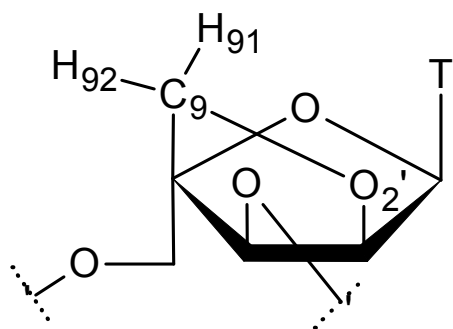


SUPPLEMENTARY MATERIAL

Table S1. RESP derived atomic charges and atom types for α -L-LNA thymine monomer



AMBER atom types differing from standard DNA

atom	atom type
H2'	H1
O2'	OS
C9	CT
H91	H1
H92	H1

atom	charge	atom	charge	atom	charge
O5'	-0.5176	H3'	0.0962	C5	0.0251
C5'	0.0176	O2'	-0.4268	C6	-0.2612
H5'	0.0974	C9	0.0613	O2	-0.5504
H5''	0.0974	H91	0.1025	H3	0.3259
C1'	-0.0166	H92	0.1025	O4	-0.5473
C2'	0.1170	O3'	-0.4991	H6	0.2118
C3'	0.1525	N1	-0.0320	C7	-0.2483
C4'	0.1439	C2	0.5645	H71	0.0876
O4'	-0.3603	N3	-0.4280	H72	0.0876
H1'	0.1856	C4	0.5639	H73	0.0876
H2'	0.1455				

Table S2. A selection of chemical shifts (in ppm) for the d(C^αL^LGA^αL^LA^αL^LGC):r(GCAUAUCAG) hybrid. Values are given at 25°C relative to DSS or inorganic phosphorous. ^a

	H6/H8	H1'	H2'	H2''	H3'	H4'	H5/H2/CH ₃	H1/H3	³¹ P
C1	7.81	6.28	2.39	2.91	4.85	4.19	5.99		
^α L ^L 2	7.41	5.71	5.00		4.82		1.78	–	–3.95
G3	7.70	5.93	2.25	2.67	4.71	4.42		11.98	–4.44
A4	8.14	6.32	2.75	3.05	5.07	4.50	7.73		–3.99
^α L ^L 5	6.79	5.41	4.68		4.77		1.55	13.59	–4.18
A6	8.02	6.27	2.55	2.97	4.78	4.45	7.25		–4.22
^α L ^L 7	6.81	5.49	4.89		4.77		1.36	13.38	–4.16
G8	7.54	6.04	2.35	2.64	4.52	4.38		12.46	–4.87
C9	7.41	6.15	2.09	2.30	4.44	4.08	5.12		–4.02
G10	7.91	5.55	4.66		4.48	4.22		–	
C11	7.80	5.55	4.66		4.57	4.47	5.29		–3.90
A12	8.01	5.97	4.48		4.64	4.52	7.14		–3.41
U13	7.48	5.43	4.62		4.42	4.45	5.07	12.89	–3.93
A14	8.09	5.93	4.40		4.62	4.54	6.81		–3.65
U15	7.59	5.45	4.35		4.39	4.39	4.85	13.44	–4.23
C16	7.98	5.61	4.52		4.58	4.43	5.67		–4.02
A17	8.00	5.83	4.30		4.70	4.45	6.93		–3.53
G18	7.34	5.77	3.99		4.21	4.47		–	–3.81

^a The protons in the C2',C4' linker (H6' and H6'') have the following chemical shift values:

^αL^L2: 4.36/4.44; ^αL^L5: 4.31/4.31; ^αL^L7: 4.24/4.24.

Table S3. A selection of helix parameters for the α -L-LNA:RNA hybrid. Values for standard A- and B-type duplexes are included for comparison

	X-disp (Å)	Y-disp (Å)	Tip (°)	Inclin (°)	Prop.Twist (°)		Twist (°)	Roll (°)	Rise (Å)
C1	-3.80	0.54	4.01	13.7	-10.5	C1-T2	23.3	-4.24	2.85
$\alpha^L T^L 2$	-3.75	0.21	2.52	15.9	-18.0	T2-G3	37.5	20.98	2.23
G3	-3.81	0.01	7.53	15.8	-20.0	G3-A4	35.5	-3.78	2.70
A4	-3.81	0.29	4.45	14.2	-15.0	A4-T5	26.3	-6.27	3.09
$\alpha^L T^L 5$	-3.56	-0.11	2.22	10.1	-11.4	T5-A6	38.4	5.25	2.65
A6	-3.55	-0.01	3.94	9.4	-20.6	A6-T7	29.4	-4.07	3.00
$\alpha^L T^L 7$	-3.48	-0.32	3.44	10.1	-19.8	T7-G8	40.4	7.19	2.85
G8	-3.46	-0.29	5.68	10.6	-16.2	G8-C9	31.1	-4.81	2.85
C9	-3.39	-0.18	3.92	11.2	-9.6				
A-type	-5.4	0	0	19.1	-13.7		32.7	0	2.56
B-type	-0.7	0	0	-5.9	-3.7		36.0	0	3.38

Table S4. Backbone torsion angles and sugar puckers (in degrees) in the *in vacuo* structure of the α -L-LNA:RNA hybrid

	α	β	γ	δ	ϵ	ζ	χ	ϕ_{\max}	P
C1			47	131	137	-63	-103	28	168
$\alpha^L T^L 2$	-172	203	182	-62	174	49	-125	67	14
G3	116	153	160	114	-176	-76	-151	18	123
A4	-78	186	48	133	-166	-89	-105	34	157
$\alpha^L T^L 5$	-85	154	172	-63	178	56	-158	61	16
A6	-161	190	80	115	-166	-81	-121	37	129
$\alpha^L T^L 7$	-101	163	172	-64	176	65	-150	61	16
G8	-135	183	50	106	-170	-78	-143	22	95
C9	-77	172	57	100			-130	29	88
G10			59 59	80	-125	-91	-171	0.377	7
C11	-66	145	65	79	-160	-66	-156	41	9
A12	-78	177	58	79	-164	-62	-159	42	6
U13	-80	176	69	78	-159	-65	-158	40	3
A14	-76	178	57	81	-173	-61	-157	40	6
U15	-79	178	80	82	-159	-62	-152	37	2
C16	-77	172	55	79	-152	-58	-146	39	9
A17	-76	186	36	76	-177	-63	-155	37	16
G18	-76	170	64	72			-160	44	12