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Biochemistry, 1996, 35(11), 3534-3544, DOI: [10.1021/bi952571n](https://doi.org/10.1021/bi952571n)

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Sequential Assignments.

The sequential assignment procedure started with the analysis of the characteristic thymine 5CH₃ and cytosine 5H resonances. Of particular value were the sequential $d_s(6,8;5H,5CH_3)$ NOEs, providing 22 (out of 30 possible) well-resolved connectivities. In addition, all $d_s(5H,5CH_3;5H,5CH_3)$ NOEs for cytidines next to a thymidine were readily observed (T₆/C₇, C₂₁/T₂₂, T₂₂/C₂₃, C₄₂/T₄₃, C₅₆/T₅₇), as were the sole $d_s(5H;5H)$ NOE (C₅₀/C₅₁) and $d_s(5CH_3;5CH_3)$ NOEs (T₃₆/T₃₇, T₆₀/T₆₁).

Careful processing of the JR-NOESY spectrum acquired at 305K, optimizing for resolution, allowed for identification of numerous of the standard B-type DNA sequential NOEs $d_s(1'H,2'H,2''H;6H,8H)$ (Figure S1 and S2). At least one such NOE was resolved for 88% of the residues (13 in strand 1, 13 in strand 2, 13 in strand 3, 14 in strand 4). Another important sequential NOE was the $d_s(6H,8H;6H,8H)$, which provided the critical connectivity for certain residues.

In the following, a rather detailed description of the sequential resonance assignment of strand 1 is given, illustrating the manner in which critical NOE connectivities are pieced together.

A good starting point for the assignment of strand 1 is the unique d-(AATC) tract. Only two pairs of adjacent adenine residues, A₄/A₅ and A₂₈/A₂₉, are found in J2P1. Both pairs were linked by readily identified $d_s(1'H;8H)$ NOEs. $d_s(8H;5CH_3)$ and $d_s(1'H,2'H,2''H;6H)$ NOEs extended the connectivity for both, to T₆ and T₃₀, respectively. Observation of a $d_s(6H;5H)$ NOE to C₇ positively identified which of the two d-(AAT) segments corresponded to strand 1 (A₂₈A₂₉T₃₀ is followed by G₃₁ in strand 3). Further evidence for the sequential connectivity to C₇ was obtained from the $d_s(1'H,2'H,2''H;6H)$ NOEs observed in the 2D plane at $\omega_1 = \delta(C_7\ 5H)$ of the 3D TOCSY/NOESY experiment. Connectivity to A₈ could be established despite the problem with the assignment of the 2''H

resonance of C₇, through $d_s(6H;8H)$ and $d_s(2'H,3'H;8H)$ NOEs together with a $d_s(1'H;8H)$ NOE identified in the NOESY spectrum recorded in D₂O. The $d_s(1'H;8H)$ NOE could not be unambiguously identified in the JR-NOESY spectra due to overlap with one of the $d_i(5H;4NH_2)$ NOEs for C₇.

T₉ was difficult to assign because the 6H resonance of T₉ is in a very crowded region, broader lineshapes were observed for the resonances of this residue, and there is distorted conformation at the junction point for this crossover strand (see main text). T₉ could only be connected to A₈ by weaker $d_s(1'H,2'H,2''H,3'H;5CH_3)$ NOEs and a very weak $d_s(8H;5CH_3)$ NOE to the 5CH₃ protons. There is a gap between T₉ and G₁₀ because sequential NOEs between these two residues were ambiguous due to chemical shift degeneracy of the 8H resonance of G₁₀ with several other guanosine 8H resonances. A₁₁ showed well resolved resonances and extension backwards to G₁₀ was possible from identification of $d_s(1'H,2'H,2''H,3'H;8H)$ and $d_s(8H,8H)$ NOEs. Connectivities on to G₁₂, C₁₃, A₁₄ and C₁₅ were obtained by identification of one or several of $d_s(1'H,2'H,2''H;6H,8H)$ NOEs and, where possible, $d_s(8H;5H)$ NOEs. These connectivities uniquely identified this tract as belonging to strand 1. Extension to G₁₆ was obtained through resolved $d_s(1'H,2'H,2''H;8H)$, even though the G₁₆ 8H resonance has the same chemical shift as five other guanosine 8H resonances. The uniquely downfield shift of the 1'H of a guanosine residue at the 3'-end of the DNA strand give further evidence for the 8H chemical shift of G₁₆.

Returning back to the 5'-end, C₁ is readily identified through its uniquely down-field shifted 6H resonance. Sequential connectivity to G₂ was established through $d_s(1'H,2'H,2''H;8H)$ NOEs, and on to C₃ through $d_s(1'H,2'H,2''H;6H)$ and $d_s(8H;5H)$ NOEs, which uniquely identifies this stretch to strand 1, and not strand 2. Observation of $d_s(1'H,2'H,2''H;8H)$ to A₄, finally completes the sequential assignment of strand 1.

In summary, the segments on either side of T₉/G₁₀ were sequentially assigned on the basis of at least one unambiguous NOE, together with observation of several other sequential

and intra-residue cross peaks. No evidence for connectivity between the two fragments was obtained, but the sequential assignments are still unambiguous as the sequences uniquely fit to strand 1.

The sequential assignment of the labile protons relies primarily on observation of connectivities between adjacent imino protons, together with NOEs corresponding to short inter-base-pair distances between the imino proton and the neighboring adenine 2H or cytosine 4NH₂ protons. More indirect pathways exist, involving various NOEs to the adenine 2H, cytosine 5H and thymine 5CH₃. Figure S3 shows a summary of all of the sequential and intra-base-pair connectivities identified for J2P1 in the various JR-NOESY spectra recorded at different temperatures.

K-3544-m4

Figure S1A

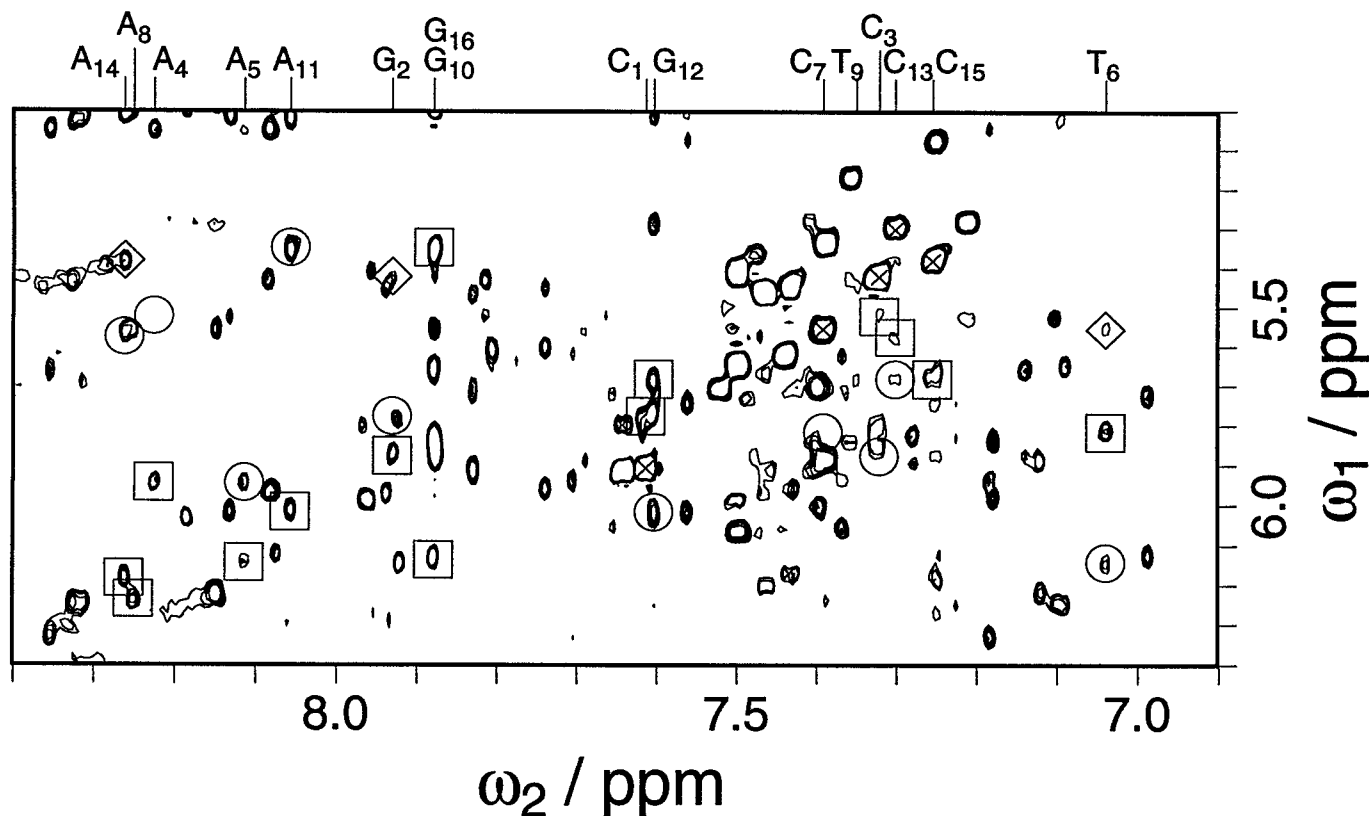
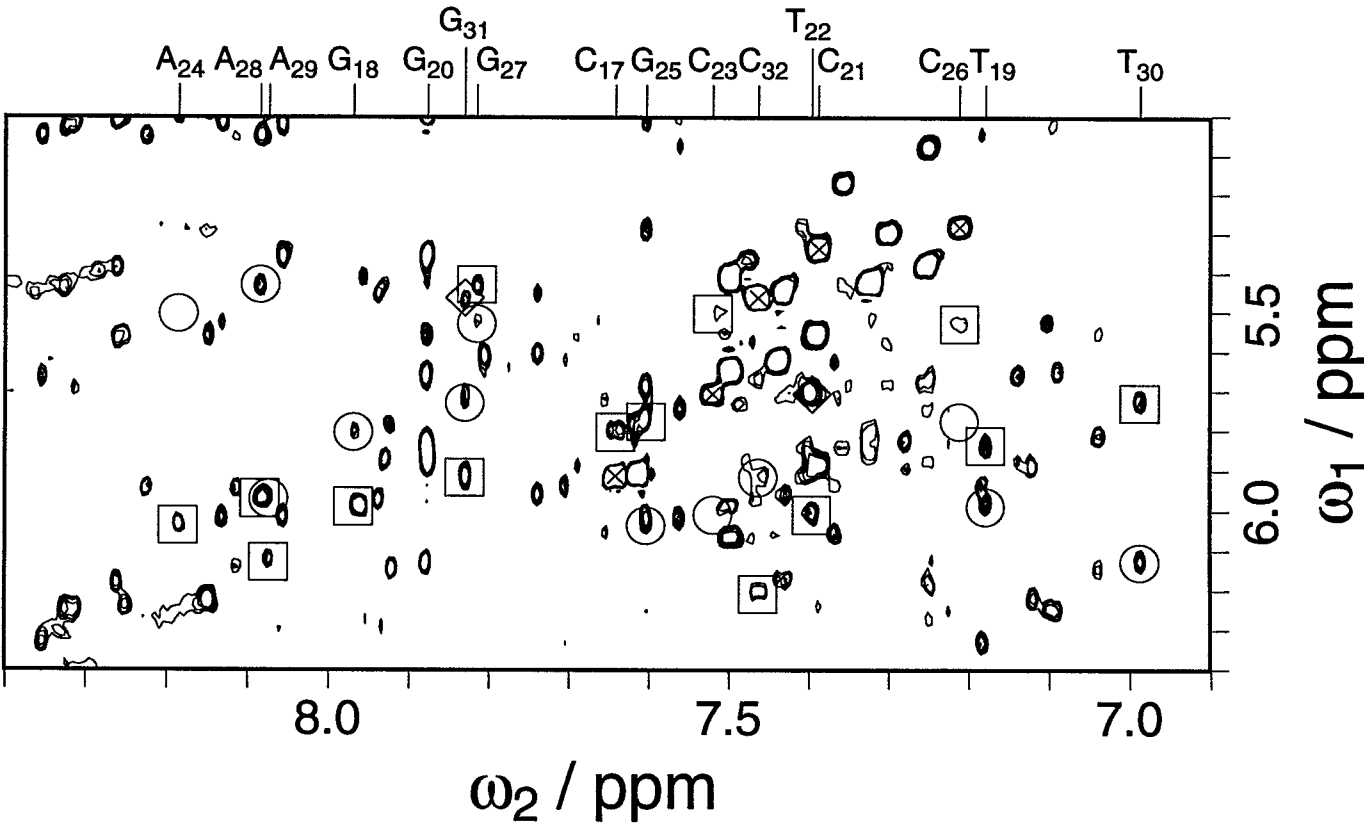


Figure S1. Region of the 600 MHz JR-NOESY spectrum of ~1 mM J2P1 containing cross peaks from 1H and 5H resonances to base proton resonances. Separate panels are shown for each of the four strands: A) 1-16; B) 17-32; C) 33-48; D) 49-64. The spectrum was acquired at 305K with a mixing time of 200 ms from a H₂O solution. The base proton frequencies are indicated at the top of each panel with unambiguously identified intraresidue and sequential NOEs identified by boxes and circles, respectively. The strong intraresidue cytosine $d_i(5H;6H)$ NOEs are marked with an X within the crosspeak, and the unambiguously identified sequential $d_s(6H,8H;5H)$ NOEs are identified by diamonds.

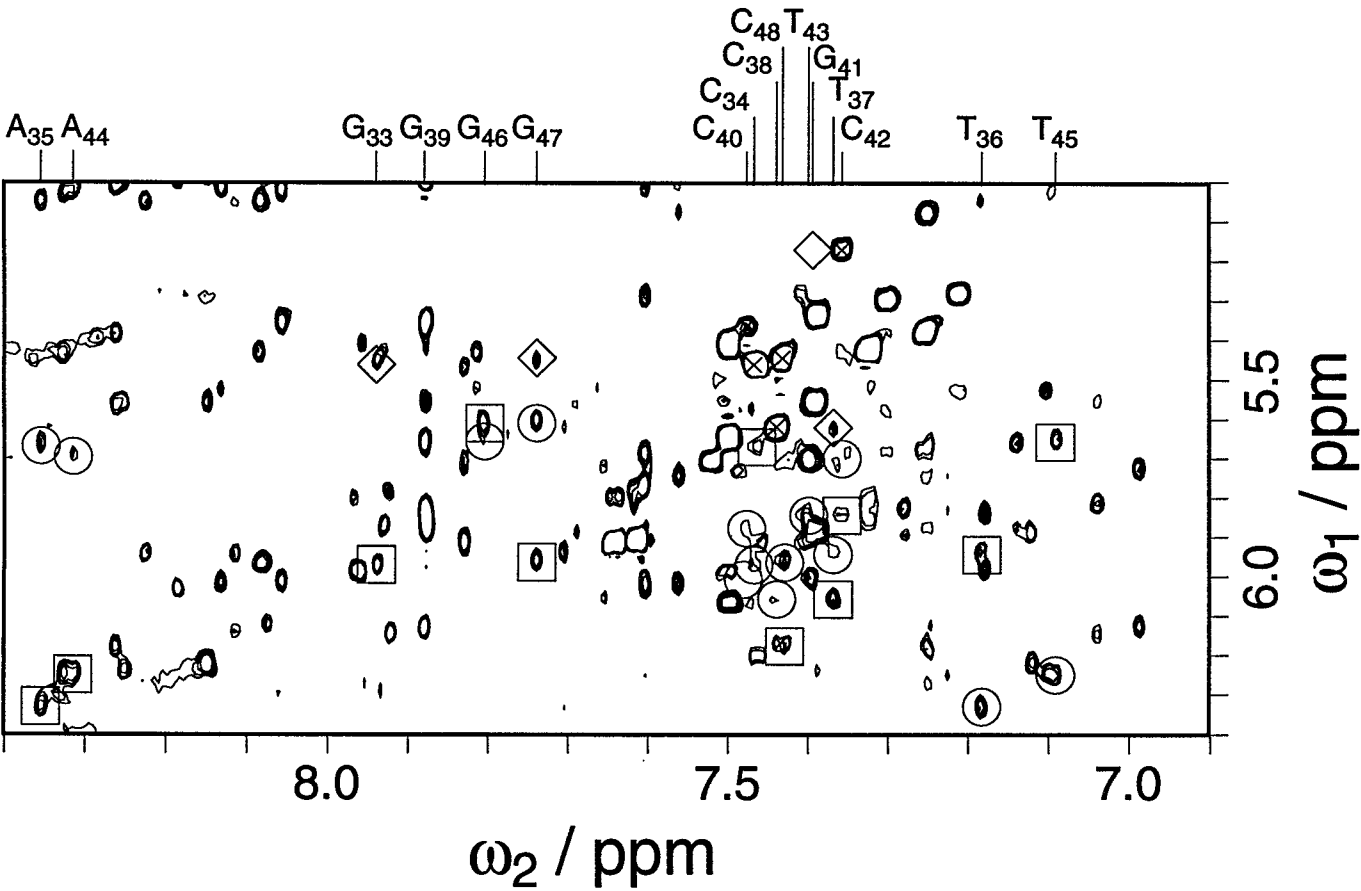
K-3544-MS

Figure S1B



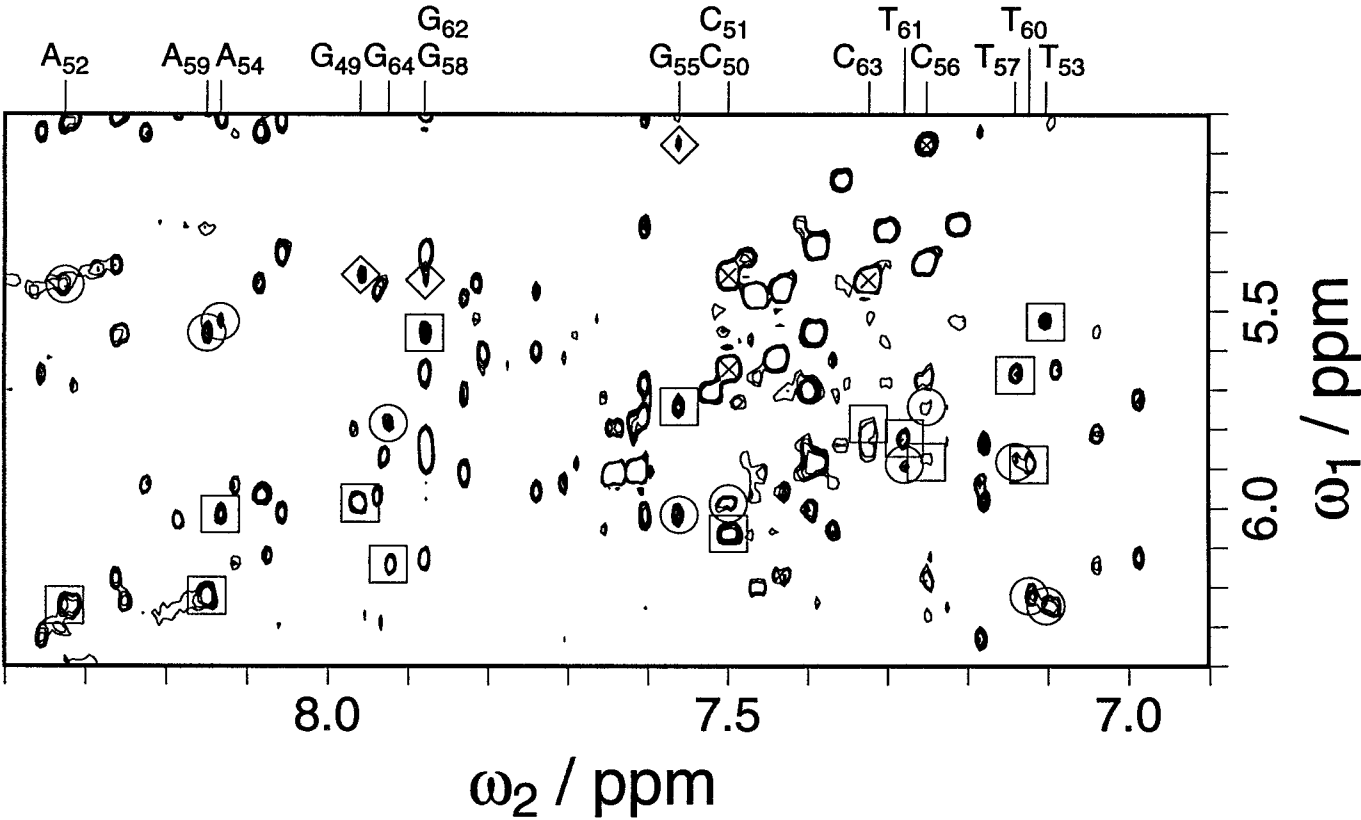
K-3S44.m6

Figure S1C



K-3S44-M7

Figure S1D



K-3544-m8

Figure S2A

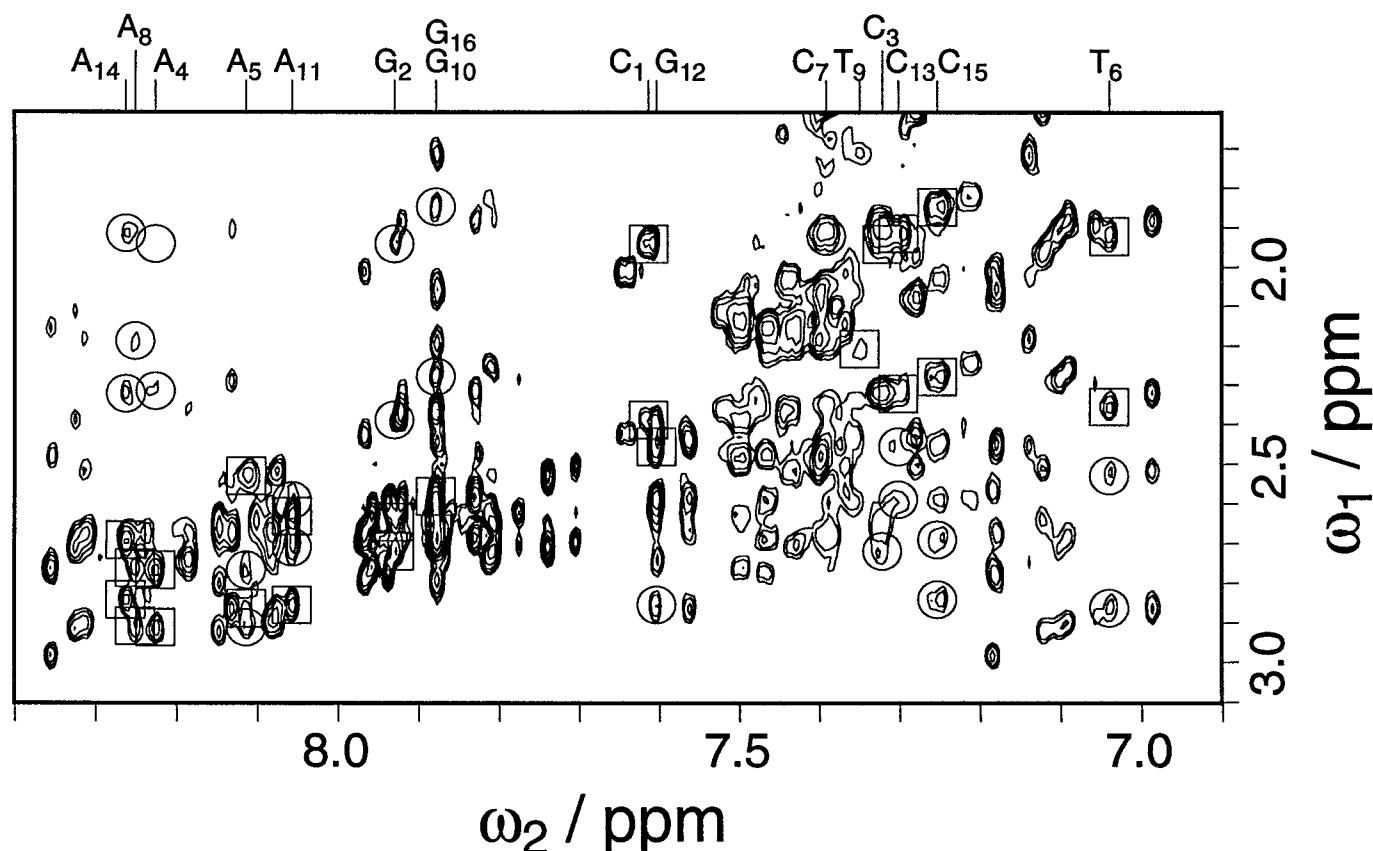
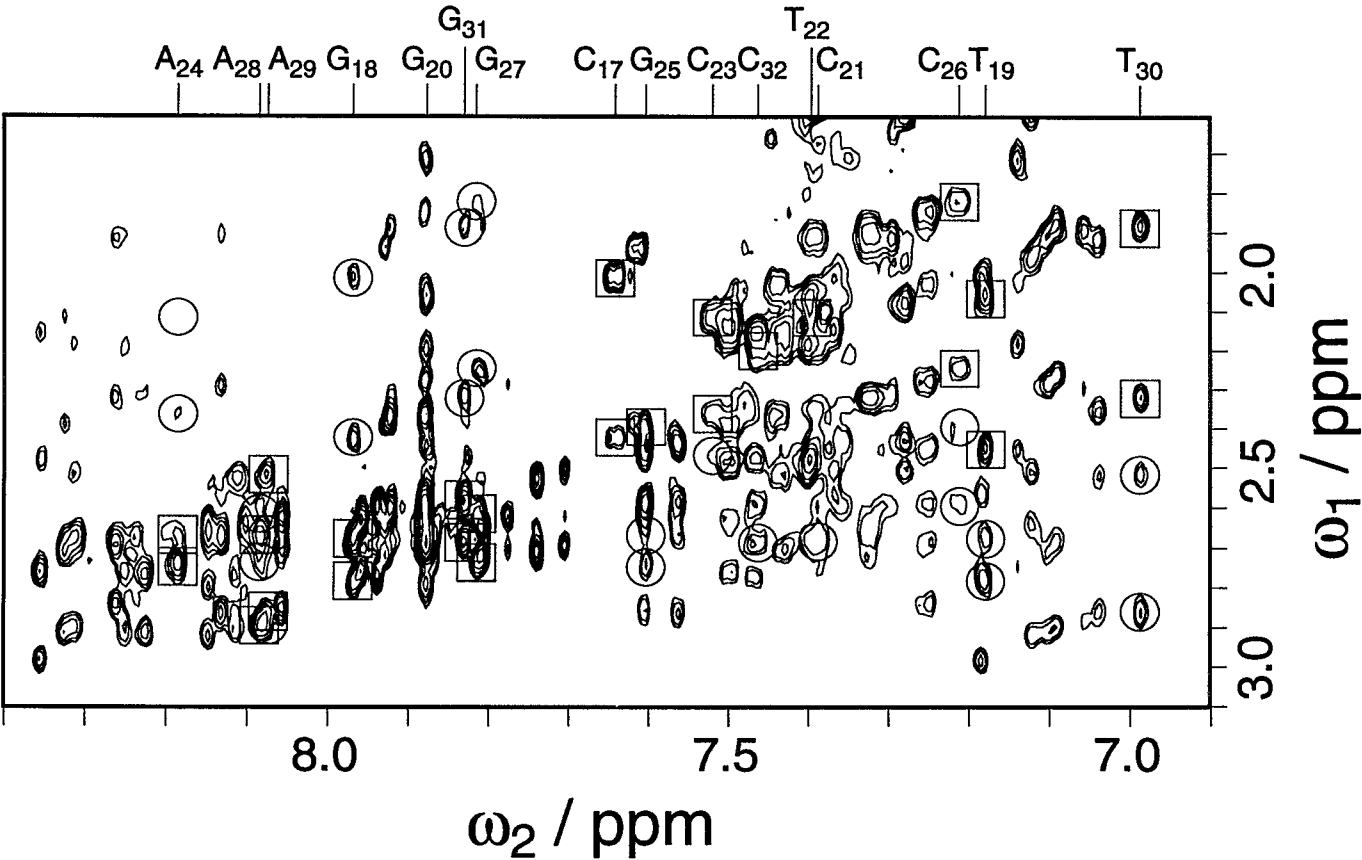


Figure S2. Region of the 600 MHz JR-NOESY spectrum of ~1 mM J2P1 containing cross peaks from 2'H and 2''H resonances to base proton resonances. Separate panels are shown for each of the four strands: A) 1-16; B) 17-32; C) 33-48; D) 49-64. The spectrum was acquired at 305 K with a mixing time of 200 ms from a H₂O solution. The base proton frequencies are indicated at the top of each panel with unambiguously identified intraresidue and sequential NOEs identified by boxes and circles, respectively.

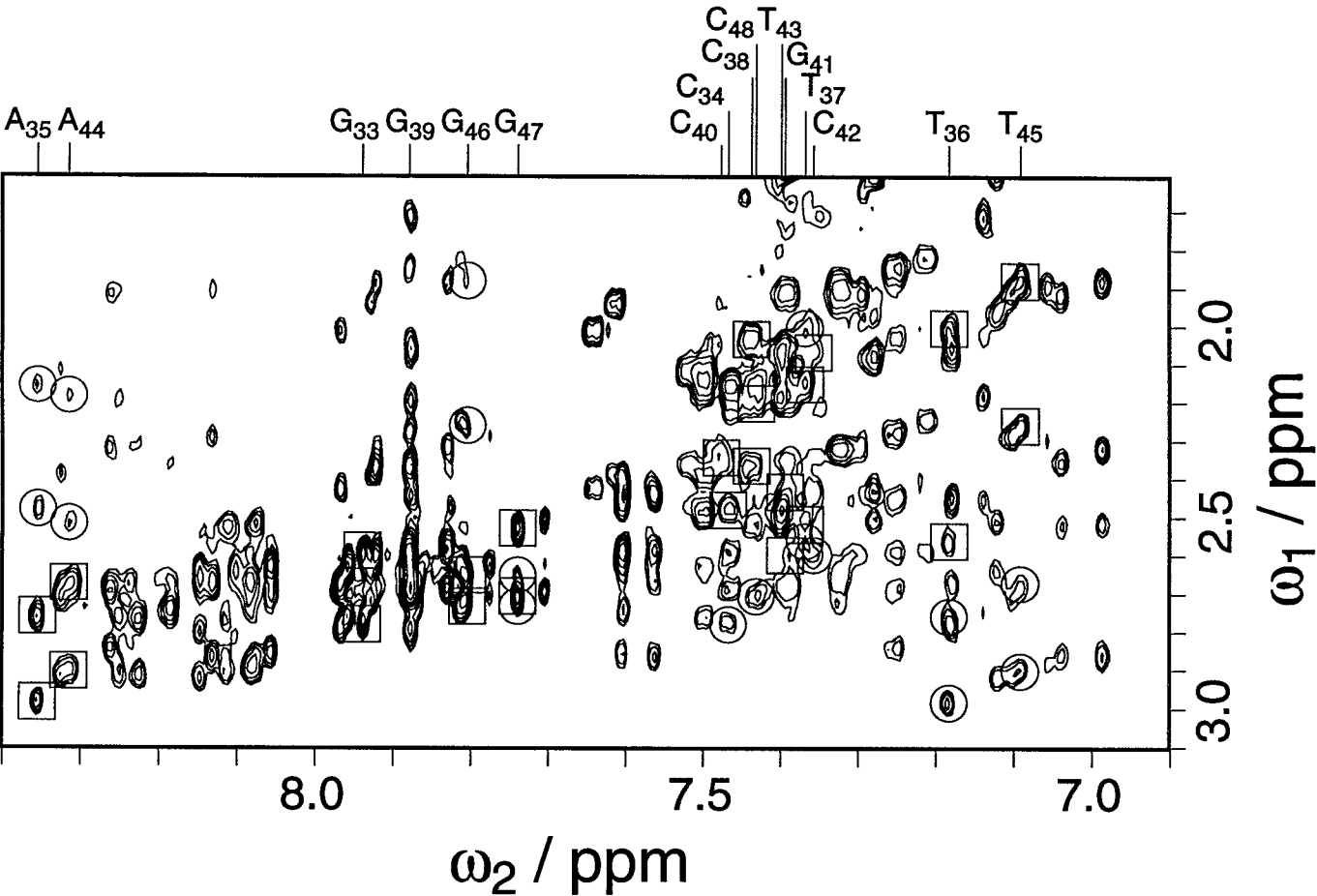
K-3544-m9

Figure S2B



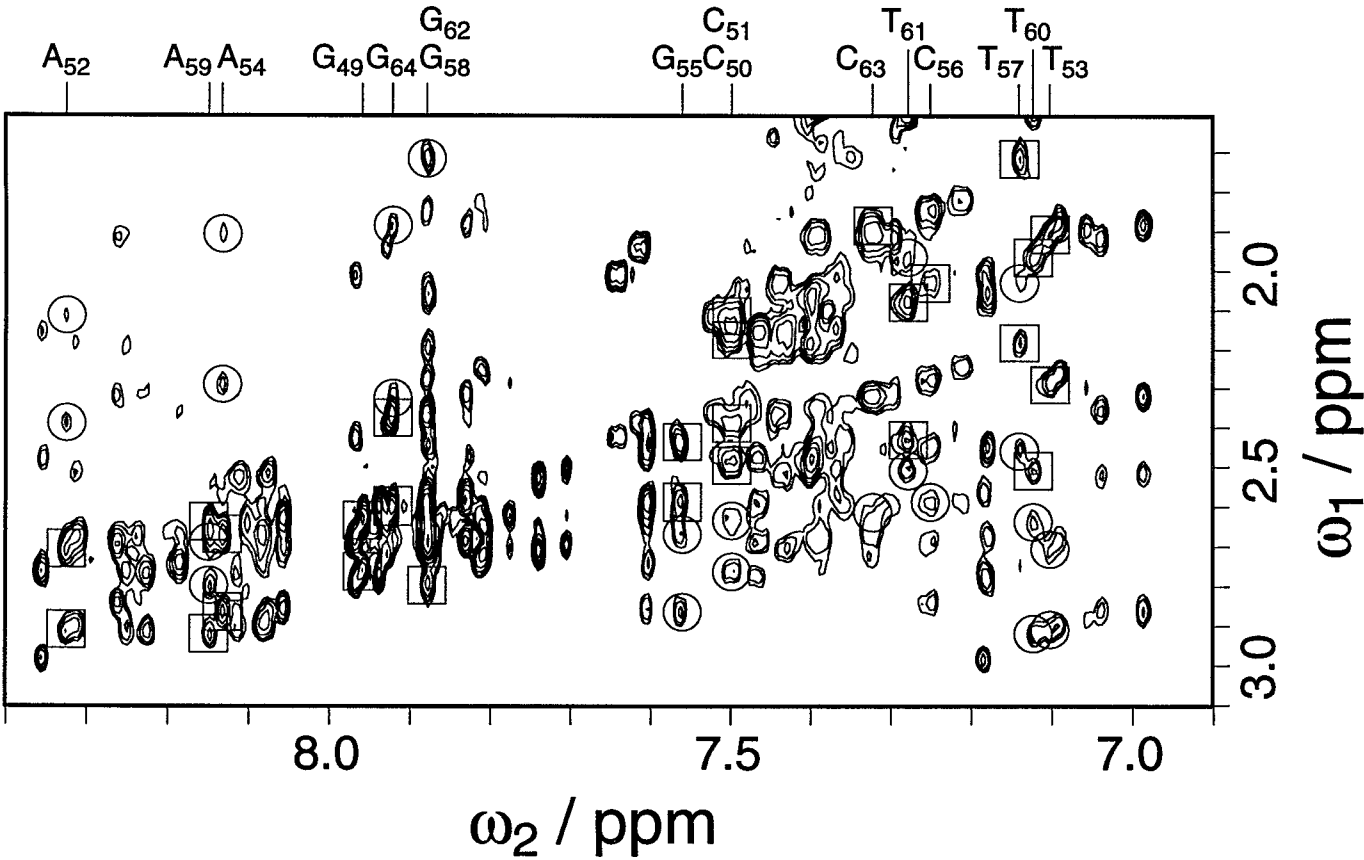
K-3544-m10

Figure S2C



K-3544-m11

Figure S2D



R-3544-m12

Figure S3

Stacking Domain	Inter-Base Pair					Intra-Base Pair			
	(NH:NH)	(NH:2)	(NH:4NH ₂)	(2:2)	(NH:5CH ₃)	(3NH:2)	(1NH:4NH ₂)	(1NH:5H)	(3NH:5CH ₃)
I + IV									
C ₁ •G ₆₄	-		-			-		-	
G ₂ •C ₆₃	•					X		X	
C ₃ •G ₆₂	•	•	•			X		X	
A ₄ •T ₆₁	•	•	•	•	-	X			X
A ₅ •T ₆₀	•	•		d	•	X			X
T ₆ •A ₅₉	•	•			-	X			Xw
C ₇ •G ₅₈	-	•	-		-		X	X	
A ₈ •T ₅₇	-	•	-		-	-			-
G ₄₁ •C ₅₆	-	-	-		-	-		Xvw	
C ₄₂ •G ₅₅	-	•	-		-	X		X	
T ₄₃ •A ₅₄	-	•		•	•	X			X
A ₄₄ •T ₅₃	-	•		•	•vw	X			X
T ₄₅ •A ₅₂	•	•		•	•	X			X
G ₄₆ •C ₅₁	•	•	-		-		X	X	
G ₄₇ •C ₅₀	•		-				X	X	
C ₄₈ •G ₄₉	-		-			-		-	
II + III									
C ₁₇ •G ₁₆	-		-			-		-	
G ₁₈ •C ₁₅	-	•	•			X		X	
T ₁₉ •A ₁₄	-	•	•		-	X			X
G ₂₀ •C ₁₃	d	•	-		•		X	X	
C ₂₁ •G ₁₂	•	•	-		•		X	X	
T ₂₂ •A ₁₁	•	•	-		•	X			X
C ₂₃ •G ₁₀	-	•	-		•		X	X	
A ₂₄ •T ₉	-	•	-		-	-			-
G ₂₅ •C ₄₀	-		-			-		-	
C ₂₆ •G ₃₉	•		-			X		X	
G ₂₇ •C ₃₈	•	•	-		•	X		X	
A ₂₈ •T ₃₇	•	•		•	•	X			X
A ₂₉ •T ₃₆	•	•		d	•	X			X
T ₃₀ •A ₃₅	•	•	•		•	X			X
G ₃₁ •C ₃₄	-	•	•		-		X	X	
C ₃₂ •G ₃₃	-		-			-		-	

Figure S3. Summary of NOE cross peaks involving the labile protons of J2P1 observed in 2D JR-NOESY spectra obtained at 285, 293, 296, 300, and 305 K. A 'd' indicates that this NOE could not be observed due to degeneracy, and a 'w' and a 'vw' indicates that the observed crosspeak was 'weak' and 'very weak', respectively.