



NANOSCIENCES

2nd YEAR PhD FINAL EXAM

Sara Chiarugi

Pisa, 17/10/2019

COURSES & OTHER ACTIVITIES

Attended courses	Hours
Seminar Cycle in Biophysical Sciences – Prof. F. Cardarelli	45
Topics in Structural Biology – Prof. A. Pastore	20

Seminars/Workshops	Hours
Seminars	31
EMBO Workshop (Hamburg): "Tools for Structural Biology of Membrane Proteins"	25







RESEARCH ACTIVITY

BioStructures Lab Dr. Gianpiero Garau

Design and engineering of a 3D protein-trapping nano-biostructure for X-ray crystallography

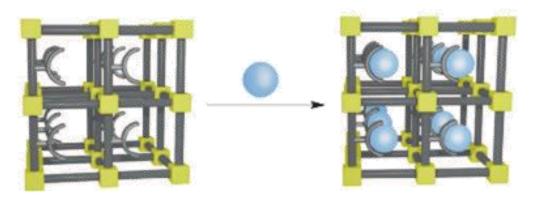
Discovery of ligands targeting the membrane protein NAPE-PLD

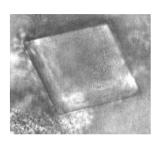
PROJECT GOAL & BACKGROUND

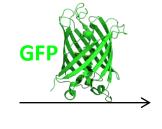
Efficient and innovative methodology for membrane protein crystallization and structure determination based on **trapping protein-nanocages**

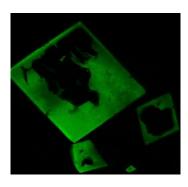
Smart protein scaffolds (gCAD) able to:

- ☐ Generate crystalline cages with internal cavities
- Crystallize easily and efficiently (known crystallization conditions)
- Trap guest target proteins







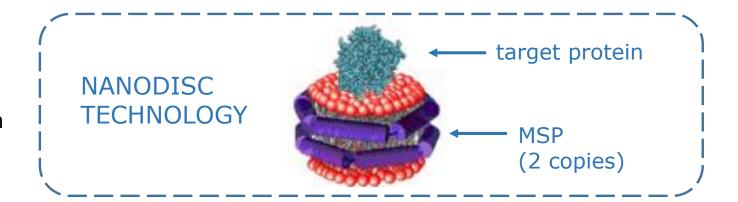


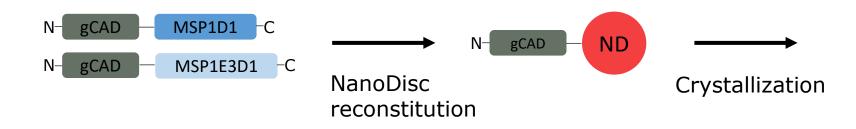
gCAD crystal

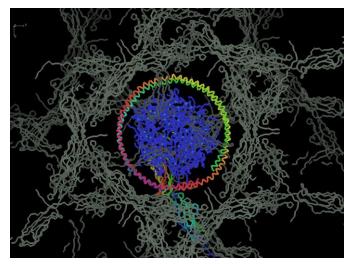
Scheme. Assembly model of the system

MEMBRANE PROTEIN TARGETS

- ☐ **Challenging** structure determination
- □ ~3% of crystal structures in the PDB

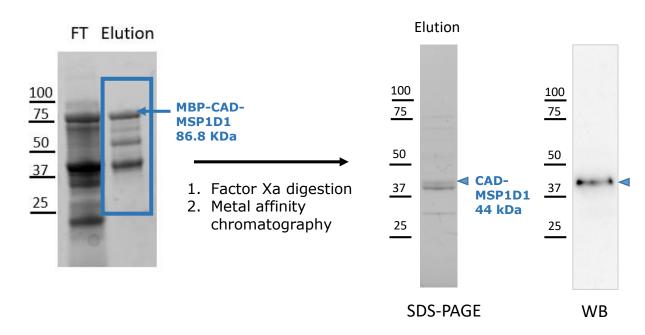






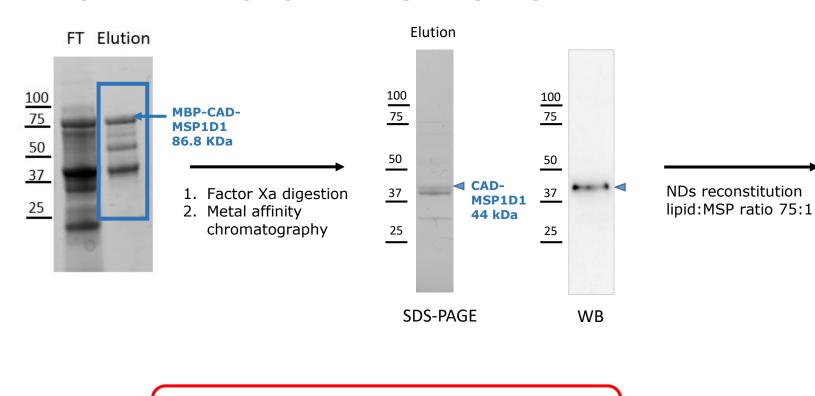
Model design

PROTEIN EXTRACTION AND PURIFICATION



X Limited purity & protein yield

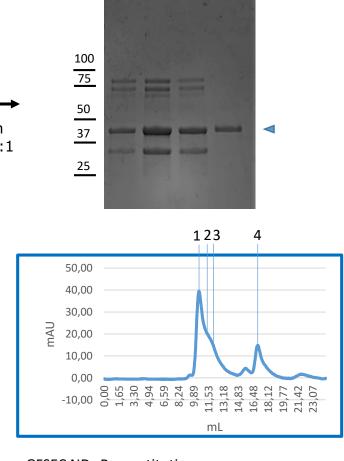
PROTEIN EXTRACTION AND PURIFICATION



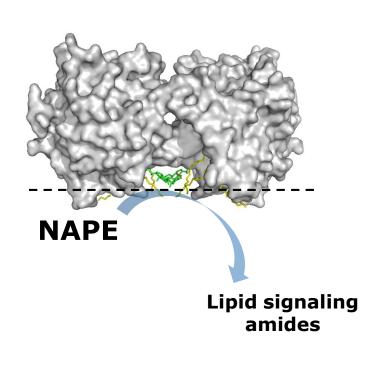
X Limited purity & protein yield

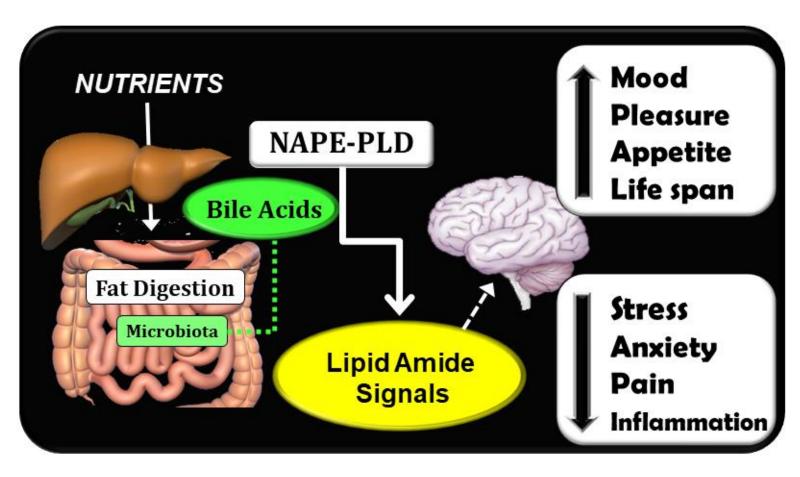
NANODISC RECONSTITUTION

1 2 3

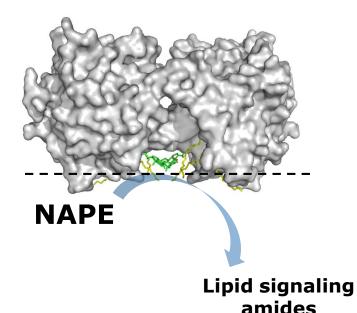


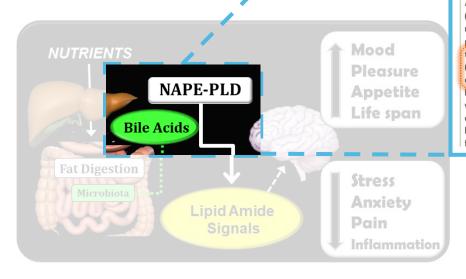
GFSEC NDs Reconstitution





Ligand Discovery with applications in metabolic and eating disorders







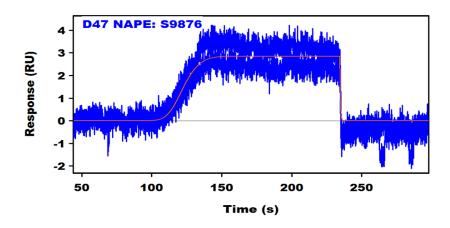
Bile acids in glucose metabolism and insulin signalling — mechanisms and research needs

Tiara R. Ahmad^{1,2} and Rebecca A. Haeusler⁶ ^{1,2}*

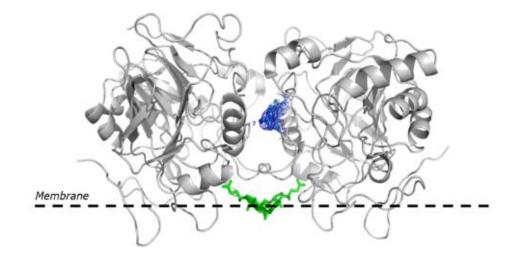
Abstract | Of all the novel glucoregulatory molecules discovered in the past 20 years, bile acids (BAs) are notable for the fact that they were hiding in plain sight. BAs were well known for their requirement in dietary lipid absorption and biliary cholesterol secretion, due to their micelle-forming properties. However, it was not until 1999 that BAs were discovered to be endogenous ligands for the nuclear receptor FXR. Since that time, BAs have been shown to act through multiple receptors (PXR, VDR, TGR5 and S1PR2), as well as to have receptor-independent mechanisms (membrane dynamics, allosteric modulation of N-acyl phosphatidylethanolamine phospholipase D). We now also have an appreciation of the range of physiological, pathophysiological and therapeutic conditions in which endogenous BAs are altered, raising the possibility that BAs contribute to the effects of these conditions on glycaemia. In this Review, we highlight the mechanisms by which BAs regulate glucose homeostasis and the settings in which endogenous BAs are altered, and provide suggestions for future research.

Ligand Discovery with applications in metabolic and eating disorders

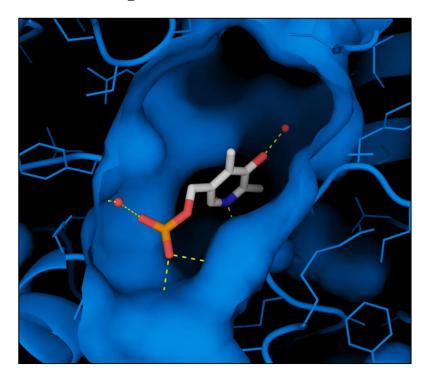
☐ Ligands screening by SPR



■ Interactions at atomic resolution by X-ray diffraction



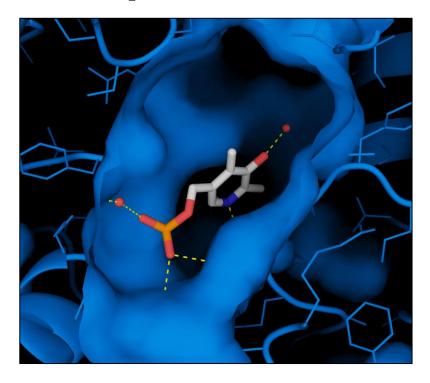
Structure of NAPE-PLD in complex with PLP



Pyridoxal-5'-phosphate

- Active form of Vitamin B6
- Coenzyme in different cellular pathways
- > Involved in **neurotransmitters generation**

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NAPE-PLD at the cross-talk between Lipid Amides and Bile Acid signaling



Hypothesis: modulation of neurotransmission

VALIDATION



Design and synthesis of PLP analogues





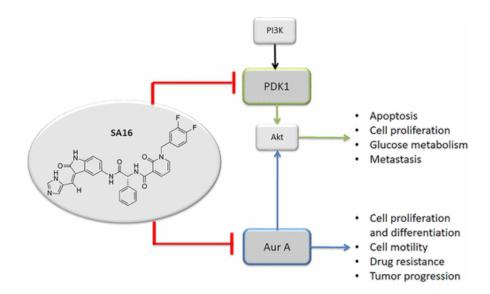
In vitro studies (binding & bioassays in living cells)



In vivo studies

Structural, Functional & Physiological basis

AurA PROJECT



ACS Chemical Neuroscience

Article

Dual inhibition of PDK1 and Aurora Kinase A: an effective strategy to induce differentiation and apoptosis of human glioblastoma multiforme stem cells

Simona Daniele, Simona Sestito, Deborah Pietrobono, Chiara Giacomelli, Grazia Chiellini, Danilo Di Maio, Luciana Marinelli, Ettore Novellino, Claudia Martini, and Simona Rapposelli

Glioblastoma Multiforme resistance & recurrence



Prof. Simona Rapposelli University of Pisa Department of Pharmacy

Crystal structure of AurA in complex with the lead compound VI8

