

Università degli Studi di Napoli Federico II

Tutor: prof.ssa Michela Varra

Co-tutor: prof. Aldo Galeone

**PROJECT TITLE:** New Molecules and Modified Aptamers for Targeting Proteins Involved in Pathological Processes

## **Project description**

The present PhD project aims at the synthesis and structural characterization of new molecules capable of selectively modulating the functional activity of proteins involved in complex pathological processes, such as cancer, neurodegenerative diseases, and chronic inflammation. Among the most relevant protein targets, the proteasome<sup>1-3</sup> will be analyzed first, due to its crucial role in protein degradation and in the regulation of cellular homeostasis. The resulting experimental approach will also be extended to other clinically relevant proteins<sup>4-6</sup>. A central element of the project will be the rational development of new molecules, such as derivatives of small organic molecules already known for their affinity towards specific protein targets, or analogues of endogenous ligands – such as proteasome regulatory proteins, miRNA, or siRNA – suitably modified to increase their resistance to proteases/nucleases, their affinity for biological targets, and to modulate the immune response. In parallel, new phosphoramidite building blocks will be synthesized to be used as nucleoside analogues in the automated solid-phase synthesis of nucleic acids, selecting appropriate sequences of therapeutic RNA and/or DNA aptamers. The modifications introduced into the selected sequences will aim to promote the bioactive conformation<sup>7-8</sup> in the recognition of the target protein and to modulate the innate immune response associated with the administration of therapeutic nucleic acids<sup>9,10</sup>, through the analysis of the correlations between the cellular response and the introduced modifications. All newly synthesized molecules will undergo indepth characterization using spectroscopic techniques (CD, UV, NMR) and spectrometric techniques (MS, HRMS, MS<sup>n</sup>), employing the advanced instrumentation available at the departmental instrumental analysis laboratories, including the NMR Laboratory and the MS Laboratory. In summary, the project aims to develop an innovative and versatile platform for the production and characterization of molecular regulators – whether small organic molecules or modified nucleic acids – with potential applications in the treatment of complex diseases and in the understanding of the involved molecular mechanisms.

Dipartimento di Farmacia

Via Domenico Montesano, 49 • 80131 Napoli, Italia angelo.izzo@unina.it • + 39 081678658 www.farmacia.unina.it



Università degli Studi di Napoli Federico II

## REFERENCES

- CHUAH, J.J.Y., et al. Minimal mechanistic component of HbYX-dependent proteasome activation that reverses impairment by neurodegenerative-associated oligomers. *Commun Biol* 6, 725 (2023). <u>https://doi.org/10.1038/s42003-023-05082-9</u>
- DI DATO, A., et al. Electrostatic map of proteasome α-rings encodes the design of allosteric porphyrin-based inhibitors able to affect 20S conformation by cooperative binding. *Sci Rep* 7, 17098 (2017). <u>https://doi.org/10.1038/s41598-017-17183-6</u>
- 3. **PERSICO, M., et al.** Modulation of the 20S Proteasome Activity by Porphyrin Derivatives Is Steered through Their Charge Distribution. *Biomolecules* **12**, 741 (2022). <u>https://doi.org/10.3390/biom12060741</u>
- 4. **BAHIT, M.C., et al.** Thrombin as target for prevention of recurrent events after acute coronary syndromes. *Thromb Res* **235**, 116–121 (2024). <u>https://doi.org/10.1016/j.thromres.2024.02.003</u>
- TONELLO, F., et al. Nucleolin: a cell portal for viruses, bacteria, and toxins. *Cell Mol Life Sci* 79, 271 (2022). <u>https://doi.org/10.1007/s00018-022-04300-7</u>
- 6. VAN DEN AVONT, A., et al. Molecular diagnostics and therapeutics. *Front Mol Biosci* 10, Article 1217769 (2023). <u>https://doi.org/10.3389/fmolb.2023.1217769</u>
- VIRGILIO, A., et al. Structural properties and anticoagulant/cytotoxic activities of heterochiral enantiomeric thrombin binding aptamer (TBA) derivatives. *Nucleic Acids Res* 48, 12556–12565 (2020). <u>https://doi.org/10.1093/nar/gkaa1109</u>
- VIRGILIO, A., et al. Improved thrombin binding aptamer analogues containing inversion of polarity sites: structural effects of extra-residues at the ends. *Org Biomol Chem* 14, 7707–7714 (2016). <u>https://doi.org/10.1039/c6ob00931j</u>
- 9. NELSON, J., et al. Impact of mRNA chemistry and manufacturing process on innate immune activation. *Sci Adv* 6, eaaz6893 (2020). <u>https://doi.org/10.1126/sciadv.aaz6893</u>
- 10. WU, K., et al. Variant SARS-CoV-2 mRNA vaccines confer broad neutralization as primary or booster series in mice. *Vaccine* **39**, 7394–7400 (2021). https://doi.org/10.1016/j.vaccine.2021.10.046

## FUNDS

PRIN 2022 "Modulating confoRmational Equilibria of prion protein and proteaSome to tune protEostasis neTwork (RESET)"-

Dipartimento di Farmacia