Research Summary

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My research interests are centered around application of organic synthesis to designing molecules with useful properties for biomedical applications. This broad definition led me to pursue research in the range of subjects from medicinal chemistry to molecular pharmacology and chemical biology of membrane proteins and non-coding RNA, and development of diagnostic fluorescent sensors.

Biased modulators of GPCRs

Biased μ -opioid receptor agonists. As a part of multi-center collaboration I synthesized newly designed biased agonists of μ -opioid receptor. My main contribution was synthesis and preparative separation of individual enantiomers of the optimized hits. The stere-ochemistry turned out to be the key parameter in SAR that defined signaling bias and affinity. The most promising molecule – PZM21 – did not recruit β -2-arrestin, while being single-digit nanomolar activator of G_i protein signaling. Animal testing demonstrated strong analgesia, lack of side effects, and improved therapeutic index¹.

BIASED NEGATIVE ALLOSTERIC MODULATORS OF CXCR3. Using homology modeling I constructed the structural model of CXCR3 based on X-ray structure of CXCR4 and docked known allosteric modulators (ChEMBL) into the model. This helped to identify amino acid residues that regulate small molecule binding and downstream signaling. Attempts to engage a nucleophilic subpocket via reversible covalent interaction with strategically placed electrophilic boronic acid led to discovery of the first biased negative allosteric modulator or CXCR3 (14, Fig. 3).²

Optimization of physico-chemical properties (cLogP and PSA) via introduction of flexible side chain led to identification of the negative allosteric modulator with the opposite signaling bias (**1b**, Fig. 3). These biased modulators exhibited opposite probe dependence – they preferentially downregulated activation of CXCR3 by CXCL10 or CXCL11 chemokines.³ Affinities of these small-molecule allosteric modulators were similar (double-digit nM) as measured in radioligand RAMX3 displacement assay.⁴

Structural model of CXCR3 that I prepared was subsequently used as is in virtual screening for identification of completely novel chemotypes of specific and dual CXCR3 and CXCR4 allosteric modulators.⁵

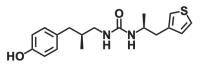


Figure 1: Structure of PZM21

¹ A. Manglik *et al. Nature*, 537(7619): 185–190, 2016

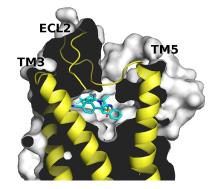


Figure 2: Homology model of CXCR₃ with docked small molecule allosteric modulator.

- ² V. Bernat *et al. ACS Chem. Biol.*, 9(11): 2664–2677, 2014
- ³ V. Bernat *et al. ChemMedChem*, 10(3): 566–574, 2015
- ⁴ V. Bernat *et al. ChemMedChem*, 7(8): 1481–1489, 2012
- ⁵ D. Schmidt *et al. ACS Chem. Biol.*, 10 (3):715–724, 2015

Chemical biology of non-coding RNA

Non-coding RNA folds into 3D-structures that can be targeted by small molecules.⁶ My research in chemical biology of RNA was about identification of new chemotypes targeting pathogenic RNA structures and development of chemical tools for studying their role in cellular markers of disease.

ELLIPTICINES AS LIGANDS FOR $r(G_4C_2)^{EXT}$. Building upon the previous work in the Disney lab, I developed fluorescent reporter 1a-TOO which served as a tool for quick SAR interrogation in vitro and led to discovery of better RNA binders from the same chemotype⁷. Key insight was that permanent positive charge was not required for potent binding to RNA and activity in cellular assays. This dramatically improved blood-brain barrier penetration in mice.

CHEMICAL BIOLOGY OF $r(CUG)^{EXP}$ AND $r(CCUG)^{EXP}$. To study the role of pathogenic non-coding RNAs in myotonic dystrophy I developed series of conjugates that disrupted interaction of r(CUG)^{exp} with proteins (GC-2H-CG, Fig. 6A) or targeted the pathogenic r(CCUG)^{exp} for degradation by latent cellular nuclease RNAse L (Fig. 6B). Further research in Disney lab built upon this work to develop microRNA degrading tool compounds.

NIR fluorescent glucose sensors

At Profusa, I developed near-infrared (NIR) fluorescent sensors for continuous sensing of glucose in vivo. The design of glucose sensors was based on known bis-boronic anthracene module that was incorporated into biocompatible hydrogel matrix. From a library of 120 dyes that I prepared⁸, I could explore structure-activity relationships. Based on that I was able to rationally improve the performance of the company's prototype dye up to 5-fold. Finally, I scaled the syn-

Figure 3: Small molecule allosteric modulators of CXCR3

⁶ V. Bernat and M. D. Disney. Neuron, 87 (1):28-46, 2015

Figure 4: Fluorescent reporter for r(G₄C₂)^{exp} binding assay ⁷ Z. F. Wang et al. Cell Chem. Biol., 26(2): 179-190.e12, 2019

Figure 5: Example of glucosesensing polymerizable NIR dye.

8 S. Gamsey et al. US Patent 10,494,385; US Pat. App. 15/855,555; US Patent App. 16/455,038, 2019

thesis of the lead dye to produce 1.8 g of material for clinical trials in humans.

DNA bioconjugation for next generation sequencing

At Quantapore, I established workflow for immobilization of DNA on flat surfaces and nanoparticles. Via rational optimization of conjugation conditions and surface chemistry I have enabled proof of principle for Quantapore's proprietary sequencing platform.

References

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A. Manglik et al. Nature, 537(7619):185-190, 2016.

D. Schmidt et al. ACS Chem. Biol., 10(3):715-724, 2015.

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Figure 6: A) Structure of peptoid-PNA-Hoechst conjugate for disruption of r(CUG)exp - MBNL1 interaction; B) Structure of peptoid-[2-5]A₄-Kanamycin conjugate for targeting r(CCUG)^{exp} for degradation by RNAse L