## **Accepted Manuscript**

Modeling, dynamics and phosphoinositide binding of the pleckstrin homology domain of two novel PLCs: n1 and n2

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PII: \$1093-3263(18)30432-7

DOI: 10.1016/j.jmgm.2018.07.012

Reference: JMG 7213

To appear in: Journal of Molecular Graphics and Modelling

Received Date: 10 June 2018
Revised Date: 23 July 2018
Accepted Date: 25 July 2018

Please cite this article as: S. Rai, P. Mohanty, S. Bhatnagar, Modeling, dynamics and phosphoinositide binding of the pleckstrin homology domain of two novel PLCs:  $\eta 1$  and  $\eta 2$ , *Journal of Molecular Graphics and Modelling* (2018), doi: 10.1016/j.jmgm.2018.07.012.

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