# **Supporting Information**

Optically Sensing Phospholipid Induced Coil-Helix Transitions in the Phosphoinositide-Binding Motif of Gelsolin

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#### 1. Supplementary Figures



**Figure S1:** Helical character (%) of the Gel150-169 peptide monitored over 100 ns in the presence of PI(4,5)P2 containing membrane.



**Figure S2.** Distance difference map representing the inter residue distance changes between the alpha carbons within Gel150-169 sequence with amide as C-terminus upon going from the helix form to the coil form. The color scale represents the change in average distances between 0-6 Å. A, B and C indicate positions with maximum distance change.



**Figure S3.** Absorption spectrum of 10  $\mu$ M IAEDANS solution in aqueous buffer (20 mM HEPES, 100 mM NaCl, pH 7.4, 0.07 % DMSO). Molar absorption coefficient ( $\epsilon$ ) of IAEDANS at 287 nm is 1339 M<sup>-1</sup>cm<sup>-1</sup>. The concentration of IAEDANS was calculated based on the reported value of  $\epsilon$  of IAEDANS which was 4500 M<sup>-1</sup>cm<sup>-1</sup> at 337 nm.<sup>(1)</sup>



**Figure S4.** CD spectra of Gel-*Q11W-G18C* peptide solution (40  $\mu$ M) in phosphate buffer (10 mM, pH 7.5) in presence of varying concentrations of PI(4,5)P2 micelles – 0  $\mu$ M (black), 10  $\mu$ M (red), 20  $\mu$ M (blue), 40  $\mu$ M (pink), 80  $\mu$ M (green). CD spectra were recorded at 25 °C in a quartz cuvette (10 mm x 2 mm) in a JASCO J-1500 CD Spectrometer.



**Figure S5.** Normalized fluorescence emission at 480 nm for IAEDANS at  $\lambda_{ex}$  340 nm (black) and at  $\lambda_{ex}$  287 nm (red). Concentration of Gel-*Q11W-G18C*-IAEDANS was 10  $\mu$ M.



**Figure S6.** Normalized fluorescence intensity at 360 nm ( $\lambda_{ex}$  287 nm) for Gel-*Q11W-G18C*-IAEDANS (10  $\mu$ M) in the presence of 100% PI(4,5)P2 micelles, 50% PI(4,5)P2 vesicles, 20% PI(4,5)P2 vesicles, PC, and in the absence of phospholipids. All phospholipid concentrations were 10  $\mu$ M.



**Figure S7.** Fluorescence emission spectra for Gel-*Q11W-G18C*-IAEDANS (10  $\mu$ M) with increasing concentrations of PC (0-20  $\mu$ M),  $\lambda_{ex}$  287 nm.



**Figure S8.** Average lifetime for tryptophan in Gel-*Q11W-G18C* (10  $\mu$ M) in the presence of increasing concentrations of a) PI(4,5)P2-PC micelles (1-20  $\mu$ M), and b) PC vesicles (1-20  $\mu$ M),  $\lambda_{ex}$  287 nm.



**Figure S9.** a) Average lifetime for tryptophan in Gel-*Q11W-G18C*-IAEDANS (10  $\mu$ M) in the presence of increasing concentrations of PC (black squares) and increasing concentrations of 20% PI(4,5)P2-PC (red circles),  $\lambda_{ex}$  287 nm. b) Average lifetime for tryptophan in Gel-*Q11W-G18C*-IAEDANS (10  $\mu$ M) in the presence of 100% PI(4,5)P2 micelles, 50% PI(4,5)P2 vesicles, 20% PI(4,5)P2 vesicles, PC, and in the absence of phospholipids. All phospholipid concentrations were 10  $\mu$ M.

#### 2. Characterization and purity of peptides

a) Gel-Q11W-G18C, KHVVPNEVVVWRLFQVKCRR-CONH<sub>2</sub>

Molecular weight: 2492.01 (Calculated)

LR-ESI/MS, profile mode: 2492

MALDI-TOF-MS: m/z, 2492.90 (M+H<sup>+</sup>)

#### LC elution trace at 280 nm:



ESI-MS (+ve mode) of peak (25 min) in LC:



## b) Gel-Q11W-G18C-IAEDANS: KHVVPNEVVVWRLFQVKC-(IAEDANS)RR

Molecular weight: 2798.35 (Calculated)

LR-ESI/MS, profile mode: 2798

MALDI-TOF-MS: m/z, 2799.07 (M+H<sup>+</sup>)

## LC elution trace at 340 nm:





#### ESI-MS (+ve mode) of peak (26 min) in LC:

**Figure S10.** Size distribution (hydrodynamic radius) of a) PC vesicles, b) 20 % PI(4,5)P2-PC vesicles, and c) 50 % PI(4,5)P2-PC vesicles measured by DLS.

## 6. <u>References</u>

(1) Saxena, A. M.; Udgaonkar, J. B.; Krishnamoorthy, G. Characterization of Intra-molecular Distances and Site-specific Dynamics in Chemically Unfolded Barstar: Evidence for Denaturant-dependent Non-random Structure. *J. Mol. Biol.* **2006**, 359, 174-189.