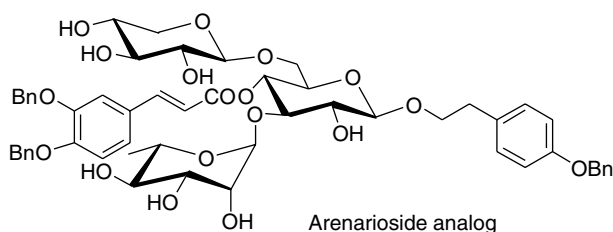


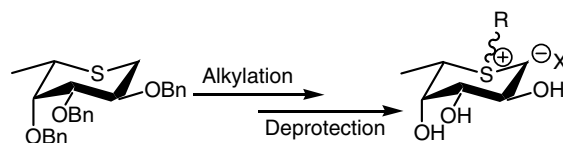
Contents

FULL PAPERS

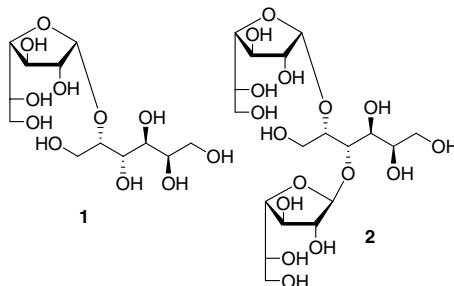
- Synthesis of a benzyl-protected analog of arenarioside, a trisaccharide phenylpropanoid glycoside** pp 2469–2477
Feng-Yan Zhou, Jin She and Yan-Guang Wang*



- Facile synthesis of sulfonium ion derivatives of 1,5-anhydro-5-thio-L-fucitol as potential α -L-fucosidase inhibitors** pp 2478–2486
Guofeng Gu, Hui Liu and B. Mario Pinto*



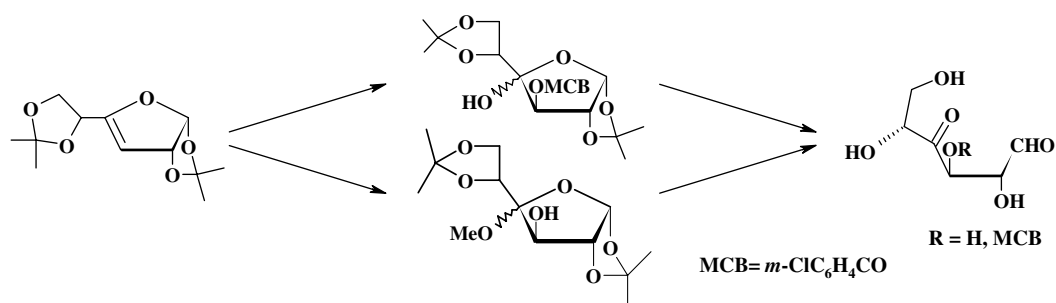
- Synthesis of α -D-Galf-(1 \rightarrow 2)-D-galactitol and α -D-Galf-(1 \rightarrow 2)[β -D-Galf-(1 \rightarrow 3)]-D-galactitol, oligosaccharide derivatives from *Bacteroides cellulosolvens* glycoproteins** pp 2487–2497
Lucía Gandolfi-Donadio, Gabriel Gola, Rosa M. de Lederkremer and Carola Gallo-Rodriguez*



A new and efficient entry to D-xyllo-hexos-4-ulose and some derivatives thereof through epoxidation of the 3,4-hexeno derivative of diacetone-D-glucose

pp 2498–2506

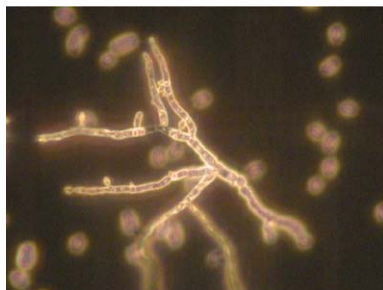
Emanuele Attolino, Giorgio Catelani, Felicia D'Andrea* and Martina Landi



Bioconversion of shellfish chitin wastes for the production of *Bacillus subtilis* W-118 chitinase

pp 2507–2515

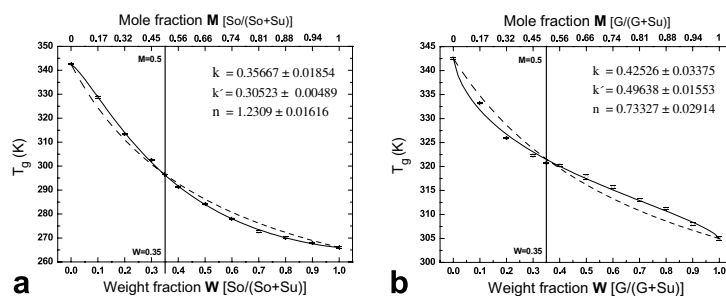
San-Lang Wang,* Tzu-Yin Lin, Yue-Horng Yen, Hui-Fen Liao and Yu-Jen Chen



The glass transition temperatures of sugar mixtures

pp 2516–2520

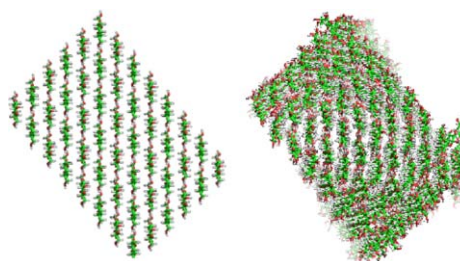
Jeong-Ah Seo, S. J. Kim, Hyun-Joung Kwon, Y. S. Yang, Hyung Kook Kim and Yoon-Hwae Hwang*



Swelling behavior of the cellulose I β crystal models by molecular dynamics

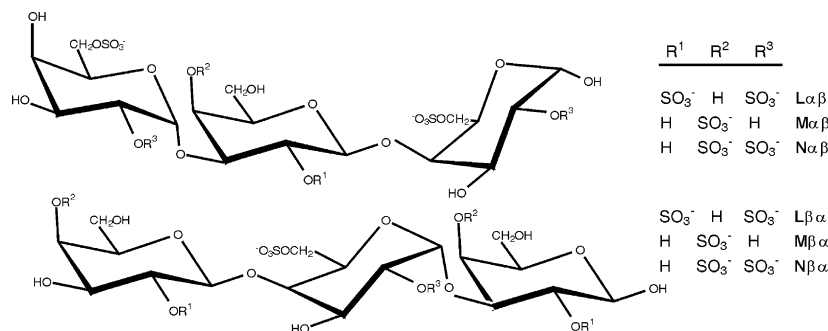
pp 2521–2530

Toshifumi Yui,* Shinya Nishimura, Shingo Akiba and Sachio Hayashi

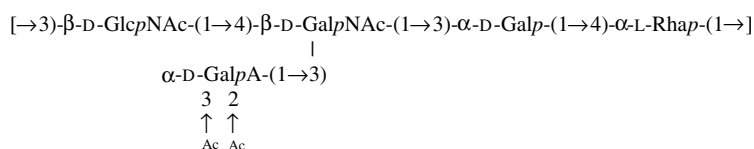


MM3 Potential energy surfaces of trisaccharide models of λ -, μ -, and ν -carrageenans**pp 2531–2542**

Carlos A. Stortz

**Structural elucidation of the O-antigenic polysaccharide from enterohemorrhagic (EHEC) *Escherichia coli* O48:H21****pp 2543–2549**

Leann L. MacLean, Ann C. Webb and Malcolm B. Perry*

**The enhancement of xylose monomer and xylotriose degradation by inorganic salts in aqueous solutions at 180°C****pp 2550–2556**

Chaogang Liu and Charles E. Wyman*

Inorganic salts such as KCl, NaCl, CaCl₂, MgCl₂, and FeCl₃, and particularly the latter, significantly increased xylose monomer and xylotriose degradation in water heated to 180°C and also caused considerable losses of xylose to unidentified compounds.

Progesterone binding to the tryptophan residues of human α_1 -acid glycoprotein**pp 2557–2564**

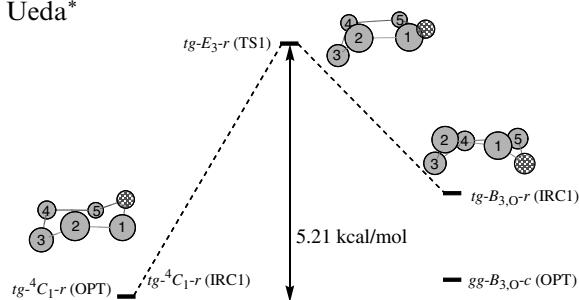
J. R. Albani

Binding studies between progesterone and α_1 -acid glycoprotein allowed us to demonstrate that the binding site of progesterone contains one hydrophobic tryptophan residue and that the structure of the protein is not altered upon binding. The data obtained at saturated concentrations of progesterone clearly reveals the type of interaction at physiological levels.

An investigation of the pyranose ring interconversion path of α -L-idose calculated using density functional theory

pp 2565–2574

Youji Kurihara* and Kazuyoshi Ueda*



B3LYP/6-311++G** study of structure and spin–spin coupling constant in methyl 2-O-sulfo- α -L-iduronate

pp 2575–2580

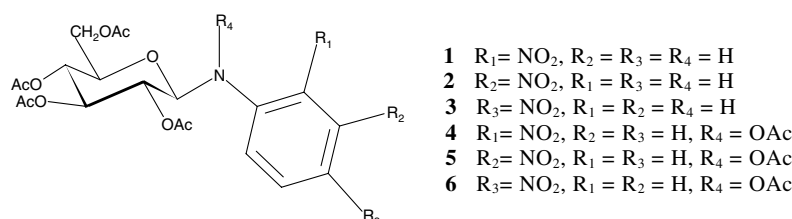
Miloš Hricovíni

Structures and spin–spin coupling constants of three most stable conformations (1C_4 , 4C_1 , 2S_0) of methyl 2-O-sulfo- α -L-iduronate monosodium salt have been analyzed by DFT at the B3LYP/6-311++G** level of theory.

Crystal structure and solid-state ^{13}C NMR analysis of *N*-*o*-, *N*-*m*- and *N*-*p*-nitrophenyl-2,3,4,6-tetra-*O*-acetyl- β -D-glucopyranosylamines, and their *N*-acetyl derivatives

pp 2581–2590

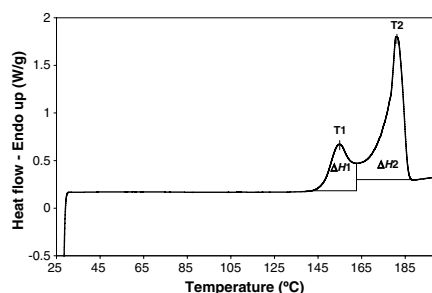
Andrzej Temeriusz,* Tomasz Gubica, Paulina Rogowska, Katarzyna Paradowska and Michał K. Cyrański



DSC study of sucrose melting

pp 2591–2599

Stephen T. Beckett, M. Grazia Francesconi, Peter M. Geary, Grahame Mackenzie* and Aurélia P. E. Maulny



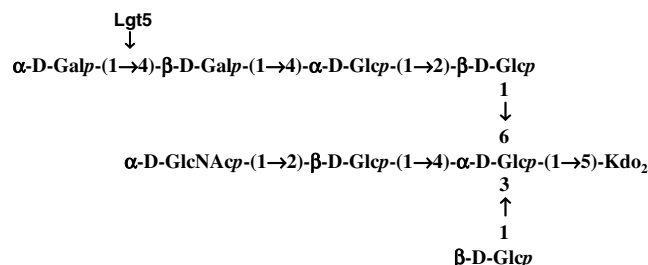
Typical DSC thermogram for recrystallised sucrose showing T1, ΔH_1 and T2, ΔH_2 .

NOTES

Identification of a novel glycosyltransferase involved in LOS biosynthesis of *Moraxella catarrhalis*

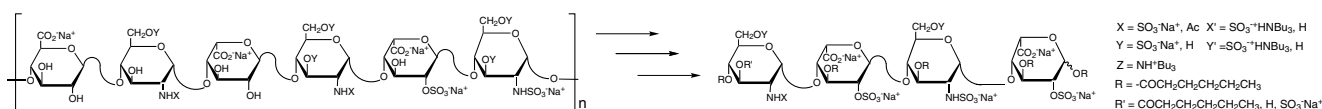
pp 2600–2606

Jennifer C. Wilson, Patrick M. Collins, Zoran Klipic, I. Darren Grice and Ian R. Peak*

**Increase in the growth inhibition of bovine pulmonary artery smooth muscle cells by an *O*-hexanoyl low-molecular-weight heparin derivative**

pp 2607–2612

Hari G. Garg,* Charles A. Hales, Lunyin Yu, Melissa Butler, Tasneem Islam, Jin Xie and Robert J. Linhardt

**The structures of the cell wall teichoic acids from the thermophilic microorganism *Geobacillus thermoleovorans* strain Fango**

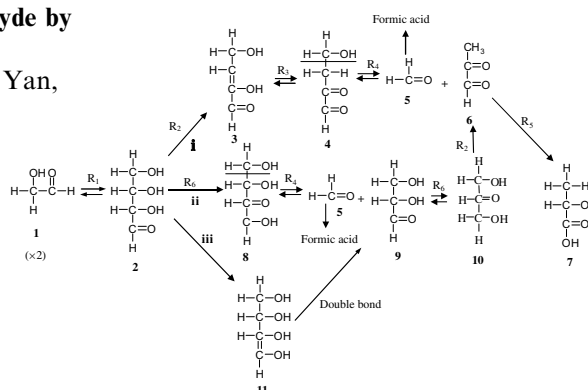
pp 2613–2618

Serena Leone, Antonio Molinaro,* Ida Romano, Barbara Nicolaus, Rosa Lanzetta, Michelangelo Parrilli and Otto Holst


**Formation of lactic acid from glycolaldehyde by alkaline hydrothermal reaction**

pp 2619–2623

Hisanori Kishida, Fangming Jin,* Xiuyi Yan, Takehiko Moriya and Heiji Enomoto



*Corresponding author

 Supplementary data available via ScienceDirect

COVER

Image represents a key process of malaria parasites multiplying in, and rupturing from the human blood cell. The parasite surface is coated with glycosylphosphatidylinositols (GPIs), which have been identified as the malaria toxin by a collaborative effort between the research groups headed by Peter Seeberger (Swiss Federal Institute of Technology (ETH) Zürich, Switzerland) and Louis Schofield (Walter and Eliza Hall Institute of Medical Research, Australia). The space filling model represents the native GPI molecule from malaria parasite that has been chemically synthesized by the Seeberger group. Professor Peter Seeberger was presented with the Carbohydrate Research Award at the 13th European Carbohydrate Symposium (Bratislava, 2005).

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