



## Atomistic insights into the Protein-functionalized nanoparticles

S.Yasami Khiabani<sup>a</sup>, H.Rahimi<sup>b</sup>, M.Golkar<sup>c</sup>, A.Karkhaneh<sup>a\*</sup>, MA.Shokrgozar<sup>d\*\*</sup>

<sup>a</sup> Department of Biomedical Engineering, AmirKabir University of Technology, Tehran, Iran

<sup>b</sup> Molecular Medicine Department, Biotechnology Research Center, Pasteur Institute of Iran, Tehran, Iran

<sup>c</sup> Molecular Parasitology Laboratory, Department of Parasitology, Pasteur Institute of Iran, Tehran, Iran

<sup>d</sup> National Cell Bank of Iran, Pasteur Institute of Iran, Tehran, Iran

\* a.karkhaneh@aut.ac.ir

\*\* mashokrgozar@pasteur.ac.ir

**Abstract:** Protein-functionalized nanoparticles can be used in various pharmaceutical approaches but the choice of nanoparticle size, shape, charge, linking chemistry, protein-labeling site, etc. remains challenging due to their effect on protein structure and function. Here molecular dynamics simulation was used as a functional way to predict atomistic insights into the charge of polystyrene nanoparticle and epidermal growth factor stability as a model of the nanoparticle-protein complex. Our result showed that the epidermal growth factor orientation regarding the conjugation is suitable and the RMSD analysis represents the complex stability during MD simulation.

**Keywords:** Molecular Dynamics, nanoparticle, conjugation, polystyrene, epidermal growth factor

### References

- [1] M. Mahmoudi, I. Lynch, M. R. Ejtehad, M. P. Monopoli, F. B. Bombelli, and S. Laurent, "Protein-nanoparticle interactions: Opportunities and challenges," *Chem. Rev.*, vol. 111, no. 9, pp. 5610–5637, 2011.
- [2] K. E. Sapsford, K. M. Tyner, B. J. Dair, J. R. Deschamps, and I. L. Medintz, "Analyzing nanomaterial bioconjugates: A review of current and emerging purification and characterization techniques," *Anal. Chem.*, vol. 83, no. 12, pp. 4453–4488, 2011.
- [3] S. J. Marrink, H. J. Risselada, S. Yefimov, D. P. Tieleman, and A. H. De Vries, "The MARTINI Force Field : Coarse Grained Model for Biomolecular Simulations The MARTINI Force Field : Coarse Grained Model for Biomolecular Simulations," *J. Phys. Chem. B*, vol. 111, no. June, pp. 7812–7824, 2007.
- [4] G. Rossi, L. Monticelli, S. R. Puisto, I. Vattulainen, and T. Ala-Nissila, "Coarse-graining polymers with the MARTINI force-field: polystyrene as a benchmark case," *Soft Matter*, vol. 7, no. 2, pp. 698–708, 2011.
- [5] D. H. De Jong, G. Singh, W. F. D. Bennett, T. a Wassenaar, L. V Sch, X. Periole, P. Tieleman, and S. J. Marrink, "Improved Parameters For F or The Martini Coarse- Coarse - Grained Protein Force Field," *J. Chem. Theory Comput.*, vol. 9, no. 1, pp. 687–697, 2012.
- [6] L. Monticelli, S. K. Kandasamy, X. Periole, R. G. Larson, D. P. Tieleman, and S. J. Marrink, "The MARTINI coarse grained force field: extension to proteins," *J. Chem. Theory Comput.*, vol. 4, no. 5, pp. 819–834, 2008.
- [7] G. Rossi, J. Barnoud, and L. Monticelli, "Polystyrene nanoparticles perturb lipid membranes," *J. Phys. Chem. Lett.*, vol. 5, pp. 241–246, 2014.