

Supporting Information for:

Protein Cross-Linking Capillary Electrophoresis for Protein-Protein Interaction Analysis

Claire M. Ouimet[†], Hao Shao[§], Jennifer N. Rauch[§], Mohamed Dawod[†], Bryce Nordhues^{||}, Chad A. Dickey^{||}, Jason E. Gestwicki[§], Robert T. Kennedy^{†,‡,*}

[§]Department of Pharmaceutical Chemistry and the Institute for Neurodegenerative Disease, 675 Nelson Rising Ln, University of California at San Francisco, San Francisco, California 94158, United States

^{||}Department of Molecular Medicine, University of South Florida, 4001 E. Fletcher Ave., MDC 36, Tampa, Florida 33613, United States

[†]Department of Chemistry, University of Michigan
930 N. University Ave, Ann Arbor, Michigan, 48109, United States

[‡]Department of Pharmacology, University of Michigan
1150 W. Medical Center Dr., Ann Arbor, Michigan, 48109, United States

*Corresponding author

E-mail: rtkenn@umich.edu

Tel: 734-615-4363

Fax: 745-615-6462

Supporting Information:

Figure S-1. Free solution electrophoresis of chaperone complexes Hsp70-488-Bag3 and Hsp90-488 homodimer.

Table S-1. Quantitation of resolution of free proteins and protein complexes.

Figure S-2. Capillary gel electrophoresis of lysozyme immunocomplex.

Table S-2. Dependence of Hsp70-488-Bag3 complex detected on concentration of formaldehyde and cross-linking reaction time.

Table S-3. Dependence of Hsp90-488 complex detected on concentration of formaldehyde and cross-linking reaction time.

Table S-4. Dependence of FITC-lysozyme-antibody complex detected on concentration of formaldehyde and cross-linking reaction time.

Figure S-3. Saturation binding assays for Hsp70-488-Bag3 at different 1% formaldehyde reaction times.

Table S-5. Dependence of measured K_d of Hsp70-488-Bag3 interaction on cross-linking reaction time using 1% formaldehyde.

Figure S-4. Calorimetric isothermal-titration measurement of FITC-lysozyme interaction with anti-lysozyme.

Figure S-5. FCPIA data of small molecule Hsp70-488-Bag3 inhibitors JG-98, JG-231 and JG-311.

Figure S-1. Free solution electrophoresis of chaperone complexes Hsp70-488-Bag3 (A) and Hsp90-488 homodimer (B). The electrophoresis buffer was 10 mM, pH 10 borate.

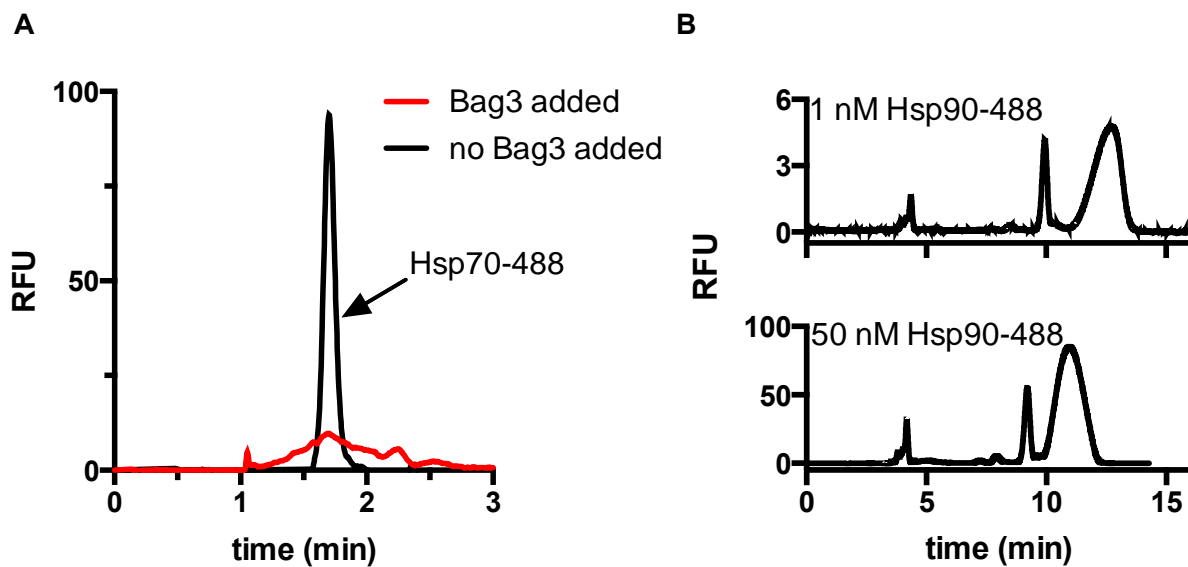


Table S-1. Resolution of free proteins from protein complexes.

Complex	Resolution
Hsp70-Bag3	1.0
Hsp90 homodimer	2.0
Lysozyme-anti-lysozyme	1.1

Figure S-2. Capillary gel electrophoresis separation of FITC-lysozyme immunocomplex.

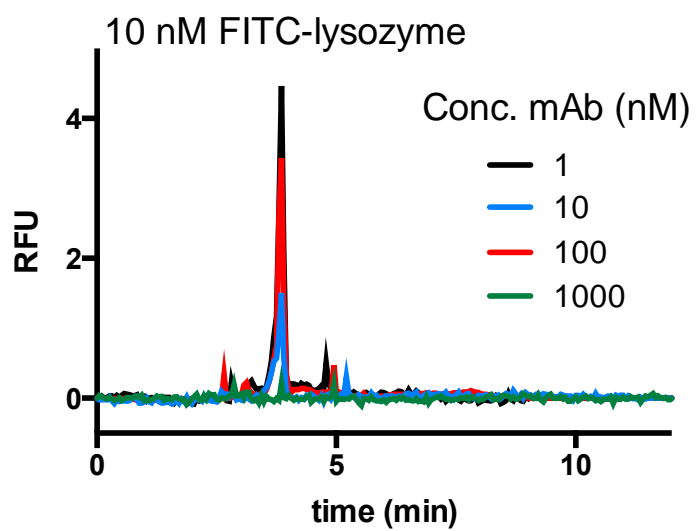


Table S-2. Dependence of 25 nM Hsp70-488 and 100 nM Bag3 on Hsp70-Bag3 complex detected on concentration of formaldehyde and cross-linking reaction time. Error is range of two trials.

Percent complex (%)					
formaldehyde concentration (%w/v)	Reaction time (min)				
	0.5	5	10	30	60
0.050	20 ± 3	21 ± 8	30 ± 17	38 ± 3	29 ± 6
0.50	19 ± 5	40 ± 13	60 ± 9	74 ± 9	46 ± 13
1.0	22 ± 9	46 ± 24	73 ± 10	73 ± 6	24 ± 3
2.5	31 ± 2	62 ± 17	73 ± 2	75 ± 8	44 ± 5

Table S-3. Dependence of 50 nM Hsp90-488 on complex detected on concentration of formaldehyde and cross-linking reaction time. Error is range of two trials.

Percent complex (%)				
formaldehyde concentration (%w/v)	Reaction time (min)			
	0.5	5	10	60
0.050	80 ± 5	77 ± 1	81.2 ± 0.4	71.4 ± 0.9
0.50	77 ± 5	74 ± 6	73.3 ± 0.6	74.6 ± 0.8
1.0	73 ± 2	79 ± 10	72 ± 3	73 ± 6.5
2.5	78 ± 6	73 ± 7	64 ± 4	58 ± 4

Table S-4. Dependence of 10 nM FITC-lysozyme and 20 nM antibody on complex detected on concentration of formaldehyde and cross-linking reaction time. Error is range of two trials.

Percent complex (%)				
formaldehyde concentration (%w/v)	Reaction time (min)			
	0.5	5	10	60
0.050	27 ± 15	39 ± 1	34 ± 7	24 ± 3
0.50	33 ± 15	37 ± 8	37 ± 12	39 ± 11
1.0	39 ± 15	35 ± 1	40 ± 9	34 ± 11
2.5	42 ± 24	27 ± 8	28 ± 12	23 ± 4

Figure S-3. Saturation binding assays for Hsp70-488-Bag3 at different 1% formaldehyde reaction times. All samples contain 25 nM Hsp70-488, error bars are standard deviation (n = 3).

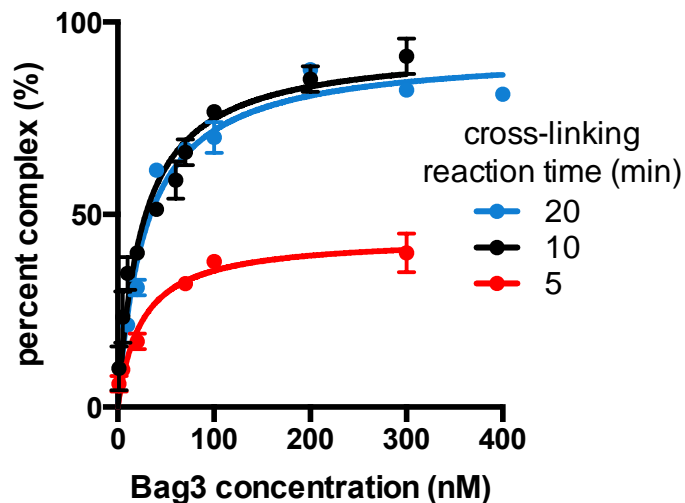


Table S-5. Dependence of measured K_d of Hsp70-488-Bag3 interaction on cross-linking reaction time using 1% formaldehyde.

Cross-linking reaction time (min)	$K_d, \text{Hsp70-Bag3}$ (nM)
5	25 ± 8
10	25 ± 5
20	29 ± 6

Figure S-4. Calorimetric isothermal-titration measurement of FITC-lysozyme interaction with anti-lysozyme.

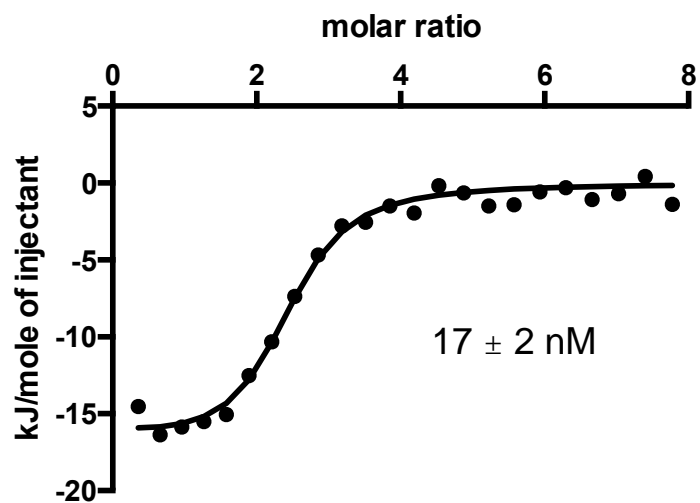


Figure S-5. FCPIA data of small molecule Hsp70-Bag3 inhibitors JG-98, JG-231 and JG-311. Error bars are range of two trials.

