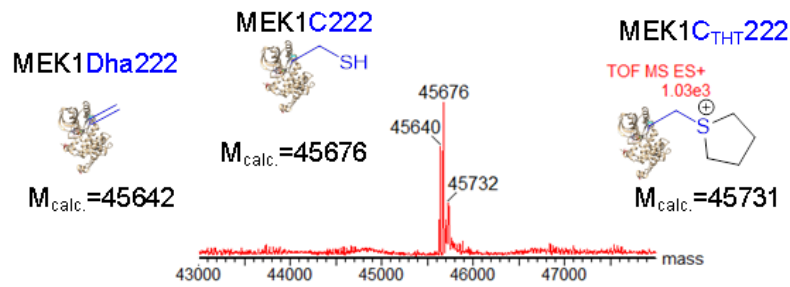


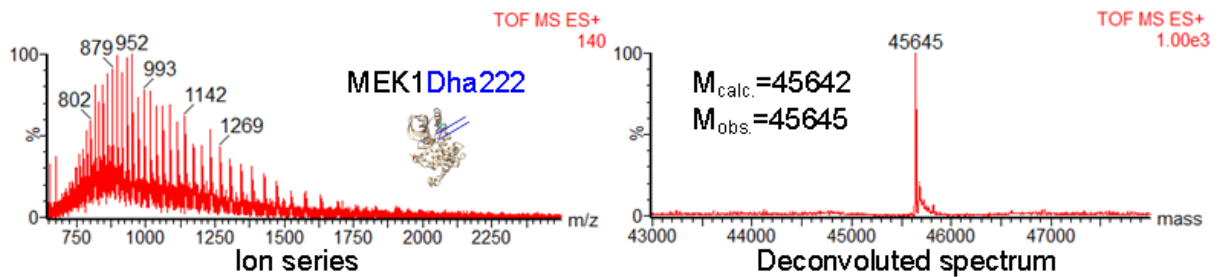
The Minimum Protein Staple? – Towards ‘bio’-Baldwin’s Rules via Inter-phosphosite Linking in the MEK1 Activation Loop

SUPPLEMENTARY DATA

Supplementary Figures

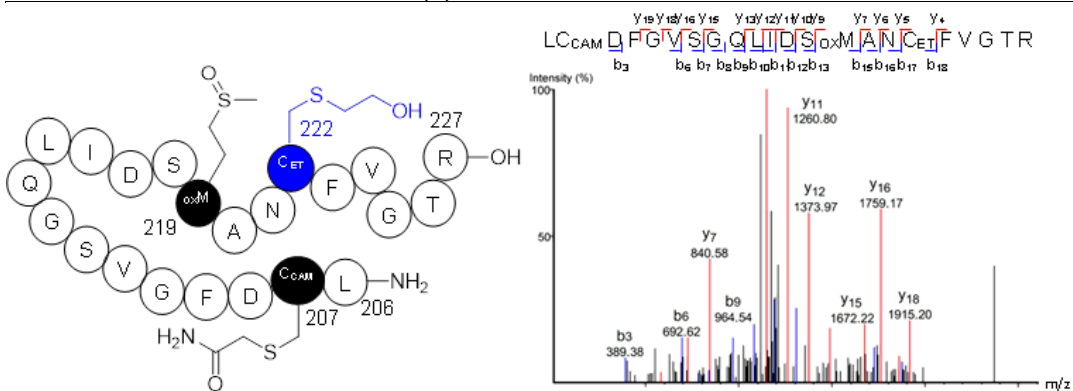


Supplementary Figure S1. Reaction of Penta-Cys MEK1-Cys222 with DIB in the presence of ADP-Mg after 6h.

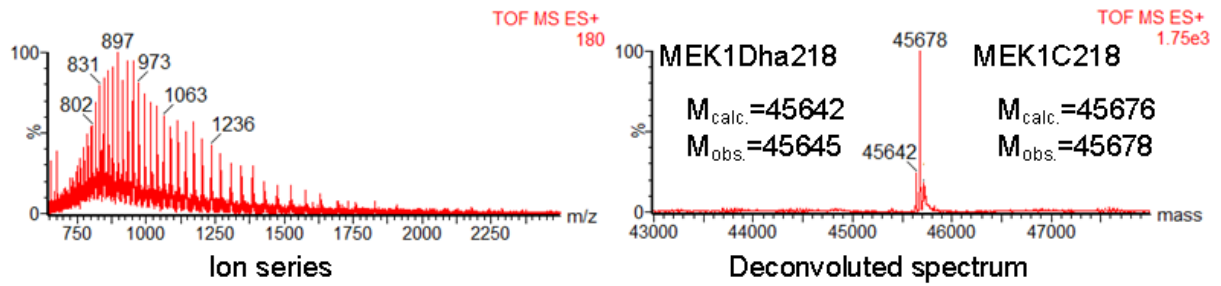


Supplementary Figure S2. Reaction of Penta-Cys MEK1-Cys222 with DIB in the presence of ADP-Mg after 20h.

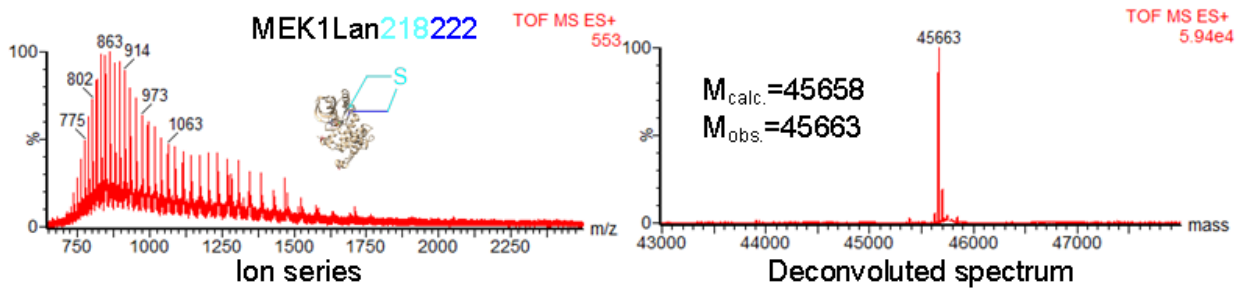
Peptide	RT/min	m/z (charge)	ppm	Modifications	-10lgP	TIC	Max Intensity
206-227	37.74	1225.5529 (2)	-6.7	C _{ET} 222, oxM219, C _{CAM} 207	98.92	1.21E4	7.74E2



Supplementary Figure S3. MSMS Analysis of Product of Reaction of Penta-Cys MEK1-Cys222 with DIB in the presence of ADP-Mg.

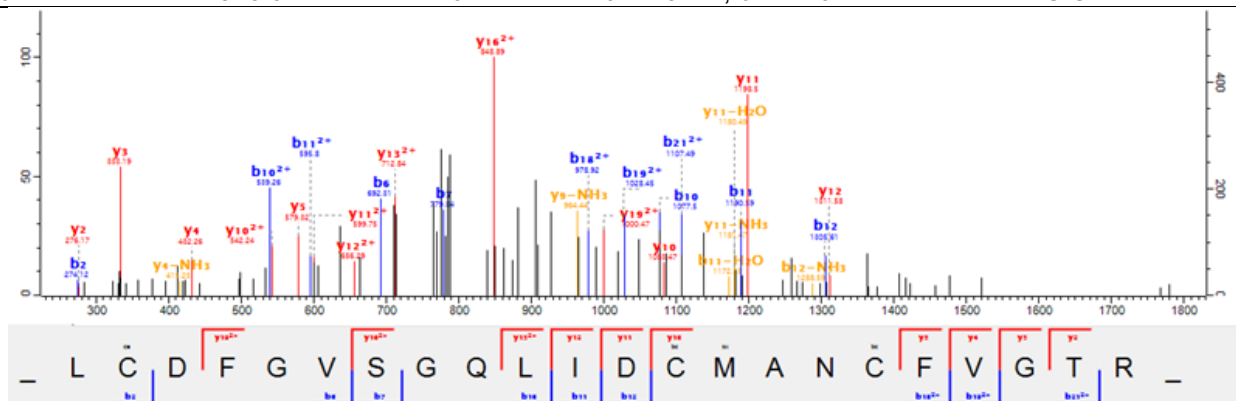


Supplementary Figure S4. Reaction of Penta-Cys MEK1-Cys218 with DIB in the presence of ADP-Mg after 26h.

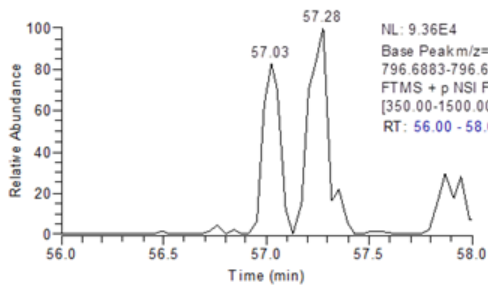


Supplementary Figure S5. Reaction of Hexa-Cys MEK1-Cys218-Cys222 with DIB in the presence of ADP-Mg after 15h.

Peptide	Mass (Da)	Score	Modifications	Method
206-227	2348.04	101	Lan218222, oxM219	ITMS CID

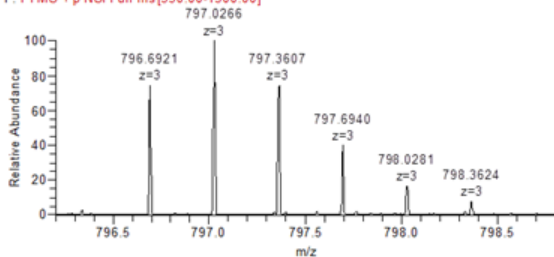


Supplementary Figure S6. MSMS Analysis of Product of Reaction of Hexa-Cys MEK1-Cys218-Cys222 with DIB in the presence of ADP-Mg.

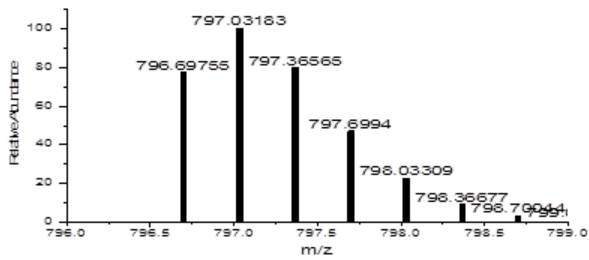


Chromatogram

#14788-14930 RT: 56.87-57.32 AV: 13 NL: 4.76E4
F: FTMS + p NSI Full ms[350.00-1500.00]

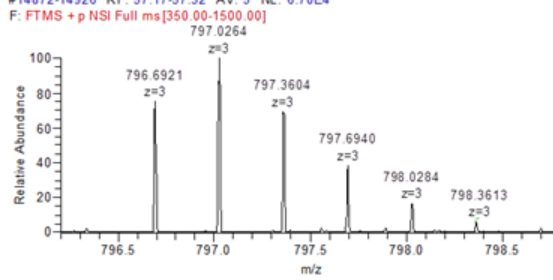


MS of RT=57.03 min



Calculated MS for desired peptide

#14872-14926 RT: 57.17-57.32 AV: 5 NL: 6.70E4



MS of RT=57.28 min

Supplementary Figure S7. LC trace of the LCMS-MS analysis of Reaction of Hexa-Cys MEK1-Cys218-Cys222 with DIB in the presence of ADP-Mg.