

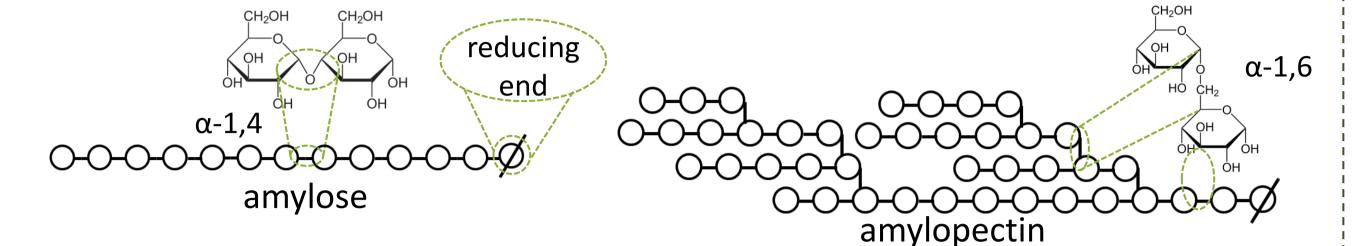
# Enzymatic fingerprinting of Isomalto/Malto-polysaccharides

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## Introduction

Starch consists of amylose & amylopectin, a-linked glucose polymers

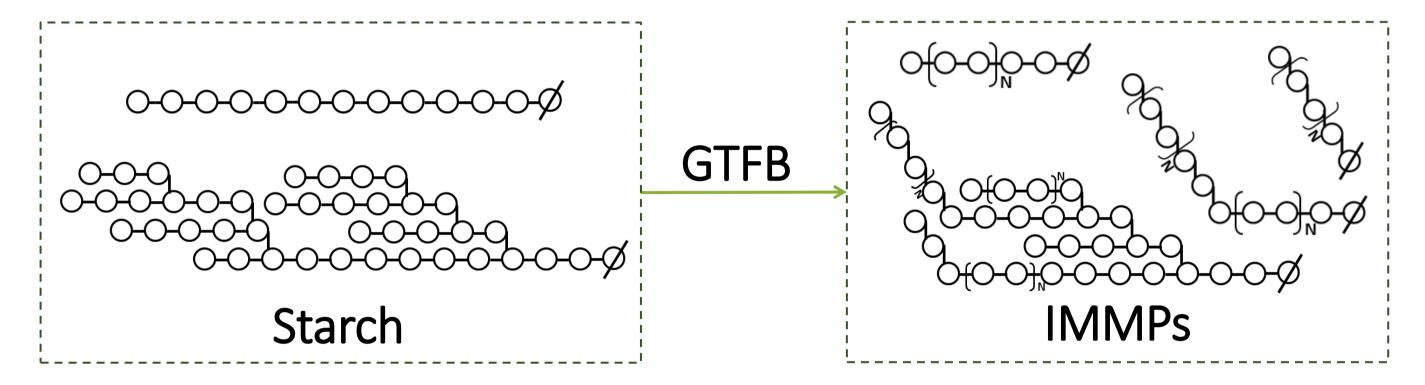


#### Results

In order to determine the presence of linear a-1,4 or linear a-1,6 linked chains in the IMMP mixture. HPAEC was used to compare a-1,4 and a-1,6 linked oligomer standards to an IMMP mixture containing

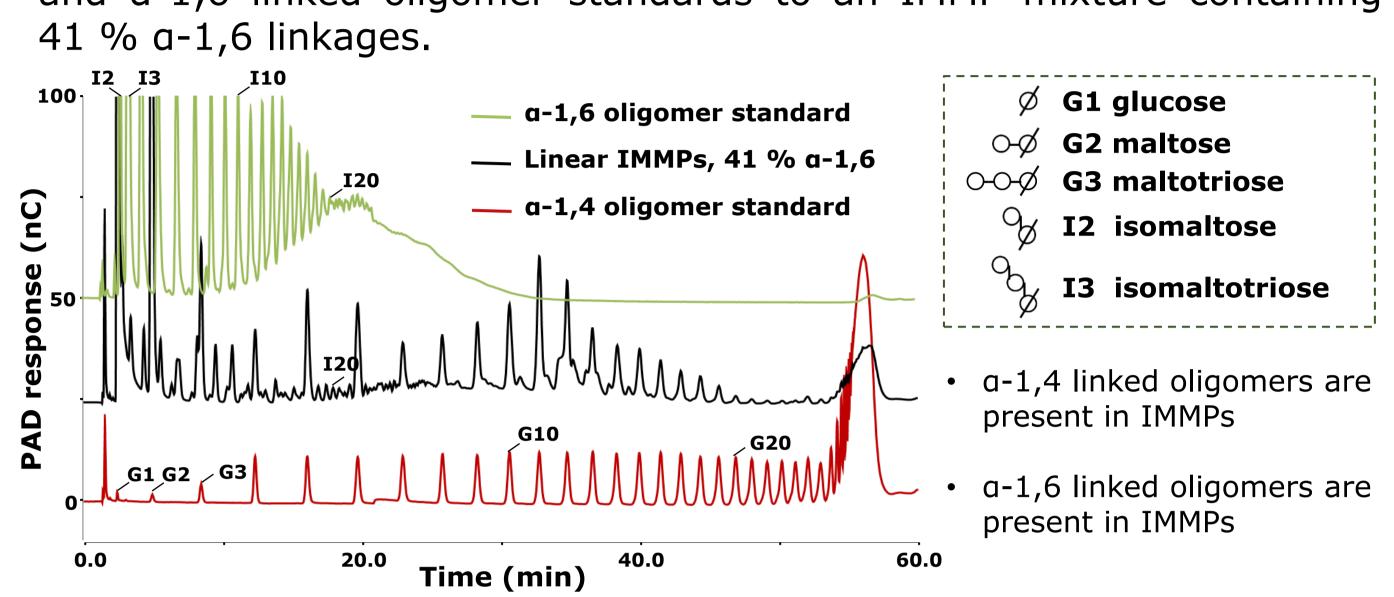
Starch can be modified in different ways to increase its technofunctional properties. The enzymatic modification of starch with 4,6-a-glucanotransferase- $\Delta N$  (GTFB) results in Isomalto/Maltopolysaccharides (IMMPs). IMMPs are molecules built of a-1,4 linked glucose chains, connected to a-1,6 linked glucose chains (Leemhuis et al., 2014).

Graphical representation of enzymatic modification of starch by 4,6-aglucanotransferase- $\Delta N$ , producing Isomalto/Malto-polysaccharides



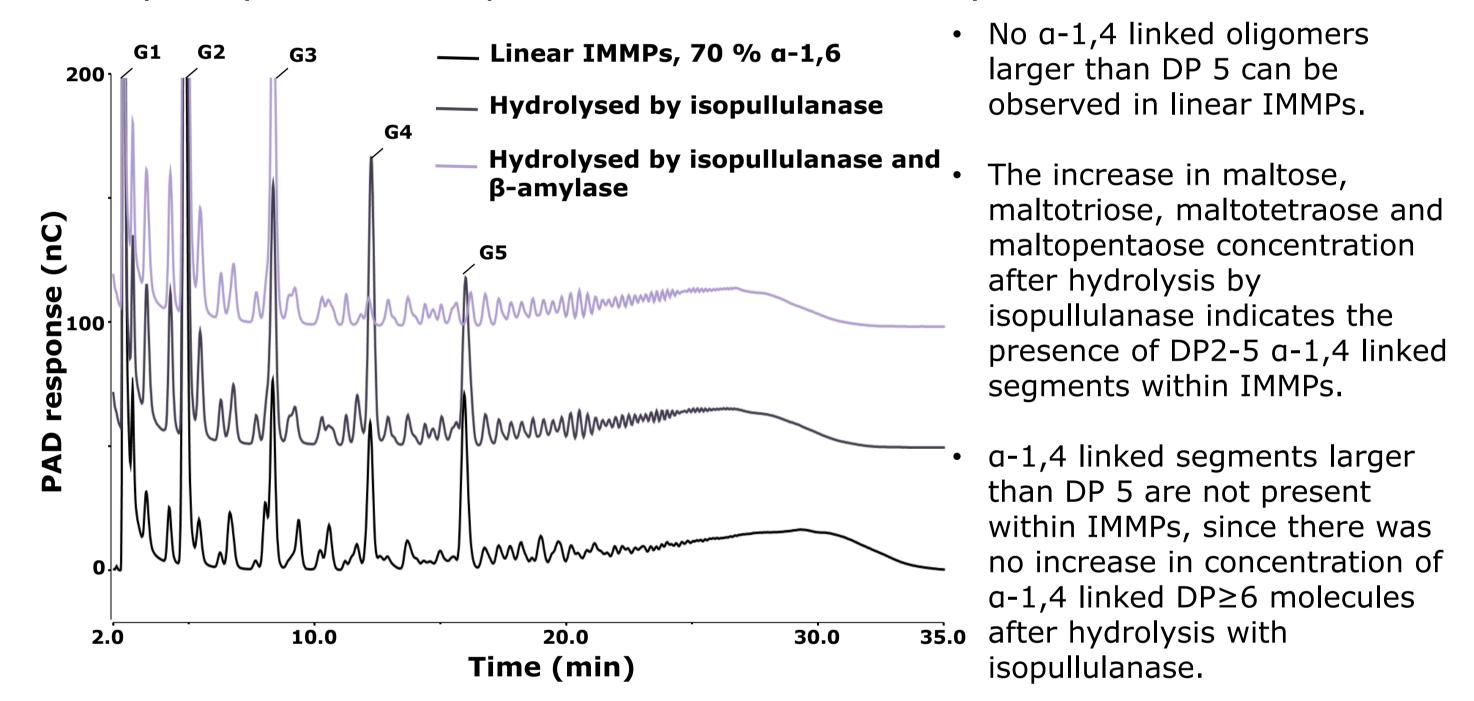
### **Objective**

"To develop an enzymatic fingerprinting method, capable to characterise linear Isomalto/Malto-polysaccharides"



The presence of completely linear a-1,6 linked oligomers indicates that GTFB is able to use glucose as an acceptor.

In order to determine the length of the a-1,4 linked chains within the IMMPs, a linear IMMP (70 % a-1,6) was incubated with isopullulanase or  $\beta$ -amylase and isopullulanase simultaneously.



#### Approach

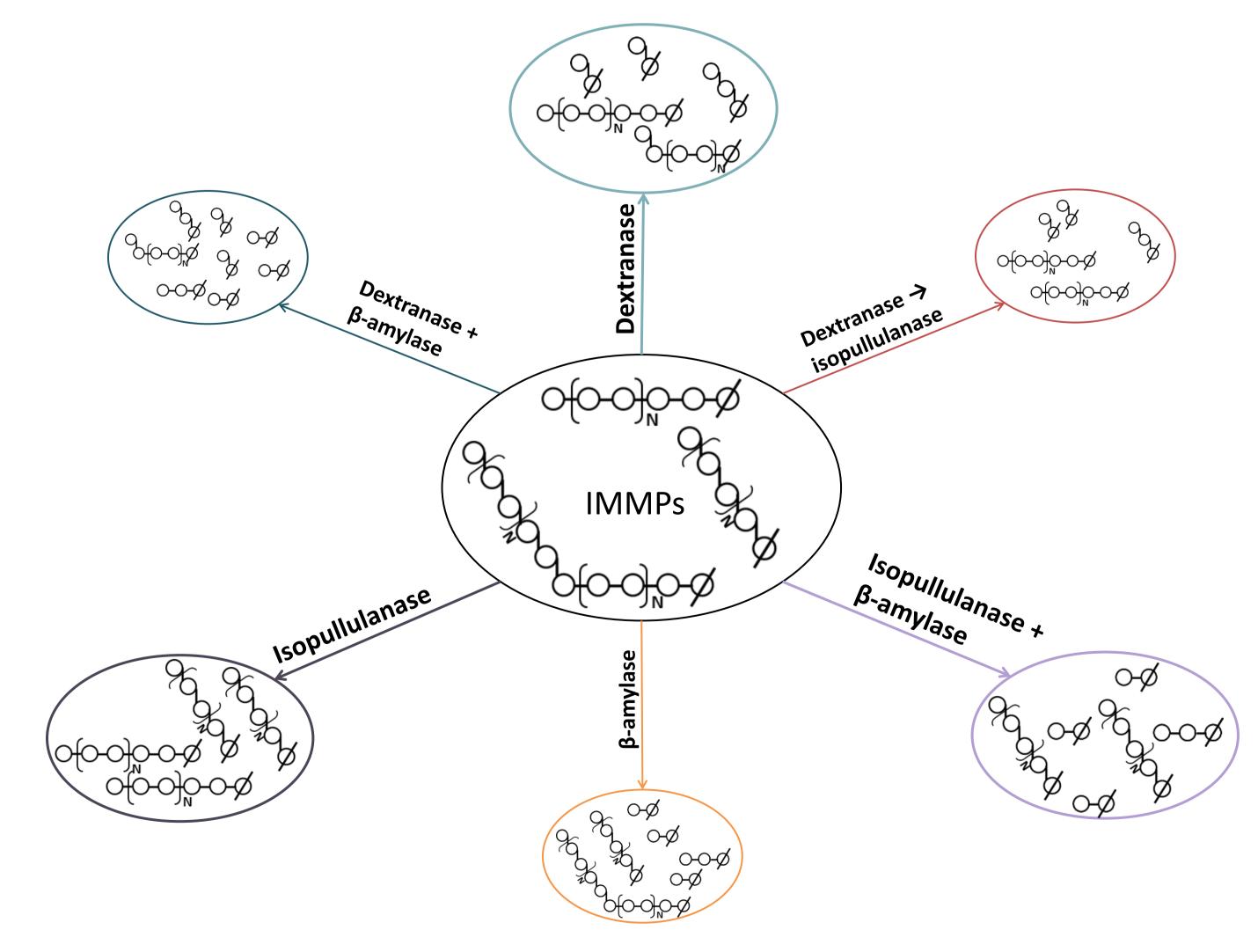
In order to characterise linear IMMPs three enzymes were used:

- $\beta$ -amylase for hydrolysis of all a-1,4 linkages from the nonreducing end
- Dextranase for the endo-hydrolysis of all a-1,6 linkages
- Isopullulanase to split hybrid a-1,4/a-1,6 linked molecules

When using these enzymes in specific orders structural information can be obtained, as shown in the figure below.

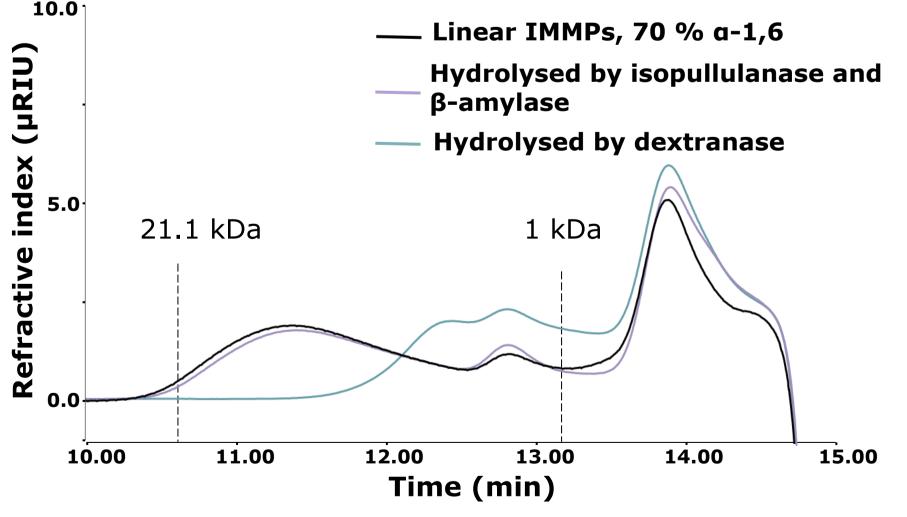
#### Characterisation of linear IMMPs with $\beta$ -amylase, dextranase and Isopullulanase.

Enzymes with a "+" are incubated simultaneously, enzymes with a " $\rightarrow$ " are incubated after each other.



GTFB uses a-1,4 linked material  $\geq$  DP 6 as donor substrate and it uses a-1,4linked material < DP 6 as acceptor substrate.

In order to determine the length of the a-1,6 linked chains within IMMPs, a linear IMMP (70 % a-1,6) was incubated with dextranase or β-amylase and isopullulanase simultaneously. HPSEC was used to analyse these samples in the high Mw range.



- After hydrolysis by isopullulanase and  $\beta$ -amylase a very slight change in Mw distribution can be observed, indicating that no long linear a-1,4 linked chains were present
- After hydrolysis by dextranase a decrease in Mw distribution is shown, indicating that many linear a-1,6 linked chains were present of around 15 kDa

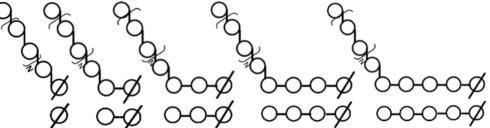
#### GTFB is able to produce IMMPs with a-1,6 linked glucose chains up to 15 kDa.

#### Conclusions

The enzymatic fingerprinting method is able to characterise IMMPs.

GTFB produces:

Fully linear a-1,6 linked chains



- IMMPs with a-1,4 linked glucose segments up to DP 5
- IMMPs with a-1,6 linked glucose segments up to 15 kDa

## Acknowledgement

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