

Supporting Information

Crystal Structure Studies of RNA Duplexes Containing s²U:A and
s²U:U Base Pairs

Jia Sheng, Aaron Larsen, Benjamin D. Heuberger, J. Craig Blain, and Jack W. Szostak*

Howard Hughes Medical Institute, Center for Computational and Integrative Biology, and Department of Molecular
Biology, Simches Research Center, Massachusetts General Hospital, Boston, MA 02114

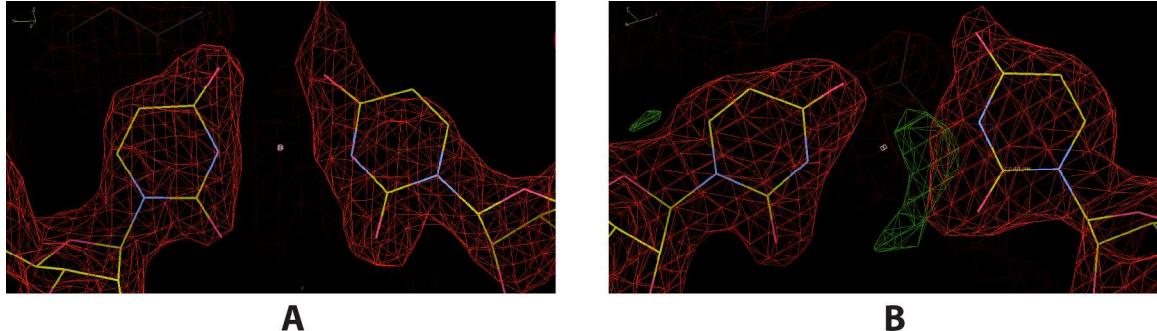


Figure S1. Experimental density map of (A) U5:U10 pair in chain EF (corresponding to Fig. 4D); and (B) U5:U10 pair in chain GH (corresponding to Fig. 4E). The red color represents the 2Fo-Fc map with σ level 1.3; the green color represents the Fo-Fc map with σ level 3.0.

Table S1. Local base-pair structural parameters in the s^2UU 7mer (s^2UU): [5'-uagc(s^2UU)cc-3'/3'-aucgUgg-5'] and the native-UU 7mer: [5'-uageucc-3'/3'-aucgugg-5'], including all four duplexes in the asymmetric unit (N-UU-1, N-UU-2, N-UU-3 and N-UU-4).

		Local Base-Pairs							
		U-A	A-U	G-C	C-G	U-U	C-G	C-G	Ave.
Shear (Å)	N-UU-1	-0.34	0.16	-0.3	0.77	-2.45	0.33	0.33	-0.21
	N-UU-2	0.02	-0.02	-0.39	0.39	2.08	0.18	0.52	0.40
	N-UU-3	-0.3	-0.1	-0.25	0.34	0.68	0.12	0.17	0.09
	N-UU-4	-1.02	-0.03	-0.36	0.03	-1.87	0.06	0.66	-0.36
	s^2UU	-0.33	0.06	-0.43	0.24	2.26	0.27	0.38	0.35
	N-UU-1	-0.07	-0.02	-0.17	-0.05	-1.66	-0.22	-0.19	-0.34
Stretch (Å)	N-UU-2	-0.22	-0.14	-0.13	-0.23	-1.8	-0.23	-0.2	-0.42
	N-UU-3	-0.24	-0.36	-0.31	-0.13	-1.36	-0.2	-0.18	-0.40
	N-UU-4	-0.14	-0.26	-0.19	-0.25	-0.74	-0.3	-0.36	-0.32
	s^2UU	-0.05	-0.14	-0.23	-0.17	-1.56	-0.16	-0.17	-0.35
	N-UU-1	-0.33	-0.11	-0.13	-0.23	0.18	-0.46	-0.27	-0.19
Stagger (Å)	N-UU-2	-0.24	-0.18	0.08	-0.27	0.42	-0.11	0.11	-0.03
	N-UU-3	0.07	0	-0.17	-0.22	0.21	0.1	-0.31	-0.05
	N-UU-4	-0.29	0.04	-0.06	0.04	-0.39	-0.17	-0.1	-0.13
	s^2UU	0.07	0.26	-0.02	-0.12	0.12	-0.12	-0.01	0.03
	N-UU-1	6.68	0.92	-1	4.15	11.53	13.57	2.86	5.53
Buckle (°)	N-UU-2	1.39	1.7	2.22	14.86	0.57	8.12	0.49	4.19
	N-UU-3	2.74	1.16	-0.6	3.32	1.81	2.81	8.14	2.77
	N-UU-4	5.57	-0.38	0.91	6.38	4.7	2.2	5.21	3.51
	s^2UU	-3.60	6.66	4.95	6.43	-6.89	5.63	-0.17	1.86
	N-UU-1	-15.02	-8.25	-12.97	-18.54	-9.36	-18.07	-1.28	-11.93
Propeller (°)	N-UU-2	-10.01	-8.84	-11	-21.05	-20.79	-16.77	-6.4	-13.55
	N-UU-3	-6.81	-13.36	-11.66	-12.73	-13.28	-11.63	-5.75	-10.75
	N-UU-4	-1.99	-10.48	-14.96	-13.86	-16.02	-15.69	-3.66	-10.95
	s^2UU	-2.51	-13.02	-11.90	-11.72	-15.51	-18.80	-1.41	-10.70
	N-UU-1	3.71	3.11	-1.54	-0.83	7.1	2.37	0.97	2.13
Opening (°)	N-UU-2	-0.43	-0.17	1.51	2.91	9.19	2.71	3.14	2.69
	N-UU-3	1.98	2.8	1.76	3.02	-4.62	3.7	-0.38	1.18
	N-UU-4	-4.66	-2.3	2.78	3.7	-11.61	2.22	2.64	-1.03
	s^2UU	0.60	4.92	2.28	-2.44	-4.86	1.58	-1.30	0.11

Table S2. Structural parameters of base pair steps in s^2 U-U 7mer (s^2 UU): [5'-uagc(s^2 U)cc-3'/3'-aucgUgg-5'] and Native-UU 7mer: [5'-uagcc-3'/3'-aucgugg-5'] including all the four duplexes in one asymmetric unit (N-UU-1, N-UU-2, N-UU-3 and N-UU-4).

		Base-Pair Steps						
		UA/UA	AG/CU	GC/GC	CU/UG	UC/GU	CC/GG	Ave.
X-disp (Å)	N-UU-1	-2.78	-4.3	-3.42	-6.86	-1.85	-4.44	-3.94
	N-UU-2	-3.67	-6.59	-3.35	-2.78	-4.4	-4	-4.13
	N-UU-3	-3.49	-3.36	-3.98	-3.88	-4.23	-4.17	-3.85
	N-UU-4	-3.74	-4.9	-3.56	-7.24	-2.67	-3.08	-4.20
	s^2 UU	-4.20	-4.45	-3.39	-3.13	-4.24	-4.38	-3.97
Inclination (°)	N-UU-1	13.16	11.47	8.35	47.25	9.49	16.55	17.71
	N-UU-2	18.49	12.23	7.65	16.74	14.44	9.5	13.18
	N-UU-3	13.59	13.76	10.3	20.78	9.74	8.71	12.81
	N-UU-4	12.18	13.6	15.11	31.96	7.11	10.68	15.11
	s^2 UU	19.27	17.69	12.79	15.25	6.61	14.24	14.31
Shift (Å)	N-UU-1	0.22	-0.55	-0.1	-0.36	-0.3	0.1	-0.17
	N-UU-2	0.1	0.42	-0.96	0.19	-0.17	0.18	-0.04
	N-UU-3	-0.01	-0.54	0.56	-0.38	0.27	0.39	0.05
	N-UU-4	0.35	-0.77	0.26	-0.55	0.47	0.18	-0.01
	s^2 UU	-0.01	-0.79	-0.37	0.3	0.68	-0.67	-0.14
Slide (Å)	N-UU-1	-0.99	-1.72	-1.55	-1.69	-0.94	-1.66	-1.43
	N-UU-2	-1.08	-2.42	-1.57	-1.1	-1.29	-1.85	-1.55
	N-UU-3	-1.33	-1.3	-1.6	-1.4	-1.71	-1.82	-1.53
	N-UU-4	-1.63	-1.86	-1.45	-1.93	-1.52	-1.36	-1.63
	s^2 UU	-1.43	-1.61	-1.35	-1.53	-1.66	-1.62	-1.53
Rise (Å)	N-UU-1	3.5	3.26	3.23	2.8	3.37	3.58	3.29
	N-UU-2	3.35	3.18	2.93	3.58	2.91	3.41	3.23
	N-UU-3	3.26	3.23	3.16	3.18	3.12	3.1	3.18
	N-UU-4	3.5	3.23	3.05	3.17	3.46	3.17	3.26
	s^2 UU	2.94	3.17	3.25	3.67	2.71	3.46	3.20
Tilt (°)	N-UU-1	-2.59	-0.57	0.65	-10.24	4.81	-1.43	-1.56
	N-UU-2	0.2	-3.21	-1.78	-2.64	9.32	-0.12	0.30
	N-UU-3	0.41	-0.33	1.67	-3.31	2.65	5.91	1.17
	N-UU-4	-2.97	-1.94	-0.26	0.03	-2.01	-0.46	-1.27
	s^2 UU	-1.89	-2.56	-0.55	2.40	7.73	-2.29	0.47
Roll (°)	N-UU-1	8.1	6.12	4.83	19.04	7.34	9.47	9.15
	N-UU-2	10.21	5.54	4.41	12.05	6.16	5.59	7.33
	N-UU-3	7.85	8.13	5.47	12.57	5.04	4.68	7.29
	N-UU-4	7.42	7.02	9.08	13.83	5.07	6.54	8.16
	s^2 UU	10.34	9.68	7.53	11.50	3.00	7.74	8.30
Twist (°)	N-UU-1	35.2	30.53	33.38	16.15	45.02	32.35	32.11
	N-UU-2	30.98	25.68	33.24	41.04	22.95	33.92	31.30
	N-UU-3	32.99	33.72	30.45	33.65	29.64	30.56	31.84
	N-UU-4	34.9	29.32	34.2	22.42	41.53	35.28	32.94
	s^2 UU	29.95	30.72	33.68	43.28	25.42	30.86	32.32

Table S3: Overlap area summary of base pair steps in s²U-U 7mer (s²UU): [5'-uagc(s²U)cc-3'/3'-aucgUgg-5'] and Native-UU 7mer: [5'-uagccu-3'/3'-aucgugg-5'] including all the four duplexes in one asymmetric unit (N-UU-1, N-UU-2, N-UU-3 and N-UU-4).

	Steps	Total Overlap Area*				
		s ² UU	N-UU-1	N-UU-2	N-UU-3	N-UU-4
1	UA/UA	2.19	1.85	2.27	2.41	1.98
2	AG/CU	1.55	1.50	3.10	2.48	1.26
3	GC/GC	13.04	13.74	11.57	12.71	12.05
4	CU/AG	2.04	2.26	3.12	1.88	0.73
5	UC/GU	4.85	10.53	6.32	5.55	7.69
6	CC/GG	4.62	3.89	3.86	3.40	5.01
Overall		28.29	33.77	30.24	28.43	28.72

*The total overlap area (in Å²) includes both intra-strand and inter-strand overlap within the four bases of each base pair step.

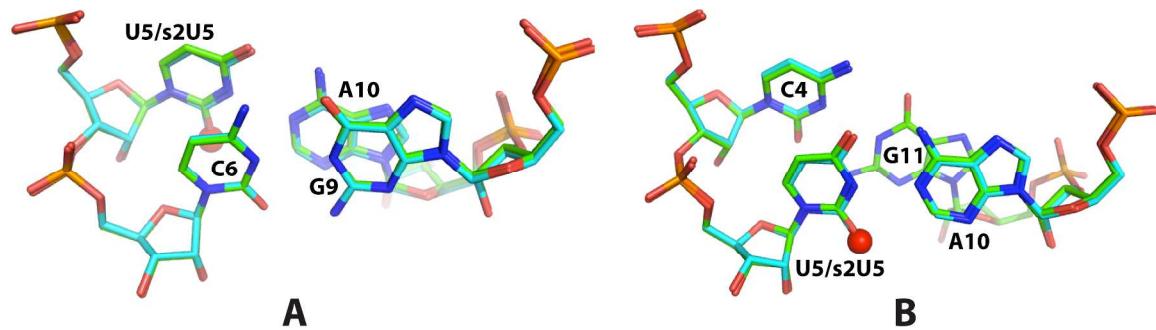


Figure S2. Comparison of base-pair steps in the native and s²U:A-containing duplex. (A) Superimposed views of the U5-C6/G9-A10 and s²U5-C6/G9-A10 steps; (B) Superimposed views of the C4-U5/A10-G11 and C4-s²U5/A10-G11 steps.

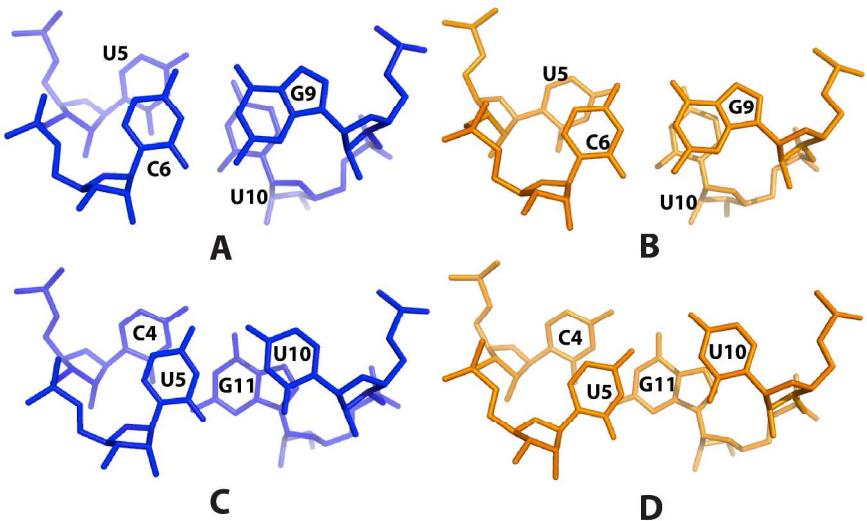


Figure S3. Stacking interactions of the two base-pair steps in the native UU-3 and UU-4 duplexes. (A) U5-C6/G9-A10 step in the native duplex UU-3. (B) U5-C6/G9-A10 step in the native duplex UU-4. (C) C4-U5/A10-G11 step in the native duplex UU-3. (D) C4-U5/A10-G11 step in the native duplex UU-4. The color code is same as in Fig. 3 and 4.

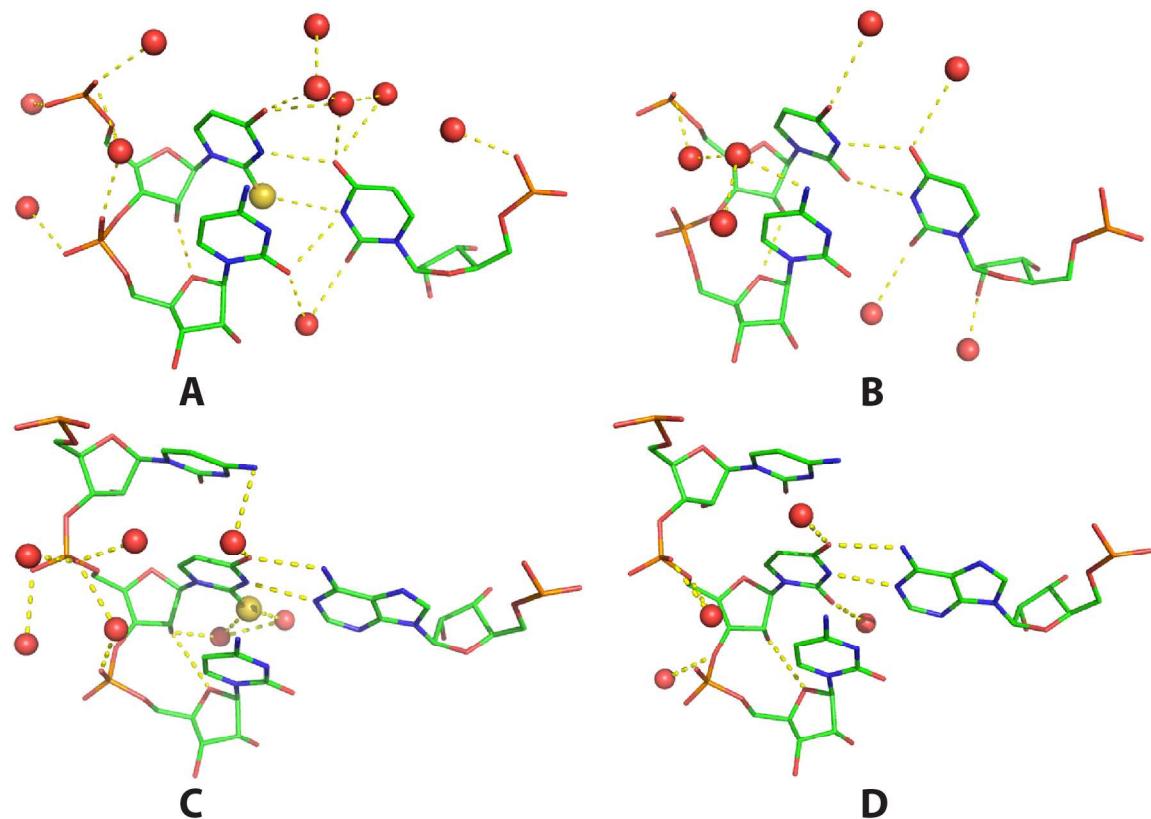


Figure S4. The hydration patterns discovered in (A) s^2U -U pair; (B) native U-U pair (in UU-1); (C) s^2U -A pair; (D) native U-A pair. (the yellow spheres represent the sulfur atoms)

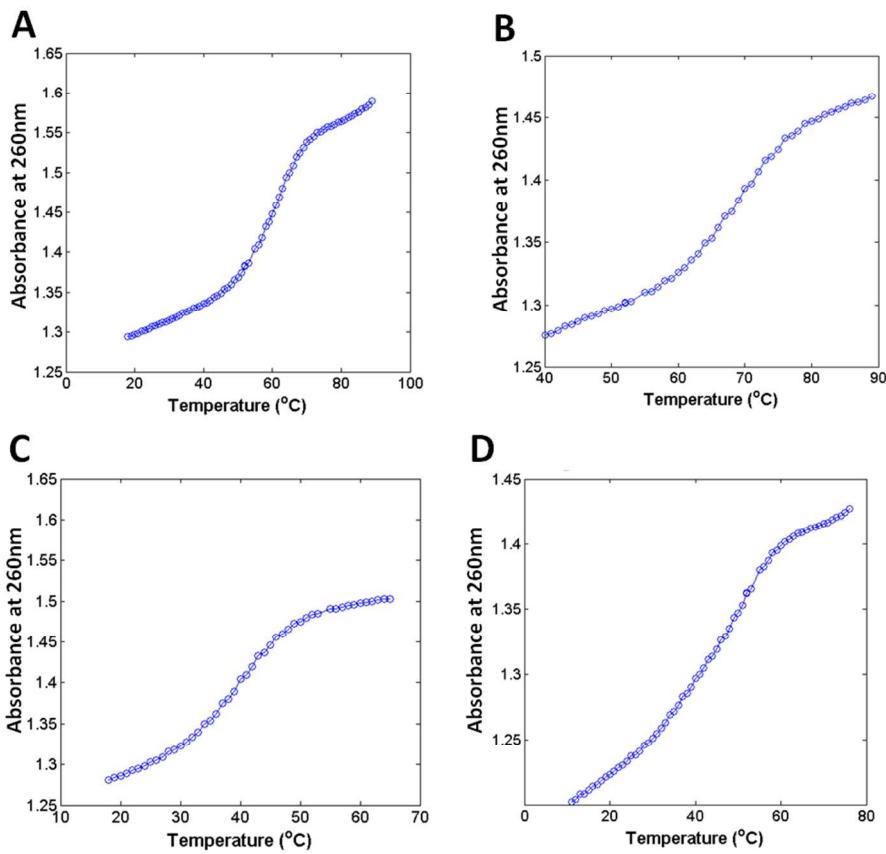


Figure S5. Thermal denaturation curves of (A) U-A 7mer; (B) s^2U -A 7mer; (C) U-U 7mer; and (D) s^2U -U 7mer. Samples were prepared by dilution of stock solutions into buffers containing 100 mM MgCl₂ and 200 mM HEPES, pH 7.5. Total strand concentration was 200 μ M (100 μ M duplex). Absorbance was monitored at 260 nm while the temperature was ramped at 1°C min⁻¹ between 4 and 89 °C. All melt data was collected in duplicate, from the first temperature ramp-up of independent samples.