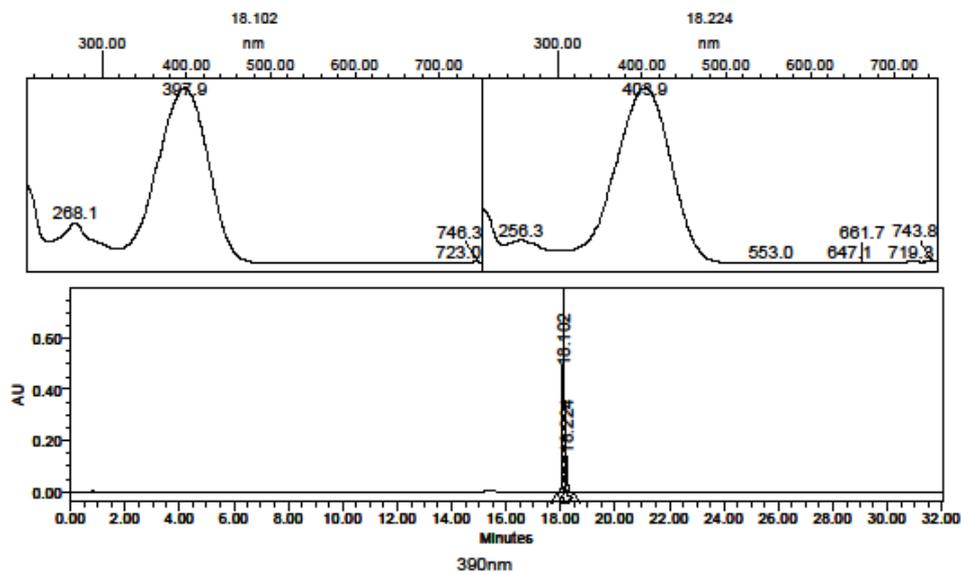
Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	15.497	281115	100.00		

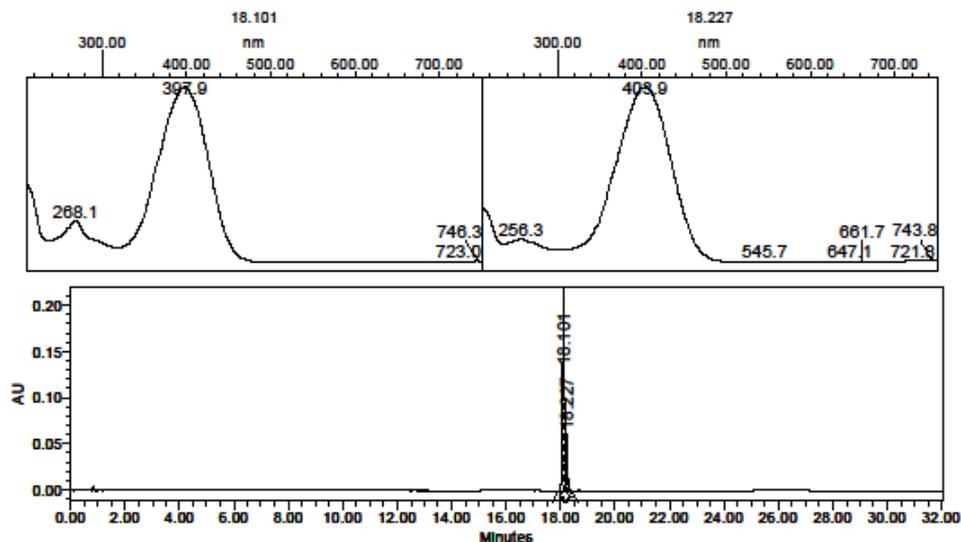
ODN5 t = 0 h, 25 °C, (detector $\lambda = 260$ nm)



ODN5 t = 0 h, 25 °C, (detector $\lambda = 390$ nm)



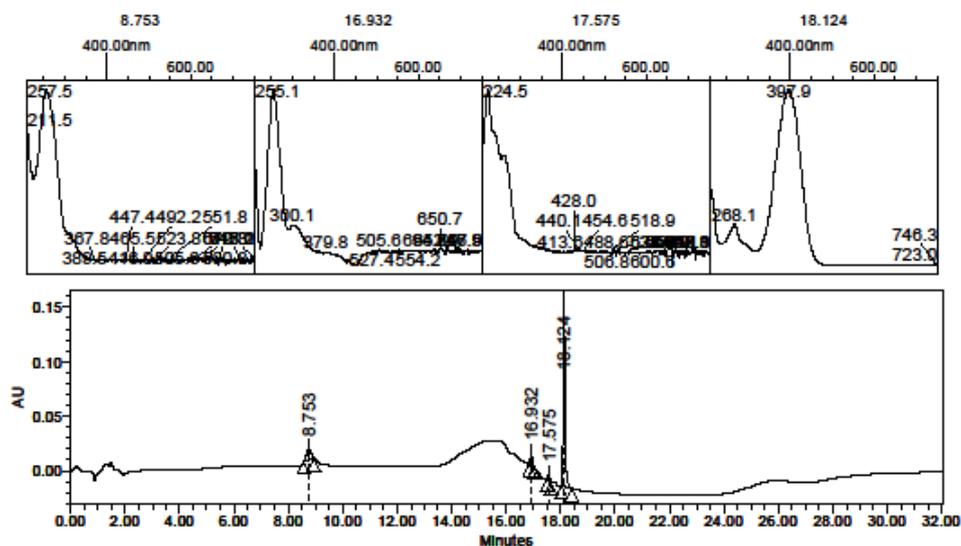
ODN5 t = 0 h, 25 °C, (detector λ = 440 nm)



440nm
Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	18.101	1046533	83.08		
2	18.227	213167	16.92		

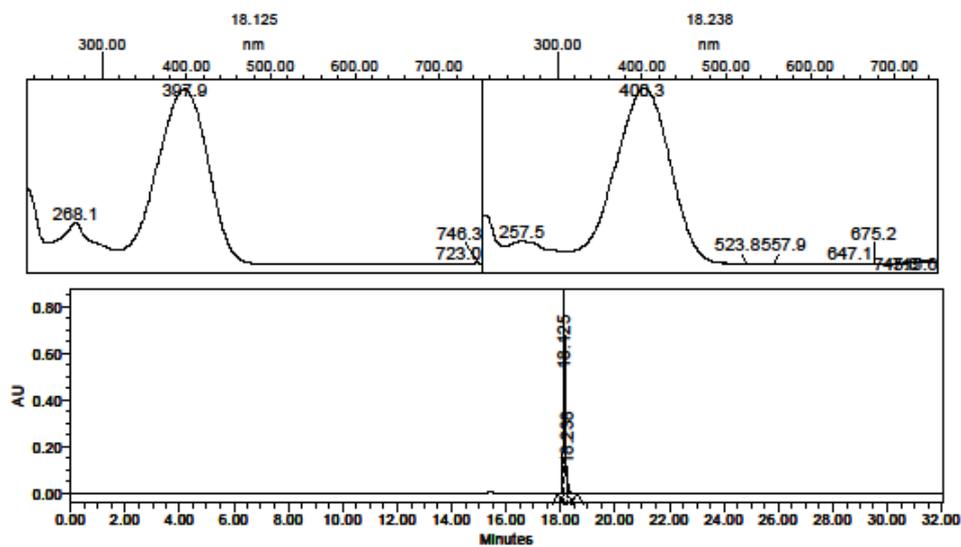
ODN5 t = 1 h, 25 °C, λ = (detector λ = 260 nm)



260nm
Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	8.753	129973	12.51		
2	16.932	45493	4.38		
3	17.575	22974	2.21		
4	18.124	840253	80.90		

ODN5 t = 1 h, 25 °C, (detector $\lambda = 390$ nm)

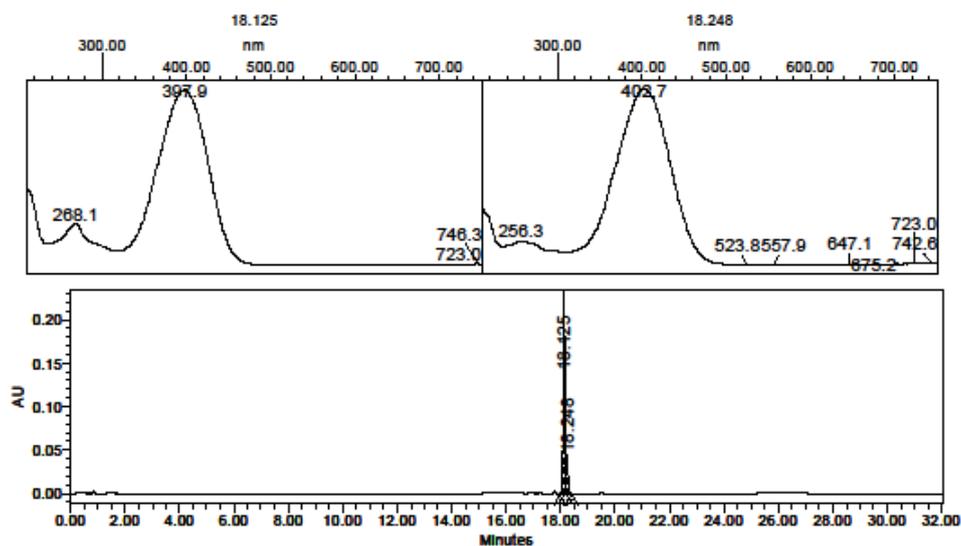


390nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	18.125	3942829	93.64		
2	18.238	267822	6.36		

ODN5 t = 1 h, 25 °C, (detector $\lambda = 440$ nm)

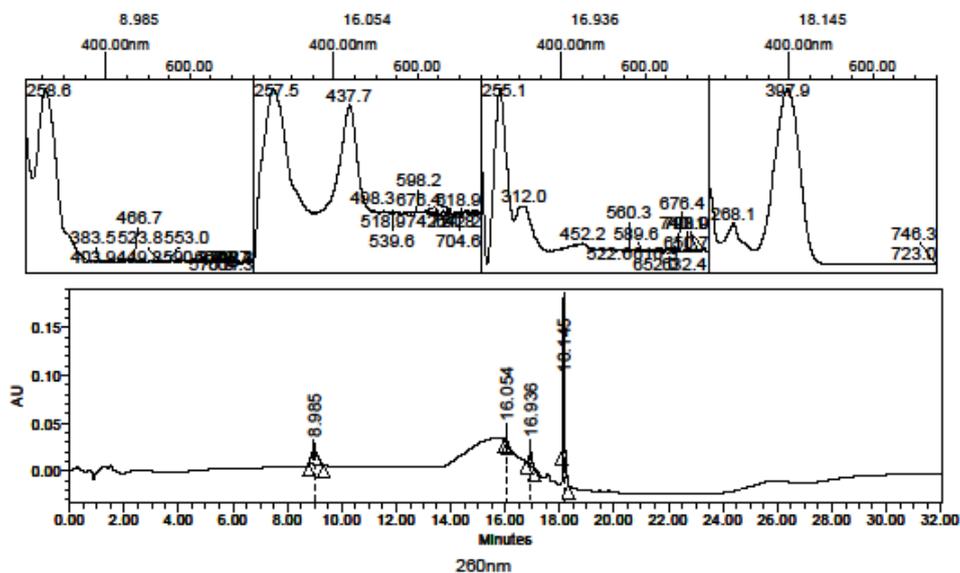


440nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	18.125	1053289	87.25		
2	18.246	153858	12.75		

ODN5 t = 2 h, 25 °C, (detector λ = 260 nm)

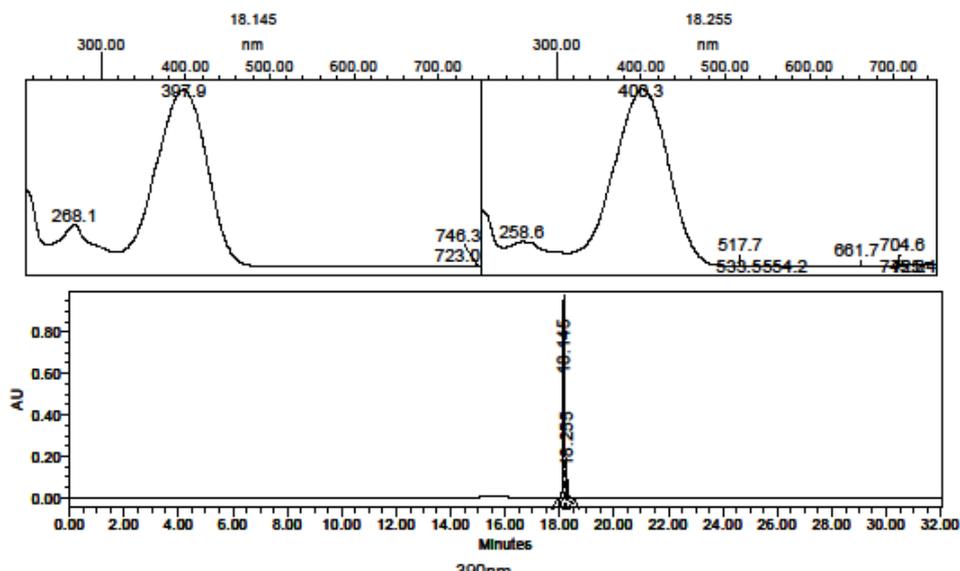


260nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	8.985	236949	22.21		
2	16.054	25005	2.34		
3	16.936	101731	9.54		
4	18.145	703055	65.91		

ODN5 t = 2 h, 25 °C (detector λ = 390 nm)

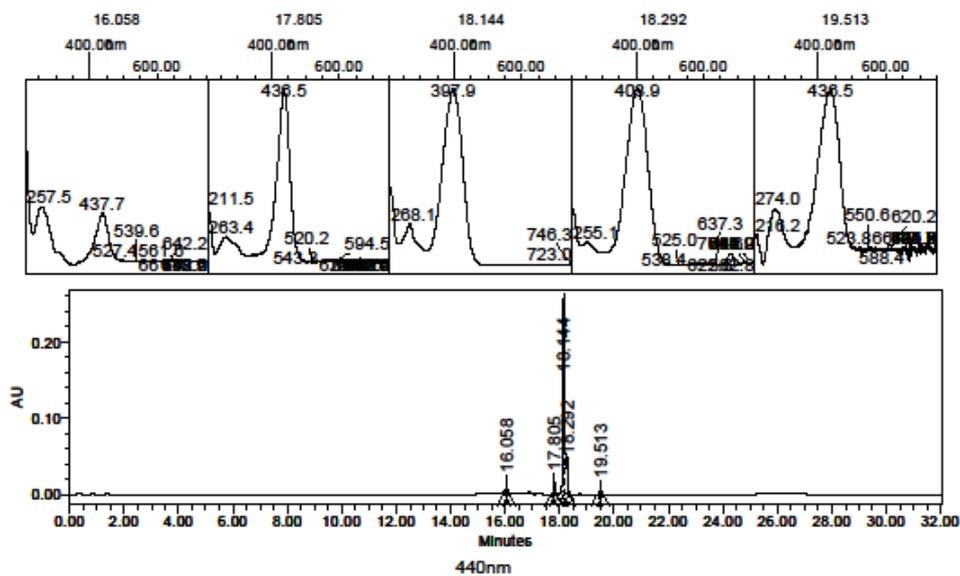


390nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	18.145	4611742	92.30		
2	18.255	384840	7.70		

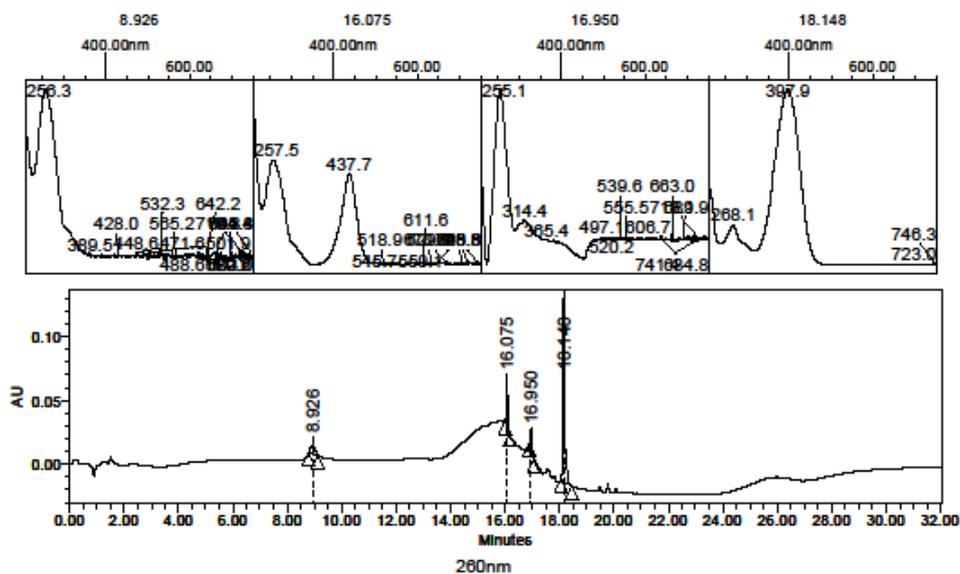
ODN5 t = 2 h, 25 °C, (detector $\lambda = 440$ nm)



440nm
Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	16.058	38109	2.51		
2	17.805	56459	3.71		
3	18.144	1345373	88.52		
4	18.292	51892	3.41		
5	19.513	28029	1.84		

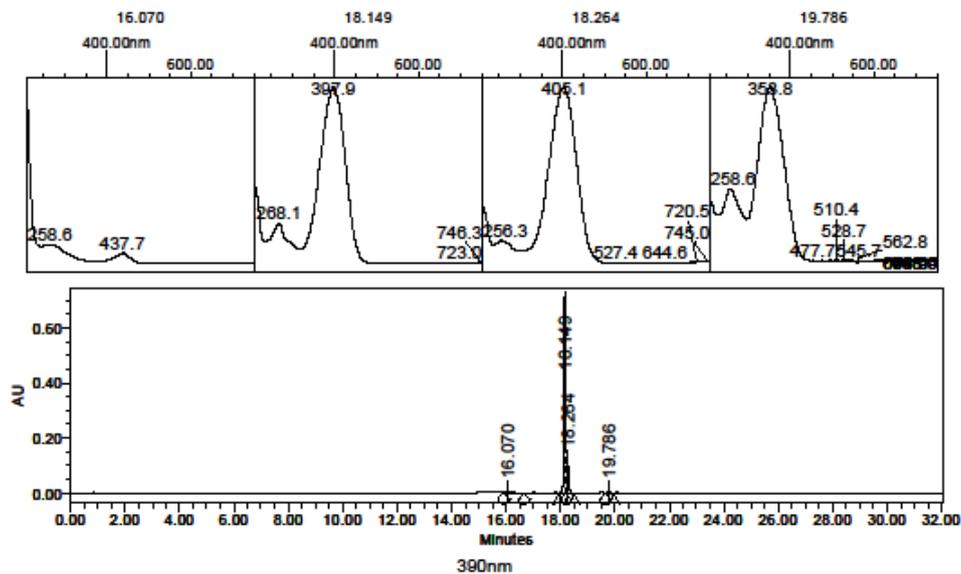
ODN5 t = 2 h at 25°C + 1h at 40°C (detector $\lambda = 260$ nm)



260nm
Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	8.926	69293	7.20		
2	16.075	131996	13.72		
3	16.950	41447	4.31		
4	18.148	719376	74.77		

ODN5 t = 2 h at 25°C + 1h at 40°C(detector λ = 390 nm)

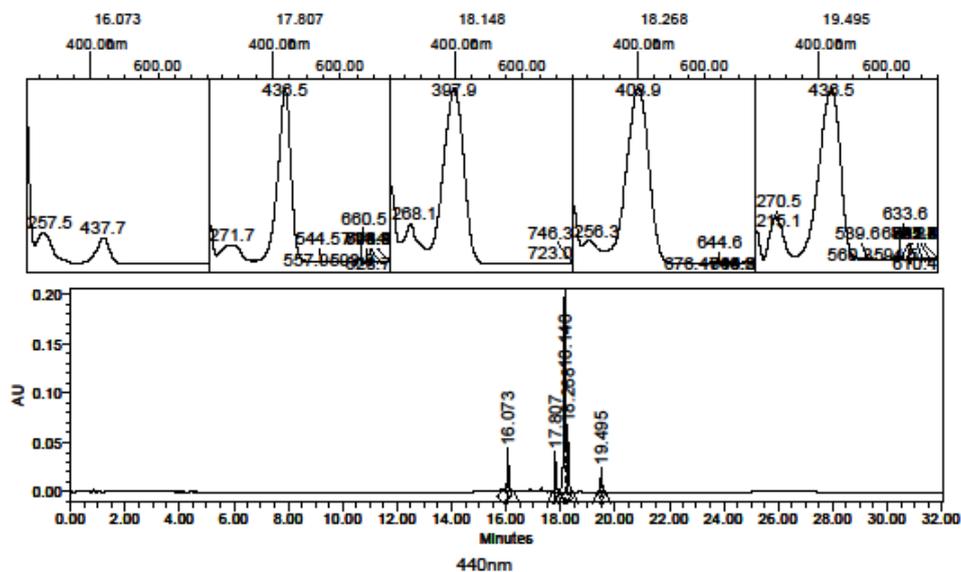


390nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	16.070	111948	2.75		
2	18.149	3364055	82.66		
3	18.264	550870	13.54		
4	19.786	43030	1.06		

ODN5 t = 2 h at 25°C + 1h at 40°C(detector λ = 440 nm)

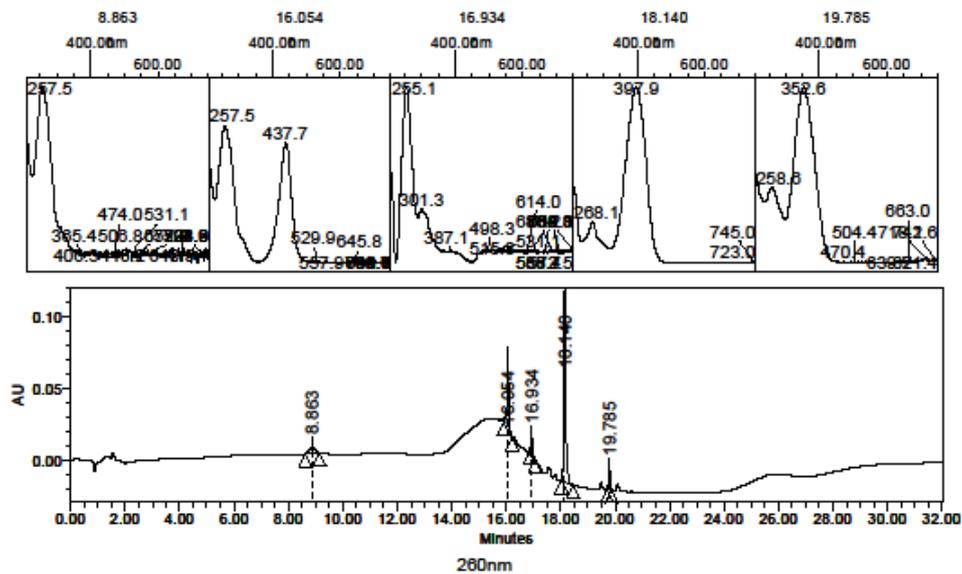


440nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	16.073	158151	9.63		
2	17.807	134906	8.22		
3	18.148	1008060	61.39		
4	18.268	262999	16.02		
5	19.495	77918	4.75		

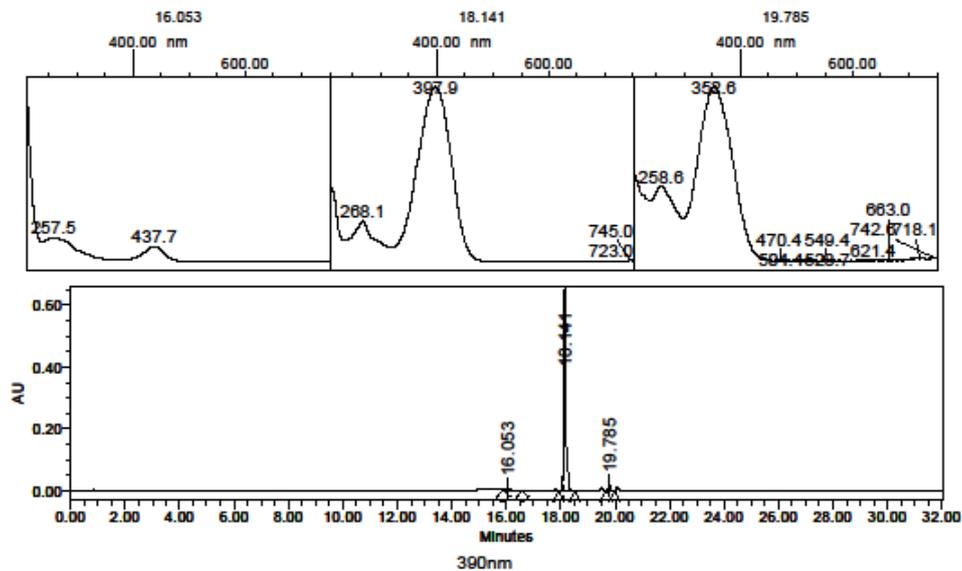
ODN5 t = 2 h at 25°C + 2h at 40°C(detector $\lambda = 260$ nm)



Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	8.863	69009	6.45		
2	16.054	231586	21.64		
3	16.934	67531	6.31		
4	18.140	635567	59.39		
5	19.785	66376	6.20		

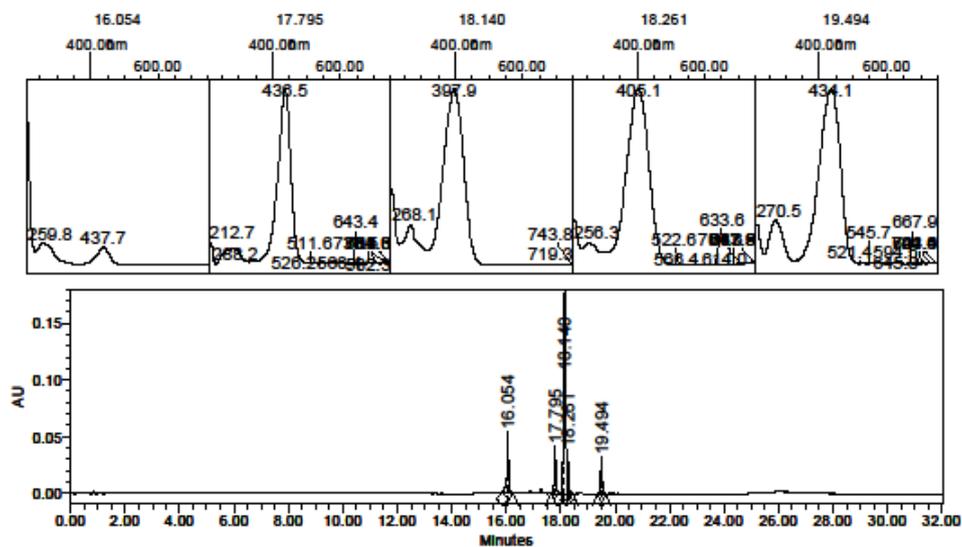
ODN5 t = 2 h at 25°C + 2h at 40°C (detector $\lambda = 390$ nm)



Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	16.053	86992	2.55		
2	18.141	3232345	94.88		
3	19.785	87547	2.57		

ODN5 t = 2 h at 25°C + 2h at 40°C (detector $\lambda = 440$ nm)

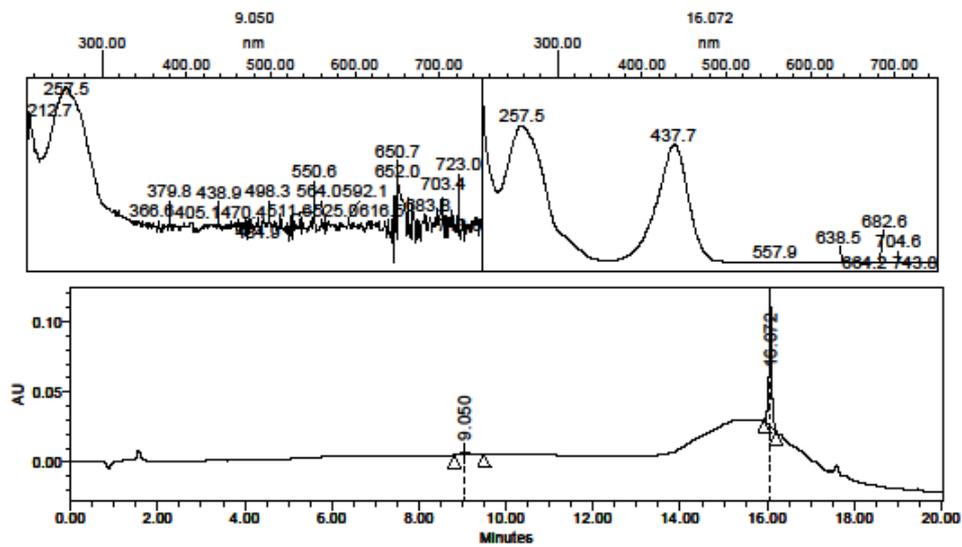


440nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	16.054	215736	14.53		
2	17.795	146316	9.86		
3	18.140	873630	58.86		
4	18.261	123395	8.31		
5	19.494	125293	8.44		

purified ODN5 (UV detector $\lambda = 260$ nm)

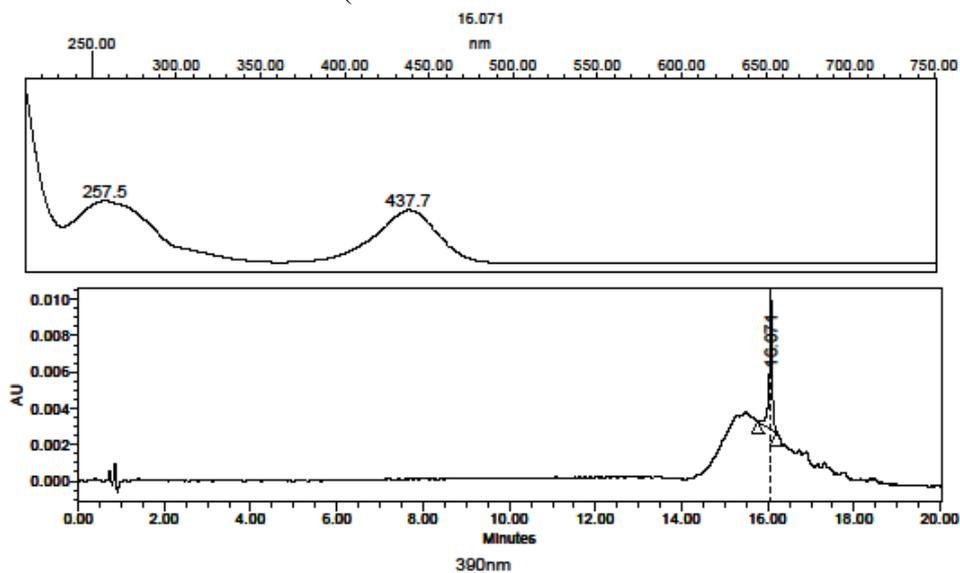


260nm

Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	9.050	27623	6.53		
2	16.072	395276	93.47		

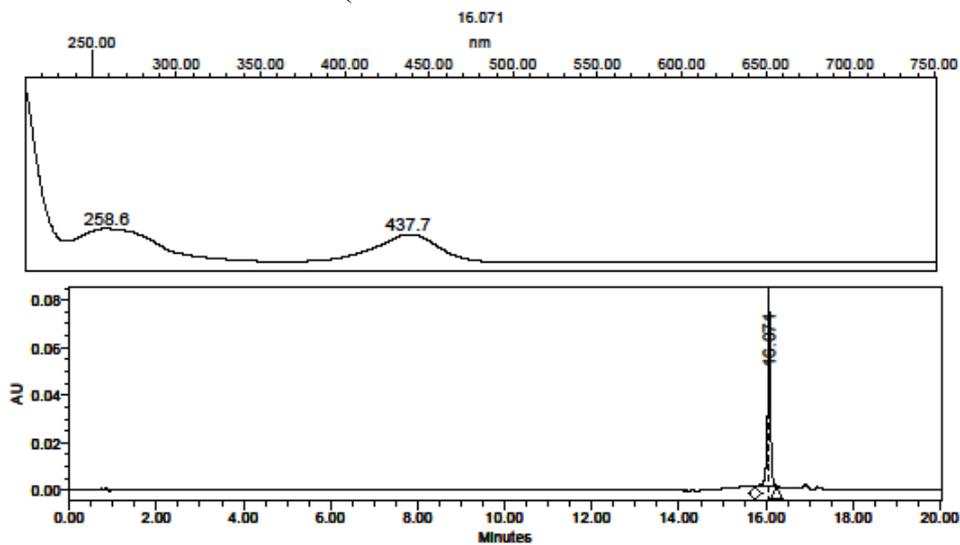
purified ODN5 (UV detector λ = 390 nm)



Pureté et comparaison avec la lirairie

Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	16.071	35279	100.00		

purified ODN5 (UV detector λ = 440 nm)

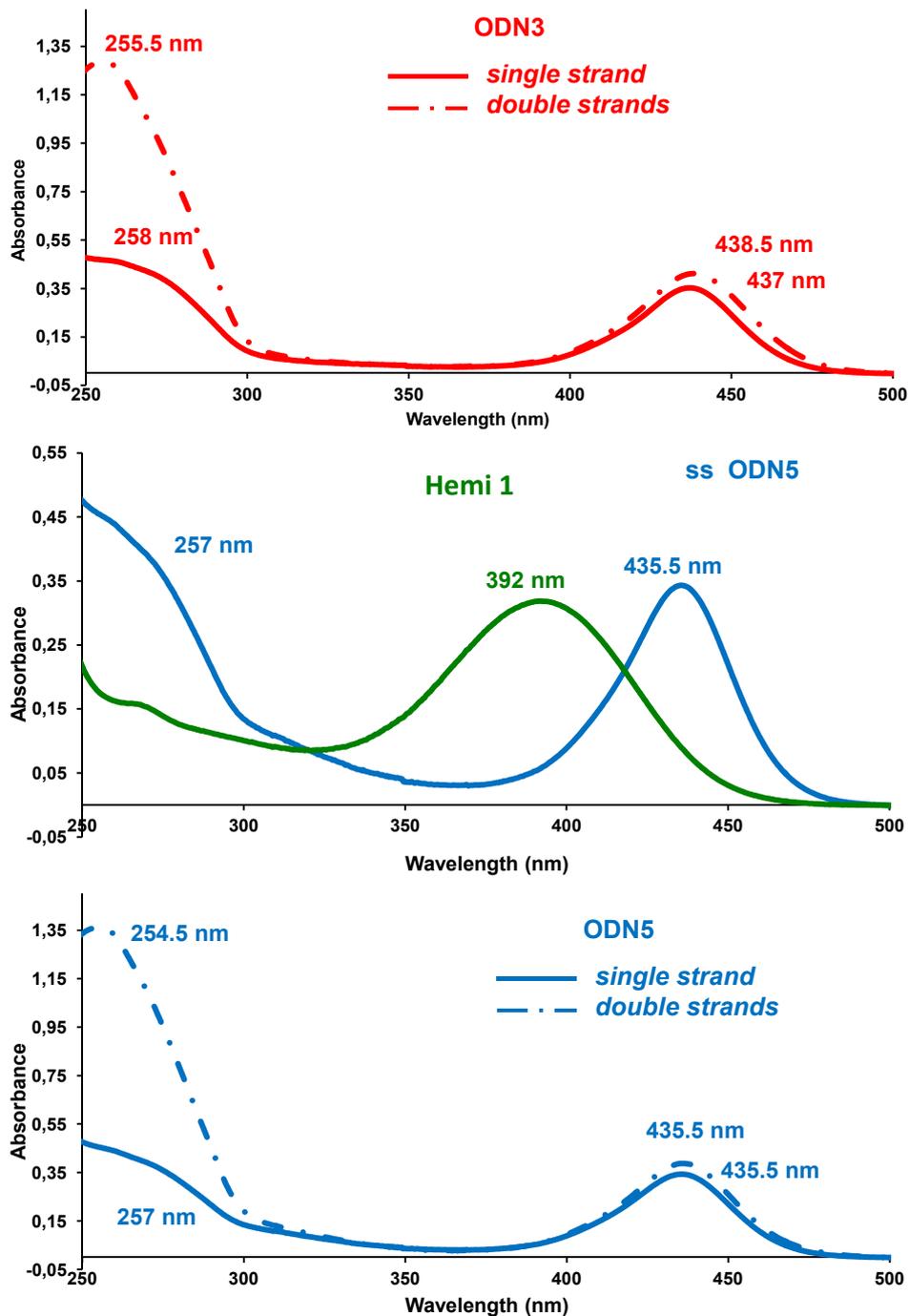


Pureté et comparaison avec la lirairie

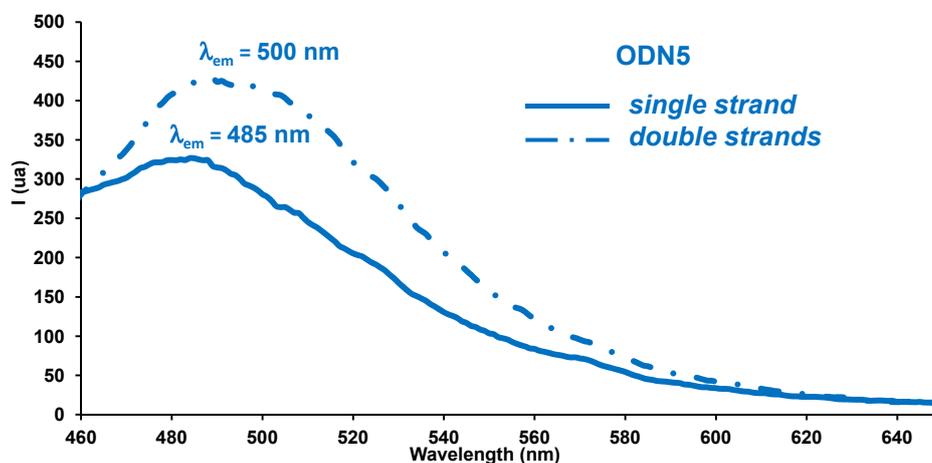
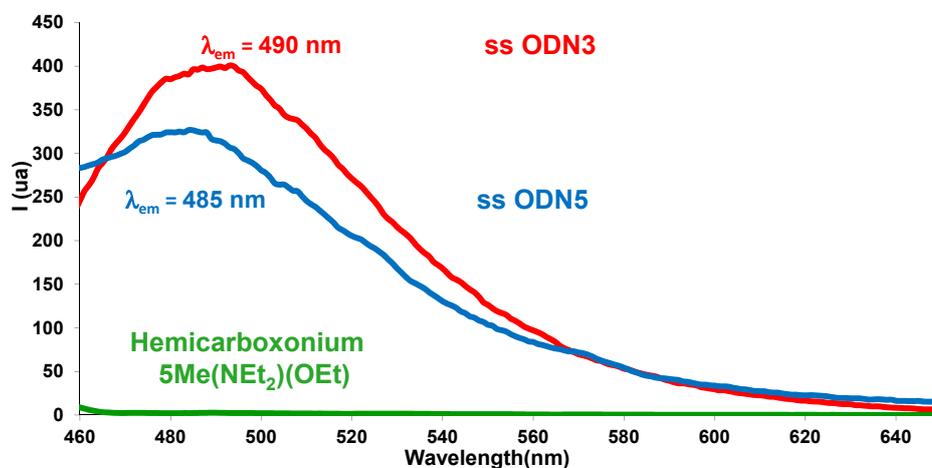
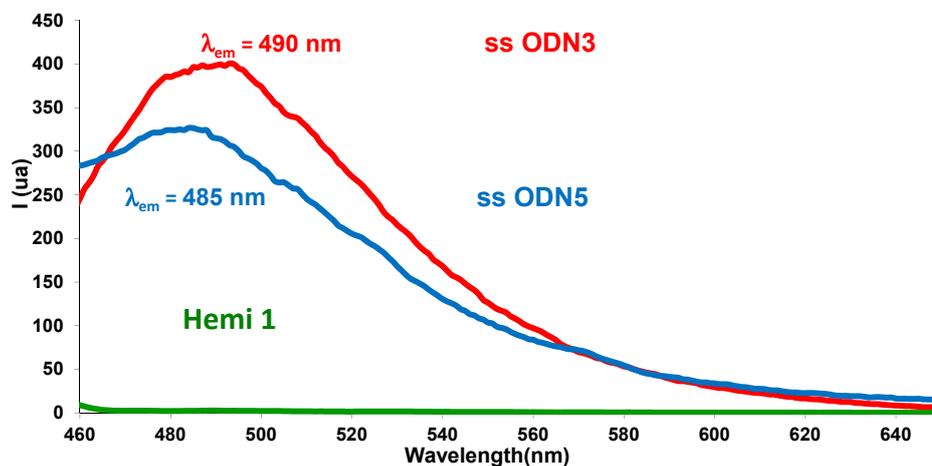
Name	RT	Area	% Area	Purity1 Angle	Purity1 Threshold
1	16.071	359302	100.00		

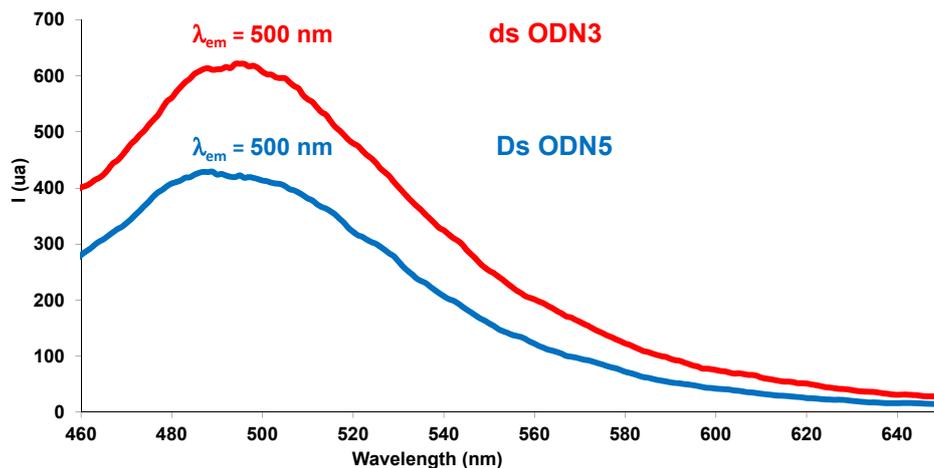
S2: UV Spectra of ODN3/5 in Single or Double Strand*Conjugates UV-Vis Properties*

[ODN] = $5.96 \cdot 10^{-6}$ M in phosphate buffer (10 mM Na_2HPO_4 , 100 mM NaCl, 1 mM EDTA, pH = 7)



Conjugates Fluorescence Properties

Emission fluorescence spectra $\lambda_{ex} = 420$ nm[ODN] = 1.49×10^{-6} M in phosphate buffer (10 mM Na_2HPO_4 , 100 mM NaCl, 1 mM EDTA, pH = 7)



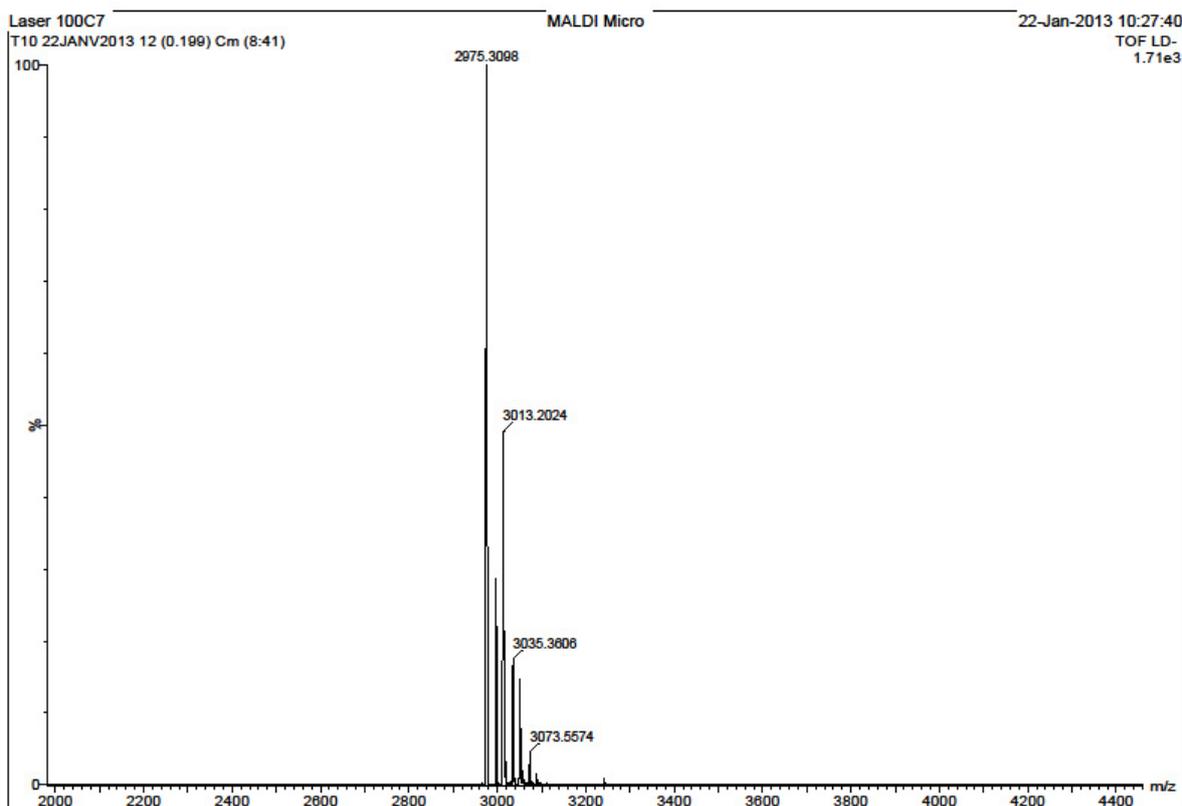
S3: Mass spectra of ODN2, ODN4, ODN3 and ODN5

ODN2

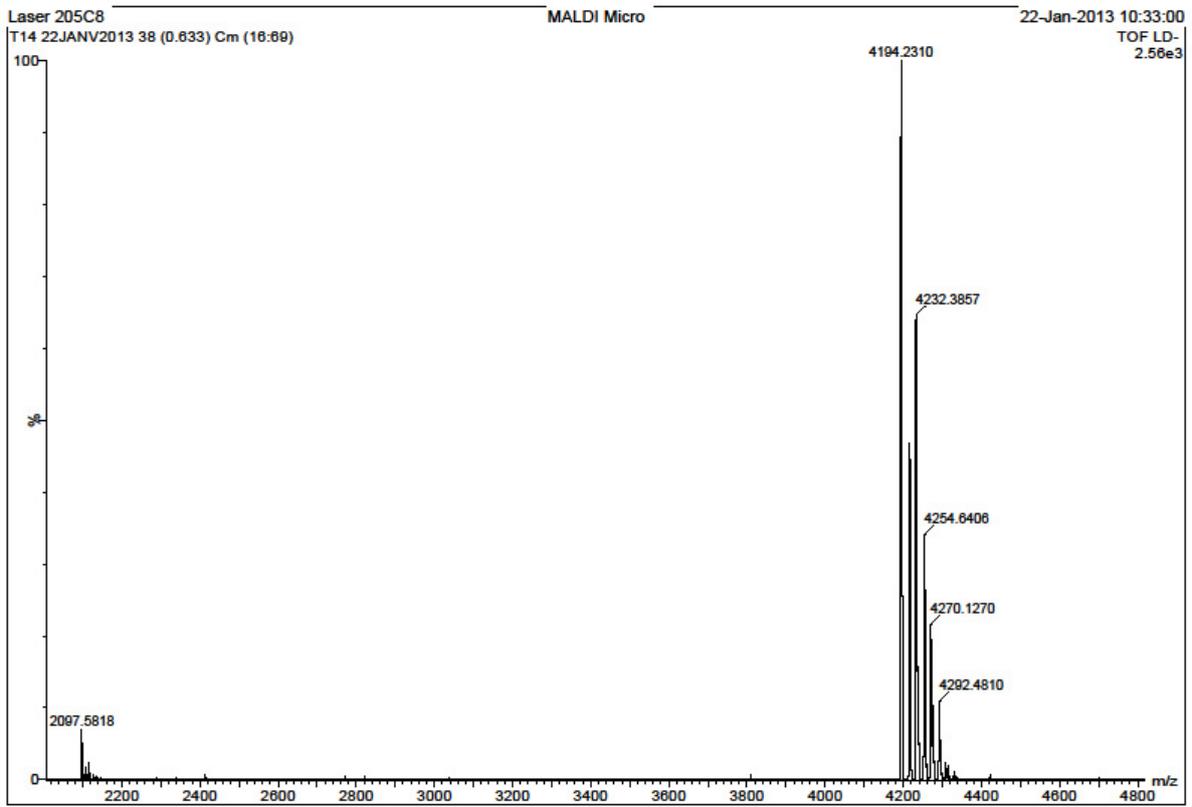
These analyses were recorded in a row with THAP matrix on a MALDI-ToF spectrophotometer WATERS Micro-MX.

	Exact mass [M]	Expected mass $m/z = [M-H]^+$	Obtained mass	% error
T10	2978.50	2977.50	2973.3	0.12%
T14	4194.69	4193.69	4192.2	0.035%
ODN2	3101.61	3100.61	3097.9	0.06%

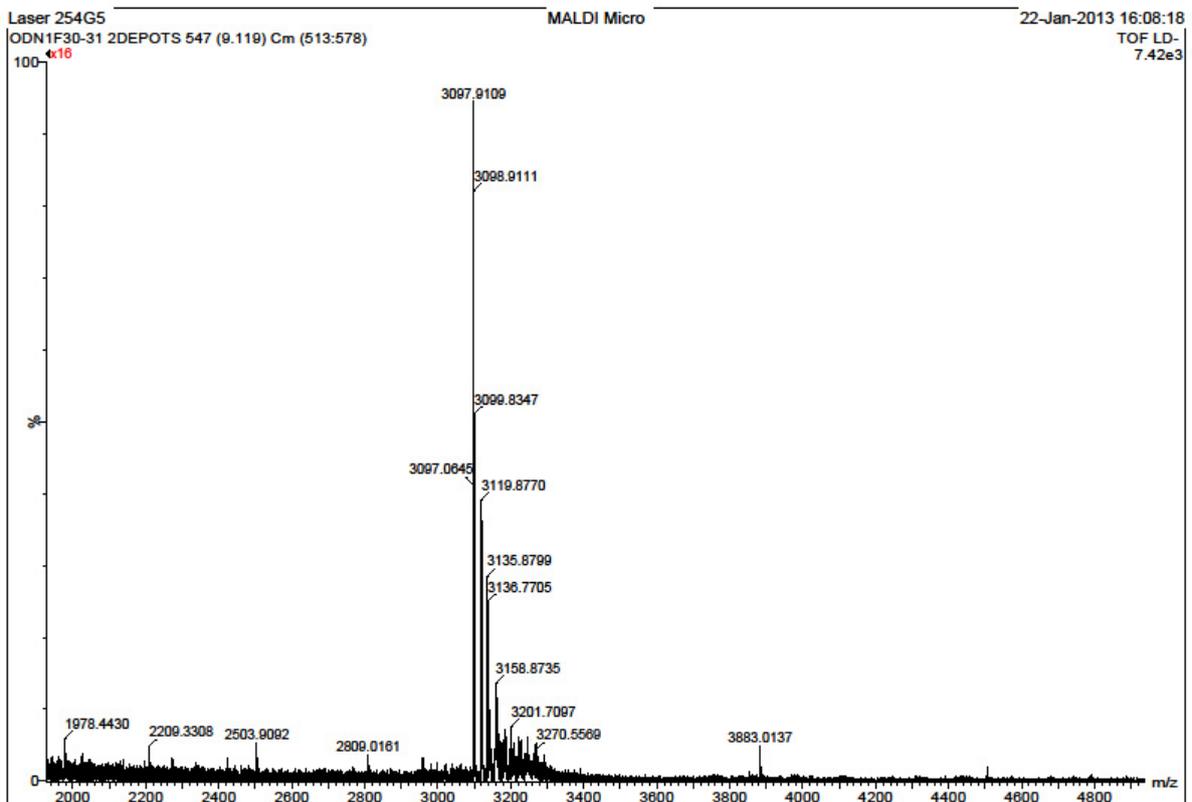
T10



T14



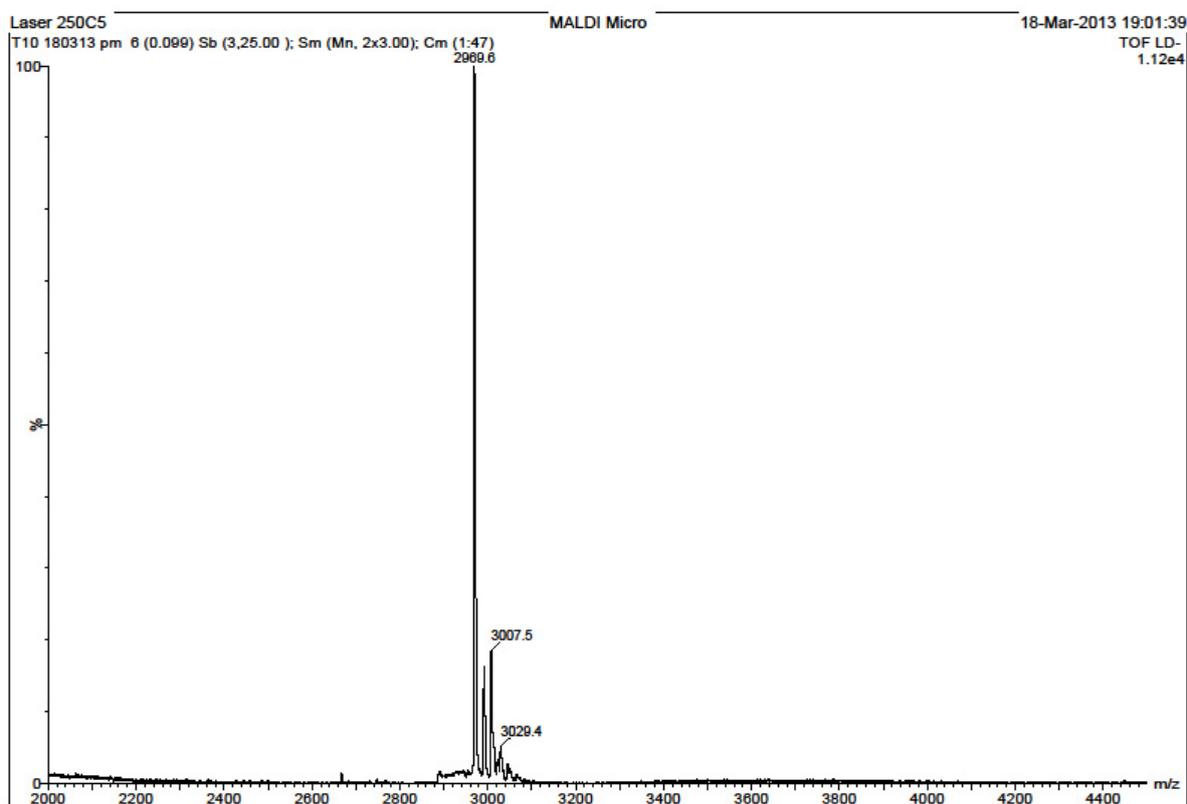
ODN2



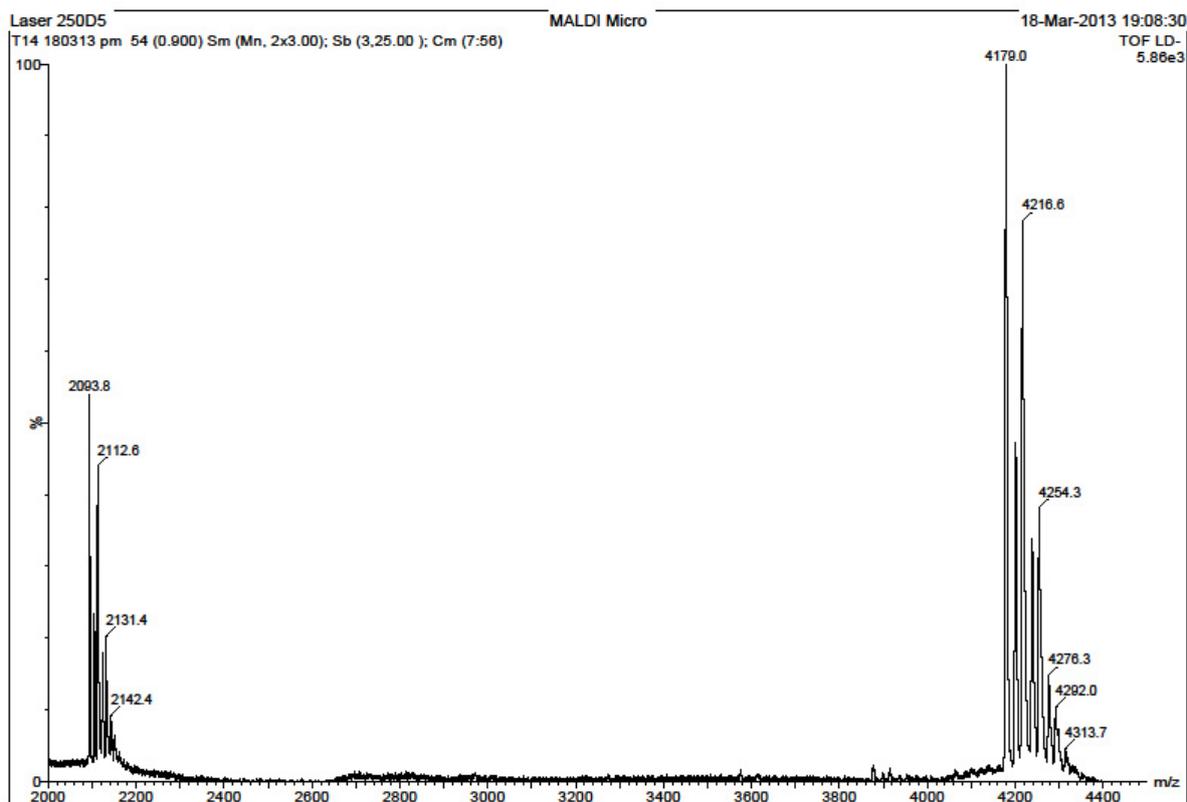
ODN4

These analyses were recorded in a row with THAP matrix on a MALDI-ToF spectrophotometer WATERS Micro-MX

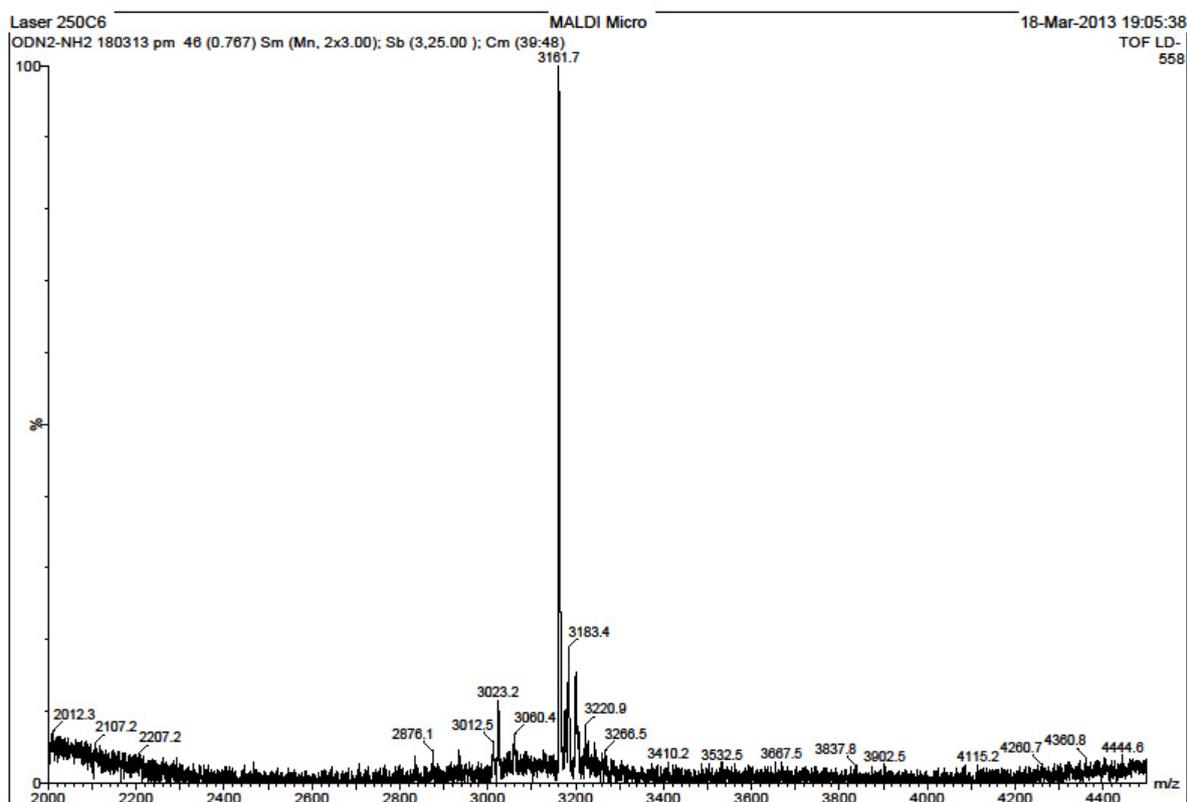
	Exact mass [M]	Expected mass $m/z = [M-H]^+$	Obtained mass	% error
T10	2978.50	2977.50	2969.6	0.26 %
T14	4194.69	4193.69	4179.0	0.35 %
ODN4	3172.64	3171.64	3161.7	0.31 %

T10

T14



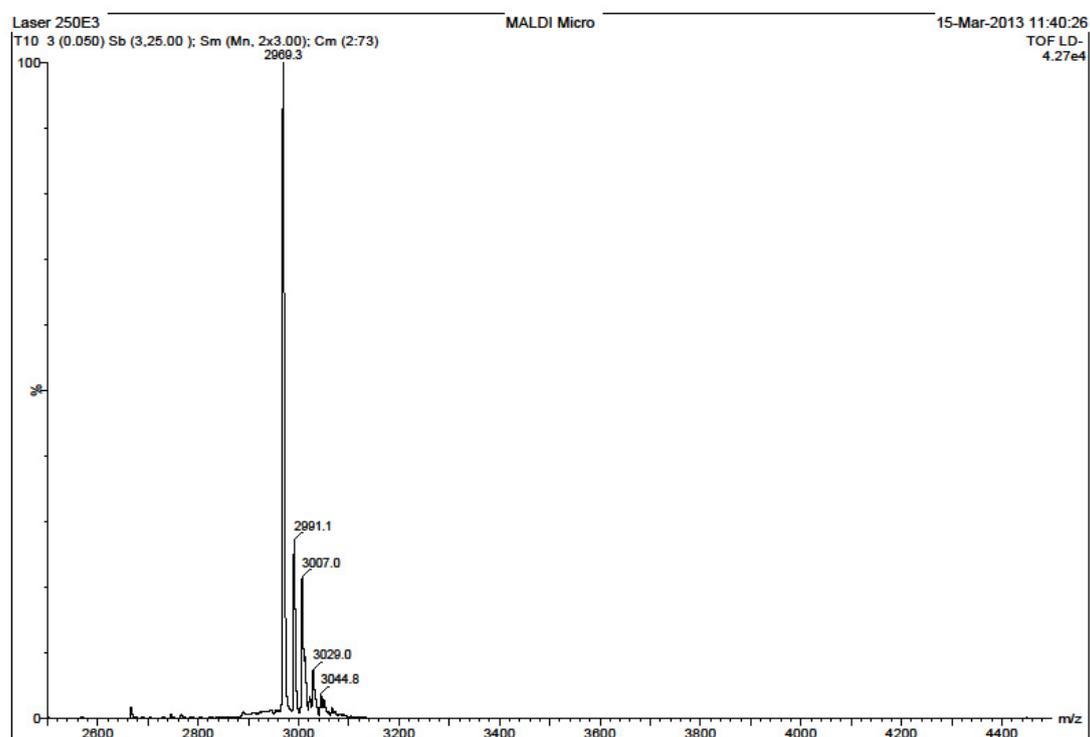
ODN4



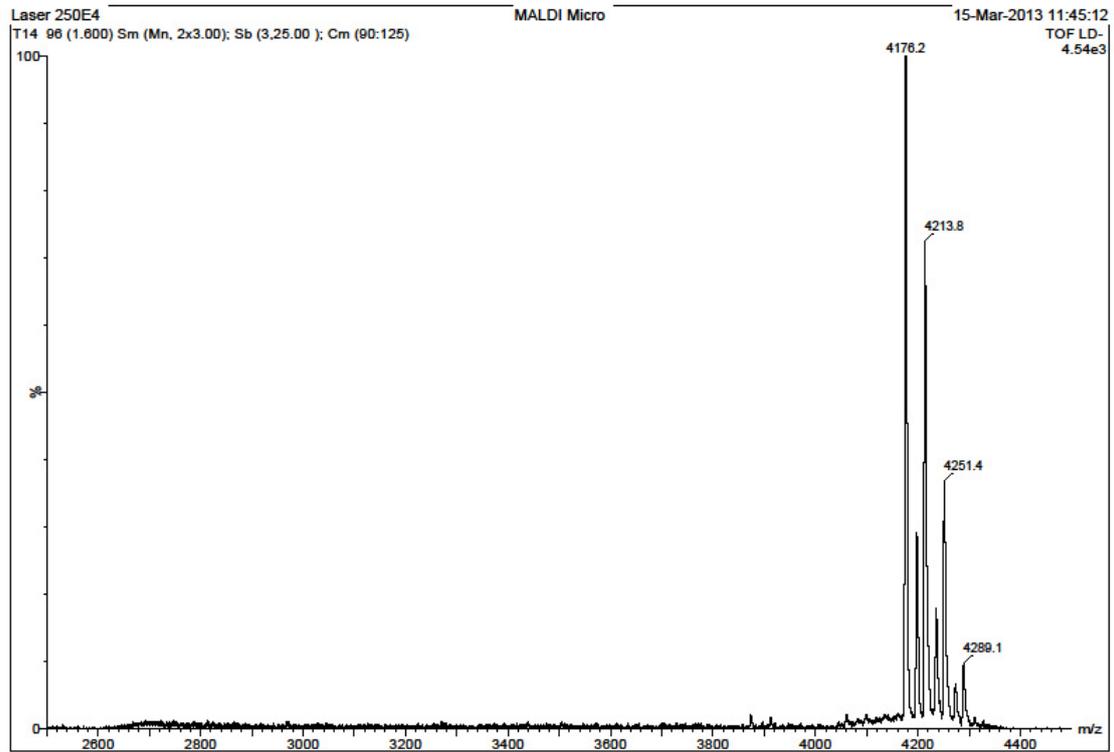
ODN3 and ODN5

These analyses were recorded in a row with THAP matrix on a MALDI-ToF spectrophotometer WATERS Micro-MX.

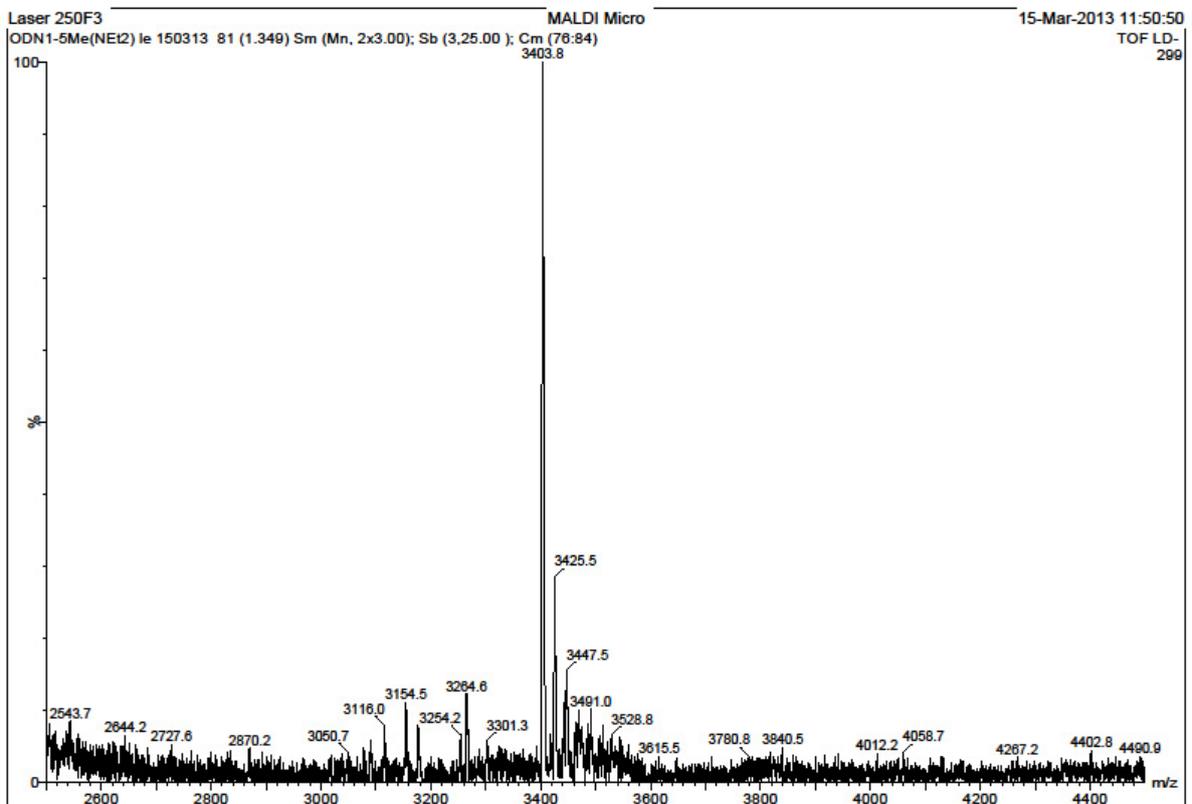
	Exact mass [M]	Expected mass m/z	Obtained mass	% error
T10	2978.50	[M-H] ⁺ 2977.50	2969.3	0.27%
T14	4194.69	[M-H] ⁺ 4193.69	4176.2	0.41%
ODN3	3417.81	[M-2H] ⁺ 3415.81	3403.8	0.35%
ODN5	3488.85	[M-2H] ⁺ 3486.85	3475.3	0.33%

T10

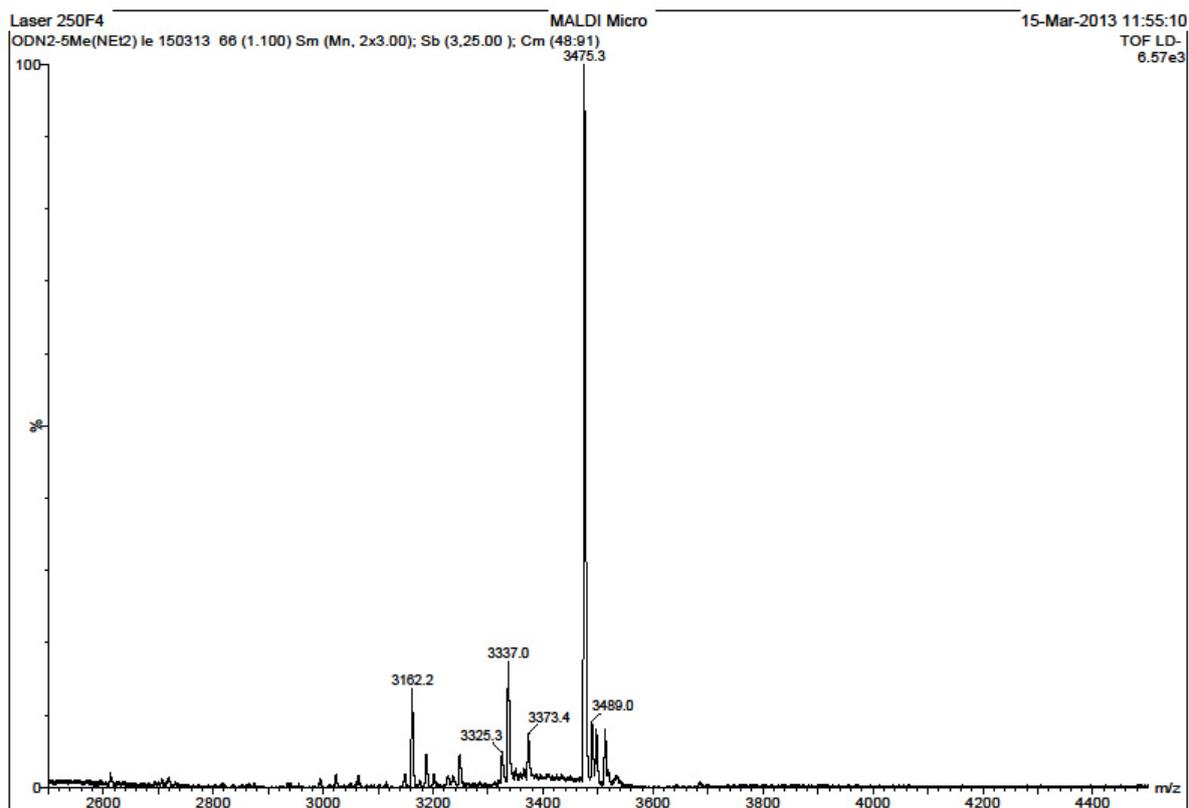
T14



ODN3



ODN5



S4: Melting Temperature of ODN3 and ODN5 within Duplex

The thermal denaturation curves were recorded on a Varian CARY-300 Bio UV-vis spectrophotometer with the Cary Win UV thermal software. All measurements were performed in 1.0 cm path-length micro cuvettes in 10 mM Na₂HPO₄, 100 mM NaCl, 1 mM EDTA, pH = 7, buffer. The concentration of the ODNs were calculated by measuring the absorbance at 260nm and at 80°C. Sample were heated at 90°C and cooled to 10°C at a rate of 0.5 °C/min and then warm up to 90°C at the same rate.

	Sequences	T _m °C (ΔT _m)
unmodified ADN/ADNc	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CAA GCG C) _{-3'}	53.0
unmodified ADN/ARNc	5'.d(GCG CTT GCC G) _{-3'} /5'.r(CGG CAA GCG C) _{-3'}	55.2
ODN3/ADNc	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CAA GCG C) _{-3'}	52.0 (-1.0)
ODN3/RcT4	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG TAA GCG C) _{-3'}	39.0 (-14.0)
ODN3/RcT5	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CTA GCG C) _{-3'}	43.8 (-10.0)
ODN3/RcT6	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CAT GCG C) _{-3'}	46.0 (-7.0)
ODN3/RcT7	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CAA TCG C) _{-3'}	Nd (nd)
ODN3/ARNc	5'.d(GCG CTT GCC G) _{-3'} /5'.r(CGC CAA GCG C) _{-3'}	50 .0 (-5.2)
ODN3/RcU4	5'.d(GCG CTT GCC G) _{-3'} /5'.r(CGG UAA GCG C) _{-3'}	34.0 (-21.2)
ODN3/RcU5	5'.d(GCG CTT GCC G) _{-3'} /5'.r(CGG CUA GCG C) _{-3'}	50.0 (-5.2)
ODN3/RcU6	5'.d(GCG CTT GCC G) _{-3'} /5'.r(CGG CAU GCG C) _{-3'}	49.0 (-6.2)
ODN3/RcU7	5'.d(GCG CTT GCC G) _{-3'} /5'. r(CGG CAA UCG C) _{-3'}	48.0 (-7.2)
ODN5/ADNc	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CAA GCG C) _{-3'}	52.0 (-1.0)
ODN5/RcT4	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG TAA GCG C) _{-3'}	38.0 (-15.0)
ODN5/RcT5	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CTA GCG C) _{-3'}	45.0 (-8.0)
ODN5/RcT6	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CAT GCG C) _{-3'}	45.0 (-8.0)
ODN5/RcT7	5'.d(GCG CTT GCC G) _{-3'} /5'.d(CGG CAA TCG C) _{-3'}	Nd (nd)
ODN5/ARNc	5'.d(GCG CTT GCC G) _{-3'} /5'. r(CGC CAA GCG C) _{-3'}	56.0 (+0.8)
ODN5/RcU4	5'.d(GCG CTT GCC G) _{-3'} /5'. r(CGG UAA GCG C) _{-3'}	40.0 (-15.2)
ODN5/RcU5	5'.d(GCG CTT GCC G) _{-3'} /5'. r(CGG CUA GCG C) _{-3'}	50.0 (-5.2)
ODN5/RcU6	5'.d(GCG CTT GCC G) _{-3'} /5'. r(CGG CAU GCG C) _{-3'}	48.0 (-7.2)
ODN5/RcU7	5'.d(GCG CTT GCC G) _{-3'} /5'. r(CGG CAA UCG C) _{-3'}	50.0 (-5.2)

Nd: no transition detected