

Single glycans shape viscoelastic properties of a mucin-inspired peptide hydrogel

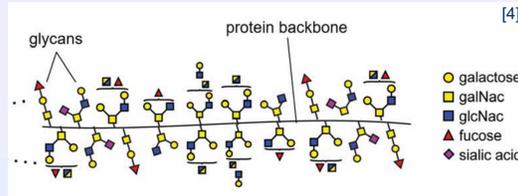
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Introduction

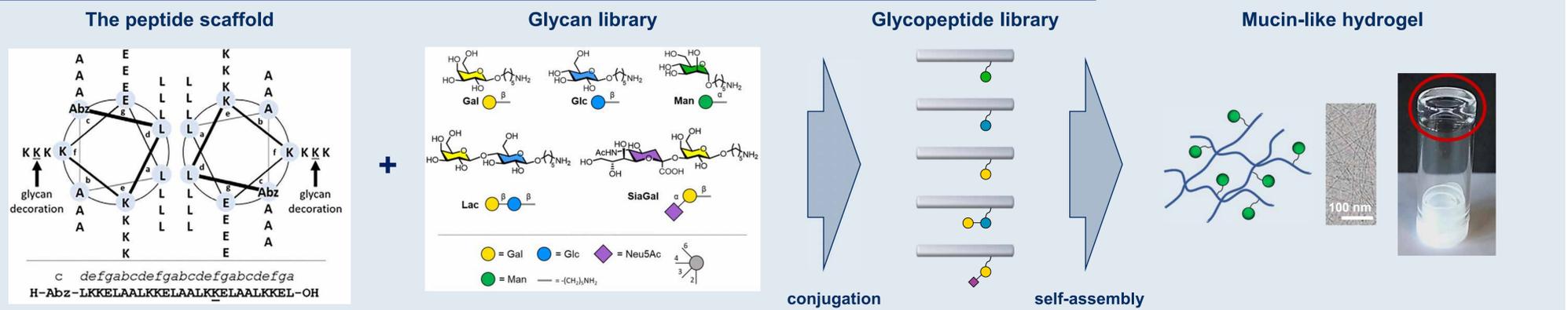
The main component of mucus are mucins, highly glycosylated proteins.^[1] Building relevant experimental models for mucins is hindered by their vast size, glycan content and structural complexity.^[2] Biologic mucus samples differ strongly depending on the donor.^[3]



Aim

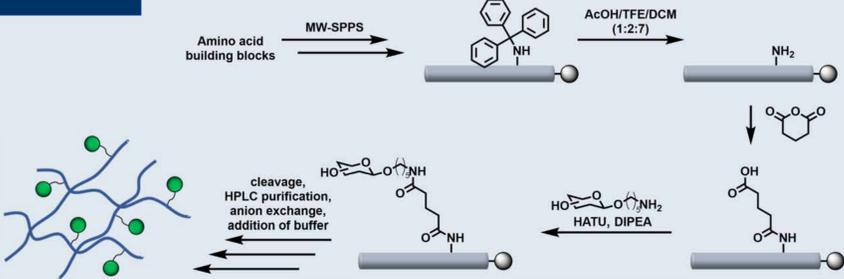
- Establish a simple and straightforward model for some key properties of mucus
- Systematically study the effect of glycan decoration on structure and viscoelastic properties
- Establish structure-property relations
- Identify key features for interactions with pathogens

Tool: A self-assembling peptide hydrogel scaffold for glycan-presentation

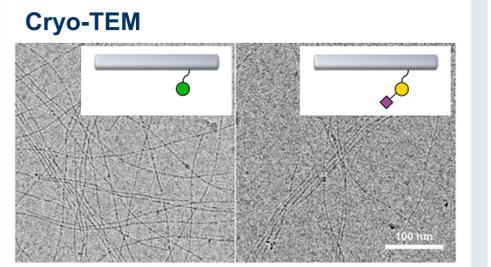
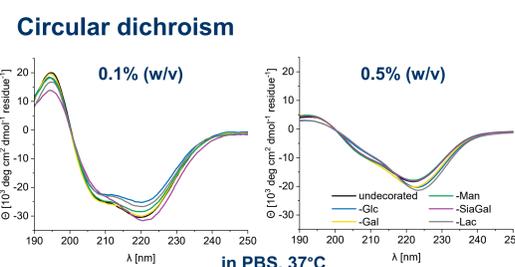


Results

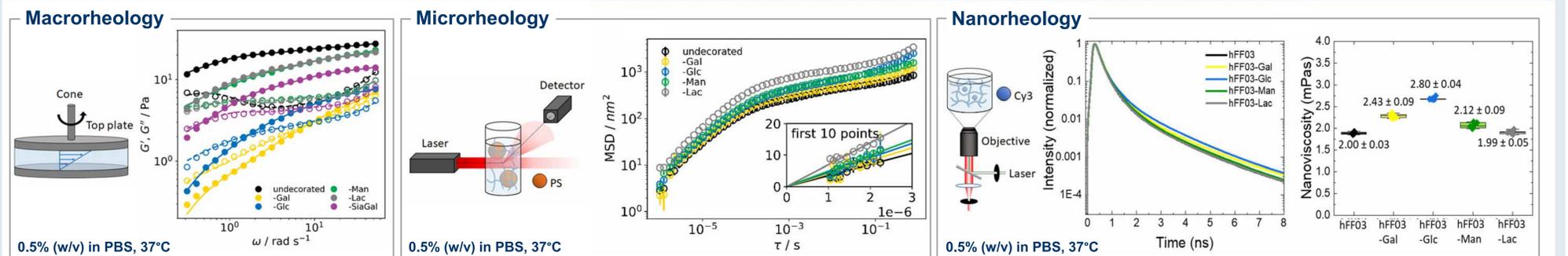
Synthesis



Structural properties

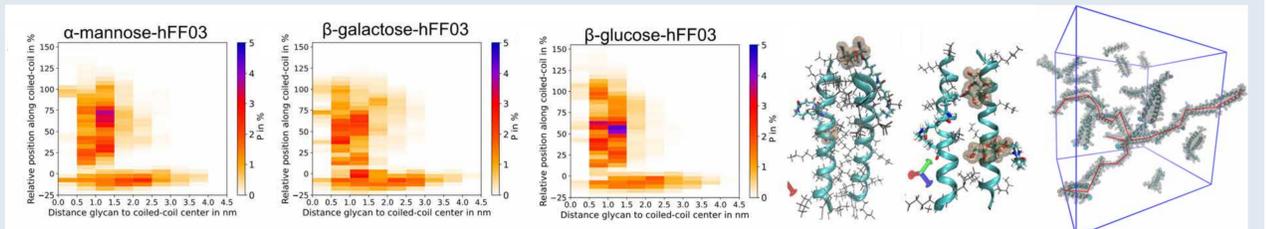


Mechanical properties



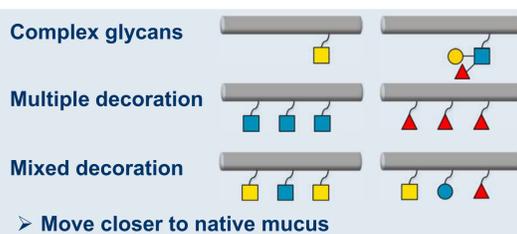
Understanding the differences

- Hydrogel structure formation robust regardless of glycan
- Differences in viscoelastic behaviour depending on the glycan
- MD suggests importance of dynamic processes and the linker



Outlook

Library expansion

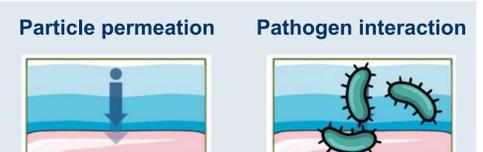


Determine the influence of the linker

➤ "Long" vs "short"

Interactions with surroundings

➤ Identify key features for interactions with pathogens



References:

- R. Bansil, B. S. Turner, *Advanced Drug Delivery Reviews*, **2018**.
- C. E. Wagner, M. Krupkin, K. B. Smith-Dupont, C. M. Wu, N. A. Bustos, J. Witten, K. Ribbeck, *Biomacromolecules*, **2023**.
- S. K. Lai, Y.-Y. Wang, D. Wirtz, J. Hanes, *Advanced Drug Delivery Reviews*, **2009**.
- B.X. Wang, C.M. Wu, K. Ribbeck, *The FEBS Journal*, **2021**.
- K. S. Hellmund, B. von Lospichl, C. Böttcher, K. Ludwig, U. Keiderling, L. Noirez, A. Weiß, D. J. Mikolajczak, M. Gradzielski, B. Kokschi, *Peptide Science*, **2021**.
- E. Zacco, J. Hütter, J. L. Heier, J. Mortier, P. H. Seeberger, B. Lepenies, B. Kokschi, *ACS Chemical Biology*, **2015**.
- J. Proksch, M. C. S. Dal Colle, F. Heinz, R. F. Schmidt, J. Gottwald, M. Delbianco, B. G. Keller, M. Gradzielski, U. Alexiev, B. Kokschi, *Journal of Peptide Science*, **2024**.
- F. Heinz, J. Proksch, R.F. Schmidt, M. Gradzielski, B. Kokschi & B. G. Keller, *Biomacromolecules*, **2024**.
- R. Bej & R. Haag, *Journal of the American Chemical Society*, **2022**.

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