

Using nanopore single-molecule sensing to identify glycans

b а С Carrier Ar unit Lac-DPE-6SL . Calindodia Composition Dwell time (ms) Cel-DPE-6SL -----Configuration 1 9 6 DPE: -3 0 and the states Mal-DPE-6SL 0.2 0.4 0.6 0.8 1.0 $I_{\rm b}/I_{\rm 0}$ d е Counts 0 0 Counts 0.348 0.323 100 LeX-DPE-6SL LeA-DPE-6SL ໌ ຮີມ 10 (sm 10

Nanopore detection of neutral glycans. a Schematic of the derivatization strategy of neutral glycan (Lac, for example). The complex label contains 6SL as a carrier and diphenyl ether (DPE) as the aromatic (Ar) unit. b Three complex label tagged disaccharide epimers (Lac-DPE-6SL, Cel-DPE-6SL, and Mal-DPE-6SL) that differ only in the C4 or C1 stereochemistry of the terminal monosaccharide. c Dot density maps of nanopore data of three disaccharide derivatives. Scatter plots and the corresponding I_b/I_0 and dwell time distributions of two complex label tagged branched trisaccharide isomers (LeA-DPE-6SL and LeX-DPE-6SL)

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(d, e) and tetrasaccharide isomers (LNnT-DPE-6SL and LNT-DPE-6SL) (f, g). All measurements were done in a 10 mM Tris-HCl buffer containing 1 mM EDTA and 1 M KCl with pH 8.0 at +100 mV voltage. All nanopore data were recorded using a 250 kHz sampling rate with a 5 kHz low-pass filtering. Each scatter plot contains at least 9,000 events. Credit: *Nature Communications* (2023). DOI: 10.1038/s41467-023-37348-5

Glycans perform varied and crucial functions in numerous cellular activities. The diverse roles of glycans are matched by their highly complex structures, which derive from differences in composition, branching, regio- and stereochemistry, and modification. This incomparable structural diversity is challenging to the structural analysis of glycans.

Recently, a joint research group led by Prof. Qing Guangyan and Prof. Liang Xinmiao from the Dalian Institute of Chemical Physics (DICP) of the Chinese Academy of Sciences (CAS) has developed a <u>glycan</u> identification method based on <u>nanopore</u> single-molecule sensing through a glycan derivatization strategy. The study was published in *Nature Communications* on March 28.

Identifying and sequencing glycans using nanopore single-molecule techniques has sparked interest; however, it has achieved little progress over the past dozen years. Only a handful of cases that focused on either high molecular weight polysaccharides or some monosaccharides were reported.

For smaller but structurally more diverse glycans with greater biological significance, single molecule detection with nanopores has not yet been achieved, largely because the fast passage of glycans through a nanopore cannot be sensed due to the <u>small size</u> and weak affinity of the glycan



with the nanopore.

To address the challenge, the researchers introduced a derivatization strategy by linking an aromatic-type tag group to small glycans via a highefficiency and facile reductive amination reaction. The resulting tagged glycan was sensed with a wild-type aerolysin nanopore by presenting strong nanopore blockage signals.

The researchers obtained a scatter plot based on blockage current and dwell time as the fingerprint map by processing the nanopore singlemolecule blockage events. They identified different glycan isomers, glycans with varying lengths, and branched simple glycans.

Moreover, they revealed that multiple cation- π interactions between the aromatic tag of glycan with K238 residues of the nanopore interface retarded the translocation of the tagged glycan and contributed to sensing.

"This study pushes the boundary of nanopore sensing beyond its traditional focus on nucleic acid and protein, and activates its power in the glycomics and glycoscience field, which might pave the way towards nanopore glycan sequencing," said Prof. Qing.

More information: Minmin Li et al, Identification of tagged glycans with a protein nanopore, *Nature Communications* (2023). DOI: 10.1038/s41467-023-37348-5

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