

Supplementary material to

**The disorderly conduct of Hsc70:
its interaction with the Alzheimer's related TAU protein as monitored by NMR.**

Dedicated to the memory of Chad A. Dickey.

Isabelle R. Taylor², Atta Ahmad¹, Taia Wu², Bryce A. Nordhues³, Anup Bhullar¹, Jason E. Gestwicki^{2*} and Erik R.P. Zuiderweg^{1*},

1) Department of Biological Chemistry, The University of Michigan Medical School,
1500 Medical Center Drive, Ann Arbor, MI 48109

2) Institute for Neurodegenerative Disease, University of California at San Francisco
675 Nelson Rising Lane, San Francisco, CA 94158

3) Department of Molecular Medicine and Byrd Alzheimer's Institute, University of South Florida,
Tampa, FL 33613.

*) corresponding authors zuiderwe@umich.edu and Jason.Gestwicki@ucsf.edu.

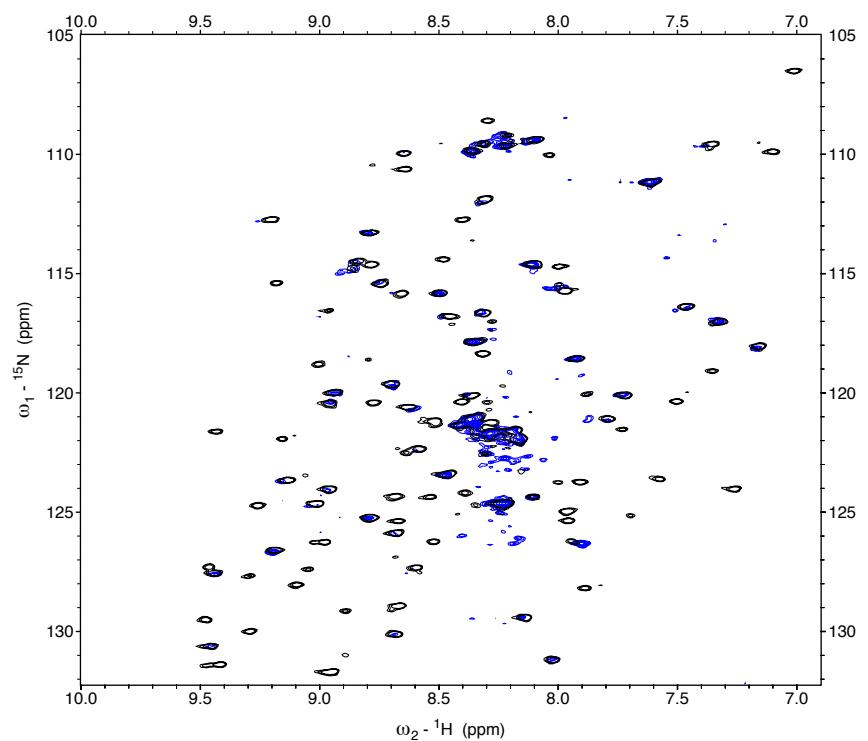
Supplemental Table 1		
	Wild type	BETA (395-508)
	EC ₅₀ (μM)	EC ₅₀ (μM)
4R0N TAU	0.9 ± 0.2	3.4 ± 2.9
TAU1	17.0 ± 9.3	32.0 ± 22
NRLLLTG	6.3 ± 3.5	28.0 ± 9.8
C-PEPTIDE	0.2 ± 1.8	1.3 ± 0.6
TAU2	> 50	> 200
Huntingtin	> 200	> 200
KFERQ	> 200	> 200

Table S1: Binding of substrates to Hsp70 (HSPA1A)

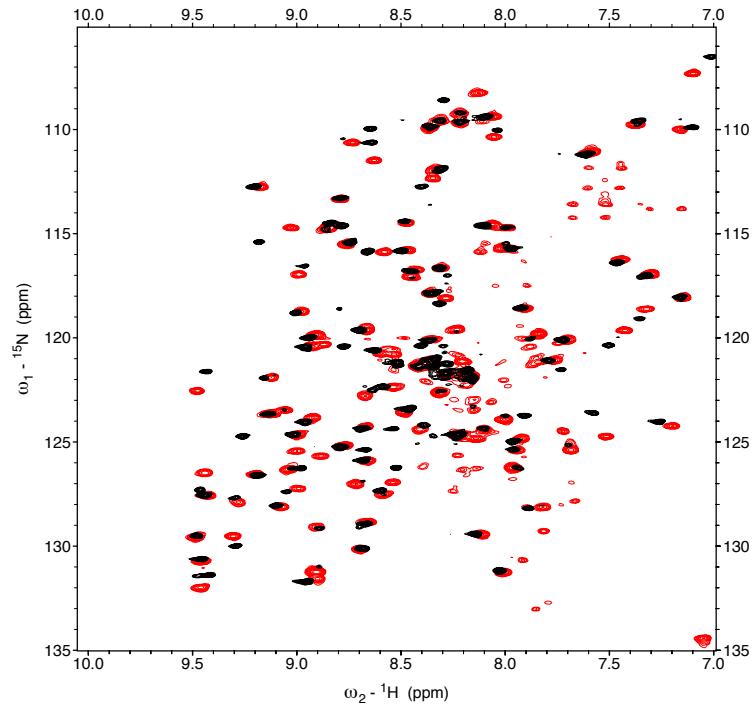
and its truncation by fluorescence polarization (FP).

TAU1: GKVQIINKKG; Insulin C-peptide (C-PEP); TAU2: VQIVYK; huntingtin: MATLEKLMKAFESLKSF.

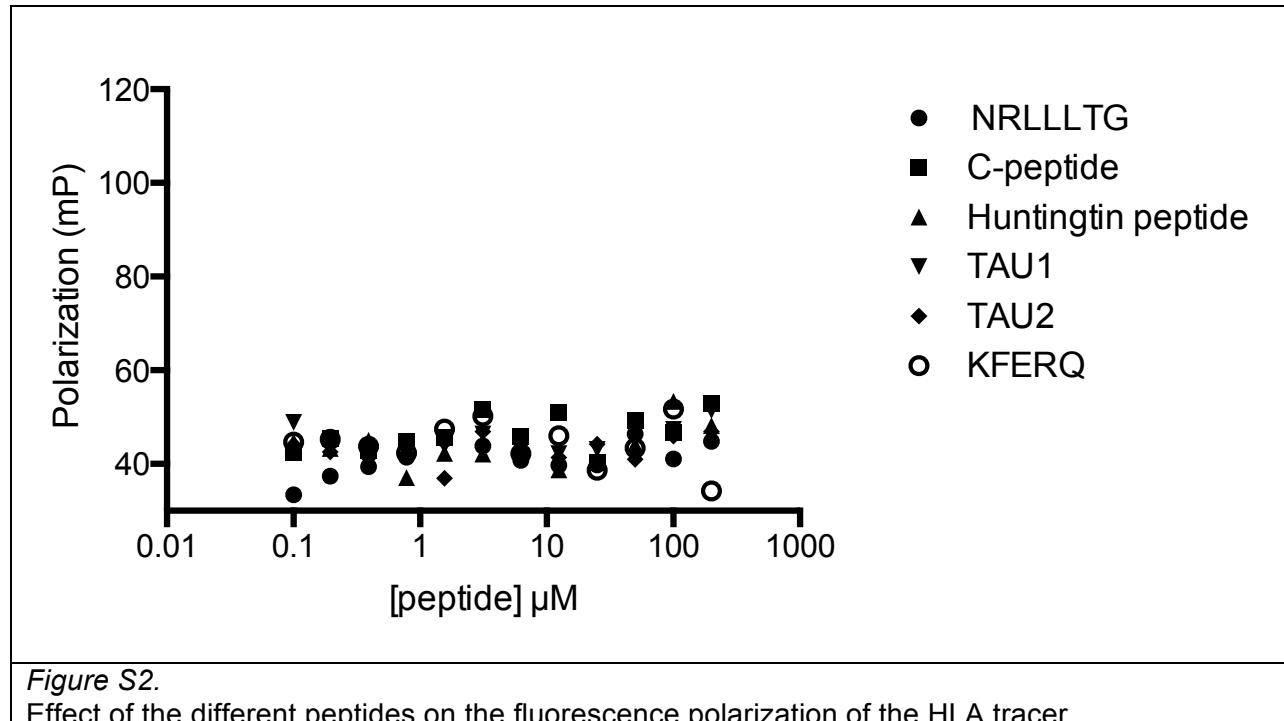
The Fam-HLA tracer was at 20 nM, wt-Hsp70 at 5 μM, Hsp70- BETA at 1 μM.



*Fig S1 A Binding of Fam-HLA
Blue 50 μM BETA apo, Black, with 60 μM Fam-HLA*



*Fig S1 B Binding of Fam-HLA compared with TAU1
Black, 50 μM BETA with 60 μM Fam-HLA
Red, 35 μM BETA + 200 μM TAU1 (GKVQIINKKG)*



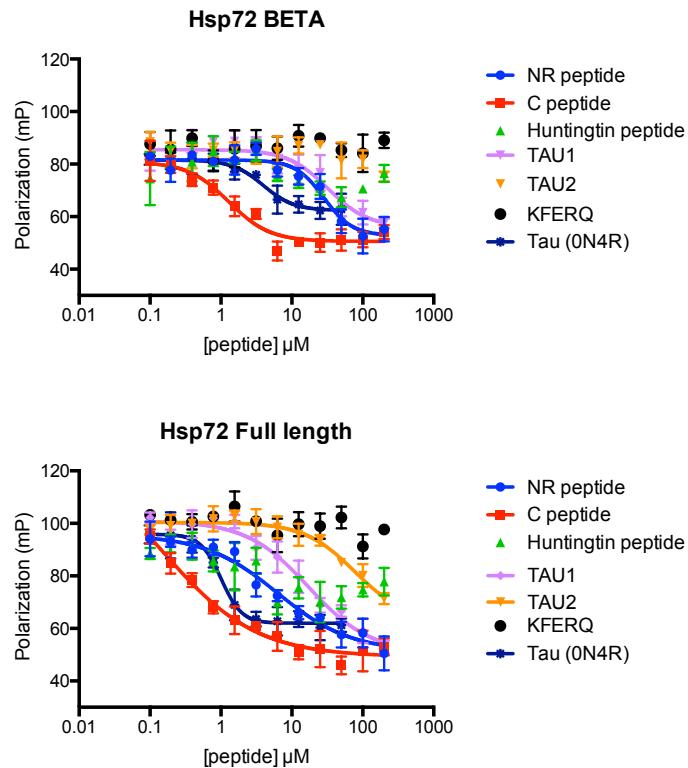


Figure S3. Interaction of peptides with inducible Hsp70 (HSPA1A) and its BETA construct.

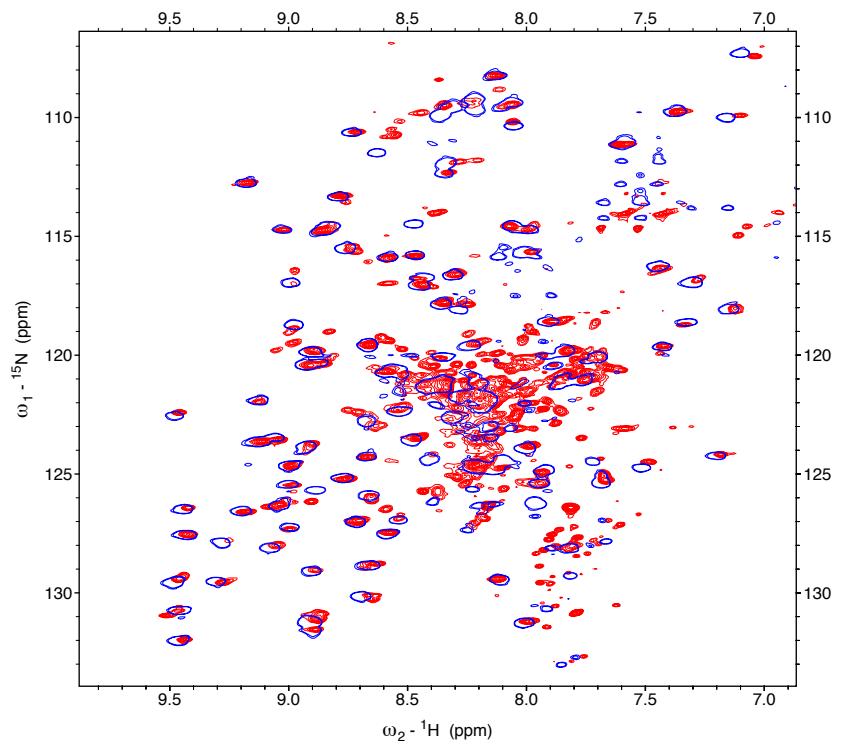


Figure S4.

$^{15}\text{N}-^1\text{H}$ TROSY spectra of BETA with unlabeled TAU1 in trans (blue; 113 residues) and the BETA-TAU1-CIS construct (red, 165 residues)

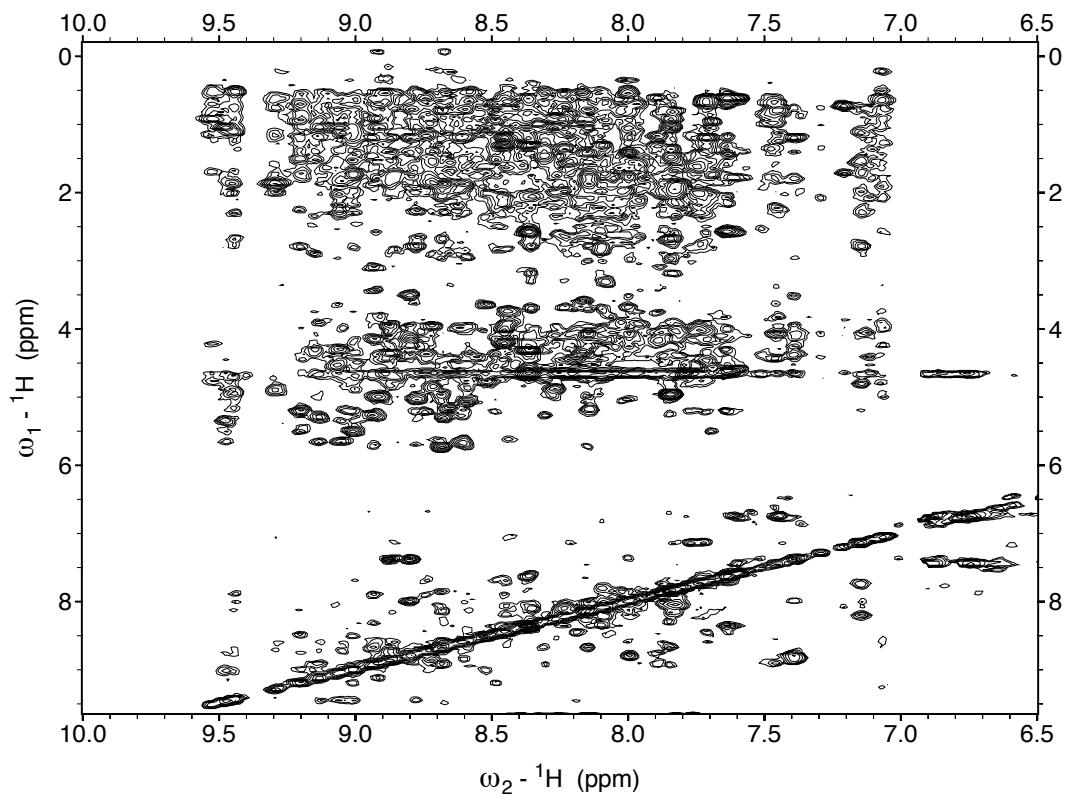


Figure S5. ${}^1\text{H}-{}^1\text{HN}$ Projection of the NUS 3D ${}^1\text{H}-\tau_M-{}^{15}\text{N}-{}^1\text{HN}$ NOESY-TROSY spectrum of BETA-TAU1-CIS. The mixing time τ_M was 200 ms.

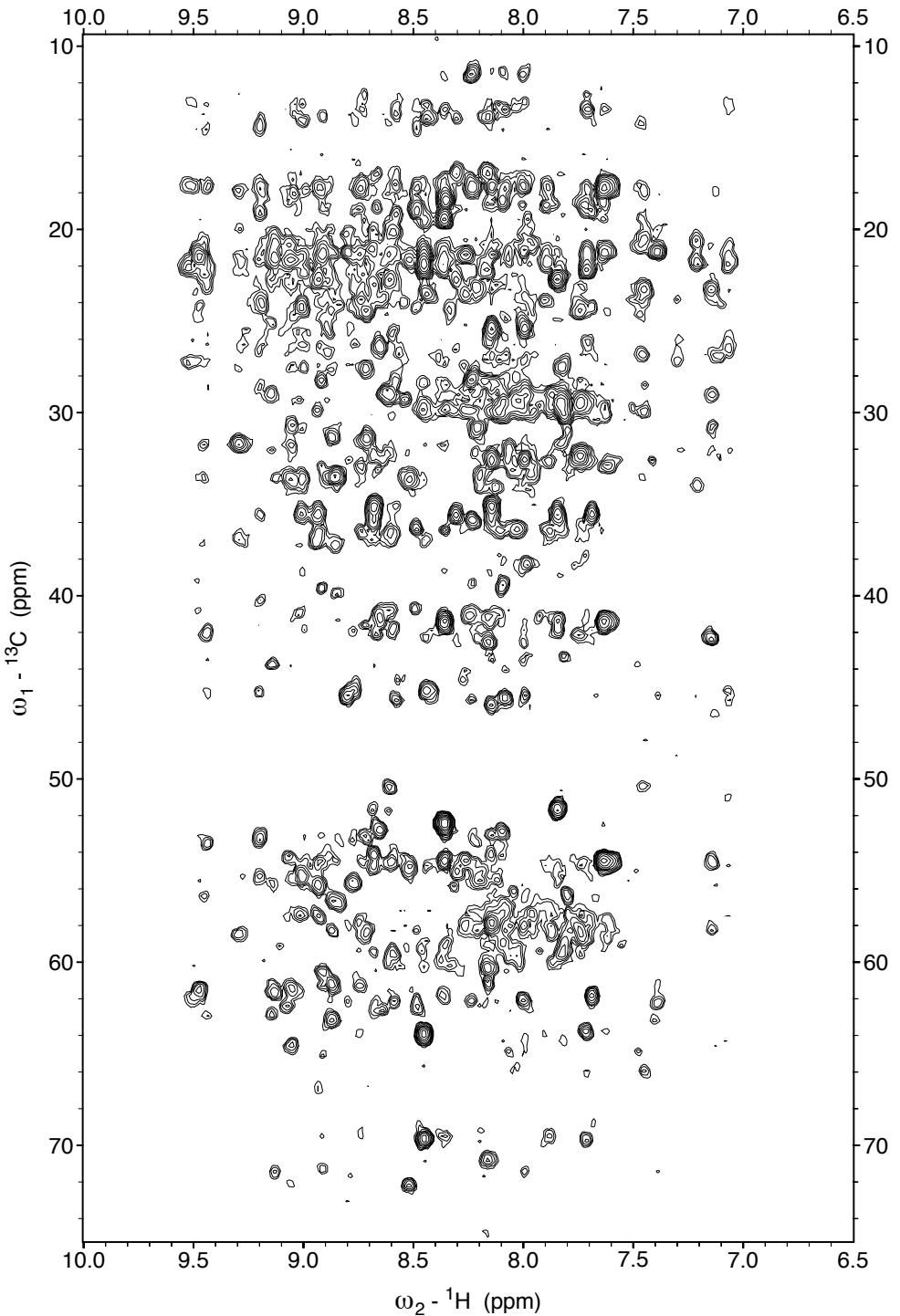


Figure S6. ^{13}C - ^1H Projection of the NUS 3D (^1H)- ^{13}C - (^1H) - τ_M - ^{15}N - ^1H HMQC-NOESY-TROSY spectrum of BETA-TAU1-CIS. The mixing time τ_M was 200 ms.

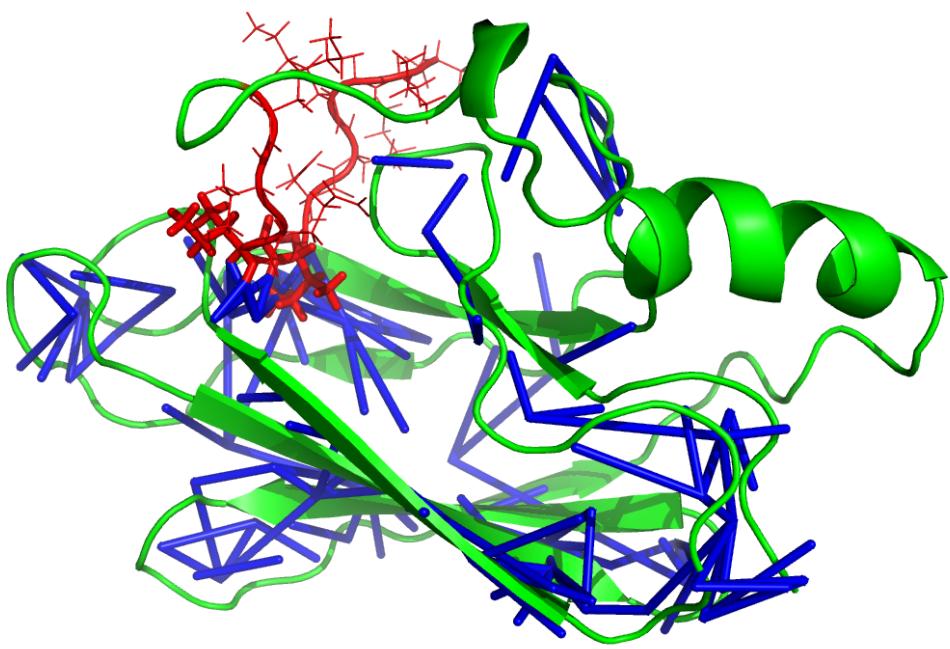
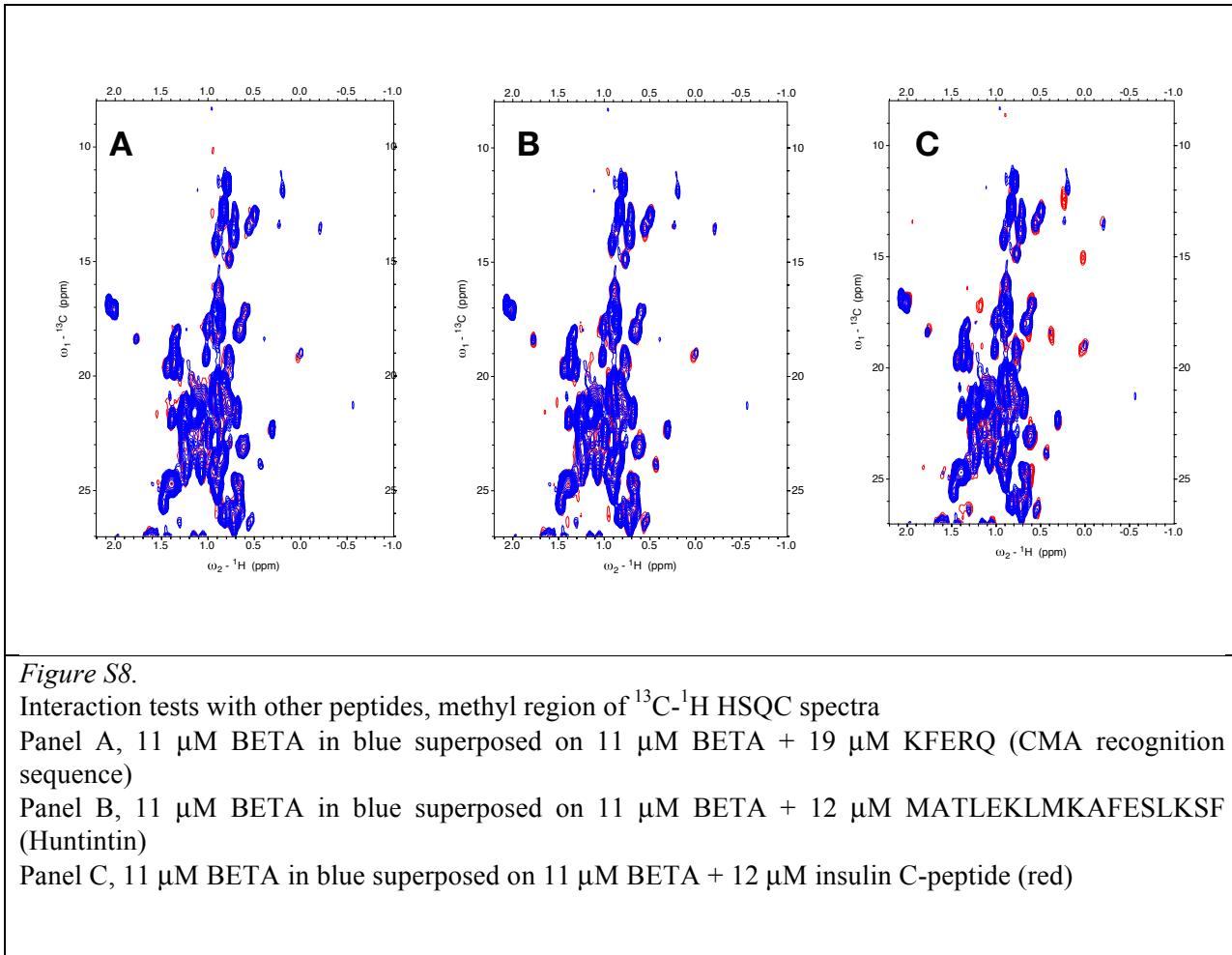


Figure S7

The molecular replacement structure for Hsc70 SBD with the sequence GKVQIINKKG in cis (red). In blue are non-sequential NOEs identified in the set of 3D NOE spectra confirming the structure of the BETA domain, and establishing details of the binding of GKVQIINKKG.



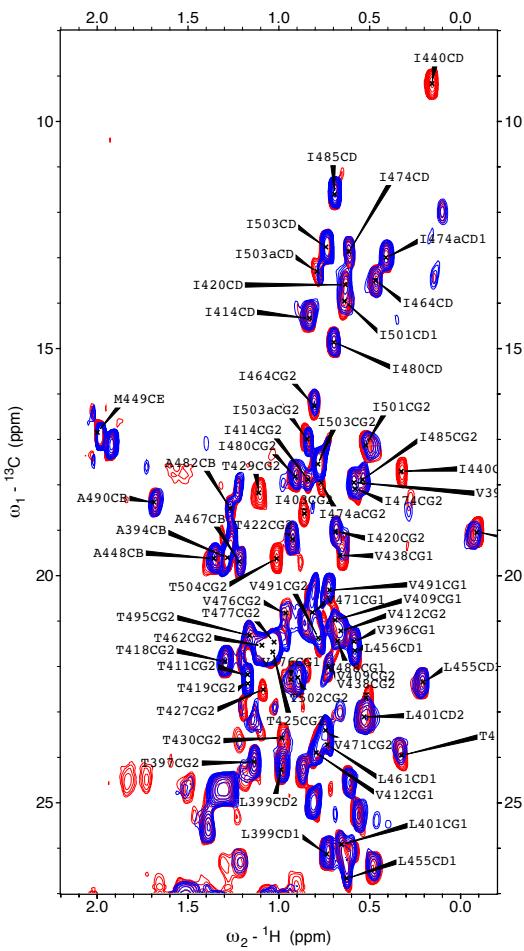


Figure 9.

Overlay of the ^1H - ^{13}C constant-time HSQC of 35 μM Hsc70 BETA (blue) and 35 μM Hsc70-BETA with 200 μM GKVQIINKKG (red). The assignments are indicated (BMRB accession code 27395)