



MicroScale Thermophoresis Experiments (2 labs)

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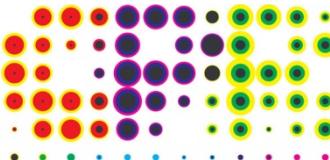
From 30/07/2017 to 04/08/2017

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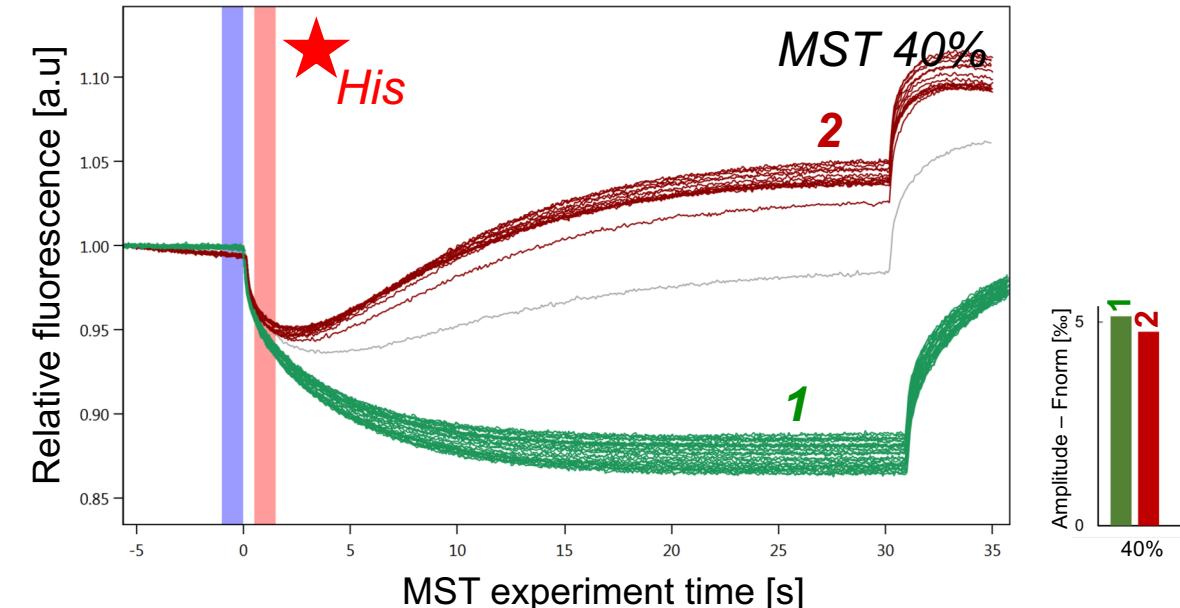
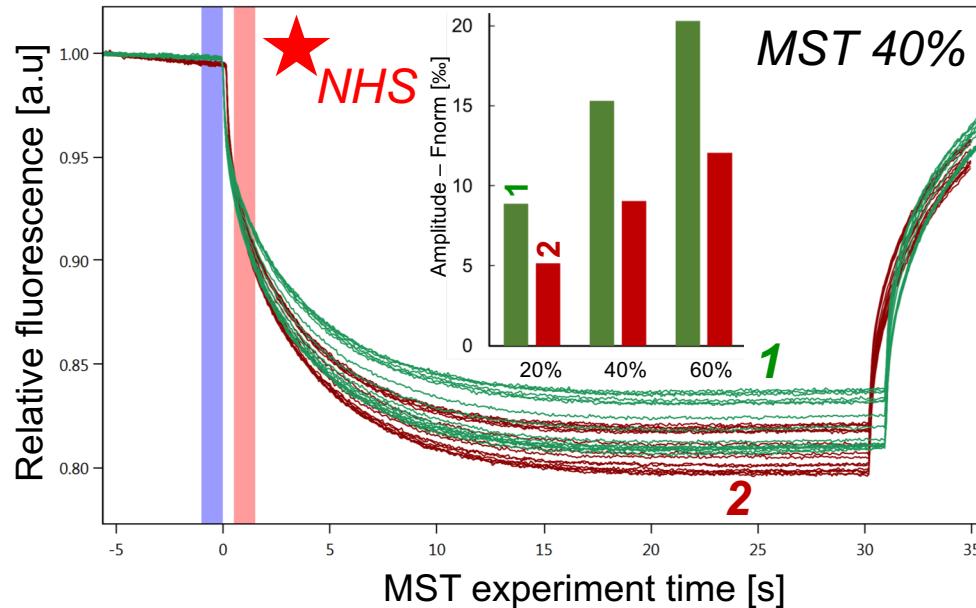
Two systems tested:

- ★ Protein/peptide interaction – low micromolar affinity
- ★ Protein/ DNA interaction – nanomolar affinity
- ★ Protein labeling efficiency - DOL:

Institution	DOL [%] – NHS chemistry	DOL [%] – MALEIMIDE chemistry
1	96	23
2	78	38

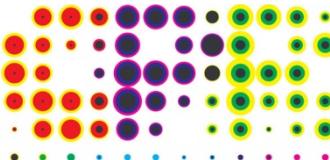


MST Experiments. ★ Protein/peptide interaction (2 labs)

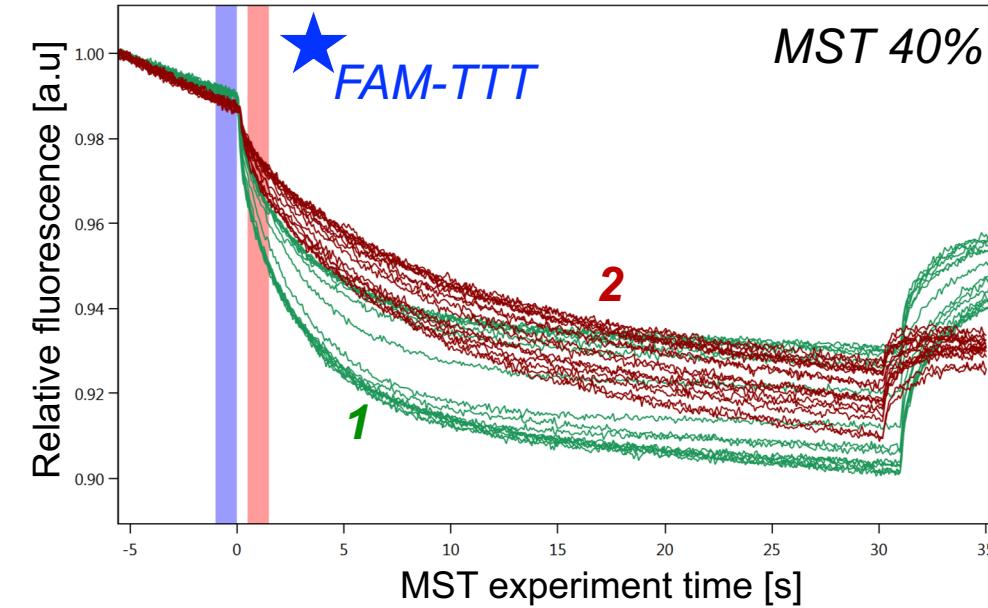
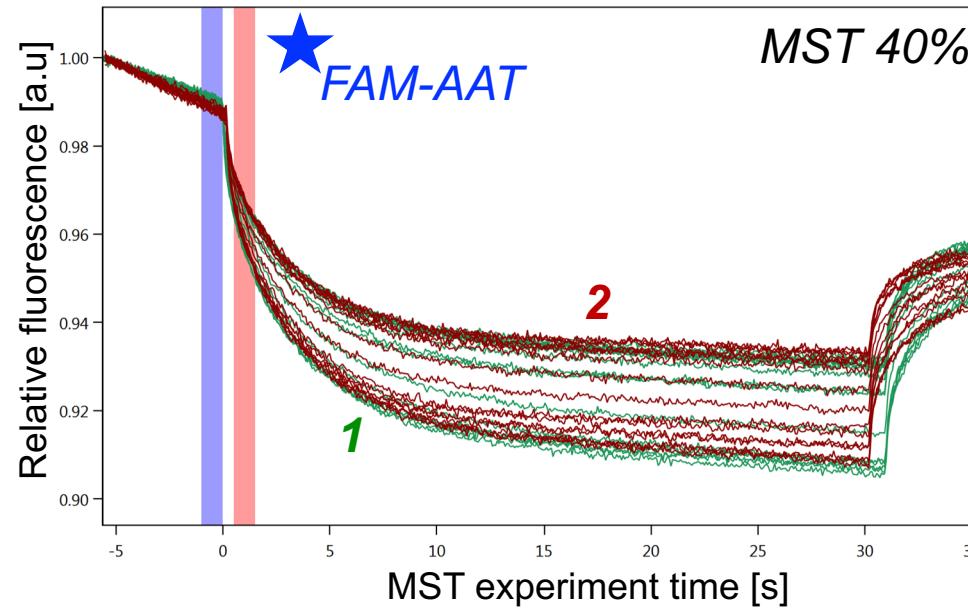


Institution | K_D [μM] at 40% MST power – (K_D) ITC values

	<i>Peptide 1</i>		<i>Peptide 2</i>
	<i>NHS labeling</i>	<i>His labeling</i>	<i>NHS labeling</i>
1	4.7 (32)	1.4 (32)	0.84 (4)
2	5.2 (32)	2.6 (32)	0.68 (4)



MST Experiments. Protein/[★]DNA interaction (2 labs)



Institution	Amplitudes at 40% MST power		K_D [nM] at 40% MST power – (K_D) ITC values	
	AAT, DNA 1	TTT, DNA 2	AAT, DNA 1	TTT, DNA 2
1	10.3	12.1	20 (40)	41.2 (450)
2	8.5	5.6	2 (40)	25.5 (450)