

residue i random coil NH shift

	A	B	C	D	E	F	G	H	I	K	L	M	N	P	Q	R	S	T	V	W	Y
A	8.08 0.44 89		7.91 0.66 12				8.22 0.74 167				8.35 1.2 57		8.23 0.57 63								
B	8.5 0.7 22				8.85 0.8 8			7.6 1.73 8		8.46 0.57 23		9.09 0.62 2			9.4 1.53 9		8.66 0.44 20				
C			8.83 0.75 25								8.45 0.56 23	9.03 0.62 4	8.65 0.57 20		8.79 1.3 7		8.7 0.41 37	8.48 0.72 27	8.98 0.82 12		
D	8.07 0.51 93				8.23 0.53 86		8.47 0.86 262										8.04 0.47 79		7.77 0.63 72		
E		8.59 0.69 18					8.51 0.61 116														
F			7.72 0.14 4	8.11 0.81 58	8.11 0.66 29				8.67 0.98 13	8.62 0.78 23	7.72 0.64 16	7.47 0.45 7				7.62 1.11 25		8.5 0.83 45			
G	8.07 0.48 125	7.91 0.68 19			8.14 0.56 142	7.83 0.72 63	8.27 0.67 212		7.93 0.69 95	7.97 0.54 165		8.02 0.62 32			8 0.48 87	8.05 0.62 113		8 0.55 127	7.83 0.52 93		7.93 0.62 67
H	7.9 0.69 24	7.15 0.16 3		8.1 0.68 30		9.16 1.06 18				7.91 1.06 31					8.56 1.35 15	7.92 0.56 18					
I	8.4 0.57 37	9.01 0.54 4		8.49 0.56 62				8.65 0.44 26		8.44 0.65 66	8.5 0.55 28					8.57 0.66 22		8.36 0.7 40	8.39 0.59 31		
K	8.34 0.62 82												8.18 0.6 75								
L				8.19 0.63 80			8.18 0.73 99											8.45 0.58 90	7.93 0.78 39		
M					8.86 0.72 11																
N				8.59 0.58 45			8.52 0.93 116											8.02 0.78 56	7.89 0.51 59		7.44 0.85 6

residue i-1

residue i random coil NH shift

	A	B	C	D	E	F	G	H	I	K	L	M	N	P	Q	R	S	T	V	W	Y		
residue i-1	P	8.35 0.55 96				8.55 0.57 82					8.48 0.67 83		8.58 0.4 27			8.51 0.61 32	8.49 0.6 54				7.6 1.06 25		
	Q	8.05 0.63 42																		8.35 0.57 35			
	R							8.18 1 99			8.48 0.62 66							8.07 0.87 62					
	S	8.35 0.56 95						8.52 0.87 158			8.01 0.55 105												
	T									8.41 0.63 38				8.23 0.76 53			8.46 0.67 45			8.39 0.84 48	8.5 0.64 17		
	V	8.57 0.66 44								8.47 0.61 33	8.41 0.67 64		7.92 0.78 10			8.5 0.51 22	8.5 0.54 30	8.42 0.61 80			9.13 0.74 7	8.53 0.48 16	
	W																	8.78 0.95 18	8.67 1 11			9.3 1.34 5	
	Y											8.45 0.82 23					8.49 0.55 18						