

The Research Group

Organic Chemistry

has the honor to invite you to the public defense of the PhD thesis of

## Morgane Mannes

to obtain the degree of Doctor of Bioengineering Sciences

Title of the PhD thesis:  
Discovery, design and synthesis of G protein peptidomimetics

Promotors:

**Prof. dr. Steven Ballet (VUB)**

**Dr. Toon Laeremans (Confo Therapeutics)**

**Prof. dr. Charlotte Martin (VUB)**

The defense will take place on  
**Friday, April 29, 2022 at 17h00**

in auditorium D0.08 on the Campus Humanities,  
Sciences and Engineering of the Vrije  
Universiteit Brussel, Pleinlaan 2, 1050 Elsenne,  
and will be followed by a reception.

### Members of the jury

Prof. dr. Jan Steyaert (VUB, chair)

Dr. Oleksandr Volkov (VUB, secretary)

Prof. dr. Sophie Hernot (VUB)

Prof. dr. Jean-Louis Banères (University of  
Montpellier)

Prof. dr. Christa Müller (University of Bonn)

Prof. dr. Antonios Kolokouris (National and  
Kapodistrian University of Athens)

### Curriculum vitae

Morgane Mannes graduated as Master in Bioengineering Sciences - Chemistry and Bioprocess Technology - at the Vrije Universiteit Brussel (VUB) in 2017. Next, she obtained an Innoviris scholarship to start a semi-industrial PhD in the Research Group of Organic Chemistry (ORGC) of the VUB in collaboration with Confo Therapeutics, a drug discovery company.

During her PhD, she focused on G protein peptidomimetics to stabilize G protein-coupled receptors in an active signaling conformation. Her research has led to a patent application on G<sub>s</sub> mimetics, a review and a first author publication in one of the most influential chemistry journals (*Angew. Chem. Int. Ed.*). Besides teaching practicals and exercise sessions, she supervised a Master and Bachelor thesis student.

### Abstract of the PhD research

G protein-coupled receptors (GPCRs) represent an important family of membrane receptors that play a central role in modern medicine. Unfortunately, it is quite challenging to gain insights into the structure and functioning of GPCRs due to their intrinsic dynamics and high instability when extracted from the cell membrane. Along these lines, conformational promiscuity hampers the full therapeutic exploitation of GPCRs, since the largest population of the receptors will adopt a basal conformation, subsequently challenging screens for agonist drug discovery programs. In past years, Confbodies (Cb) were discovered to lock GPCRs in active signaling states and these have been recognized to be of extreme value in the search of specific orthosteric GPCR agonist ligands. Yet, even though Cbs are crucial tools for structural biology and drug discovery, their use shows some limitations in terms of development costs, including the discovery and purification of the Cbs.

To overcome the drawbacks associated to Cbs, a set of peptidomimetic ligands stabilizing the active state conformation of the  $\beta_2$  adrenergic receptor ( $\beta_2$ AR) and the dopamine 1 receptor (D1R), both signaling through the G<sub>s</sub> protein, was developed. Of note, the developed peptidomimetics equalled the stabilization capacity of Cb80, a reported Cb of  $\beta_2$ AR. During fragment-based screening efforts, the (un)constrained peptide analogues of the  $\alpha_5$  helix in G<sub>s</sub> proteins allowed to, gratifyingly, identify agonism pre-imprinted fragments for the examined GPCRs. Having the proof-of-concept that the designed G<sub>s</sub> peptidomimetics act as a generic toolset to stabilize active conformations of G<sub>s</sub> GPCRs, their fusion to the receptors represents a promising approach to standardize ligand screenings for all G<sub>s</sub>-mediated receptors. To this end, different methods were considered to covalently conjugate a G<sub>s</sub> peptidomimetic to the  $\beta_2$ AR. However, despite the promising results obtained for the covalent linkage of the G<sub>s</sub> peptidomimetics to the membrane-embedded  $\beta_2$ AR, application of the envisaged receptor-peptidomimetics conjugates in drug discovery programs needs further research.