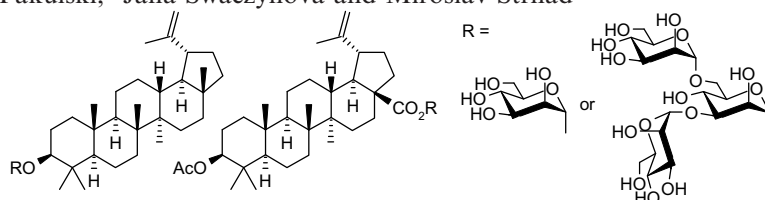


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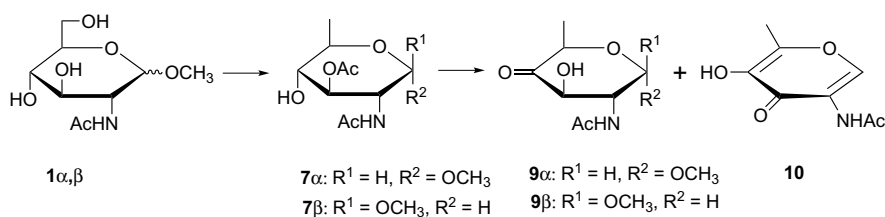


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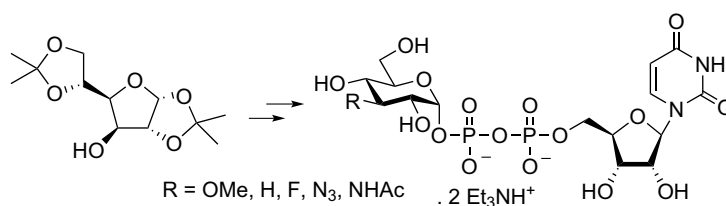
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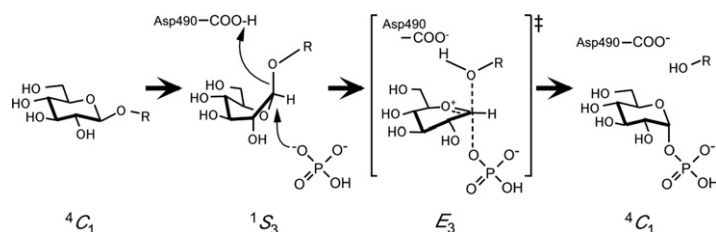
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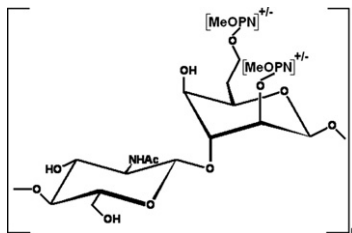
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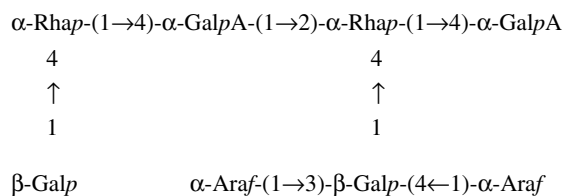
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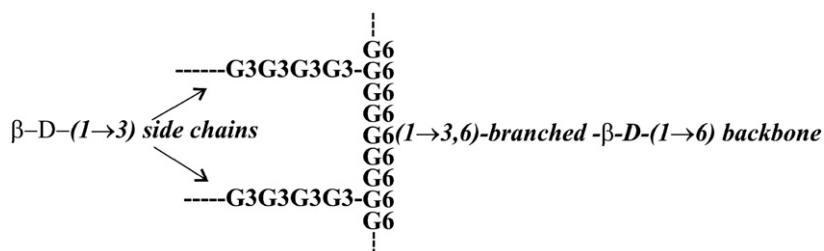
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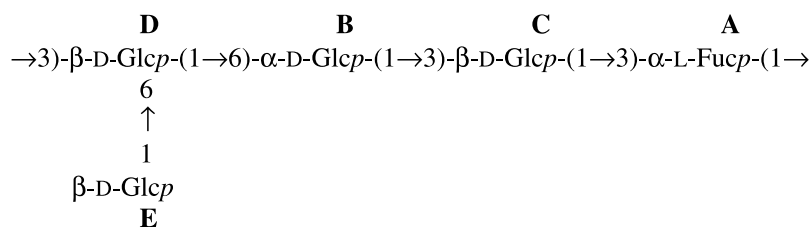
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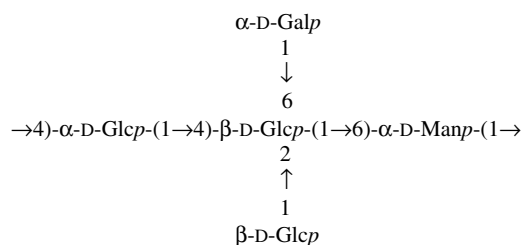
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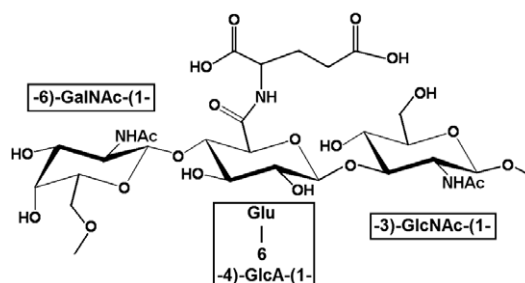
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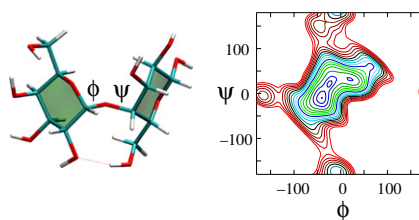
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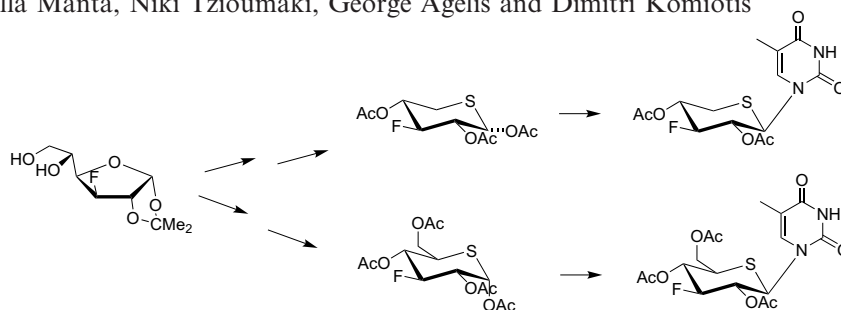


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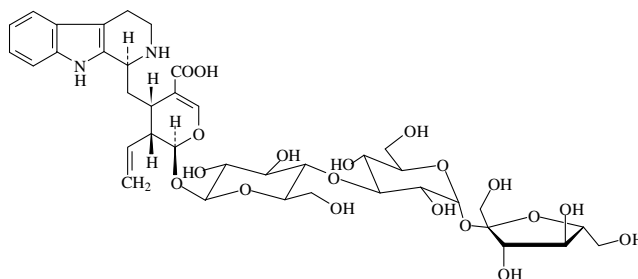
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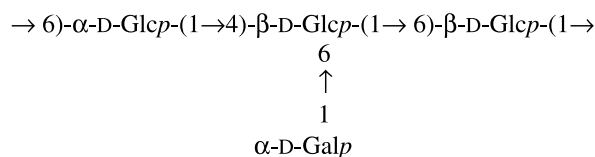
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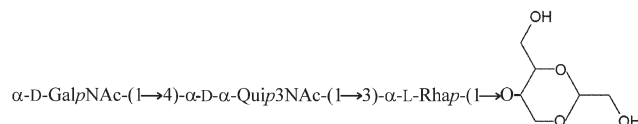
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A ^1H NMR structural-reporter-group concept, based on the analysis of a series of linear $\alpha\text{-D-glucopyranose}$ di- and trisaccharide structures, has been formulated to function as a tool in the structural analysis of bacterial $\alpha\text{-D-glucans}$.

Smith degradation of the O-antigenic polysaccharide of *Salmonella* Dakar: structural studies of the products pp 1120–1125

Jolanta Kumirska,* Janusz Szafrank, Małgorzata Czerwiczka, Marek Gołębiowski, Monika Paszkiewicz, Halina Dziadziuszko, Danuta Kunikowska and Piotr Stepnowski

Two different oligosaccharides: α -D-GalpNAc-(1→4)- α -D-Quip3NAc-(1→3)- α -L-Rhap-(1→2)-threitol and



where Quip3NAc is 3-acetamido-3,6-dideoxyglucose, were obtained from the Smith degradation of the O-polysaccharide isolated from the lipopolysaccharide of *Salmonella* Dakar. This paper presents practical methods for isolating and determining the full structure of such compounds.

*Corresponding author

 Supplementary data available via ScienceDirect

COVER

The graphic represents a molecular dynamics simulation of water density around the disaccharide α -D-Araf-(1→5)- α -D-Araf-OCH₃, highlighting the interglycosidic linkage. The red clouds represent regions where the probability of finding an oxygen atom is high while the gray clouds are for hydrogen atoms. This work is the result of a collaboration in the Alberta Ingenuity Centre for Carbohydrate Science and Department of Chemistry at the University of Alberta between the groups of Pierre-Nicolas Roy and Todd L. Lowary (Castillo, N.; Roy, P. N.; Lowary, T. L. Manuscript in Preparation).

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Abstracted/Indexed in: Chem. Abstr.: Curr. Contents: Phys., Chem. & Earth Sci. Life Sci. Current Awareness in Bio. Sci. (CABS). Science Citation Index. Full texts are incorporated in CJELSEVIER, a file in the Chemical Journals Online database which is available on STN® International. Also covered in the abstract and citation database SCOPUS®. Full text available on ScienceDirect®



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ISSN 0008-6215