



Hiver 2018  
**Conférence**  
au Département de chimie  
présentée conjointement avec  
PROTEO

CONFÉRENCIER

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DATE

Lundi, 12 février 2018

TITRE

## **Discovery, design and development of human amylase inhibitors: from nM to pM**

RÉSUMÉ

Carbohydrates play important roles in biological systems, not only in the form of energy storage materials such as starch, but also as «recognition elements» on cell surfaces. The degradation of such sugar structures is achieved using enzymes known as glycoside hydrolases (glycosidases). Specific enzyme inhibitors are not only useful tools for understanding enzyme mechanisms, but also can play important roles as therapeutics if inhibition suppresses unwanted reactions. In this talk I shall discuss our efforts to discover new inhibitors of human pancreatic alpha-amylase (HPA) that have potential in the control of blood sugar levels, thus in the treatment of diabetes and possibly obesity.

High-throughput screening of natural product extract libraries from terrestrial and marine sources, in conjunction with my colleague Ray Andersen, has yielded two new classes of potent ( $K_i = 8 \text{ nM}$  and  $10 \text{ pM}$ ) inhibitors of human  $\alpha$ -amylase. The first compound, Montbretin A, was isolated from the beautiful Crocosmia plant shown below, and its inhibition characterized through degradation studies and through X-ray crystallographic analysis of its complex with HPA in conjunction with Gary Brayer<sup>1</sup>. Subsequent animal studies revealed, good control of blood glucose levels in diabetic rats when administered orally<sup>2</sup>. The second screen revealed helianthamide as a highly potent peptidic inhibitor of HPA, derived from a sea anemone<sup>3</sup>. Structural studies with these inhibitors reveal a new paradigm for glycosidase inhibition in which pairs of aromatic moieties joined via a short linker provide the key inhibitory motif. Synthesis of simple mimics of this core structure has yielded inhibitors with  $K_i$  values ranging down to  $50 \text{ nM}$  to date, while screening of very large libraries of cyclic peptides through ribosome display methodologies in conjunction with the Suga group has produced two sets of inhibitors with  $K_i$  values between  $1$  and  $10 \text{ nM}$ , with similar modes of inhibition<sup>4</sup>.

1. Williams, L. et al (2015) **Nature Chemical Biology** 11, 691-696.

2. Yuen, V.G. et al (2016) **Mol. Cell. Biochem.** 411, 373-381.

3. Tysoe et al (2016) **ACS Central Science** 2, 154-161.

4. Jongkees et al (2017) **Cell Chemical Biology** 24, 381-390.

La conférence aura lieu à **14h30 local 2830** du **Pav. Alexandre-Vachon**  
*Cordiale invitation à toutes et à tous !*

Professeur hôte : **Prof. Denis Giguère**

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