

# Membrane curvature sensing by model biomolecular condensates

Midhun Mohan Anila<sup>1</sup>, Rikhia Ghosh<sup>2</sup>, Paweł Rogowski<sup>1</sup>, Bartosz Różycki<sup>1</sup>

<sup>1</sup>Institute of Physics, Polish Academy of Sciences, Aleja Lotników 32/46, 02-668 Warsaw, Poland.

<sup>2</sup>Icahn School of Medicine at Mount Sinai, 1 Gustave L. Levy Pl, New York, NY 10029, USA.

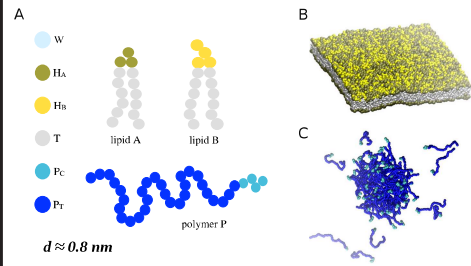
Email: midhun@ifpan.edu.pl



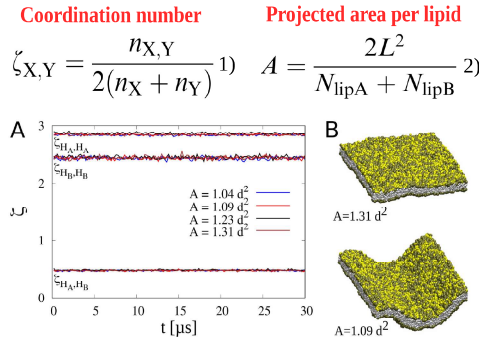
## Abstract

Biomolecular condensates (BCs), comprised mainly of intrinsically disordered proteins, form through liquid-liquid phase separation within biological cells. Despite considerable research on BCs in the cytosol and nucleus, their behavior at cellular membranes remains largely unexplored. Galectin-3, a mixed-folded protein consisting of an intrinsically disordered N-terminal domain (NTD) and a carbohydrate recognition domain (CRD), plays pivotal roles in various biological processes such as immune responses, cell migration, and signaling. It has been demonstrated that galectin-3 interacts with glycosphingolipids on the cell membrane, facilitating clathrin-independent endocytosis [1]. Using dissipative particle dynamics (DPD) simulations, we explore how polymer models resembling galectin-3 sense and respond to membrane curvature. Our findings suggest a generic mechanism by which BCs sense membrane curvature, potentially influencing such cellular processes as endocytosis [2].

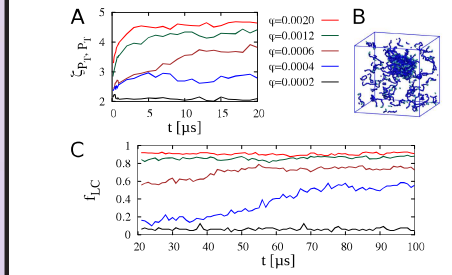
## DPD Model



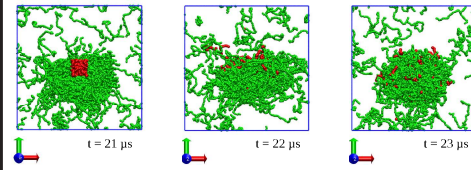
## Absence of polymers



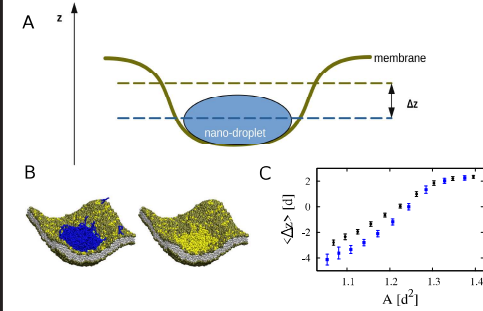
## Absence of lipids



## Fluidity of nanodroplets



## Immersion depth



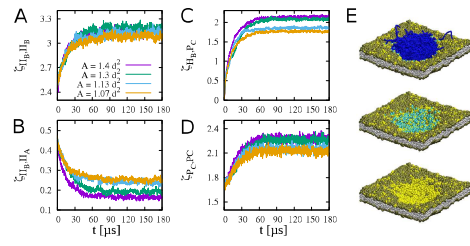
## References

- [1] R. Lakshminarayanan et al. Nature Cell Biology 16(6), 592-603 (2014).
- [2] M. M. Anila et al. Soft Matter 19(20), 3723-3732 (2023).

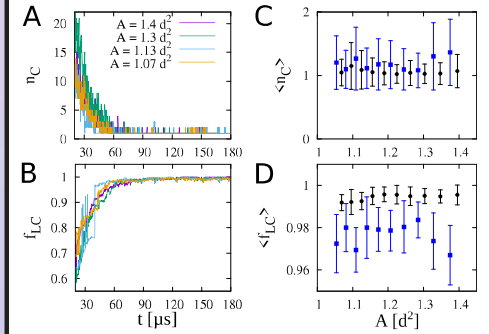
## Acknowledgements

This research has received support from the National Science Centre (grant No. 2020/39/B/NZ1/00377). We acknowledge the Polish high-performance computing infrastructure PLGrid for awarding this project access to the LUMI supercomputer (grant No. PLL/2023/04/016485).

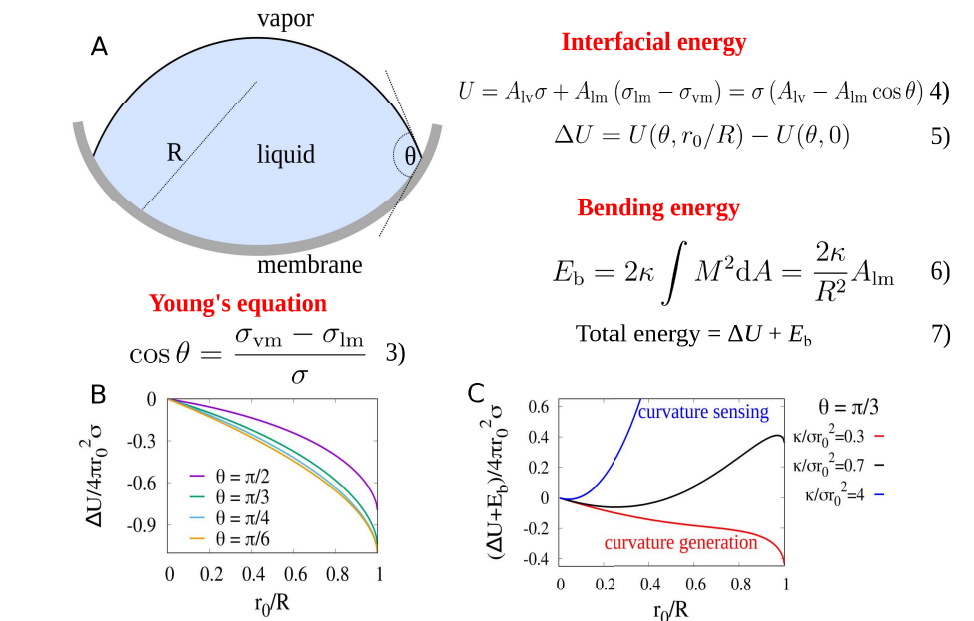
## With lipids and polymers



## Clusterization of polymers



## Toy-model



## Conclusion

Our results of the coarse-grained simulations and analytical calculations indicate together towards a generic mechanism of membrane curvature sensing by biomolecular condensates.

