VaporSPOT: Parallel Synthesis of Oligosaccharides on Membranes

Alexandra Tsouka,† Pietro Dallab Bernardino,† Marco Mende, Eric T. Sletten, Sabrina Leichnitz, Klaus Bienert, Kim Le Mai Hoang, Peter H. Seeberger, and Felix F. Loeffler*†

Cite This: J. Am. Chem. Soc. 2022, 144, 19832−19837

ABSTRACT: Automated chemical synthesis has revolutionized synthetic access to biopolymers in terms of simplicity and speed. While automated oligosaccharide synthesis has become faster and more versatile, the parallel synthesis of oligosaccharides is not yet possible. Here, a chemical vapor glycosylation strategy (VaporSPOT) is described that enables the simultaneous synthesis of oligosaccharides on a cellulose membrane solid support. Different linkers allow for flexible and straightforward cleavage, purification, and characterization of the target oligosaccharides. This method is the basis for the development of parallel automated glycan synthesis platforms.

INTRODUCTION

Oligosaccharides are the most abundant biopolymers in nature and play a fundamental role in many biological functions. Despite their relevance in many processes of life, their role is still not sufficiently understood.1−3 Their structural heterogeneity, complexity, and diversity often make their isolation from natural sources laborious and their synthesis a cumbersome process. Therefore, different automated platforms for enzymatic,5−7 chemoenzymatic,6,7 and chemical oligosaccharide1−12 synthesis have been developed, giving access to complex and biologically valuable structures. In the last two decades, the development of automated glycan assembly (AGA) has enabled the successful synthesis of defined and complex synthetic oligosaccharide libraries of biological and medical interest,6,15−21 as well as very long polysaccharides including complex branching, up to 100-mers.22,23 Currently, AGA is used to prepare one single oligosaccharide at a time. Parallel oligosaccharide synthesis would be more cost- and time-efficient. Compared to the well-established parallel peptide and oligonucleotide synthesis methods,24,25 the parallel synthesis of oligosaccharides remains a major challenge. Early work by the Kahne group relied mainly on acylations to generate diversity and study the interaction of disaccharides with carbohydrate-binding proteins.26 Mrksich and colleagues demonstrated the on-chip synthesis of a collection of different disaccharides and subsequent enzymatic modification.27,28 Recently, Heo et al. reported their on-chip sequential enzymatic glycosylation strategy for the synthesis of several Globo H-related oligosaccharides on a DNA linker, enabling characterization and purification of the synthetic structures.28

None of these methods were advanced beyond the proof-of-principle stage, since the chemical synthesis requires inert and temperature-controlled glycosylation conditions, making automation a challenging process. Enzymatic synthesis overcomes these issues, but still, only a limited number of glycosyltransferases is available.29

Here, we present the VaporSPOT method for parallel oligosaccharide synthesis that overcomes these limitations. SPOT synthesis, initially developed by Frank et al.,30 can be performed manually or automated31−34 and is commonly used to simultaneously generate peptide, small-molecule, or glycopeptide libraries.35−37 The original SPOT method follows the 9-fluorenylmethoxycarbonyl (Fmoc)-based solid-phase synthesis protocol under ambient conditions, using cellulose membranes as a solid support. Cleavage of the products can be achieved after treatment with strong bases or acids, giving access to a library of soluble and/or cellulose-tethered peptides. However, SPOT synthesis is incompatible with chemical carbohydrate synthesis, since the conditions are neither inert nor temperature-controlled. To solve this problem, we devised a novel method and designed a setup to ensure controlled conditions suitable for glycosylation reactions. With this, we show the parallel synthesis of six different oligosaccharides and up to four residues in length in the micromolar scale (~1 µmol). In contrast to other solid-phase approaches, it is a simple setup that saves time and reagents by parallelization.

Received: July 11, 2022
Published: October 21, 2022
RESULTS AND DISCUSSION

The VaporSPOT process under inert argon conditions was designed for building block delivery at room temperature and subsequent chemical vapor glycosylation at low temperature. The synthesis begins with the functionalization and preparation of the cellulose membrane (Figure 1), which was preloaded with a base-labile mannopyranoside linker (Sections C and E in the Supporting Information), bearing an Fmoc group on the C-6 position, after deprotection, serving as the temporary protecting group on the C-6 position, after deprotection, giving the best glycosylation outcome in comparison to galactoside 10 and glucose 12, due to its higher reactivity (arming/disarming effect) at these temperatures. Moreover, no glycosylation was observed with more reactive BBs 5 and 6 with two electron-donating groups. This is likely due to the currently limited minimum temperature (−15 °C) achievable by the setup.

However, a significantly higher yield was obtained with building block 8, where the Fmoc group on the C-6 position is replaced with the more electron-withdrawing Lev group, yielding 48% of the desired di-mannopyranoside 15 under the exact same conditions.

Next, trimannoside 18 and tetramannoside 19 were synthesized, using mannopyranosides 1, 3, and 4 for chain elongation. While BB 8 resulted in a higher yield, the Lev deprotection would require extensive optimization on the cellulose membrane. Thus, we selected the already established Fmoc deprotection strategy to synthesize longer structures. In the first experiments, partial decomposition of the membrane and reproducibility issues were observed during the synthesis of 18 under the reported glycosylation conditions. Reduction of the activator amount to 4% provided trisaccharide 18, with two electron-donating groups. This is likely due to the higher reactivity (arming/disarming effect) at these temperatures. Moreover, no glycosylation was observed with more reactive BBs 5 and 6 with two electron-donating groups. This is likely due to the currently limited minimum temperature (−15 °C) achievable by the setup.

Figure 1. Schematic representation of VaporSPOT synthesis: (A) reagents and conditions for membrane functionalization: (i) capping 10% Ac2O, 2% MsOH in DCM, rt, 30 min; (ii) 20% piperidine in DMF, rt, 20 min; (iii) attachment of the linker, rt, overnight; and (iv) capping 10% Ac2O, 20% DIPEA in DMF, rt, 30 min. (B) Modules of the VaporSPOT process: (i) preparation of the membrane; (ii) acidic wash of the membrane; (iii) spotting of the building block; (iv) chemical vapor glycosylation; (v) removal of tPG; (vi) deprotection of pPGs and release of oligosaccharides from the solid support; and (vii) purification and characterization of synthesized structures.

of the first building block and then drying under high vacuum. One or several spotted membranes were transferred to the bottom of the custom-built instrument (Figure 2) and cooled to −15 °C. Activation of the glycosyl donor, similar to batch or solid-phase syntheses, was achieved by delivery and condensation of TMSOTf and dichloromethane vapor inside the glycosylation chamber (2 min). Then, the temperature was slowly increased to rt and maintained for 30 min. After completion, the remaining condensate was removed from the glycosylation chamber under high vacuum. The membrane(s) were transferred to a Petri dish and washed with dichloromethane and dimethylformamide. For higher coupling efficiencies with poorly reactive and/or sterically hindered building blocks, coupling can be repeated. Removal of the temporary protecting group with piperidine unmasks the nucleophile for the next synthesis cycle. These steps are repeated until the target structure(s) are formed (Figure 1B, modules ii–v).

Simultaneous deprotection of the ester protecting groups and release of the product(s) from the surface was achieved by sodium methoxide, followed by purification and characterization.

For successful glycosylations, two main parameters had to be optimized: the temperature inside the glycosylation chamber and the amount and concentration of the activator solution to be delivered. Both affect the reactivity of the building blocks and the yield. For the preparation of different oligosaccharides, 14 building blocks (BBs, Figure 3) were either synthesized following established protocols or prepared from commercially available precursors (trichloroacetimidates 3, 5, and 9 and phosphates 6 and 8, see Section B in the Supporting Information). To optimize the coupling conditions, glycosyl donors 1–6 and 8–12 were screened, using 8% TMSOTf in dichloromethane (Figure 4A), to furnish the corresponding dimers 15–17 in different yields (Figure 4B). While perbenzoylated building blocks (1, 9, 11) show relatively similar results, mannose 3, with an Fmoc temporary protecting group on the C-6 position, gave the best glycosylation outcome in comparison to galactoside 10 and glucose 12, due to its higher reactivity (arming/disarming effect) at these temperatures. Moreover, no glycosylation was observed with more reactive BBs 5 and 6 with two electron-donating groups. This is likely due to the currently limited minimum temperature (−15 °C) achievable by the setup.

For the preparation of different oligosaccharides, 14 building blocks (BBs, Figure 3) were either synthesized following established protocols or prepared from commercially available precursors (trichloroacetimidates 3, 5, and 9 and phosphates 6 and 8, see Section B in the Supporting Information). To optimize the coupling conditions, glycosyl donors 1–6 and 8–12 were screened, using 8% TMSOTf in dichloromethane (Figure 4A), to furnish the corresponding dimers 15–17 in different yields (Figure 4B). While perbenzoylated building blocks (1, 9, 11) show relatively similar results, mannose 3, with an Fmoc temporary protecting group on the C-6 position, gave the best glycosylation outcome in comparison to galactoside 10 and glucose 12, due to its higher reactivity (arming/disarming effect) at these temperatures. Moreover, no glycosylation was observed with more reactive BBs 5 and 6 with two electron-donating groups. This is likely due to the currently limited minimum temperature (−15 °C) achievable by the setup.

For successful glycosylations, two main parameters had to be optimized: the temperature inside the glycosylation chamber and the amount and concentration of the activator solution to be delivered. Both affect the reactivity of the building blocks and the yield. For the preparation of different oligosaccharides, 14 building blocks (BBs, Figure 3) were either synthesized following established protocols or prepared from commercially available precursors (trichloroacetimidates 3, 5, and 9 and phosphates 6 and 8, see Section B in the Supporting Information). To optimize the coupling conditions, glycosyl donors 1–6 and 8–12 were screened, using 8% TMSOTf in dichloromethane (Figure 4A), to furnish the corresponding dimers 15–17 in different yields (Figure 4B). While perbenzoylated building blocks (1, 9, 11) show relatively similar results, mannose 3, with an Fmoc temporary protecting group on the C-6 position, gave the best glycosylation outcome in comparison to galactoside 10 and glucose 12, due to its higher reactivity (arming/disarming effect) at these temperatures. Moreover, no glycosylation was observed with more reactive BBs 5 and 6 with two electron-donating groups. This is likely due to the currently limited minimum temperature (−15 °C) achievable by the setup.

However, a significantly higher yield was obtained with building block 8, where the Fmoc group on the C-6 position is replaced with the more electron-withdrawing Lev group, yielding 48% of the desired di-mannopyranoside 15 under the exact same conditions.

Figure 1. Schematic representation of VaporSPOT synthesis: (A) reagents and conditions for membrane functionalization: (i) capping 10% Ac2O, 2% MsOH in DCM, rt, 30 min; (ii) 20% piperidine in DMF, rt, 20 min; (iii) attachment of the linker, rt, overnight; and (iv) capping 10% Ac2O, 20% DIPEA in DMF, rt, 30 min. (B) Modules of the VaporSPOT process: (i) preparation of the membrane; (ii) acidic wash of the membrane; (iii) spotting of the building block; (iv) chemical vapor glycosylation; (v) removal of tPG; (vi) deprotection of pPGs and release of oligosaccharides from the solid support; and (vii) purification and characterization of synthesized structures.
obtained in 34% yield with $\beta$-(1-6) and $\beta$-(1-4) linkage, starting from disaccharide 14 (Scheme 1A). All final structures 15−20 were characterized using mass spectrometry (MS) and nuclear magnetic resonance spectroscopy (NMR). Both purity and anomeric purity of the final structures were analyzed by high-performance liquid chromatography (HPLC) and de-coupled $^1$H−$^{13}$C heteronuclear single quantum coherence spectroscopy (HSQC NMR).

Encouraged by these results, the parallel synthesis of oligosaccharides was investigated (Scheme 1B). To assess, whether diffusion or contamination can occur between different membrane pieces placed in close proximity inside the setup, a photolabile linker was designed and synthesized (Sections D and E in the Supporting Information). With the current set of BBs, the previously used base-labile linker would deliver partially deprotected compounds of mostly indistinguishable molecular weight, since cleavage from the solid support and deprotection of the ester and carbonyl protecting groups occur simultaneously. In contrast, the photolabile linker will deliver protected compounds with distinguishable molecular weight. Using the same experimental setup, reaction time, and the 4% activator solution, six different glycosyl...
SPOT synthesis is widely used for the parallel synthesis of peptides. Here, we developed the VaporSPOT synthesis method for the parallel synthesis of oligosaccharides on cellulose membranes under inert and temperature-controlled chemical vapor conditions. This method offers a flexible and cost-efficient way to rapidly screen the glycosylation outcome of different glycosyl donors in parallel and synthesize oligosaccharides in good purity on the micromolar scale. In a parallel reaction approach, diffusion or contamination between the different spotted glycosyl donors was ruled out. Future optimization of the methodology, including different protecting groups and linkers, will lead to more complex structures with higher yields. For example, the replacement of the Fmoc protecting group with the more electron-withdrawing Lev group on the C-6 position should lead to higher glycosylation yields. Besides, technical improvements (e.g., minimum achievable temperature) may enable more flexible chemical strategies.

Furthermore, other solid supports, such as cross-linked cellulose, polypropylene, or Teflon-patterned membranes, may further improve the spot density, substrate stability, and synthesis yield. The same VaporSPOT approach may be further expanded to high-throughput glycan array synthesis on functionalized glass slides or even beyond glycochemistry, for precisely controlled polymerization or cross-coupling reactions. Together with automated spotting of building blocks, this should enable even higher parallelization. Such oligosaccharide collections are ideal for microarray production, to drastically accelerate the screening of glycan–glycan binding protein interactions.

**ASSOCIATED CONTENT**

Supporting Information

The Supporting Information is available free of charge at https://pubs.acs.org/doi/10.1021/jacs.2c07285.

All experimental details regarding the VaporSPOT method; building block synthesis; and characterization (PDF)

**AUTHOR INFORMATION**

**Corresponding Author**

Felix F. Loeffler — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany; orcid.org/0000-0002-8227-2522; Email: felix.loeffler@mpikg.mpg.de

**Authors**

Alexandra Tsouka — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany; Institute of Chemistry and Biochemistry, Freie Universität Berlin, 14195 Berlin, Germany

Pietro Dallabernardina — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany; Present Address: Aptuit (Verona) Srl, an Evotec company, Via Alessandro Fleming 4, 37135 Verona, Italy

Marco Mende — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany; Present Address: Bachem AG, Hauptstrasse 144, 4416 Bubendorf, Switzerland

Eric T. Sletten — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany; orcid.org/0000-0003-2177-2969

Sabrina Leichnitz — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany; Institute of Chemistry and Biochemistry, Freie Universität Berlin, 14195 Berlin, Germany
Klaus Bienert — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany
Kim Le Mai Hoang — GlycoUniverse GmbH & Co. KGaA, 14476 Potsdam, Germany
Peter H. Seeberger — Department of Biomolecular Systems, Max Planck Institute of Colloids and Interfaces, 14476 Potsdam, Germany; Institute of Chemistry and Biochemistry, Freie Universität Berlin, 14195 Berlin, Germany

Complete contact information is available at: https://pubs.acs.org/10.1021/jacs.2c07285

Author Contributions
A.T. and P.D. contributed equally to this work.

Funding
Open access funded by Max Planck Society.

Notes
The authors declare the following competing financial interest(s): P.I.H.S and K.L.M.H. declare a significant financial interest in GlycoUniverse GmbH & Co. KGaA, the company that commercialized AGA synthesis instruments, building blocks, and other reagents. The other authors declare no competing financial interests. F.F.L., A.T., M.M., and P.H.S. are named on a patent related to the presented technology for parallel glycan synthesis, while all other authors declare no conflict of interest.

■ ACKNOWLEDGMENTS

The authors thank the department of Biomolecular Systems for the (technical) support, as well as the electronics workshop of the MBMF, grant number 13XPS050A), the MPG-FhG cooperation (Glyco3Display), and the Max Planck Society.

■ REFERENCES

(1) Variki, A.; Cummings, R. D.; Esko, J. D.; Stanley, P.; Hart, G. W.; Aebl, M.; Mohnen, D.; Kinoshita, T.; Packer, N. H.; Prestegard, J. H.; Schnaar, R. L.; Seeberger, P. H. Essentials of Glycobiology, 4th ed.; Cold Spring Harbor Laboratory Press, 2022.
(2) Variki, A. Biological Roles of Glycans. Glyobiology 2017, 27, 3–49.
(3) Matsushita, T.; Nagashima, I.; Fumoto, M.; Ohta, T.; Yamada, K.; Shimizu, H.; Hinou, H.; Naruchi, K.; Ito, T.; Kondo, H.; Nishimura, S.-I. Artificial Golgi Apparatus: Globular Protein-like Dendrimer Facilities Fully Automated Enzymatic Glycan Synthesis. J. Am. Chem. Soc. 2010, 132, 16651–16656.
(4) Zhang, J.; Chen, C.; Gadi, M. R.; Gibbons, C.; Guo, Y.; Cao, X.; Edmunds, G.; Wang, S.; Liu, D.; Yu, J.; Wen, L.; Wang, P. G. Machine-Driven Enzymic Oligosaccharide Synthesis by Using a Peptide Synthesizer. Angew. Chem., Int. Ed. 2018, 57, 16638–16642.
(5) Li, T.; Liu, L.; Wei, N.; Yang, J.-Y.; Chapla, D. G.; Moremen, K. W.; Boons, G.-J. An Automated Platform for the Enzyme-Mediated Assembly of Complex Oligosaccharides. Nat. Chem. 2019, 11, 229–236.
(6) Liu, L.; Prudden, A. R.; Capicciotti, C. J.; Bosman, G. P.; Yang, J.-Y.; Chapla, D. G.; Moremen, K. W.; Boons, G.-J. Streamlining the Chemoenzymatic Synthesis of Complex N-Glycans by a Stop and Go Strategy. Nat. Chem. 2019, 11, 161–169.
(7) Wu, C. Y.; Wong, C. H. Automated Programmable One-Pot Synthesis of Glycans. In Glycoscience: Biology and Medicine; Springer: Tokyo, 2015; pp 45–52.
(8) Plante, O. J.; Palmacci, E. R.; Seeberger, P. H. Automated Solid-Phase Synthesis of Oligosaccharides. Science 2001, 291, 1523–1527.
(9) Jaipuri, F. A.; Pohl, N. L. Toward Solution-Phase Automated Iterative Synthesis: Fluorour-Tag Assisted Solution-Phase Synthesis of Linear and Branched Mannose Oligomers. Org. Biomol. Chem. 2008, 6, 2686–2691.
(10) Machida, K.; Hirose, Y.; Fuse, S.; Sugawara, T.; Takahashi, T. Development and Application of a Solution-Phase Automated Synthesizer, ‘ChemKonzert’. Chem. Pharm. Bull. 2010, 58, 87–93.
(11) Ganesh, N. V.; Fujikawa, K.; Tan, Y. H.; Stine, K. J.; Demchenko, A. V. HPLC-Assisted Automated Oligosaccharide Synthesis. Org. Lett. 2012, 14, 3036–3039.
(12) Nakami, T.; Hayashi, R.; Saigusa, Y.; Shimizu, A.; Liu, C.-Y.; Mong, K.-T. J.; Yoshida, J. Automated Solution-Phase Synthesis of Oligosaccharides via Iterative Electrochemical Assembly of Thioglycosides. Org. Lett. 2013, 15, 4520–4523.
(13) Liu, M.; Qin, X.; Ye, X.-S. Glycan Assembly Strategy: From Concept to Application. Chem. Rec. 2021, 21, 3256–3277.
(14) Panza, M.; Pistorio, S. G.; Stine, K. J.; Demchenko, A. V. Automated Chemical Oligosaccharide Synthesis: Novel Approach to Traditional Challenges. Chem. Rev. 2018, 118, 8105–8150.
(15) Guberman, M.; Seeberger, P. H. Automated Glycan Assembly: A Perspective. J. Am. Chem. Soc. 2019, 141, 5581–5592.
(16) Seeberger, P. H. The Logic of Automated Glycan Assembly. Acc. Chem. Res. 2015, 48, 1450–1463.
(17) Hahn, H. S.; Schlegel, M. K.; Hurevich, M.; Eller, S.; Schuhmacher, F.; Hofmann, J.; Pagel, K.; Seeberger, P. H. Automated Glycan Assembly Using the Glyconeer 2.1 Synthesizer. Proc. Natl. Acad. Sci. U.S.A. 2017, 114, E3385–E3389.
(18) Hahn, H. S.; Broecker, F.; Kawasaki, F.; Mietzsch, M.; Heilbronn, R.; Fukuda, M.; Seeberger, P. H. Automated Glycan Assembly of Oligo-N-Acetylactosamine and Keratan Sulfate Polypeptide Chains to Study Virus-Glycan Interactions. Chem 2017, 2, 114–124.
(19) Hahn, H. S.; Hurevich, M.; Seeberger, P. H. Automated Assembly of Oligosaccharides Containing Multiple Cis-Glycosidic Linkages. Nat. Commun. 2016, 7, No. 12482.
(20) Tyrikos-Ergas, T.; Sletten, E. T.; Huang, J.-Y.; Seeberger, P. H.; Delbianco, M. On Resolution of Sulfated Oligosaccharides. Chem. Sci. 2022, 13, 2115–2120.
(21) Dallabernardina, P.; Benazzi, V.; Laman, J. D.; Seeberger, P. H.; Loeffler, F. F. Automated Glycan Assembly of Peptidoglycanc Backbone Fragments. Org. Biomol. Chem. 2021, 19, 9829–9832.
(22) Joseph, A. A.; Pardo-Vargas, A.; Seeberger, P. H. Total Synthesis of Polysaccharides by Automated Glycan Assembly. J. Am. Chem. Soc. 2020, 142, 8561–8564.
(23) Zhu, Y.; Delbianco, M.; Seeberger, P. H. Automated Assembly of Starch and Glycogen Polysaccharides. J. Am. Chem. Soc. 2021, 143, 9758–9768.
(24) Mattes, D. S.; Jung, N.; Weber, L. K.; Bräse, S.; Breitling, F. Miniaturized and Automated Synthesis of Biomolecules—Overview and Perspectives. Adv. Mater. 2019, 31, No. 1806656.
(25) Paris, G.; Heidepriem, J.; Tsouka, A.; Liu, Y.; Mattes, D. S.; Pinzón Martín, S.; Dallabernardina, P.; Mende, M.; Lindner, C.; Waszinek, R.; Bademacher, C.; Seeberger, P. H.; Breitling, F.; Bischoff, F. R.; Wolf, T.; Loeffler, F. F. Automated Laser-Transfer Synthesis of High-Density Microarrays for Infectious Disease Screening. Adv. Mater. 2022, 34, No. 2200359.
(26) Liang, R.; Yan, L.; Loebach, J.; Ge, M.; Uozumi, Y.; Sekanina, K.; Horan, N.; Gildersleeve, J.; Thompson, C.; Smith, A.; Biswas, K.; Still, W. C.; Kahne, D. Parallel Synthesis and Screening of a Solid Phase Carbohydrate Library. Science 1996, 274, 1520–1522.
(27) Ban, L.; Mrksich, M. On-Chip Synthesis and Label-Free Assays of Oligosaccharide Arrays. Angew. Chem., Int. Ed. 2008, 47, 3396–3399.
(28) Heo, H. R.; Joo, K.; Il; Seo, J. H.; Kim, C. S.; Cha, H. J. Glycan Chip Based on Structure-Switchable DNA Linker for on-Chip Biosynthesis of Cancer-Associated Complex Glycans. Nat. Commun. 2021, 12, No. 1395.

(29) Krasnova, L.; Wong, C.-H. Oligosaccharide Synthesis and Translational Innovation. J. Am. Chem. Soc. 2019, 141, 3735–3754.

(30) Frank, R. Spot-Synthesis: An Easy Technique for the Positionally Addressable, Parallel Chemical Synthesis on a Membrane Support. Tetrahedron 1992, 48, 9217–9232.

(31) Ay, B.; Volkmmer, R.; Boisguerin, P. Synthesis of Cleavable Peptides with Authentic C-Terminal: An Application for Fully Automated SPOT Synthesis. Tetrahedron Lett. 2007, 48, 361–364.

(32) Winkler, D. F. H.; Hilpert, K.; Brandt, O.; Hancock, R. E. W. Synthesis of Peptide Arrays Using SPOT-Technology and the CelluSpots-Method. In Methods in Molecular Biology; Humana Press, 2009; Vol. 570, pp 157–174.

(33) Deiss, F.; Yang, Y.; Matochko, W. L.; Derda, R. Heat-Enhanced Peptide Synthesis on Teflon-Patterned Paper. Org. Biomol. Chem. 2016, 14, 5148–5156.

(34) Hilpert, K.; Winkler, D. F. H.; Hancock, R. E. W. Peptide Arrays on Cellulose Support: SPOT Synthesis, a Time and Cost Efficient Method for Synthesis of Large Numbers of Peptides in a Parallel and Addressable Fashion. Nat. Protoc. 2007, 2, 1333–1349.

(35) Blackwell, H. Hitting the SPOT: Small-Molecule Macroarrays Advance Combinatorial Synthesis. Curr. Opin. Chem. Biol. 2006, 10, 203–212.

(36) Winkler, D. F. H.; Campbell, W. D. The Spot Technique: Synthesis and Screening of Peptide Macroarrays on Cellulose Membranes. In Methods in Molecular Biology; Humana Press, 2008; Vol. 494, pp 47–70.

(37) Mehta, A. Y.; Veeraiah, R. K. H.; Dutta, S.; Goth, C. K.; Hanes, M. S.; Gao, C.; Stavenhagen, K.; Kardish, R.; Matsumoto, Y.; Heimbburg-Molinaro, J.; Boyce, M.; Pohl, N. L. B.; Cummings, R. D. Parallel Glyco-SPOT Synthesis of Glycopeptide Libraries. Cell Chem. Biol. 2020, 27, 1207–1219.

(38) Andreana, P. R.; Crich, D. Guidelines for O -Glycoside Formation from First Principles. Methods in Molecular Biology. 2019, 161, 47–56.

(39) Moon, S.; Chatterjee, S.; Seeberger, P. H.; Gilmore, K. Predicting Glycosylation Stereoselectivity Using Machine Learning. Chem. Sci. 2021, 12, 2931–2939.

(40) Chatterjee, S.; Moon, S.; Hentschel, F.; Gilmore, K.; Seeberger, P. H. An Empirical Understanding of the Glycosylation Reaction. J. Am. Chem. Soc. 2018, 140, 11942–11953.

(41) Tuck, O. T.; Sletten, E. T.; Danglad-Flores, J.; Seeberger, P. H. Towards a Systematic Understanding of the Influence of Temperature on Glycosylation Reactions. Angew. Chem., Int. Ed. 2022, 61, No. e202115433.

(42) Bien, F.; Ziegler, T. Chemoenzymatic Synthesis of Glycosylated Enantiomerically Pure 4-Pentene 1,2- and 1,3-Diol Derivatives. Tetrahedron: Asymmetry 1998, 9, 781–790.

(43) Patel, M. K.; Vijayakrishnan, B.; Koeppe, J. R.; Chalker, J. M.; Doores, K. J.; Davis, B. G. Analysis of the Dispersity in Carbohydrate Loading of Synthetic Glycoproteins Using MALDI-TOF Mass Spectrometry. Chem. Commun. 2010, 46, 9119–9121.

(44) Calin, O.; Eller, S.; Seeberger, P. H. Automated Polysaccharide Synthesis: Assembly of a 30mer Mannoside. Angew. Chem., Int. Ed. 2013, 52, 5862–5865.

(45) Danglad-Flores, J.; Leichnitz, S.; Sletten, E. T.; Abramag Joseph, A.; Bienert, K.; Le Mai Hoang, K.; Seeberger, P. H. Microwave-Assisted Automated Glycan Assembly. J. Am. Chem. Soc. 2021, 143, 8893–8901.

(46) Zhang, T.; Li, X.; Song, H.; Yao, S. Ionic Liquid-Assisted Catalysis for Glycosidation of Two Triterpenoid Sapogenins †. New J. Chem. 2019, 43, 16881–16888.

(47) Sandbhór, M. S.; Søya, N.; Albohy, A.; Zheng, R. B.; Cartmell, J.; Bundle, D. R.; Klassen, J. S.; Cairo, C. W. Substrate Recognition of the Membrane-Associated Stalidase NEU3 Requires a Hydrophobic Aglycone. Biochemistry 2011, 50, 6753–6762.