

Supporting Information

for

One-Pot Multienzyme-Catalyzed Assembly of Pentasaccharide Repeating Units from Group B *Streptococcus* Capsular Polysaccharides and Their Related Oligosaccharide Derivatives

Min Liang,^{a,b} Wei Gong,^c Jielin Zhao,^a Hong Wang,^d Chongzhen Sun,^a Xianwei Liu,^{a,*} and Guofeng Gu^{a,*}

^a National Glycoengineering Research Center and Shandong Center of Technology Innovation for Carbohydrate, Shandong University,
72 Binhai Road, Qingdao 266237, China

^b Public Health Clinical Center Affiliated to Shandong University, 2999 Gangxing West Road, Jinan, China

^c Shandong Institute for Food and Drug Control, 2749 Xinluo Road, Jinan, China

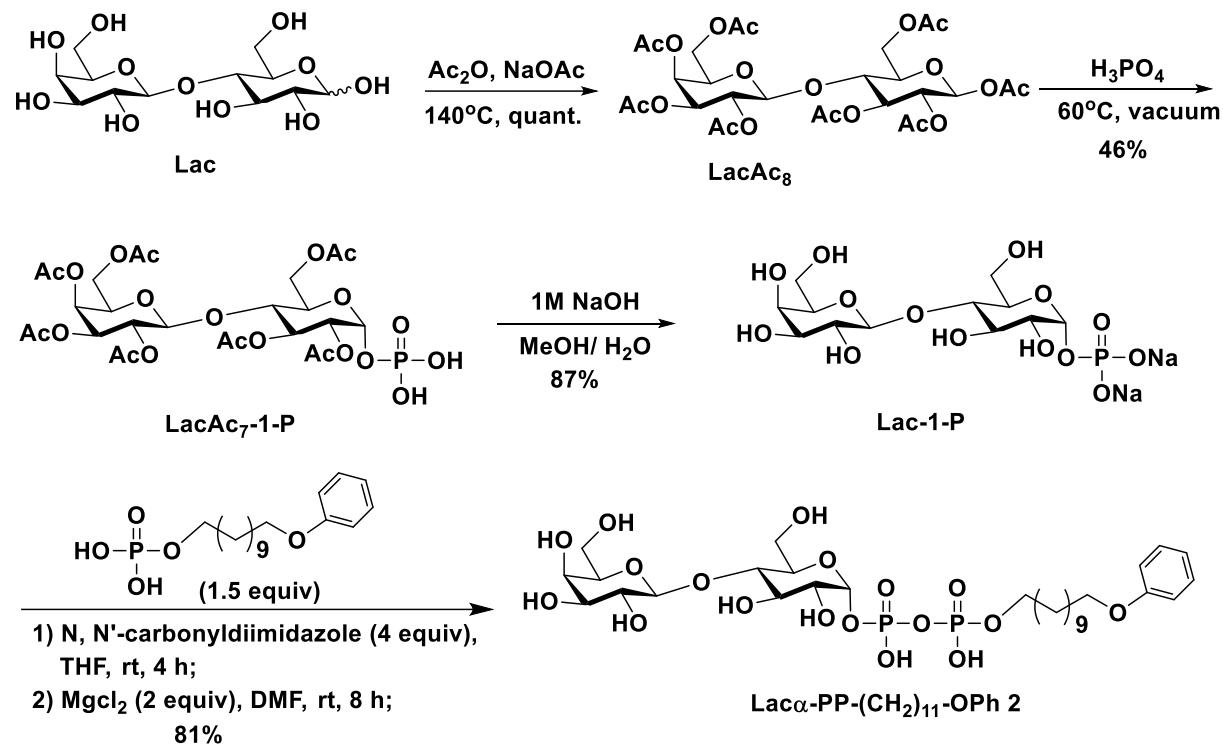
^d College of Bioengineering, Beijing Polytechnic University, No. 9 Liangshuihe 1st Street, Beijing 100176, China

Corresponding authors. Tel/Fax: +86-532-5863 1405; E-mails: xianweiliu@sdu.edu.cn (X. L); guofenggu@sdu.edu.cn (G.G)

Table of Contents

I. Supplementary Scheme and Figures.....	Page S2-S27
II. Experimental Section.....	Page S28-39
III. NMR and ESI-HRMS Spectra.....	Page S40-125
IV. References.....	Page S126

I. Supplementary Scheme and Figures



Scheme S1. Chemical synthesis of Lac α -PP-(CH₂)₁₁-OPh **2**.

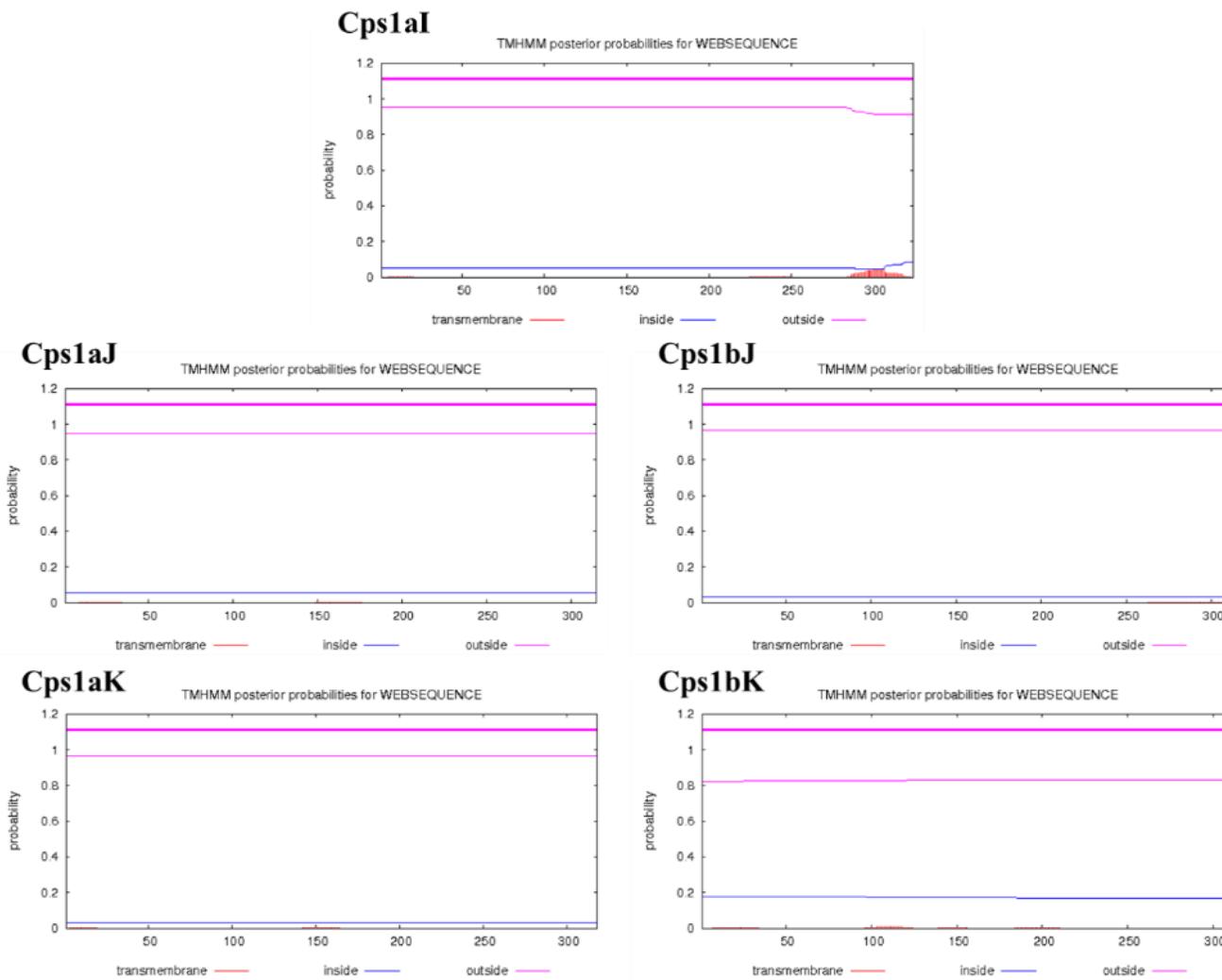


Figure S1. Prediction of transmembrane helices in five target glycosyltransferases by TMHMM Server v. 2.0 software.

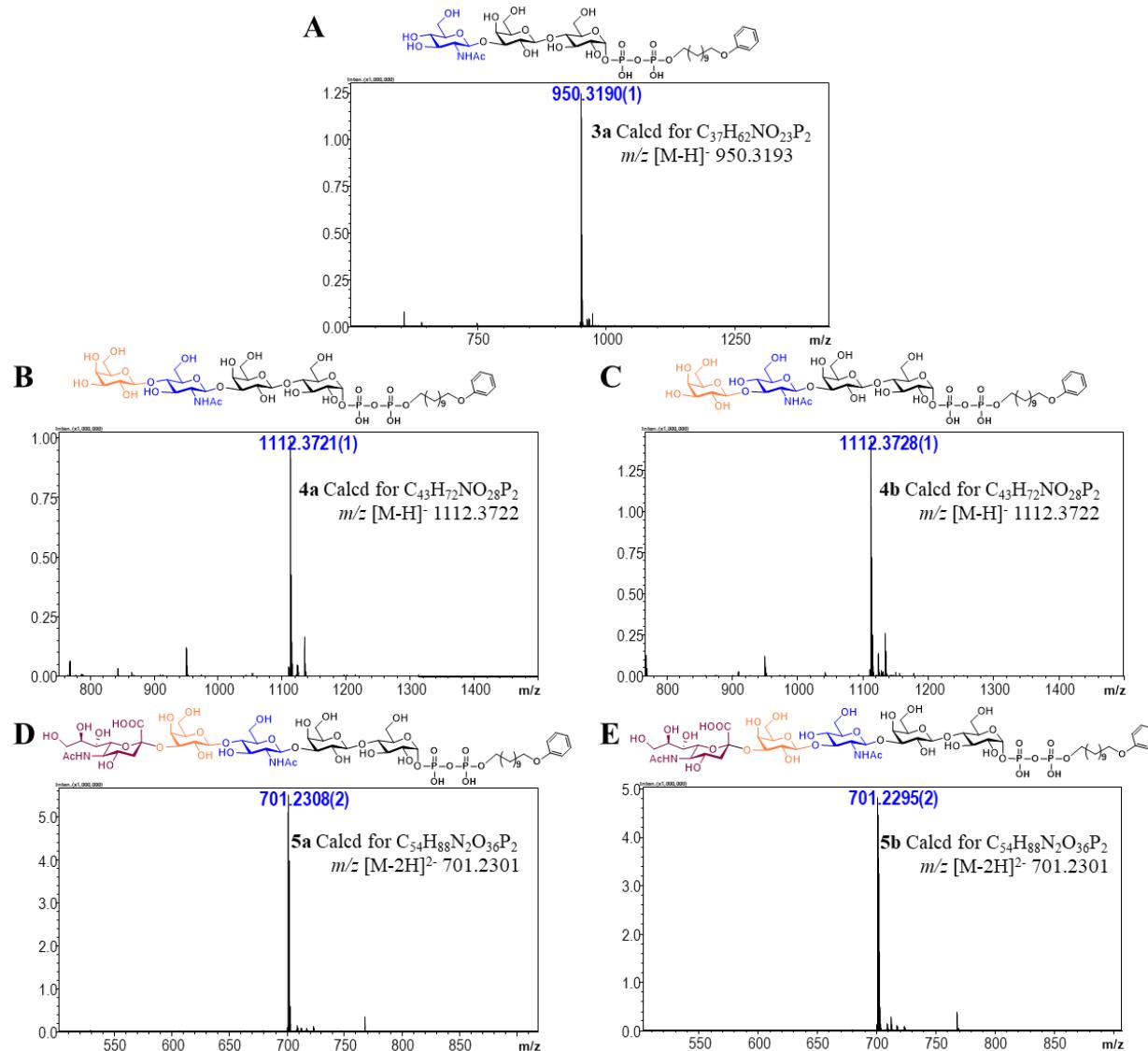


Figure S2. ESI-HRMS analysis of reaction products catalyzed by Cps1aI (A), Cps1aJ (B), Cps1bJ (C), Cps1aK (D) and Cps1bK (E), respectively.

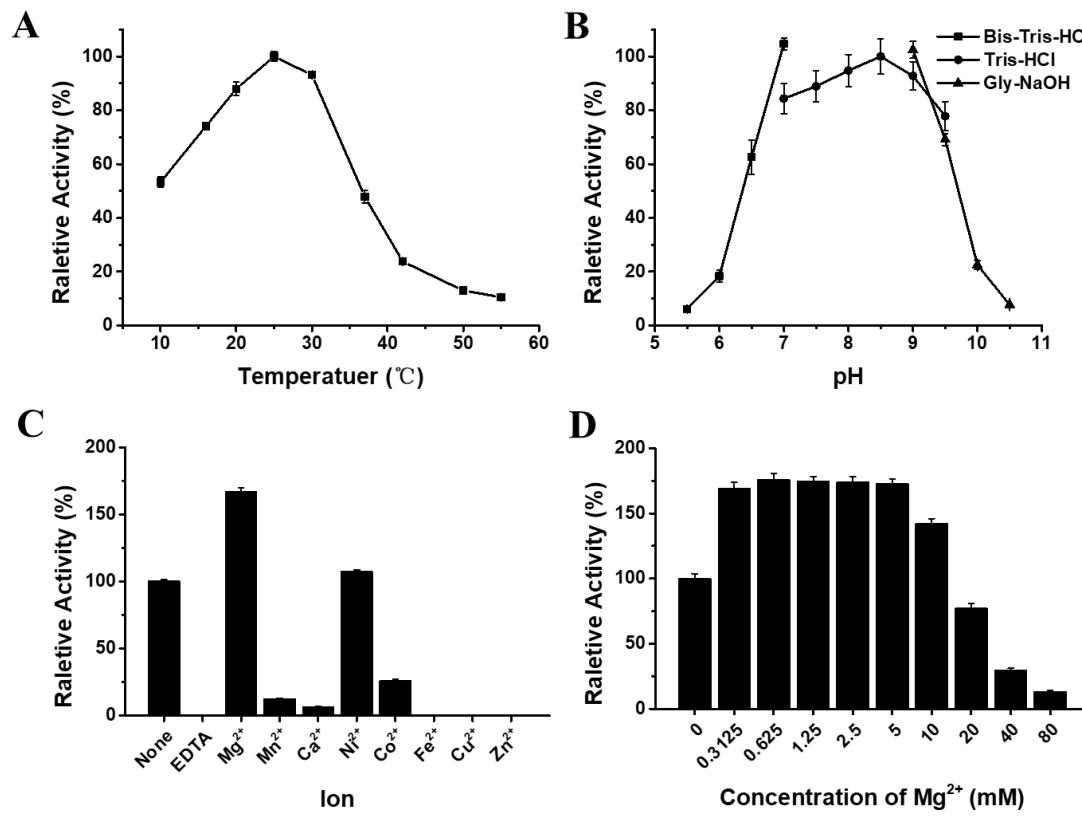


Figure S3. The influences of temperature (A), pH (B), ion (C) and concentration of Mg²⁺ (D) on the relative enzymatic activities of N-acetylglucosaminyltransferase Cps1ai to catalyze the reaction of Lac α -PP-(CH₂)₁₁-OPh 2 with UDP-GlcNAc. Each error bar represents the standard deviation of three experiments.

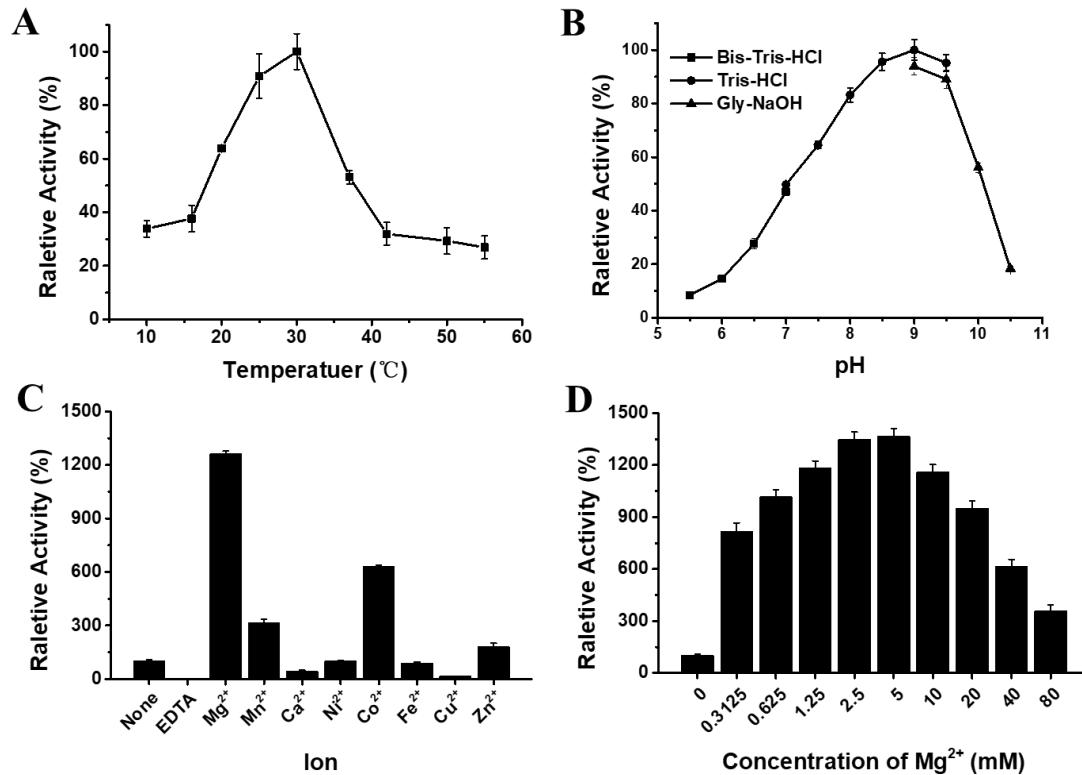


Figure S4. The influences of temperature (A), pH (B), ion (C) and concentration of Mg²⁺ (D) on the relative enzymatic activities of β -1,4-galactosyltransferase Cps1aJ to catalyze the reaction of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** with UDP-Gal. Each error bar represents the standard deviation of three experiments.

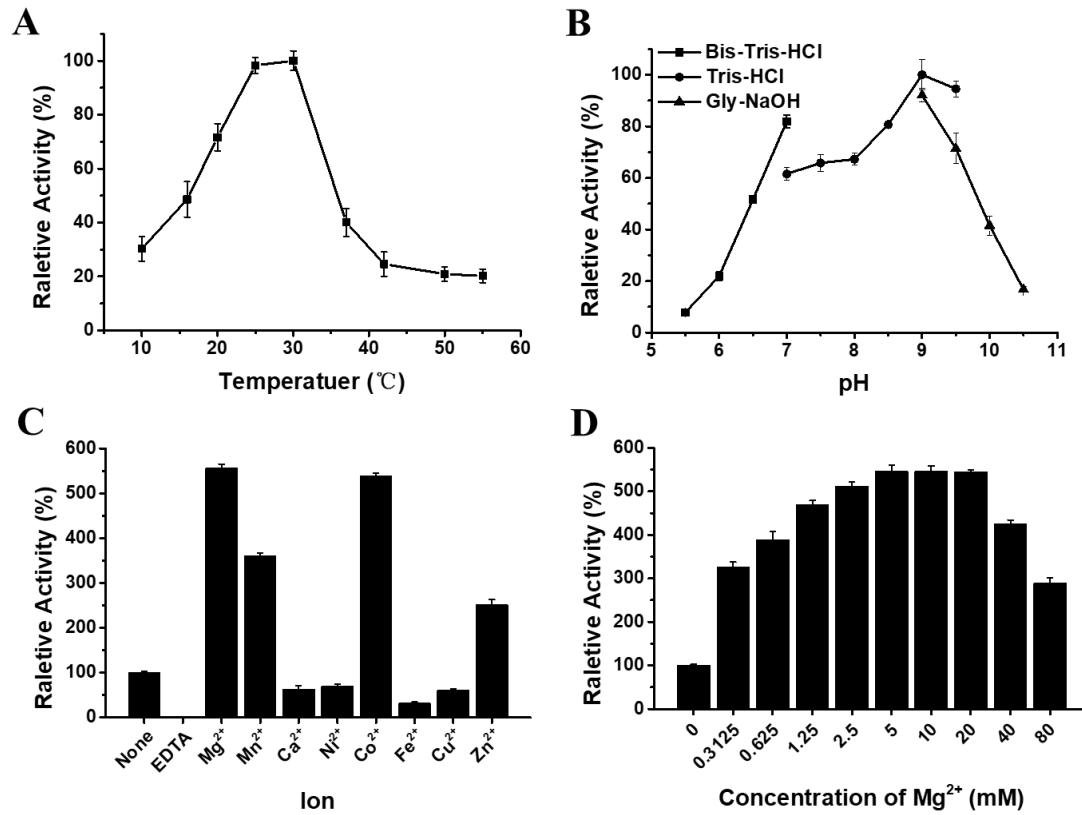


Figure S5. The influences of temperature (A), pH (B), ion (C) and concentration of Mg²⁺ (D) on the relative enzymatic activities of β-1,3-galactosyltransferase Cps1bJ to catalyze the reaction of GlcNAcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **3a** with UDP-Gal. Each error bar represents the standard deviation of three experiments.

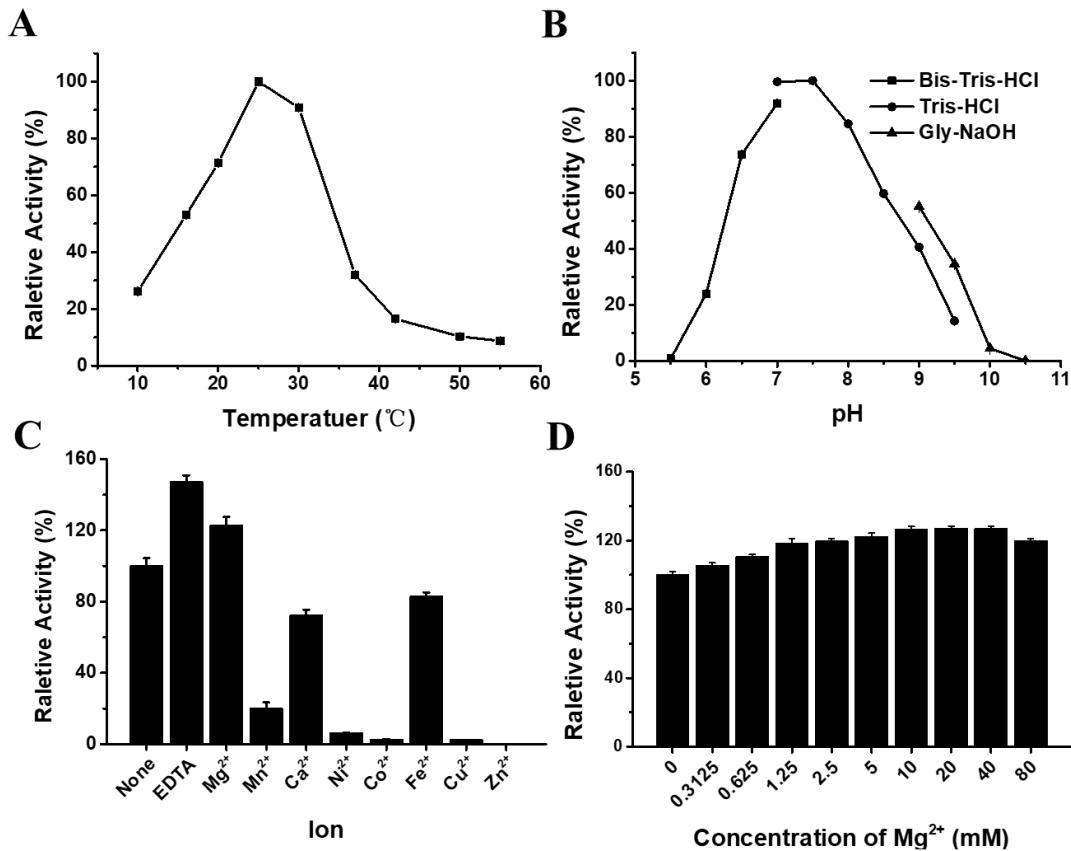


Figure S6. The influences of temperature (A), pH (B), ion (C) and concentration of Mg²⁺ (D) on the relative enzymatic activities of α -2,3-sialyltransferase Cps1aK to catalyze the reaction of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a** with CMP-NeuNAc. Each error bar represents the standard deviation of three experiments.

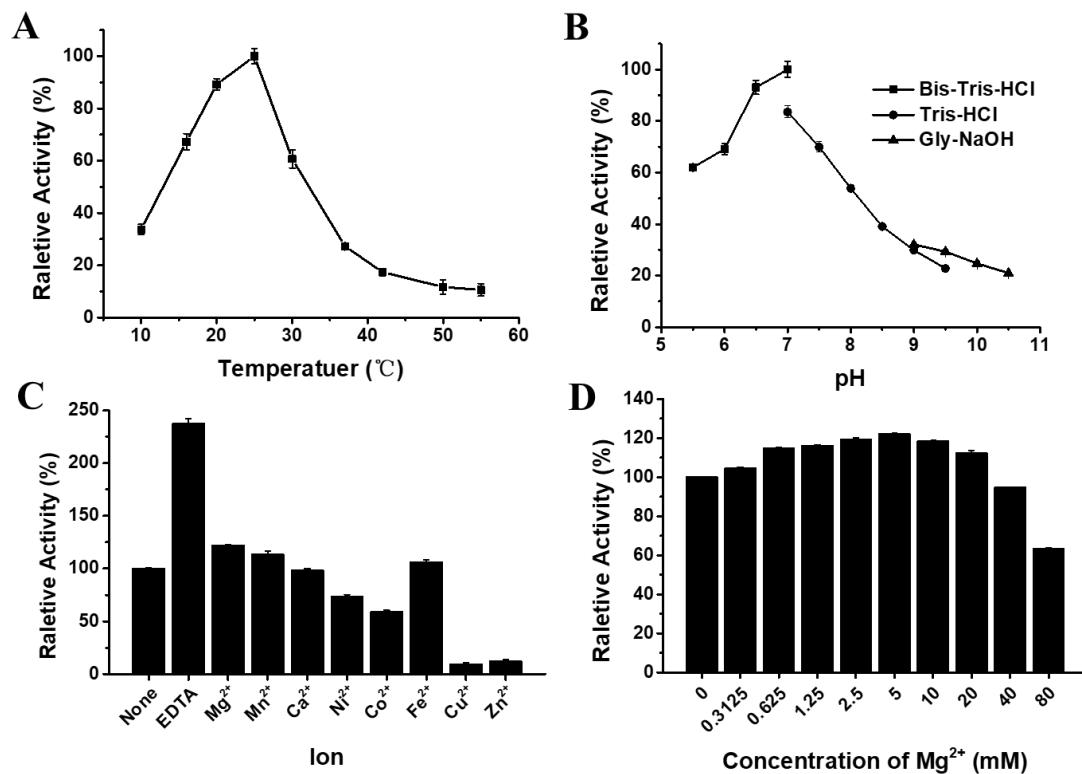


Figure S7. The influences of temperature (A), pH (B), ion (C) and concentration of Mg²⁺ (D) on the relative enzymatic activities of α -2,3-sialyltransferase Cps1bK to catalyze the reaction of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh**4b** with CMP-NeuNAc. Each error bar represents the standard deviation of three experiments.

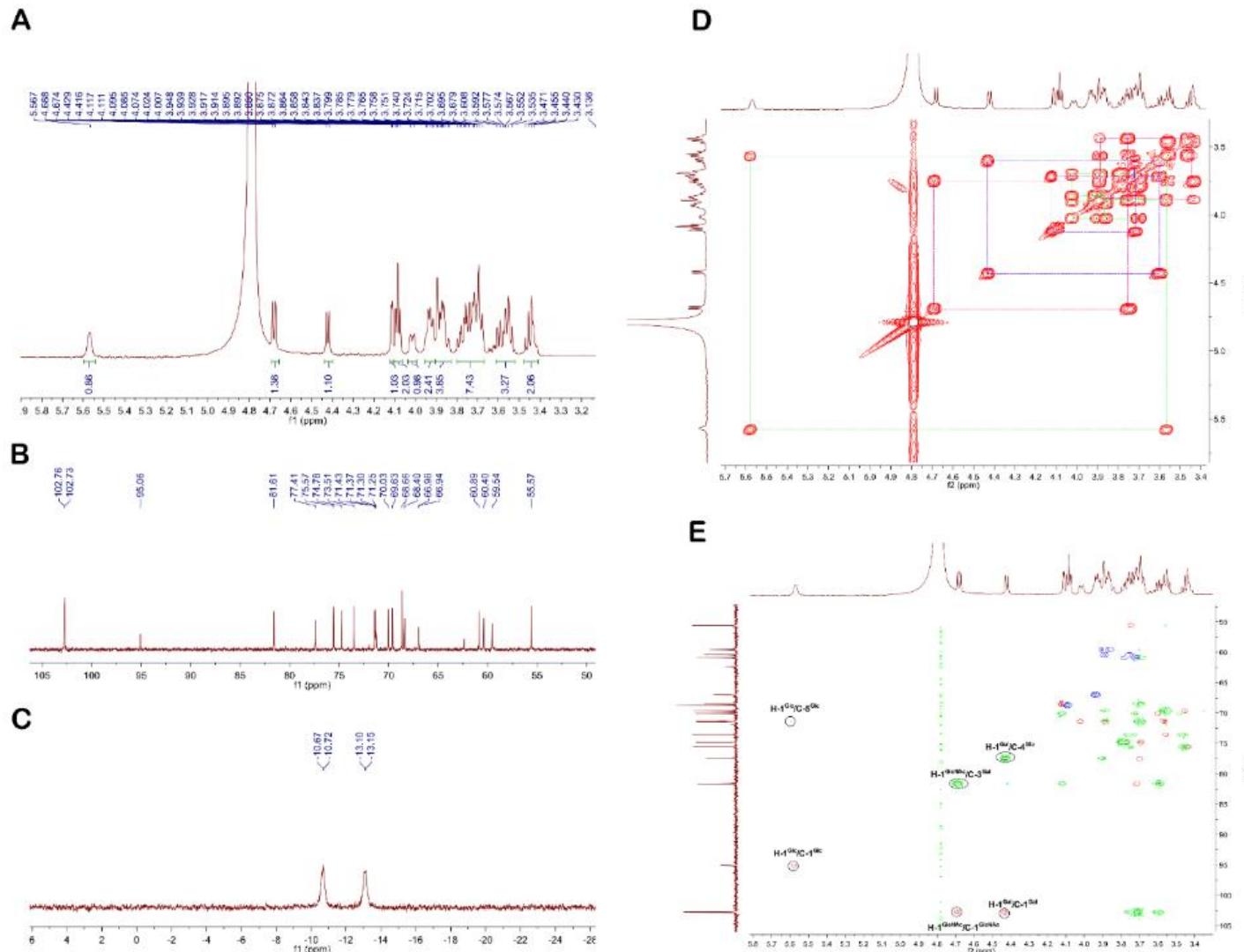


Figure S8. Structural characterization of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** via NMR analysis. Selected ¹H NMR (A), ¹³C NMR (B), ³¹P NMR (C), gCOSY (D) and combined gHSQC (red-blue singals) and gHMBC (green singals) spectra of trisaccharide **3a**.

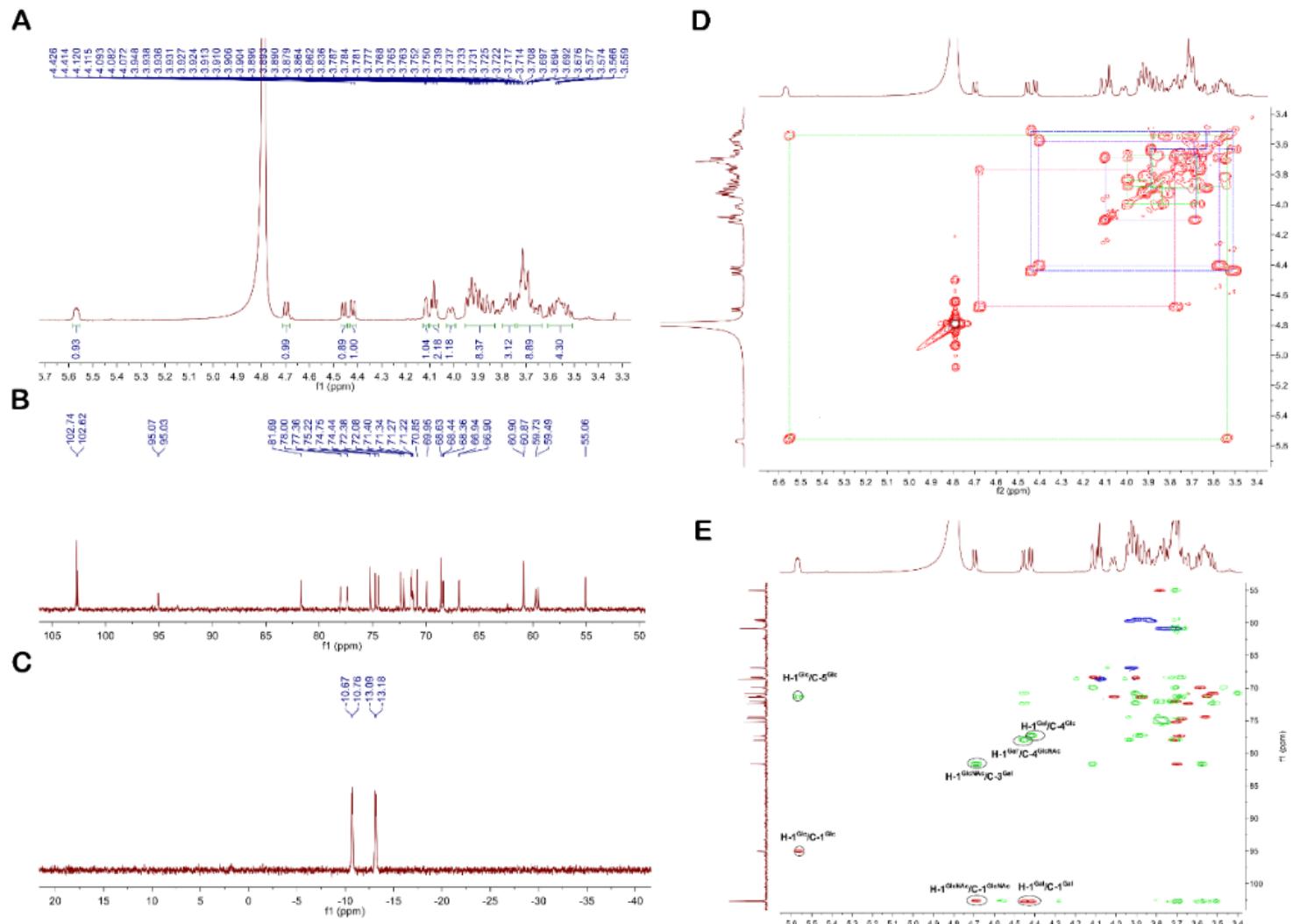


Figure S9. Structural characterization of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a** via NMR analysis. Selected ¹H NMR (A), ¹³C NMR (B), ³¹P NMR (C), gCOSY (D) and combined gHSQC (red-blue singals) and gHMBC (green singals) spectra of tetrasaccharide **4a**.

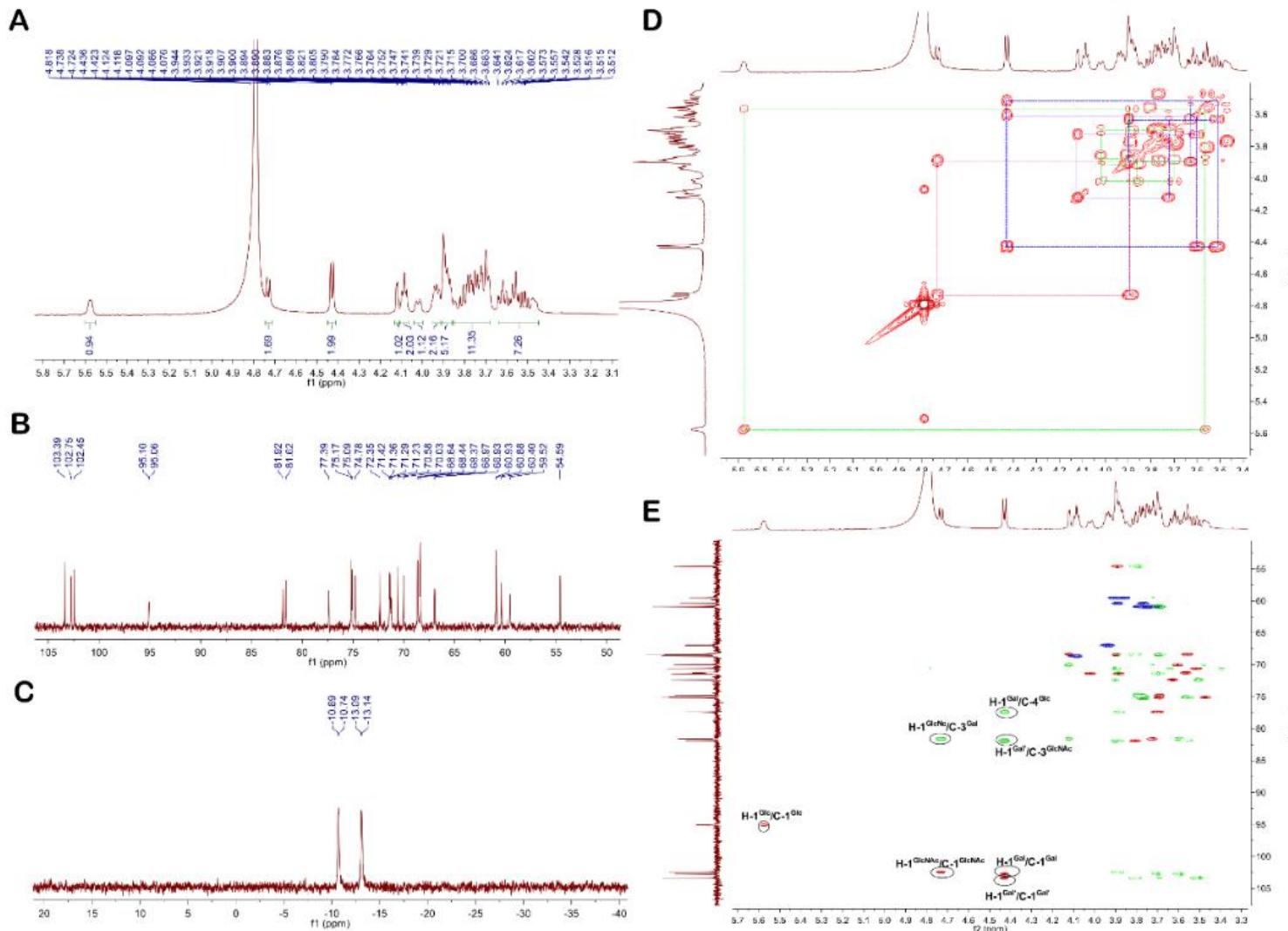


Figure S10. Structural characterization of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b** via NMR analysis. Selected ¹H NMR (A), ¹³C NMR (B), ³¹P NMR (C), gCOSY (D) and combined gHSQC (red-blue singals) and gHMBC (green singals) spectra of tetrasaccharide **4b**.

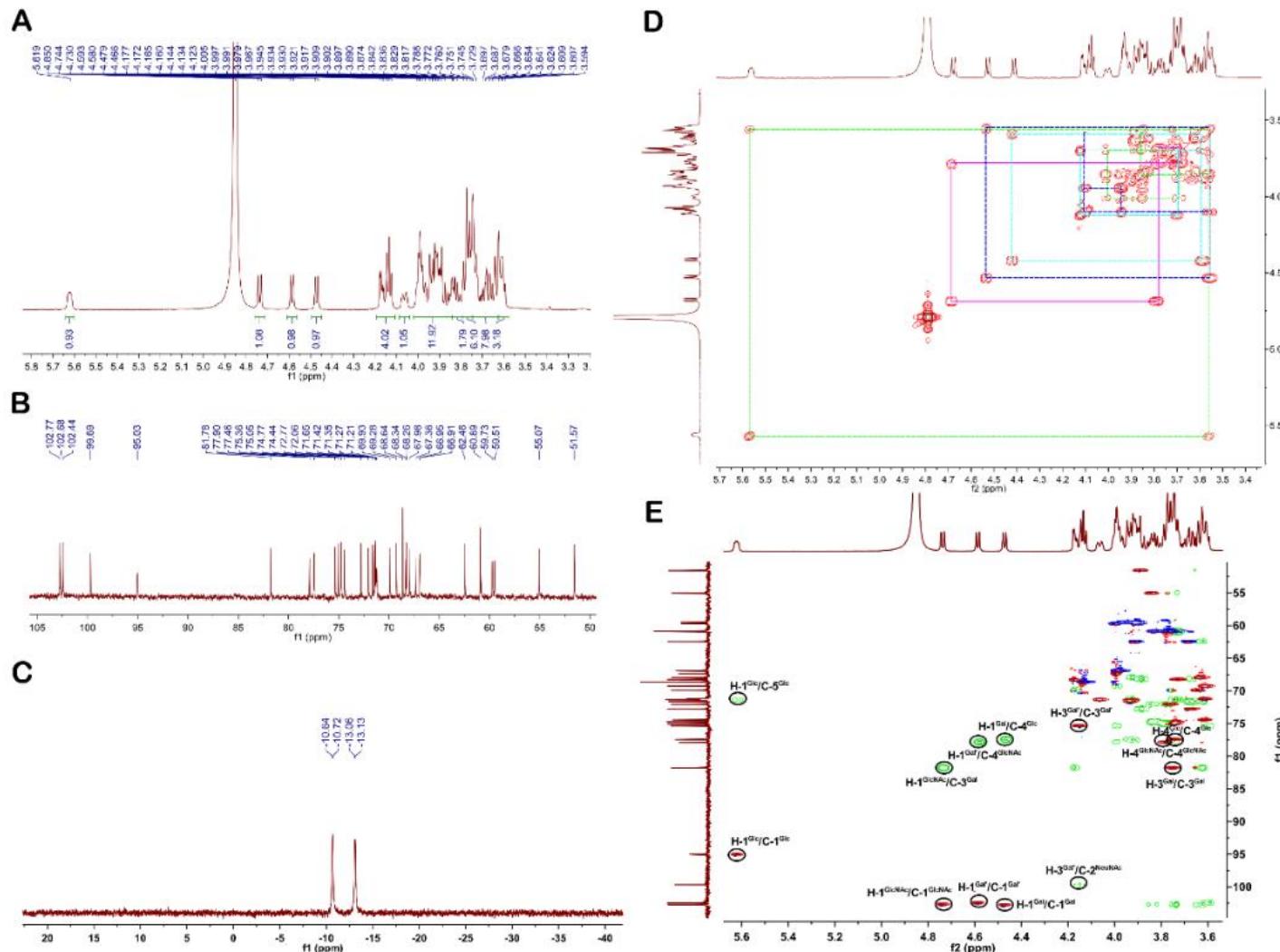


Figure S11. Structural characterization of NeupNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5a** via NMR analysis. Selected ¹H NMR (A), ¹³C NMR (B), ³¹P NMR (C), gCOSY (D) and combined gHSQC (red-blue singals) and gHMBC (green singals) spectra of pentasaccharide **5a**.

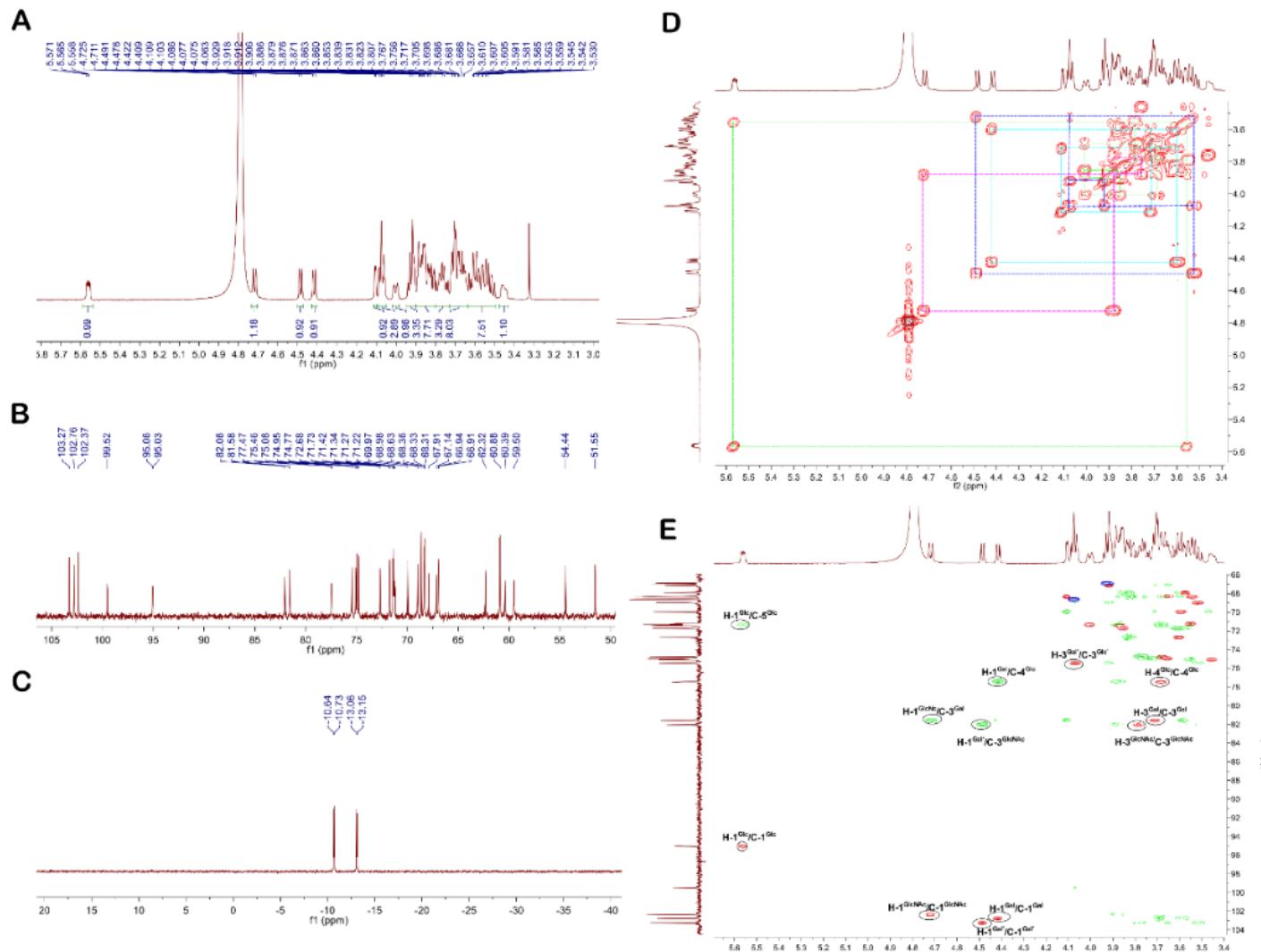


Figure S12. Structural characterization of NeupNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5b** via NMR analysis. Selected ¹H NMR (A), ¹³C NMR (B), ³¹P NMR (C), gCOSY (D) and combined gHSQC (red-blue singals) and gHMBC (green singals) spectra of pentasaccharide **5b**.

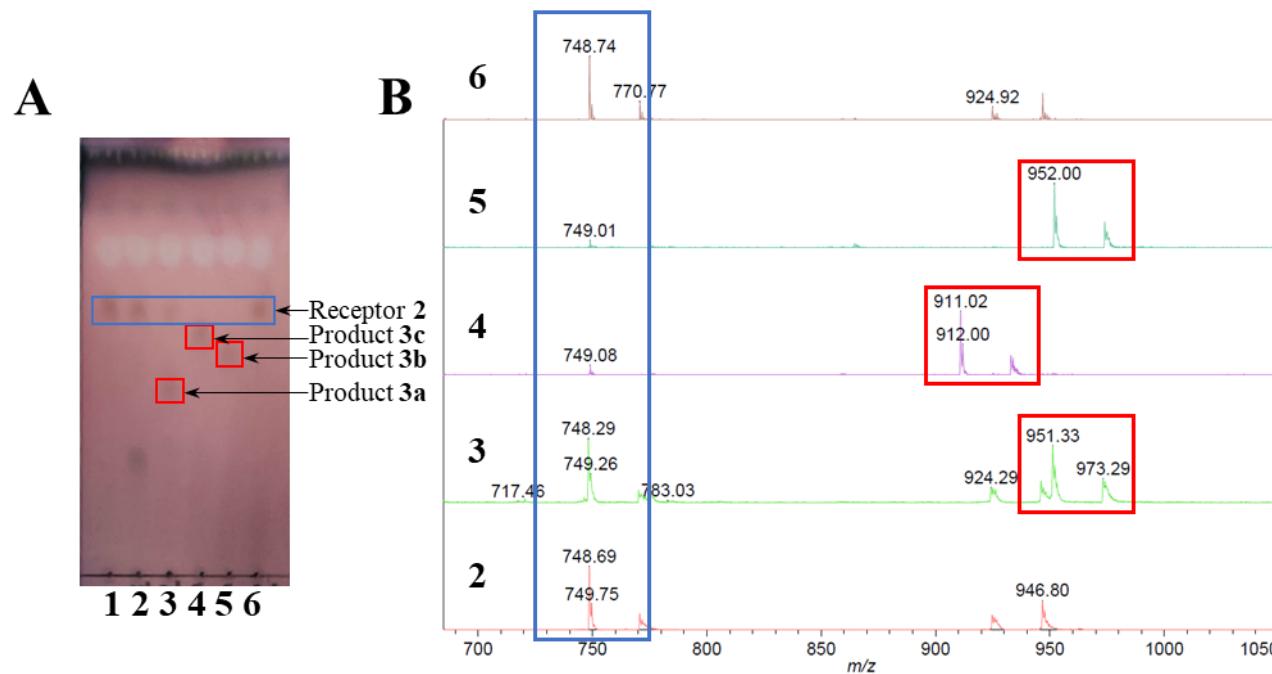


Figure S13. TLC (A) and MALDI-TOF-MS (B) traces of donor substrates recognition of Cps1aI. **1**, negative control reaction with heat-deactivated Cps1aI; **2-6**, enzyme reactions of Lac α -PP-(CH₂)₁₁-OPh **2** with UDP-Gal, UDP-GalNAc, UDP-Glc, UDP-GlcNAc and UDP-GlcAc catalyzed by Cps1aI. The signal peaks of Lac α -PP-(CH₂)₁₁-OPh **2** and trisaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively (MALDI-TOF-MS error range ± 1 Da).

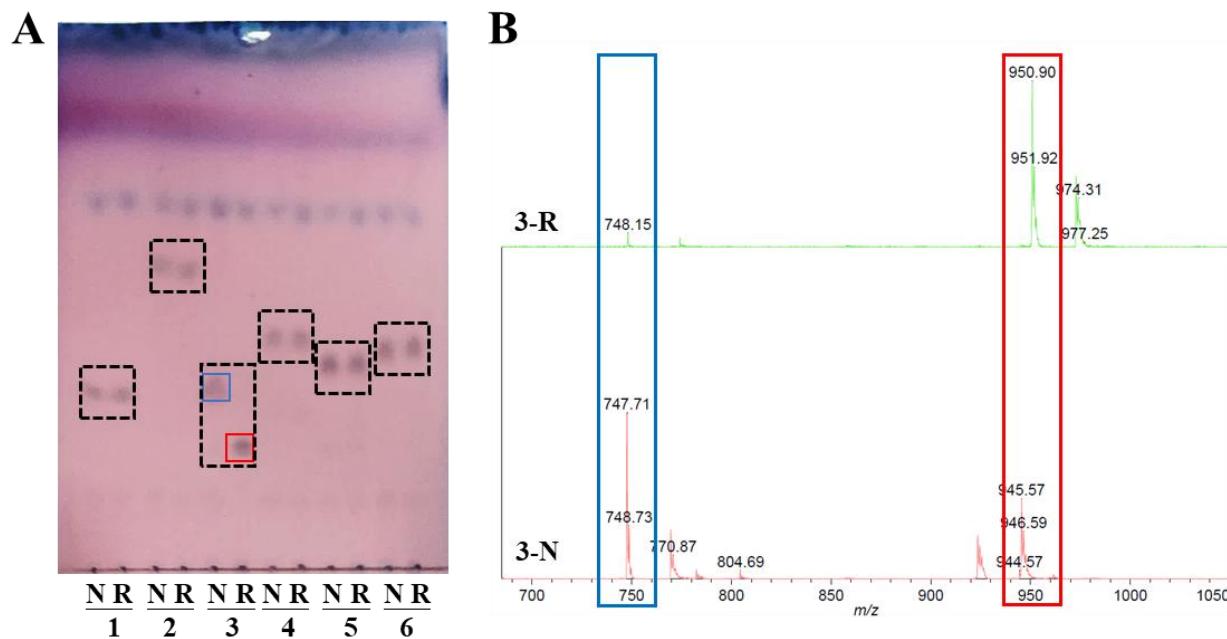


Figure S14. TLC (A) and MALDI-TOF-MS (B) traces of receptor substrates recognition of Cps1al. *N*, negative control reaction with heat-deactivated Cps1al; *R*, enzyme reactions of Cps1al. **1-6**, enzyme reactions of UDP-GlcNAc with Lac, Lac β -O(CH₂)₂N₃, Lac α -PP-(CH₂)₁₁-OPh **2**, Rha β -1,4-Glc α -PP-(CH₂)₁₁-OPh, Gal α -1,3-Glc α -PP-(CH₂)₁₁-OPh and Glc α -1,3-Glc α -PP-(CH₂)₁₁-OPh catalyzed by Cps1al. Positive results are marked with red dotted boxes in TLC analysis; the signal peaks of Lac α -PP-(CH₂)₁₁-OPh **2** and trisaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively (MALDI-TOF-MS error range ± 1 Da).

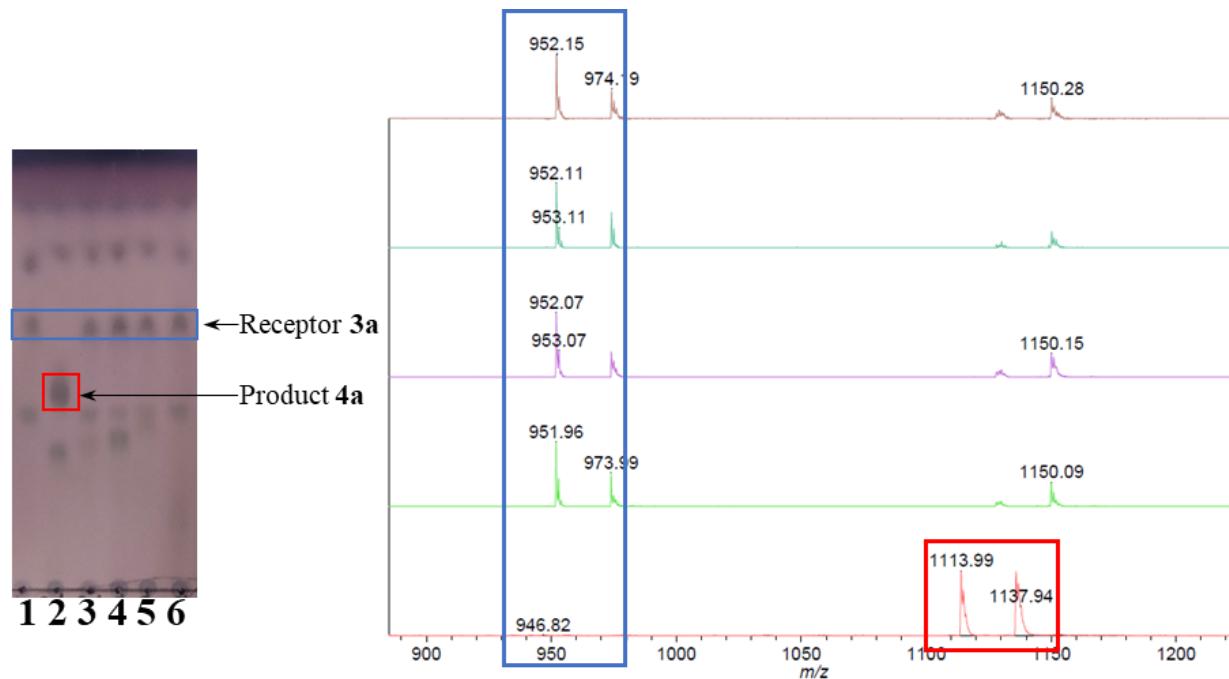


Figure S15. TLC (A) and MALDI-TOF-MS (B) traces of donor substrates recognition of Cps1aJ. **1**, negative control reaction with heat-deactivated Cps1aJ; **2-6**, enzyme reactions of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** with UDP-Gal, UDP-GalNAc, UDP-Glc, UDP-GlcNAc and UDP-GlcA catalyzed by Cps1aJ. Positive results are marked with red dotted boxes in TLC analysis; the signal peaks of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** and tetrasaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively (MALDI-TOF-MS error range ± 1 Da).

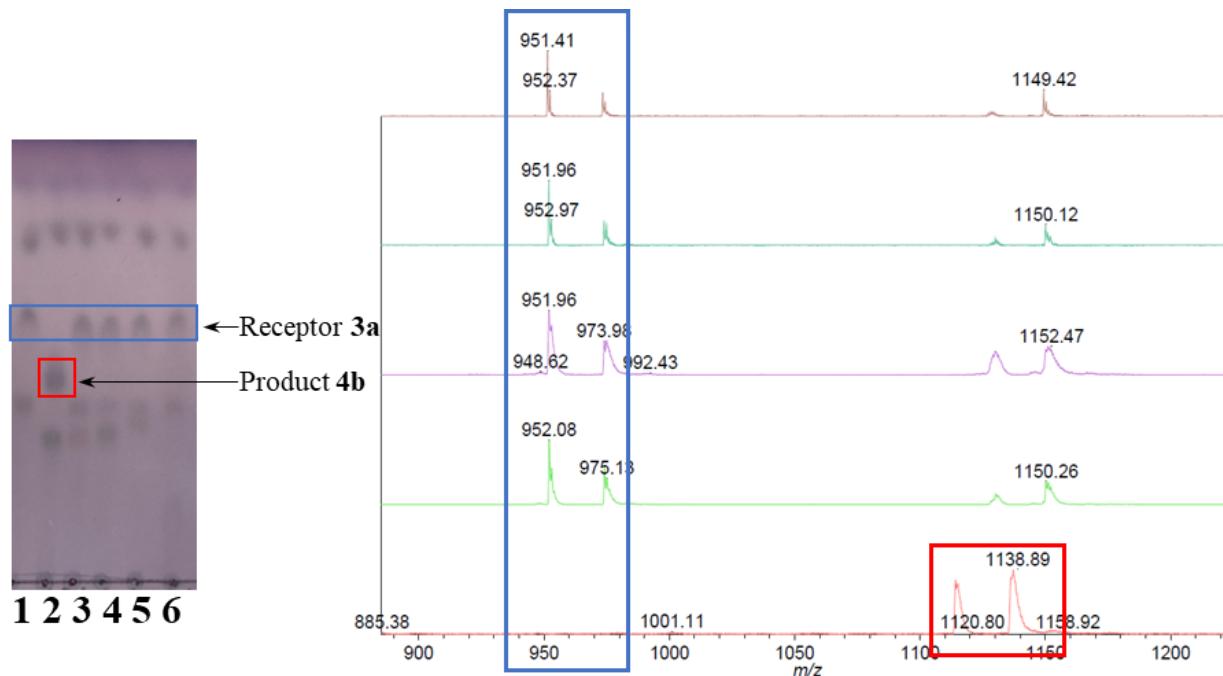


Figure S16. TLC (A) and MALDI-TOF-MS (B) traces of donor substrates recognition of Cps1bJ. **1**, negative control reaction with heat-deactivated Cps1bJ; **2-6**, enzyme reactions of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** with UDP-Gal, UDP-GalNAc, UDP-Glc, UDP-GlcNAc and UDP-GlcA catalyzed by Cps1bJ. Positive results are marked with red dotted boxes in TLC analysis; the signal peaks of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** and tetrasaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively (MALDI-TOF-MS error range ± 1 Da)

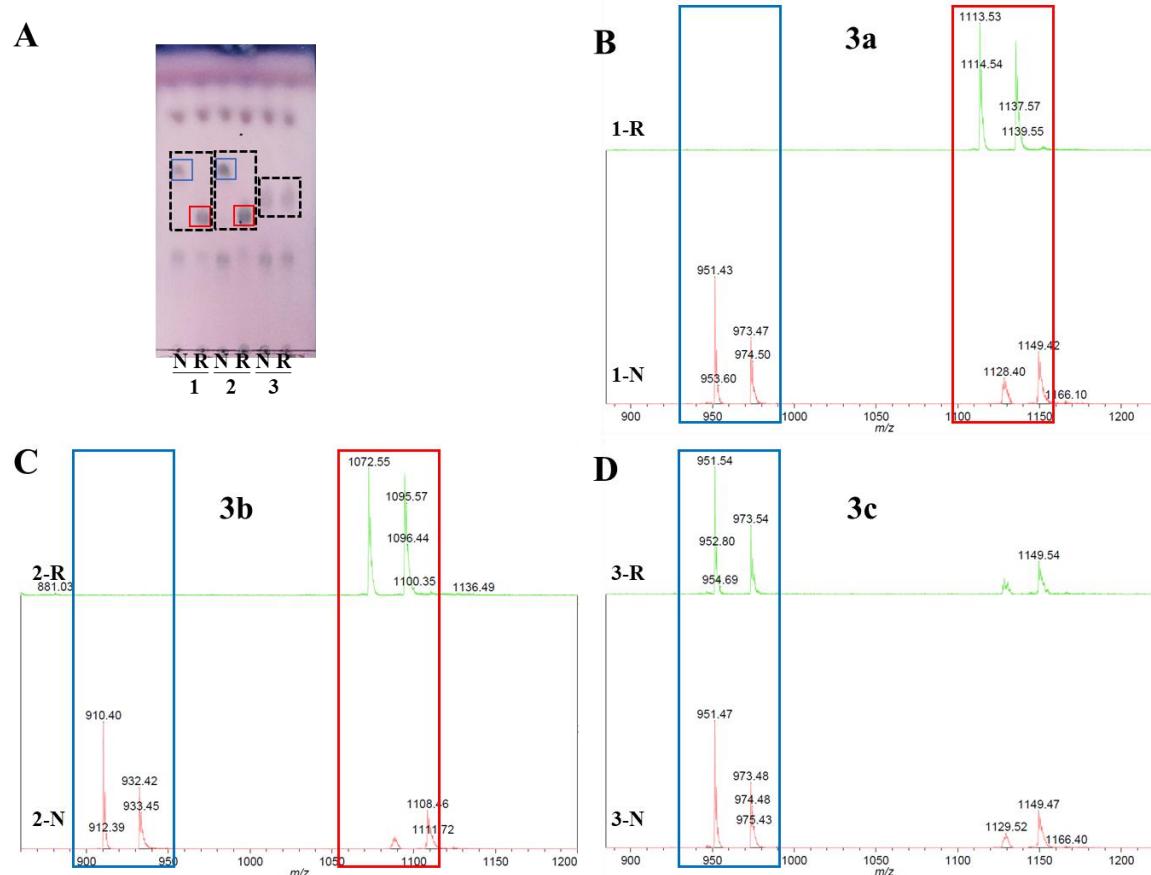


Figure S17. TLC (A) and MALDI-TOF-MS (B-D) traces of receptor substrates recognition of Cps1aJ. *N*, negative control reaction with heat-deactivated Cps1aJ; *R*, enzyme reactions of Cps1aJ. **1-3**, enzyme reactions of UDP-Gal with GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a**, Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3b** and GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3c** catalyzed by Cps1aJ. Positive results are marked with red dotted boxes in TLC analysis; the signal peaks of trisaccharide receptor substrates and tetrasaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively (MALDI-TOF-MS error range ± 1 Da)

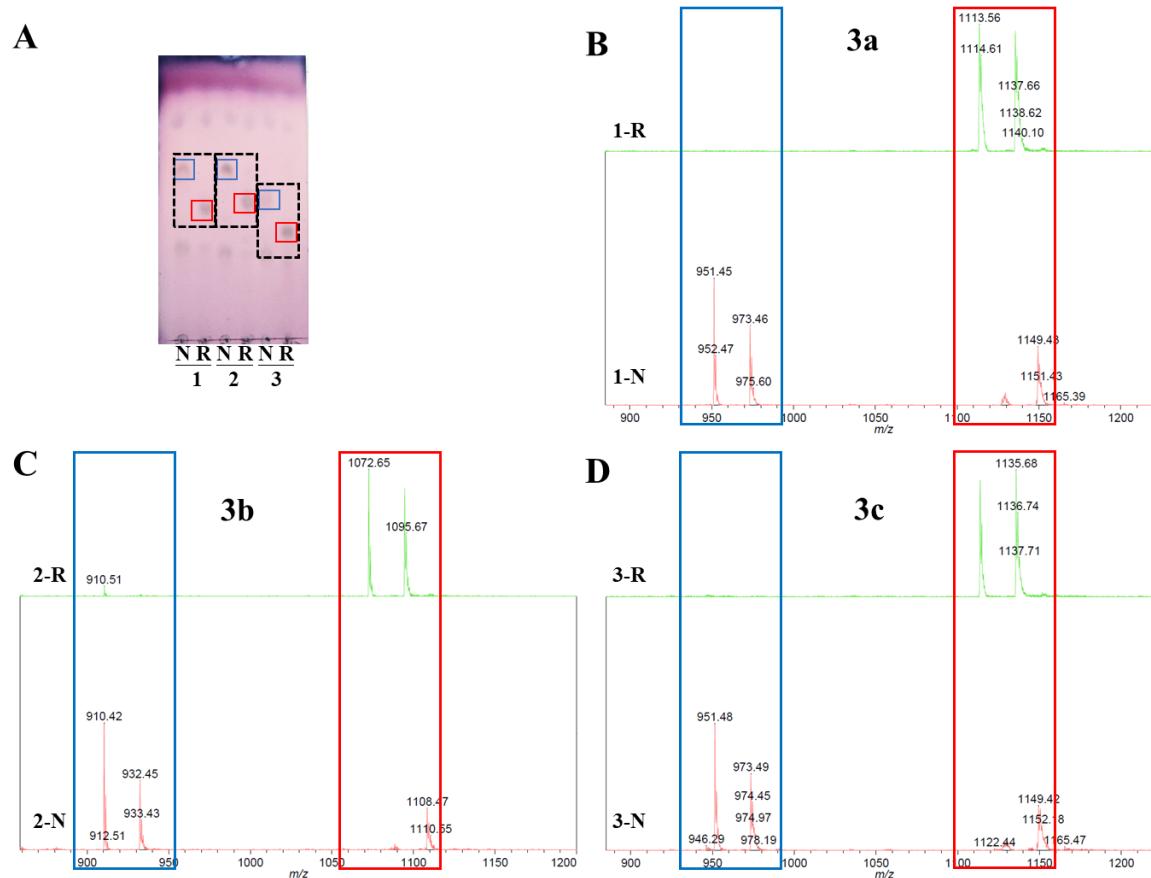


Figure S18. TLC (A) and MALDI-TOF-MS (B-D) traces of receptor substrates recognition of Cps1bJ. *N*, negative control reaction with heat-deactivated Cps1bJ; *R*, enzyme reactions of Cps1bJ. **1-3**, enzyme reactions of UDP-Gal with GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a**, Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3b** and GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3c** catalyzed by Cps1bJ. Positive results are marked with red dotted boxes in TLC analysis; the signal peaks of trisaccharide receptor substrates and tetrasaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively. (MALDI-TOF-MS error range ± 1 Da)

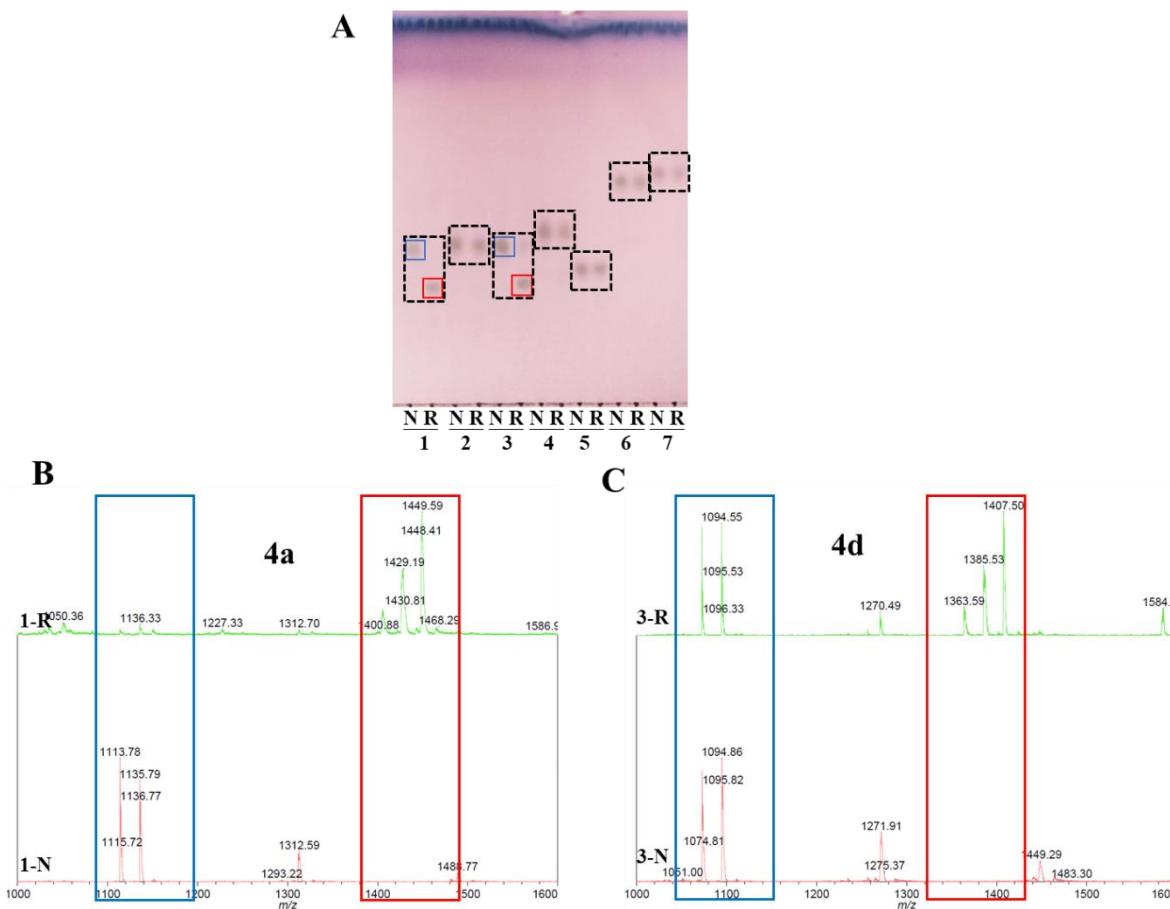


Figure S19. TLC (A) and MALDI-TOF-MS (B-C) traces of receptor substrates recognition of Cps1aK. *N*, negative control reaction with heat-deactivated Cps1aK; *R*, enzyme reactions of Cps1aK. **1-7**, enzyme reactions of CMP-NeuNAc with Galβ-1,4-GlcNAcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **4a**, Galβ-1,3-GlcNAcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **4b**, Galβ-1,4-Glcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **4d**, Galβ-1,3-Glcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **4e**, Galβ-1,3-GalNAcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **4c**, Galβ-1,4-Rhaβ-1,4-Glcα-PP-(CH₂)₁₁-OPh, and Lacα-PP-(CH₂)₁₁-OPh **2** catalyzed by Cps1aK. Positive results are marked with red dotted boxes in TLC analysis; the signal peaks of tetrasaccharide receptor substrates and pentasaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively. (MALDI-TOF-MS error range ± 1 Da).

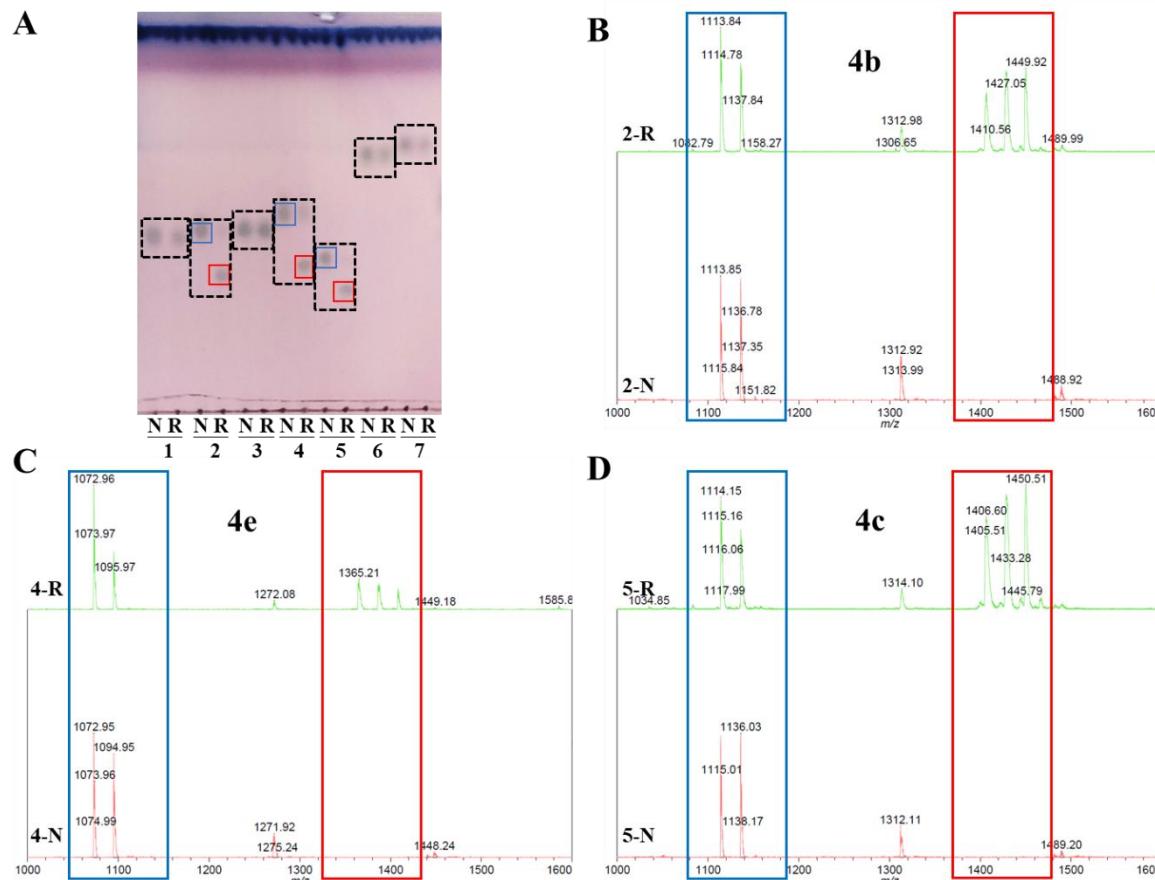


Figure S20. TLC (A) and MALDI-TOF-MS (B-D) traces of receptor substrates recognition of Cps1bK. *N*, negative control reaction with heat-deactivated Cps1bK; *R*, enzyme reactions of Cps1bK. **1-7**, enzyme reactions of CMP-NeuNAc with Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a**, Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b**, Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4d**, Gal β -1,3-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4e**, Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4c**, Gal β -1,4-Rha β -1,4-Glc α -PP-(CH₂)₁₁-OPh, and Lac α -PP-(CH₂)₁₁-OPh **2** catalyzed by Cps1bK. Positive results are marked with red dotted boxes in TLC analysis; the signal peaks of tetrasaccharide receptor substrates and pentasaccharide products in the MALDI-TOF-MS spectra are shown in blue and red solid boxes, respectively. (MALDI-TOF-MS error range ± 1 Da).

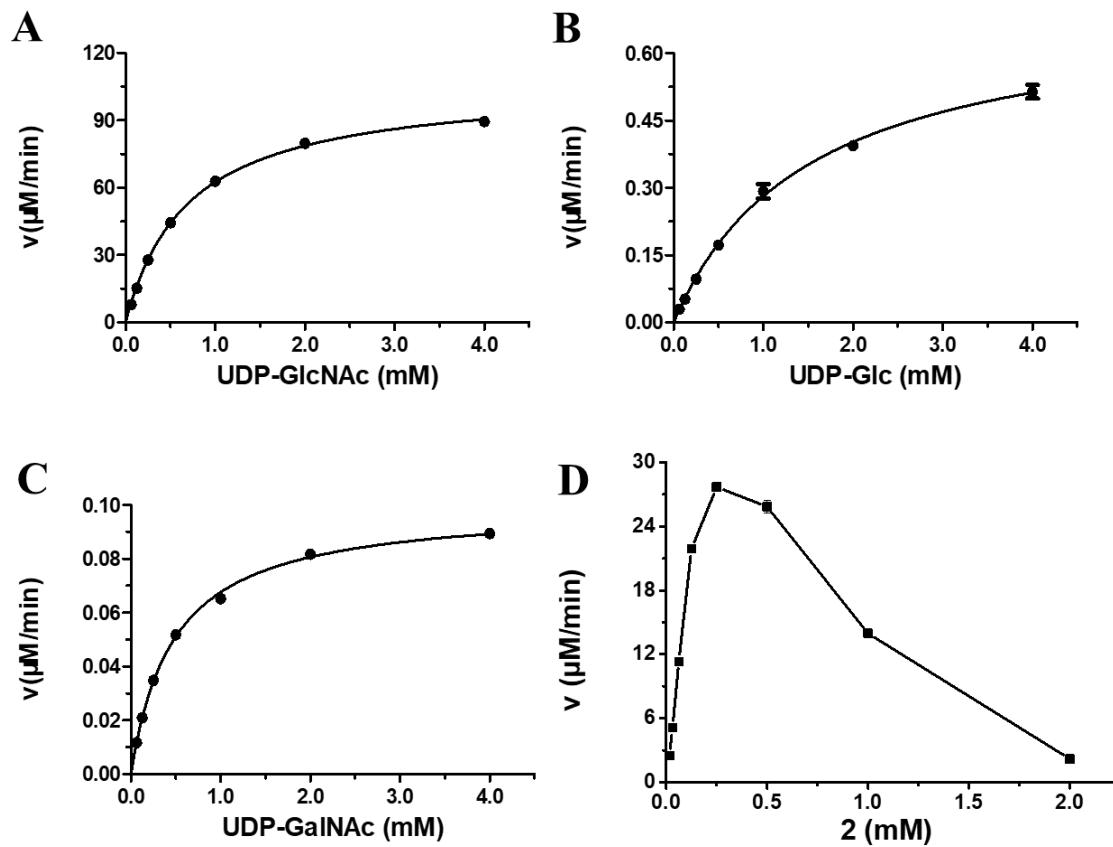


Figure S21. Enzyme kinetics studies of Cps1aI for UDP-GlcNAc (A), UDP-Glc (B) and UDP-GalNAc (C) and the influences of concentration of receptor substrate Lac α -PP-(CH₂)₁₁-OPh **2** (D) on the relative enzymatic activities of Cps1aI. Each error bar represents the standard deviation of three experiments.

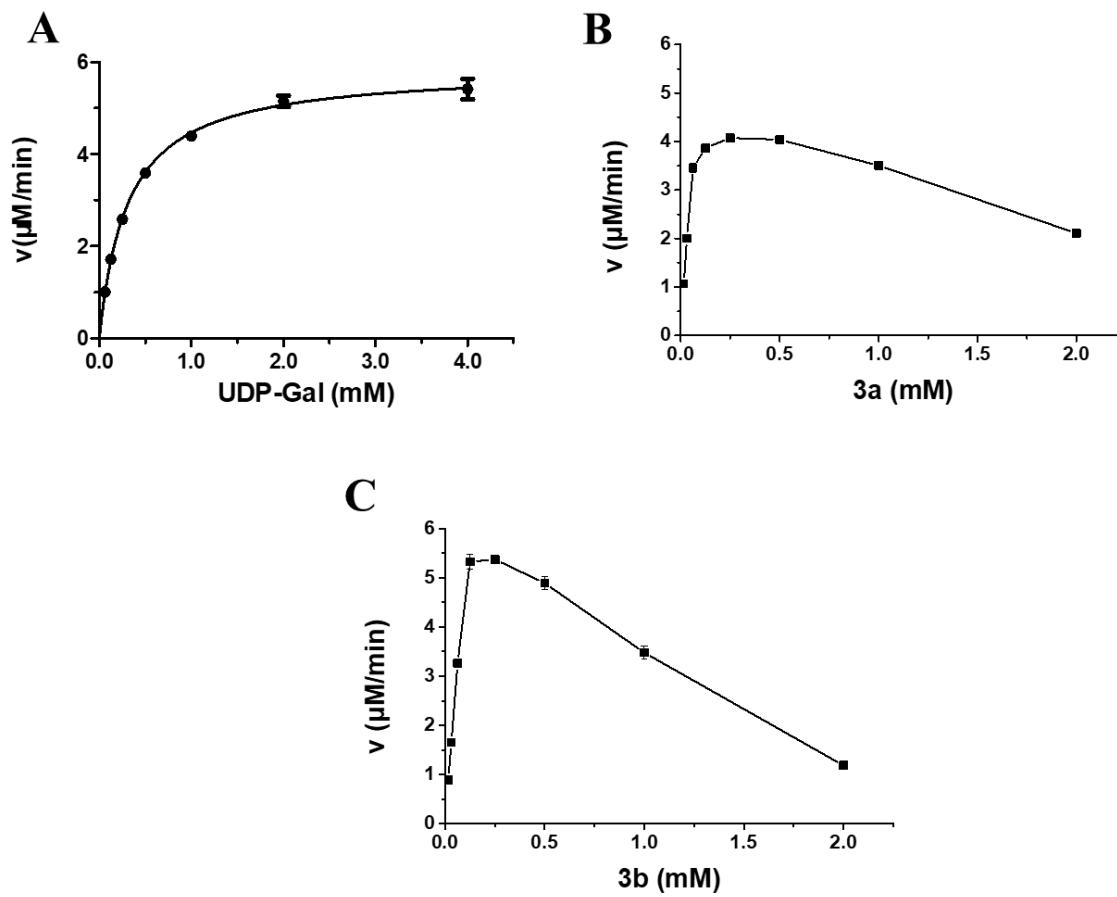


Figure S22. Enzyme kinetics studies of Cps1aJ for UDP-Gal (A) and the influences of concentration of trisaccharide receptor substrate **3a** (B) and **3b** (C) on the relative enzymatic activities of Cps1aJ. Each error bar represents the standard deviation of three experiments.

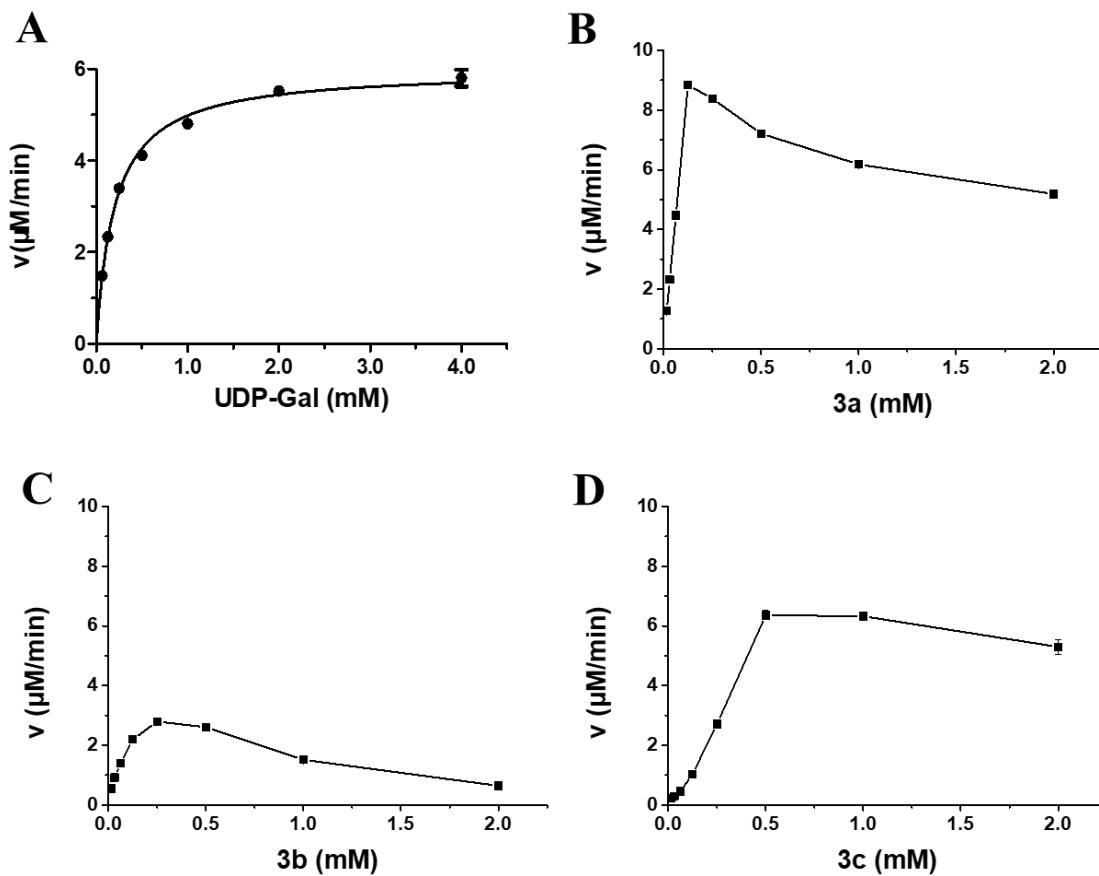


Figure S23. Enzyme kinetics studies of Cps1bJ for UDP-Gal (A) and the influences of concentration of trisaccharide receptor substrate 3a (B), 3b (C) and 3c (D) on the relative enzymatic activities of Cps1bJ. Each error bar represents the standard deviation of three experiments.

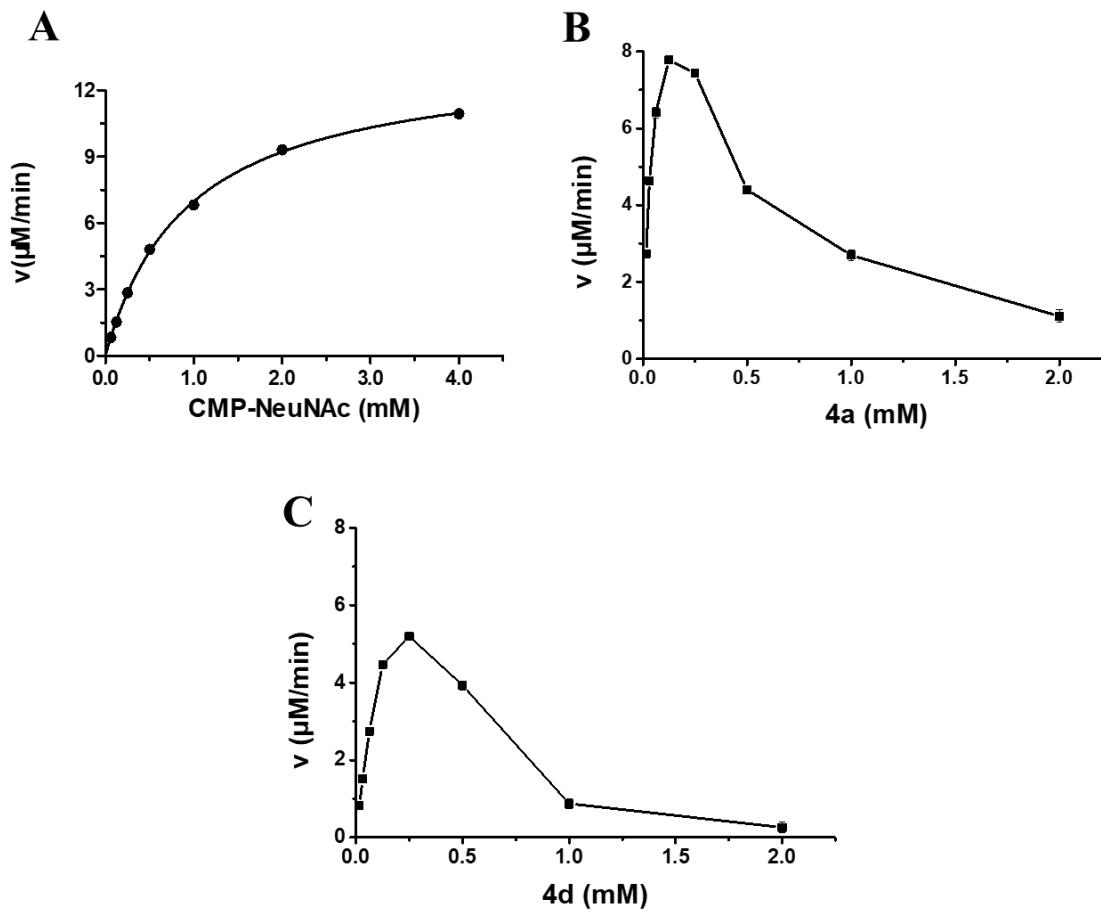


Figure S24. Enzyme kinetics studies of Cps1aK for CMP-NeuNAc (A) and the influences of concentration of tetrasaccharide receptor substrates **4a** (B) and **4d** (C) on the relative enzymatic activities of Cps1aK. Each error bar represents the standard deviation of three experiments.

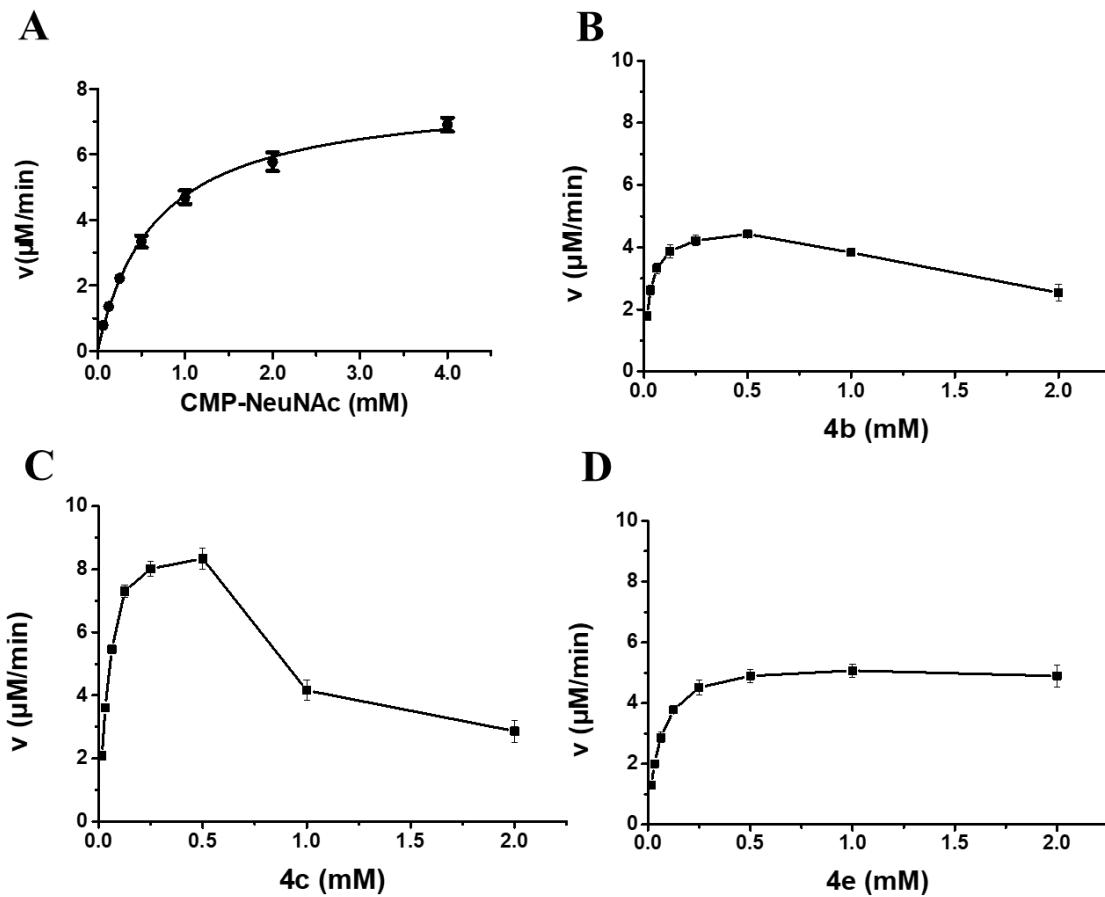


Figure S25. Enzyme kinetics studies of Cps1bK for CMP-NeuNAc (A) and the influences of concentration of tetrasaccharide receptor substrates **4b** (B), **4c** (C) and **4e** (D) on the relative enzymatic activities of Cps1bK. Each error bar represents the standard deviation of three experiments.

II. Experiments Section

Materials. *Escherichia coli* Top10 and BL21(DE3) were used as hosts for transformation and expression, respectively. Both strains were grown in Luria-Bertani (LB) medium supplemented with 100 µg/mL ampicillin at 37 °C. The restriction enzymes were obtained from NEB. The T4 DNA Ligase was from TransGen Company. Yeast extract, agar and tryptone were purchased from OXOID. The C18-SepPak column was from Waters. Other reagents were obtained from Sangon Biotech. Chemicals and solvents were from Sigma-Aldrich.

Preparation of sugar nucleotides. UDP-Glc was prepared using thymidylyltransferase Cps23FL according to the reported method.^[1] UDP-GlcNAc, UDP-Gal, and CMP-NeuNAc were prepared using fusion enzymes GalK-USP-ET64, GlmU-NahK-ET64, and NanA-CSS-ET64, respectively, according to the reported protocols.^[2]

Cloning, overexpression and purification of Cps1aI, Cps1aJ, Cps1aK, Cps1bJ and Cps1bK. DNA sequences of complete *cps1aI*, *cps1aJ*, *cps1aK*, *cps1bJ* and *cps1bK* genes of GBS Ia and Ib were derived from the GenBank (CP000114.1, *cps1aI*: 1225745-1226719, 975 bp, *cps1aJ*: 1224764-1225711, 948 bp and *cps1aK*: 1223724-1224680, 957 bp; AB050723.1, *cps1bJ*: 5008-5949, 942 bp and *cps1bK*: 5942-6880, 939 bp) and synthesized by Sangon Biotech. The amplified fragments were digested with the appropriate restriction endonucleases (*Sma* I and *Xho* I for *cps1aI*, *cps1aJ* and *cps1aK*, *Bam*H I and *Xho* I for *cps1bJ* and *cps1bK*), cloned in plasmid pGEX-4T-1, transferred in *E. coli* BL21(DE3) for protein overexpression. Plasmid DNA sequencing was performed by Sangon Biotech. *E. coli* BL21 (DE3) harboring the recombinant plasmids were allowed to grow until OD₆₀₀ reached 0.6-0.8 in Luria-Bertani (LB) medium supplemented with 100 µg/mL ampicillin (37 °C, 200 rpm). After induction by 0.3 mmol/L isopropyl-1-thio-β-D-galactopyranoside (IPTG) for another 20 h (16 °C, 110 rpm), cells were harvested by centrifugation (8,000 rpm, 10 min, 4 °C). The harvested cell pellets which expressed target proteins were re-suspended in phosphate-buffered saline (PBS, pH 7.4), disrupted by sonication and then cleared by centrifugation (12,000 rpm, 30 min, 4 °C). The resultant supernatants of Cps1aI, Cps1aJ, Cps1aK, Cps1bJ and Cps1bK were subjected to a Glutathione Sepharose 4 Fast Flow column (GE Healthcare) which was pre-equilibrated with PBS buffer (pH 7.4). The GST-tagged proteins were eluted with the same buffer containing 10 mmol/L glutathione reductase (GSH) (pH 7.4). The purified fractions were pooled, concentrated and then desalting using an Amicon Ultra 30-kDa centrifugal filter (Millipore). Finally, the purified protein was re-suspended in PBS buffer (pH 7.4) containing 20% glycerol (v/v) and stored at -80 °C. The purity and homogeneity of five proteins were identified by sodium dodecyl sulfate polyacrylamide gel electrophoresis (SDS-PAGE), and their concentrations were determined using the Thermo Scientific™ NanoDrop One calibrated with the extinction coefficient predicted by ExPASy (<http://web.expasy.org/protparam/>).

Preliminary identification of enzyme activities of Cps1aI, Cps1aJ, Cps1aK, Cps1bJ and Cps1bK. The enzymatic activities of purified Cps1aI, Cps1aJ, Cps1aK, Cps1bJ, and Cps1bK were preliminarily examined in the Tris-HCl buffer (50 mmol/L, pH 7.5) containing 1.2 mmol/L nucleotide sugar (UDP-GlcNAc for Cps1aI, UDP-Gal for Cps1aJ and Cps1bJ, or CMP-NeuNAc for Cps1aK and Cps1bK), 1.0 mmol/L acceptor substrate (Lacα-PP-(CH₂)₁₁-OPh **2** for Cps1aI, GlcNAcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **3a** for

Cps1aJ and Cps1bJ, Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a** for Cps1aK or Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b** for Cps1bK), 5 mmol/L MgCl₂, and 200 μ g/mL purified enzyme in a total volume of 20 μ L. The reaction was incubated at 25 °C using heat-deactivated corresponding enzyme as the negative control, and then detected at 2 h for Cps1aI, 4 h for Cps1aJ and Cps1bJ, or 1 h for Cps1aK and Cps1bK, respectively, by thin-layer chromatography (TLC) (EtOAc:CH₃OH:H₂O:HOAc = 6:3:2:0.5 (v/v/v/v)) or matrix-assisted laser desorption/ionization time of flight mass spectrometry (MALDI-TOF-MS) analysis.

Biochemical characterization of β -1,3-N-acetylglucosaminyltransferase Cps1aI. The reaction system for the detail characterization of purified Cps1aI contained 1 mmol/L UDP-GlcNAc, 1 mmol/L Lac α -PP-(CH₂)₁₁-OPh **2** and 24 μ g/mL of purified enzyme in 50 mmol/L buffer system. Reactions were allowed to proceed for 10 min, terminated by boiling at 100 °C for 30 s, and finally cleared by centrifugation (12,000 rpm, 10 min). Then, the supernatant was analyzed with HPLC (DionexCarboPac™ PA-100 column, 4 × 250 mm, 0.1 mol/L ammonium acetate buffer eluent). The reaction was monitored by generation of the byproduct UDP, which had the strong UV absorption at 260 nm and could be easily quantitated by HPLC.

Investigating the influences of temperatures, pH values and metal cations on the activity of Cps1aI. For determination of temperature effect, the reactions were carried out at different temperatures (10, 16, 20, 25, 30, 37, 42, 50, 55 °C) in 50 mmol/L Tris-HCl buffer (pH 8.5). For determination of pH effect on enzymatic activity, pH values were varied from 5.5 to 10.5 with three different buffer systems at 25 °C. The buffer systems were as follows: Bis-Tris-HCl (50 mmol/L, pH 5.5, 6.0, 6.5, and 7.0), Tris-HCl (50 mmol/L, pH 7.0, 7.5, 8.0, 8.5, 9.0 and 9.5), and Gly-NaOH (50 mmol/L, pH 9.0, 9.5, 10.0 and 10.5). The effects of various metal ions on enzyme activity were assessed in the presence of the following metal salts including 5 mmol/L ethylenediamine tetraacetic acid (EDTA), MgCl₂, MnCl₂, CaCl₂, NiSO₄, CoSO₄, FeSO₄, CuSO₄ and ZnSO₄ in Tris-HCl buffer (pH 8.5) at 25 °C. To obtain the optimized Mg²⁺ concentration, the enzymatic reactions were carried out under varied concentrations of Mg²⁺ (0.3125-80 mM). Negative controls were performed in parallel under the same conditions using heat-deactivated Cps1aI. Relative activity concluded from pH and temperature test was defined as the relative value to the maximum enzyme activity, and the effect of metal ions on enzyme activity was determined using the activity measured without adding ions as the reference value.

Substrate specificity study of Cps1aI. For acceptor substrate specificity study, Lac α -PP-(CH₂)₁₁-OPh **2**, Lac, Lac-O(CH₂)₂N₃, Rha β -1,4-Glc α -PP-(CH₂)₁₁-OPh^[3], Gal α -1,3-Glc α -PP-(CH₂)₁₁-OPh^[4] and Glc α -1,3-Glc α -PP-(CH₂)₁₁-OPh^[4] were examined with UDP-GlcNAc as nucleotide donor, respectively. For donor substrate specificity study, UDP-Gal, UDP-Glc, UDP-GalNAc, UDP-GlcNAc and UDP-GlcA were examined with Lac α -PP-(CH₂)₁₁-OPh **2** as acceptor substrate, respectively. The reaction was performed in the optimized condition for Cps1aI.

Enzyme kinetics of Cps1aI. The enzymatic reactions were carried out under above-optimized conditions, i.e. in Tris-HCl buffer (50 mmol/L, pH 8.5) containing 12 μ g/mL of Cps1aI and 5 mmol/L MgCl₂ with varied concentrations of UDP-sugar (UDP-GlcNAc, UDP-Glc or UDP-GalNAc) and Lac α -PP-(CH₂)₁₁-OPh **2** at 25 °C. Then, reactions were performed for 10 min using saturated UDP-GlcNAc

(2 mmol/L) and varied concentrations of Lac α -PP-(CH₂)₁₁-OPh **2** (0.015625-2.0 mmol/L) or using Lac α -PP-(CH₂)₁₁-O-Ph **2** (1 mmol/L) and varied concentrations of UDP-sugar (0.0625-4.0 mmol/L). The results obtained were used to calculate the initial reaction velocities and to determine the Michaelis constant (K_m and k_{cat}) values using the GraphPad Prism 6.04 program.

Biochemical characterization of β -1,4-galactosyltransferase Cps1aJ and β -1,3-galactosyltransferase Cps1bJ

The reaction system for the detail characterization of purified Cps1aJ and Cps1bJ contained 1 mmol/L UDP-Gal, 1 mmol/L GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** and 100 μ g/mL of purified enzyme in 50 mmol/L buffer system. Reactions and detected methods followed those as mentioned above.

Investigating the influences of temperatures, pH values and metal cations on the activity of Cps1aJ and Cps1bJ. For determination of temperature effect, the reactions were carried out at different temperatures (10, 16, 20, 25, 30, 37, 42, 50, 55 °C) in 50 mmol/L Tris-HCl buffer (pH 9.0). For determination of pH effect on enzymatic activity, pH values were varied from 5.5 to 10.5 with the foregoing three different buffer systems at 30 °C. The effects of various metal ions on enzyme activity and the optimized Mg²⁺ concentrations were assessed in Tris-HCl buffer (pH 9.0) at 30 °C. Negative controls were performed in parallel under the same conditions using heat-deactivated Cps1aJ or Cps1bJ. Relative activity concluded from pH and temperature test was defined as the relative value to the maximum enzyme activity, and the effect of metal ions on enzyme activity was determined using the activity measured without adding ions as the reference value.

Substrate specificity study of Cps1aJ and Cps1bJ. For acceptor substrate specificity study of the two galactosyltransferases, GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a**, Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3b** and GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3c** were examined with UDP-Gal as nucleotide donor, respectively. For donor substrate specificity study, UDP-Gal, UDP-Glc, UDP-GalNAc, UDP-GlcNAc and UDP-GlcA were examined with GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** as acceptor substrate, respectively. The reactions were performed in the optimized condition for Cps1aJ or Cps1bJ.

Enzyme kinetics of Cps1aJ and Cps1bJ. The enzymatic reactions were carried out under above-optimized conditions, i.e. in Tris-HCl buffer (50 mmol/L, pH 9.0) containing 50 μ g/mL of Cps1aJ or Cps1bJ and 5 mmol/L MgCl₂ with varied concentrations of UDP-Gal and acceptor substrate (GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a**, Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3b**, and GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3c**) at 30 °C. Then, reactions were performed for 10 min using saturated UDP-Gal (2 mmol/L) and varied concentrations of receptor substrates (0.015625-2.0 mmol/L) or using GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** (1 mmol/L) and varied concentrations of UDP-Gal (0.0625-4.0 mmol/L). The results obtained were used to calculate the initial reaction velocities and determine the Michaelis constant (K_m and k_{cat}) values using the GraphPad Prism 6.04 program.

Biochemical characterization of α -2,3-sialyltransferases Cps1aK and Cps1bK. The reaction system for the detail characterization of purified Cps1aK and Cps1bK contained 1 mmol/L CMP-NeuNAc, 1 mmol/L Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh

4a for Cps1aK (20 μ g/mL) or Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b** for Cps1bK (200 μ g/mL) in 50 mmol/L buffer system. Reactions were allowed to proceed for 10 min, terminated by boiling at 100 °C for 30 s, and finally cleared by centrifugation (12,000 rpm, 10 min). Then, the supernatants were analyzed with HPLC to quantitate formation of pentasaccharide NeuNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5a** for Cps1aK or NeuNAc α -2,3-Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5b** for Cps1bK that had the strong UV absorption at 211 nm (C18 reverse phase column, 4.6 \times 250 mm, 10-100% methanol in water containing 10 mmol/L NH₄HCO₃ as gradient eluent).

Investigating the influences of temperatures, pH values and metal cations on the activity of Cps1aK and Cps1bK. For determination of temperature effect, the reactions were carried out at different temperatures in 50 mmol/L Tris-HCl buffer (pH 7.0). For determination of pH effect on enzymatic activity, pH values were varied from 5.5 to 10.5 with the aforementioned three different buffer systems at 25 °C. The effects of various metal ions on enzyme activity and the optimized Mg²⁺ concentrations were assessed in Tris-HCl buffer (pH 7.0) at 25 °C. Negative controls were performed in parallel under the same conditions using heat-deactivated Cps1aK or Cps1bK. Relative activity concluded from pH and temperature test was defined as the relative value to the maximum enzyme activity, and the effect of metal ions on enzyme activity was determined using the activity measured without adding ions as the reference value.

Acceptor substrate specificity study of Cps1aK and Cps1bK. For acceptor substrate specificity study of purified enzymes, Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a**, Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b**, Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4c**, Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4d**, Gal β -1,3-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4e**, Gal β -1,4-Rha β -1,4-Glc α -PP-(CH₂)₁₁-OPh^[31] and Lac α -PP-(CH₂)₁₁-OPh **2** were examined with CMP-NeuNAc as nucleotide donor, respectively.

Enzyme kinetics of Cps1aK and Cps1bK. The enzymatic reactions were carried out under above-optimized conditions, i.e. in Tris-HCl buffer (50 mmol/L, pH 7.0) containing 10 μ g/mL of Cps1aK or 100 μ g/mL of Cps1bK with varied concentrations of CMP-NeuNAc and acceptor substrate (**4a** and **4d** for Cps1aK, or **4c**, **4d**, and **4e** for Cps1bK) at 25 °C. Then, reactions were performed for 10 min using saturated CMP-NeuNAc (2 mmol/L) and varied concentrations of acceptor substrates (0.015625-2.0 mmol/L) or using corresponding acceptor substrate (1 mmol/L) and varied concentrations of CMP-NeuNAc (0.0625-4.0 mmol/L). The results obtained were used to calculate the initial reaction velocities and determine the Michaelis constant (K_m and k_{cat}) values using the GraphPad Prism 6.04 program.

Chemical synthesis of Lac α -PP-(CH₂)₁₁-OPh **2**.

Acetylation. A solution of Ac₂O (30 mL) and NaOAc (5 g) was heated up to 140°C (reflux), and lactose (5 g) was added in portion. The resulting mixture was stirred at 140°C for 1 h and poured into ice water (100 mL). The mixed solution was extracted with EtOAc (100 mL), washed by NaHCO₃ (aq) and NaCl (aq), dried over Na₂SO₄, filtered and concentrated. The yellow-oil product was recrystallized from MeOH/CH₂Cl₂ (550mL, v/v 10:1) to give 5.5 g white solid peracetylated lactose (LacAc₈).

Phosphorylation.^[5,6] Crystalline phosphoric acid (3 g) was dried *in vacuo* over phosphorous pentoxide for 12 h. The peracetylated lactose (3.4 g) was added and the mixture was heated at 60°C *in vacuo*. After 2 h, heating was ceased and the resulting dark black mixture was dissolved in anhydrous tetrahydrofuran (THF) (25 mL). The solution was cooled to 0°C and concentrated ammonium hydroxide (2.5 mL) was added until ~pH 7. The precipitate of ammonium phosphate was filtered off and washed with THF (100 mL). The combined filtrates were evaporated to give a syrupy residue that was purified by flash column chromatography (EtOAc 100% to EtOAc/MeOH 50/50). The peracetylated lactose-1-phosphate (LacAc7-1-P, 1.7 g) was obtained as a white solid.

Deacetylation. Sodium hydroxide (0.1 mol/L) was slowly dripped into MeOH (50 mL) containing LacAc7-1-P (1.0 g). The reaction was stirred at room temperature until white turbidity appeared, then 50 mL water and 1M sodium hydroxide was added to the reaction solution until ~pH 7. Fractions containing crude products in methanol and water mixed solvent were collected and concentrated, and the resultant supernatant was performed by gel filtration with Bio-Gel P2 and resultant product was collected and then lyophilized. The product lactose-1-phosphate disodium salt (Lac-1-P, 550 mg) was obtained as white powder.

MgCl₂-catalyzed diphosphate bond formation.^[3,7] A mixture of 11-phenoxyundecyl dihydrogen phosphate (550 mg, 1.6 mmol) and *N,N'*-carbonyldiimidazole (1.0 g, 6.4 mmol) in anhydrous THF (5 mL) was stirred at room temperature for 4 h, and then dry methanol (0.5 mL) was added. The resulting solution was stirred for another 1 h, and then concentrated to yield the crude 11-phenoxyundecyl dihydrogen phosphorimidazolide which was directly used for next step. The above generated product was added to a vigorously stirred suspension of Lac-1-P (500 mg, 1.05 mmol) and MgCl₂ (250 mg, 2.6 mmol) in *N,N*-dimethylformamide (DMF) (5.0 mL), and the resulting reaction mixture was then stirred for 8 h at room temperature, at which time TLC analysis (EtOAc:CH₃OH:H₂O:HOAc = 8:3:2:0.5 (*v/v/v/v*)) indicated the completion of reaction. Then, the reaction mixture was filtered and loaded on a Sephadex LH-20 column for purification using methanol as an eluent. Fractions containing the desired product were collected, concentrated, and further purified on a silica gel flash column chromatography using 10:3:2 EtOAc-CH₃OH-H₂O as eluents. The desired product Lac α -PP-(CH₂)₁₁-OPh **1** (650 mg, 81%) was obtained as white solid after lyophilization. ¹H NMR (600 MHz, D₂O): δ 7.36–7.32 (m, 2H, Ph), 7.02–6.67 (m, 3H, Ph), 5.50 (dd, *J* = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.41 (d, *J* = 7.8 Hz, 1H, H-1^{Gal}), 4.08–4.02 (m, 2H, -CH₂CH₂OPh), 3.99 (br d, *J* = 10.2, 1H, H-5^{Glc}), 3.91 (q, *J* = 6.6 Hz, 2H, -OCH₂CH₂-), 3.89–3.81 (m, 4H, H-3^{Glc}, H-6a,b^{Glc}, H-4^{Gal}), 3.78 (dd, *J* = 11.4, 7.8 Hz, 1H, H-6a^{Gal}), 3.73–3.65 (m, 3H, H-4^{Glc}, H-5,6a^{Gal}), 3.62 (dd, *J* = 10.2, 3.6 Hz, 1H, H-3^{Gal}), 3.56–3.50 (m, 2H, H-2^{Glc}, H-2^{Gal}), 1.77–1.69 (m, 2H, -CH₂CH₂-), 1.64–1.57 (m, 2H, -CH₂CH₂-), 1.44–1.37 (m, 2H, -CH₂CH₂-), 1.35–1.25 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 158.0, 129.7 (2C), 121.3, 114.8 (2C), 102.8 (C-1^{Gal}), 95.0 (d, *J*_{C,P} = 6.0 Hz, C-1^{Glc}), 77.5 (C-4^{Glc}), 75.2 (C-5^{Gal}), 72.4 (C-3^{Gal}), 71.4 (C-3^{Glc}), 71.3 (C-5^{Glc}), 71.2 (d, *J*_{C,P} = 7.5 Hz, C-2^{Glc}), 70.8 (C-2^{Gal}), 68.6 (-OCH₂CH₂OPh), 68.5 (C-4^{Gal}), 66.9 (d, *J*_{C,P} = 6.0 Hz, -OCH₂CH₂-), 60.9 (C-6^{Gal}), 59.5 (C-6^{Glc}), 29.7 (d, *J*_{C,P} = 7.5 Hz, -OCH₂CH₂-), 28.5 (3C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh); ³¹P NMR (243 MHz, D₂O): δ -10.70 (d, *J* = 19.4 Hz), -13.13 (d, *J* = 19.4 Hz); ESI-(-)-TOF HRMS *m/z*: Calcd for C₂₉H₅₀O₁₈P₂ 747.2400 [M - H]⁻; Found 747.2404.

Enzymatic synthesis of trisaccharides **3a, **3b** and **3c**.** Milligram-scale production of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a**, Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3b** and GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3c** were performed in a 10 mL

reaction system containing 2.4 mmol/L UDP-sugar (UDP-GlcNAc, UDP-Glc or UDP-GalNAc), 2 mmol/L Lac α -PP-(CH₂)₁₁-OPh **2**, 5 mmol/L MgCl₂ and 50 μ g/mL purified Cps1aI in 50 mmol/L Tris-HCl buffer (pH 8.5). The solutions were incubated at 25 °C for 2 h, 5 h or 12 h until the substrate **2** was completely converted to trisaccharide product as monitored by TLC analysis, and then quenched by boiling for 30 s. The mixtures were vortexed and centrifuged to remove the formed precipitate and the resultant supernatants were freeze-dried, re-suspended with methanol (5 mL) and filtered to remove the precipitation again. The filtrates were loaded on a semi-preparative HPLC (C18 reverse phase column: 10 \times 250 mm, 10-100% methanol in water containing 10 mmol/L NH₄HCO₃ as gradient eluent) in batches for purification. The fractions containing trisaccharide product were pooled and concentrated to produce **3a** (16.5 mg, 87%), **3b** (14.6 mg, 80%) and **3c** (15.8 mg, 83%) as white solid.

3a: ¹H NMR (600 MHz, D₂O): δ 7.37 (t, J = 7.8 Hz, 2H, Ph), 7.06-7.01 (m, 3H, Ph), 5.57 (br s, 1H, H-1^{Glc}), 4.70 (d, J = 8.4 Hz, 1H, H-1^{GlcNAc}), 4.42 (d, J = 7.8 Hz, 1H, H-1^{Gal}), 4.11 (d, J = 3.6 Hz, 1H, H-4^{Gal}), 4.09 (t, J = 6.0 Hz, 2H, -CH₂CH₂OPh), 4.01 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.96-3.90 (m, 2H, -OCH₂CH₂-), 3.90-3.83 (m, 4H, H-3^{Glc}, H-6a,b^{Glc}, H-6a^{GlcNAc}), 3.81-3.66 (m, 7H, H-4^{Glc}, H-3^{Gal}, H-5^{Gal}, H-6a,b^{Gal}, H-2^{GlcNAc}, H-6b^{GlcNAc}), 3.62-3.52 (m, 3H, H-2^{Glc}, H-2^{Gal}, H-3^{GlcNAc}), 3.32-3.27 (m, 2H, H-4,5^{GlcNAc}), 2.02 (s, 3H, CH₃CO), 1.78-1.72 (m, 2H, -CH₂CH₂-), 1.66-1.59 (m, 2H, -CH₂CH₂-), 1.46-1.40 (m, 2H, -CH₂CH₂-), 1.37-1.26 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 174.9 (CH₃CO), 158.0, 129.8 (2C), 121.3, 114.9 (2C), 102.8 (C-1^{Gal}), 102.7 (C-1^{GlcNAc}), 95.07 (d, $J_{C,P}$ = 6.0 Hz, C-1^{Glc}), 81.6 (C-3^{Gal}), 77.4 (C-4^{Glc}), 75.6 (C-5^{GlcNAc}), 74.8 (C-5^{Gal}), 73.5 (C-3^{GlcNAc}), 71.43 (C-3^{Glc}), 71.37 (C-5^{Glc}), 71.28 (d, $J_{C,P}$ = 7.5 Hz, C-2^{Glc}), 70.0 (C-2^{Gal}), 69.6 (C-4^{GlcNAc}), 68.7 (-OCH₂CH₂OPh), 68.4 (C-4^{Gal}), 66.9 (d, $J_{C,P}$ = 6.0 Hz, -OCH₂CH₂-), 60.9 (C-6^{Gal}), 60.4 (C-6^{GlcNAc}), 59.5 (C-6^{Glc}), 55.6 (C-2^{GlcNAc}), 29.7 (d, $J_{C,P}$ = 6.0 Hz, -OCH₂CH₂-), 28.57, 28.54, 28.52, 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 22.1 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.69 (d, J = 12.1 Hz), -13.12 (d, J = 12.1 Hz); ESI(-)-TOF HRMS *m/z*: Calcd for C₃₇H₆₂NO₂₃P₂ 950.3193 [M - H]⁻; Found 950.3190.

3b: ¹H NMR (600 MHz, D₂O): δ 7.35 (t, J = 7.8 Hz, 2H, Ph), 7.03-6.99 (m, 3H, Ph), 5.56 (br s, 1H, H-1^{Glc}), 4.63 (d, J = 7.8 Hz, 1H, H-1^{Glc'}), 4.47 (d, J = 7.8 Hz, 1H, H-1^{Gal}), 4.14 (d, J = 3.2 Hz, 1H, H-4^{Gal}), 4.06 (t, J = 6.6 Hz, 2H, -CH₂CH₂OPh), 3.99 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.94-3.89 (m, 2H, -OCH₂CH₂-), 3.89-3.81 (m, 4H, H-3^{Glc}, H-6a,b^{Glc}, H-6a^{Glc'}), 3.80-3.74 (m, 2H, H-3^{Gal}, H-6a^{Gal}), 3.73-3.66 (m, 5H, H-4^{Glc}, H-2,5,6b^{Gal}, H-6b^{Glc'}), 3.54 (br d, J = 9.6 Hz, 1H, H-2^{Glc}), 3.47 (t, J = 9.0 Hz, 1H, H-3^{Glc'}), 3.43-3.36 (m, 2H, H-4,5^{Glc'}), 3.20-3.16 (m, 1H, H-2^{Glc'}), 1.77-1.71 (m, 2H, -CH₂CH₂-), 1.64-1.57 (m, 2H, -CH₂CH₂-), 1.45-1.37 (m, 2H, -CH₂CH₂-), 1.36-1.24 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 158.0, 129.8 (2C), 121.3, 114.8 (2C), 103.6 (C-1^{Gal}), 102.4 (C-1^{Glc}), 95.1 (d, J = 6.0 Hz, C-1^{Glc}), 81.8 (C-3^{Gal}), 77.3 (C-4^{Glc}), 75.6 (C-5^{Glc}), 75.4 (C-3^{Glc'}), 74.8 (C-5^{Gal}), 73.1 (C-2^{Glc'}), 71.4 (C-3^{Glc}), 71.32 (C-5^{Glc}), 71.27 (d, $J_{C,P}$ = 7.5 Hz, C-2^{Glc}), 70.0 (C-2^{Gal}), 69.3 (C-4^{Glc'}), 68.6 (-OCH₂CH₂OPh), 68.2 (C-4^{Gal}), 66.9 (d, J = 6.0 Hz, -OCH₂CH₂-), 60.9 (C-6^{Gal}), 60.3 (C-6^{Glc'}), 59.5 (C-6^{Glc}), 29.7 (d, J = 7.5 Hz, -OCH₂CH₂-), 28.6, 28.5 (2C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh); ³¹P NMR (243 MHz, D₂O): δ -10.68 (d, J = 21.8 Hz), -13.11 (d, J = 21.8 Hz); ESI(-)-TOF HRMS *m/z*: Calcd for C₃₅H₅₉O₂₃P₂ 909.2928 [M - H]⁻; Found 909.2934.

3c: ¹H NMR (600 MHz, D₂O): δ 7.30 (t, J = 7.8 Hz, 2H, Ph), 6.99-6.94 (m, 3H, Ph), 5.50 (dd, J = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.54 (dd, J = 8.4, 1H, H-1^{GlcNAc}), 4.35 (dd, J = 7.8, 1H, H-1^{Gal}), 4.06 (d, J = 3.0 Hz, 1H, H-4^{Gal}), 4.01 (t, J = 6.6 Hz, 2H, -CH₂CH₂OPh), 3.95 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.90-3.76 (m, 7H, H-3^{Glc}, H-6a,b^{Glc}, H-2,4^{GlcNAc}, -OCH₂CH₂-), 3.74-3.60 (m, 8H, H-4^{Glc}, H-3,5^{Gal}, H-6a,b^{Gal},

H-3^{GalNAc}, H-6a,b^{GalNAc}), 3.60–3.56 (m, 1H, H-5^{GalNAc}), 3.54–3.48 (m, 2H, H-2^{Glc}, H-2^{Gal}), 1.96 (s, 3H, CH_3CO), 1.70–1.65 (m, 2H, - CH_2CH_2 -), 1.60–1.52 (m, 2H, - CH_2CH_2 -), 1.40–1.32 (m, 2H, - CH_2CH_2 -), 1.31–1.19 (m, 12H, - CH_2CH_2 -); ^{13}C NMR (150 MHz, D_2O): δ 175.0 (CH_3CO), 157.9, 129.7 (2C), 121.2, 114.7 (2C), 103.2 (C-1^{GalNAc}), 102.7 (C-1^{Gal}), 95.0 (d, J = 6.4 Hz, C-1^{Glc}), 81.3 (C-3^{Gal}), 77.2 (C-4^{Glc}), 74.8 (C-5^{GalNAc}), 74.7 (C-5^{Gal}), 71.3 (C-3^{Glc}), 71.3 (C-5^{Glc}), 71.2 (d, J = 9.0 Hz, C-2^{Glc}), 70.6 (C-3^{GalNAc}), 70.0 (C-2^{Gal}), 68.53 (- $\text{OCH}_2\text{CH}_2\text{OPh}$), 68.45 (C-4^{Gal}), 67.6 (C-4^{GalNAc}), 66.9 (d, J = 6.0 Hz, - OCH_2CH_2 -), 60.89 (C-6^{Gal}), 60.85 (C-6^{GalNAc}), 59.4 (C-6^{Glc}), 52.4 (C-2^{GalNAc}), 29.6 (d, J = 6.0 Hz, - OCH_2CH_2 -), 28.52, 28.49, 28.48, 28.32, 28.25, 28.15, 25.0, 24.7 (8C, - $\text{OCH}_2\text{CH}_2(\text{CH}_2)_8\text{CH}_2\text{OPh}$), 22.1 (CH_3CO); ^{31}P NMR (243 MHz, D_2O): δ -10.76 (d, J = 19.4 Hz), -13.17 (d, J = 19.4 Hz); ESI(-)-TOF HRMS m/z : Calcd for $\text{C}_{37}\text{H}_{62}\text{NO}_{23}\text{P}_2$ 950.3193 [M - H]⁺; Found 950.3194.

Enzymatic synthesis of tetrasccharides **4a**, **4b**, **4c**, **4d** and **4e**

Procedure A. Milligram-scale production systems of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH_2)₁₁-OPh **4a** and Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH_2)₁₁-OPh **4d** were 2 mL reaction mixture of 50 mmol/L Tris-HCl buffer (pH 8.5) containing 2.4 mmol/L UDP-Gal, 2 mmol/L **3a** or **3b**, 5 mM MgCl₂ and 200 $\mu\text{g}/\text{mL}$ Cps1aJ. Milligram-scale production systems of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH_2)₁₁-OPh **4b**, Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH_2)₁₁-OPh **4c** and Gal β -1,3-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH_2)₁₁-OPh **4e** were 2 mL reaction mixture of 50 mmol/L Tris-HCl buffer (pH 8.5) containing 2.4 mmol/L UDP-Gal, 2 mmol/L **3a**, **3b**, or **3c**, 5 mM MgCl₂ and 200 $\mu\text{g}/\text{mL}$ Cps1bJ. All of the reaction systems were incubated at 30 °C for 12 h, at which time TLC indicated the complete consumption of acceptor substrate, and then quenched by boiling for 30 s. After purification protocol as mentioned above, the fractions containing desired product were collected, then concentrated and lyophilized to afford **4a** (3.6 mg, 81%), **4d** (3.4 mg, 79%), **4b** (3.5 mg, 79%), **4c** (3.2 mg, 72%) and **4e** (3.3 mg, 77%) as white powders.

Procedure B. The one-pot two-enzyme synthesis of **4a**, **4b**, **4c**, **4d** and **4e** was achieved with Cps1aI and Cps1aJ or Cps1bJ using Lac α -PP-(CH_2)₁₁-OPh **2** as the starting acceptor substrate. A 5 mL reaction mixture of 50 mmol/L Tris-HCl buffer (pH 8.5) containing 2.4 mmol/L UDP-GlcNAc/UDP-Glc, 2.4 mmol/L UDP-Gal, 2 mmol/L **2**, 5 mmol/L MgCl₂, 50 $\mu\text{g}/\text{mL}$ Cps1aI and 200 mg/mL Cps1aJ to producing **4a/4d**. A 5 mL reaction mixture of 50 mmol/L Tris-HCl buffer (pH 8.5) containing 2.4 mmol/L UDP-GlcNAc, UDP-Glc or UDP-GalNAc, 2.4 mmol/L UDP-Gal, 2 mmol/L **2**, 5 mmol/L MgCl₂, 50 $\mu\text{g}/\text{mL}$ Cps1aI and 200 mg/mL Cps1bJ to producing **4b/4c/4e**. All of the reaction systems were incubated at 25 °C for 12 h, at which time TLC indicated the complete consumption of acceptor substrate, and then quenched by boiling for 30 s. After purification protocol as mentioned above, the fractions containing desired product were collected, then concentrated and lyophilized to afford **4a** (8.9 mg, 80%), **4d** (8.3 mg, 77%), **4b** (9.2 mg, 84%), **4c** (7.8 mg, 70%) and **4e** (7.9 mg, 74%) as white powders.

4a: ^1H NMR (600 MHz, D_2O): δ 7.37 (t, J = 7.8 Hz, 2H), 7.06–7.00 (m, 3H), 5.57 (dd, J = 6.6, 3.0 Hz, 1H, H-1^{Glc}), 4.69 (d, J = 8.4 Hz, 1H, H-1^{GalNAc}), 4.46 (d, J = 7.8 Hz, 1H, H-1^{Gal}'), 4.40 (d, J = 7.2 Hz, 1H, H-1^{Gal}), 4.12 (d, J = 3.0 Hz, 1H, H-4^{Gal}'), 4.08 (t, J = 6.4 Hz, 2H, - $\text{CH}_2\text{CH}_2\text{OPh}$), 4.01 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.96–3.81 (m, 8H, H-3^{Glc}, H-6a,b^{Glc}, H-4^{Gal}', H-6a,b^{GalNAc}, - OCH_2CH_2 -), 3.81–3.62 (m, 12H, H-4^{Glc}, H-3,5^{Gal}, H-6a,b^{Gal}, H-2,4,5^{GlcNAc}, H-3,5^{Gal}', H-6a,b^{Gal}'), 3.61–3.50 (m, 4H, H-2^{Glc}, H-2^{Gal}, H-3^{GlcNAc}, H-2^{Gal}'), 2.02 (s, 3H, CH_3CO), 1.79–1.72 (m, 2H, - CH_2CH_2 -), 1.66–1.59 (m, 2H, - CH_2CH_2 -), 1.46–1.40 (m, 2H, - CH_2CH_2 -), 1.36–1.25 (m, 12H, - CH_2CH_2 -); ^{13}C NMR (150 MHz, D_2O): δ 174.8 (CH_3CO), 158.0, 129.8 (2C), 121.3, 114.8 (2C), 102.7 (2C, C-1^{Gal}, C-1^{Gal}'), 102.6 (C-

¹GlcNAc), 95.0 (d, J = 6.0 Hz, C-1^{Glc}), 81.7 (C-3^{Gal}), 78.0 (C-4^{GlcNAc}), 77.4 (C-4^{Glc}), 75.2 (C-5^{GlcNAc}), 74.7 (C-5^{Gal}), 74.4 (C-3^{GlcNAc}), 72.4 (C-3^{Gal}'), 72.1 (C-5^{Gal}'), 71.4 (C-3^{Glc}), 71.34 (C-5^{Glc}), 71.25 (d, J = 7.5 Hz, C-2^{Glc}), 70.8 (C-2^{Gal}'), 69.9 (C-2^{Gal}), 68.6 (-OCH₂CH₂OPh), 68.44 (C-4^{Gal}'), 68.36 (C-4^{Gal}), 66.9 (d, J = 6.0 Hz, -OCH₂CH₂-), 60.9 (C-6^{Gal}), 60.87 (C-6^{Gal}'), 59.7 (C-6^{GlcNAc}), 59.5 (C-6^{Glc}), 55.1 (C-2^{GlcNAc}), 29.7 (d, J = 7.5 Hz, -OCH₂CH₂-), 28.5, 28.5 (2C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 22.1 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.70 (d, J = 21.8 Hz), -13.13 (d, J = 21.8 Hz); ESI-(-)-TOF HRMS *m/z*: Calcd for C₄₃H₇₂NO₂₈P₂ 1112.3722 [M - H]⁻; Found 1112.3721.

4b: ¹H NMR (600 MHz, D₂O): δ 7.37 (t, J = 7.8 Hz, 2H, Ph), 7.06–7.01 (m, 3H, Ph), 5.57 (br s, 1H, H-1^{Glc}), 4.73 (d, J = 8.4 Hz, 1H, H-1^{GlcNAc}), 4.43 (d, J = 7.8 Hz, 2H, H-1^{Gal}, H-1^{Gal}'), 4.12 (d, J = 3.6 Hz, 1H, H-4^{Glc}), 4.09 (t, J = 6.6 Hz, 2H, -CH₂CH₂OPh), 4.02 (br d, J = 10.8 Hz, 1H, H-5^{Glc}), 3.97–3.83 (m, 8H, H-3^{Glc}, H-6a,b^{Glc}, H-2,6a^{GlcNAc}, H-4^{Gal}'), -OCH₂CH₂-), 3.83–3.67 (m, 10H, H-4^{Glc}, H-3,5^{Gal}, H-6a,b^{Gal}, H-3,6b^{GlcNAc}, H-5^{Gal}', H-6a,b^{Gal}'), 3.65–3.45 (m, 6H, H-2^{Glc}, H-2^{Gal}, H-4,5^{GlcNAc}, H-2,3^{Gal}'), 2.02 (s, 3H, CH₃CO), 1.79–1.73 (m, 2H, -CH₂CH₂-), 1.67–1.60 (m, 2H, -CH₂CH₂-), 1.47–1.41 (m, 2H, -CH₂CH₂-), 1.39–1.26 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 174.9 (CH₃CO), 158.0, 129.8 (2C), 121.3, 114.9 (2C), 103.4 (C-1^{Gal}'), 102.7 (C-1^{Gal}), 102.4 (C-1^{GlcNAc}), 95.1 (d, J = 6.0 Hz, C-1^{Glc}), 81.9 (C-3^{GlcNAc}), 81.6 (C-3^{Gal}), 77.4 (C-4^{Glc}), 75.2 (C-5^{Gal}'), 75.1 (C-5^{GlcNAc}), 74.8 (C-5^{Gal}), 72.3 (C-3^{Gal}'), 71.42 (C-3^{Glc}), 71.36 (C-5^{Glc}), 71.3 (d, J = 9.0 Hz, C-2^{Glc}), 70.6 (C-2^{Gal}'), 70.0 (C-2^{Gal}), 68.6 (-OCH₂CH₂OPh), 68.43 (C-4^{Gal}'), 68.37 (2C, C-4^{Gal}, C-4^{GlcNAc}), 66.9 (d, J = 6.0 Hz, -OCH₂CH₂-), 60.97 (C-6^{Gal}), 60.93 (C-6^{Gal}'), 60.4 (C-6^{GlcNAc}), 59.5 (C-6^{Glc}), 54.6 (C-2^{GlcNAc}), 29.7 (d, J = 7.5 Hz, -OCH₂CH₂-), 28.5 (3C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 22.1 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.70 (d, J = 12.1 Hz), -13.12 (d, J = 12.1 Hz); ESI-(-)-TOF HRMS *m/z*: Calcd for C₄₃H₇₂NO₂₈P₂ 1112.3722 [M - H]⁻; Found 1112.3728.

4c: ¹H NMR (600 MHz, D₂O): δ 7.30 (t, J = 7.8 Hz, 2H, Ph), 7.00–6.93 (m, 3H, Ph), 5.50 (dd, J = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.60 (dd, J = 8.4, 1H, H-1^{GlcNAc}), 4.38–4.34 (m, 2H, H-1^{Gal}, H-1^{Gal}'), 4.10 (d, J = 3.6 Hz, 1H, H-4^{GlcNAc}), 4.05 (d, J = 3.0 Hz, 1H, H-4^{Gal}), 4.01 (t, J = 6.0 Hz, 2H, -CH₂CH₂OPh), 4.00–3.92 (m, 2H, H-5^{Glc}, H-2^{GlcNAc}), 3.90–3.75 (m, 7H, H-3^{Glc}, H-6a,b^{Glc}, H-3^{GlcNAc}, H-4^{Gal}', -OCH₂CH₂-), 3.74–3.59 (m, 10H, H-4^{Glc}, H-3,5^{Gal}, H-6a,b^{Gal}, H-6a,b^{GlcNAc}, H-5^{Gal}'), H-6a,b^{Gal}'), 3.59–3.47 (m, 4H, H-2^{Glc}, H-2^{Gal}, H-5^{GlcNAc}, H-3^{Gal}'), 3.44 (t, J = 9.0 Hz, 1H, H-2^{Gal}'), 1.95 (s, 3H, CH₃CO), 1.72–1.66 (m, 2H, -CH₂CH₂-), 1.60–1.52 (m, 2H, -CH₂CH₂-), 1.40–1.33 (m, 2H, -CH₂CH₂-), 1.31–1.19 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 175.0 (CH₃CO), 157.9, 129.7 (2C), 121.2, 114.7 (2C), 104.7 (C-1^{Gal}'), 102.9 (C-1^{GlcNAc}), 102.7 (C-1^{Gal}), 95.0 (d, J = 6.4 Hz, C-1^{Glc}), 81.2 (C-3^{Gal}), 79.3 (C-3^{GlcNAc}), 77.2 (C-4^{Glc}), 74.8 (C-5^{GlcNAc}), 74.7 (C-5^{Gal}), 74.5 (C-5^{Gal}'), 72.2 (C-3^{Gal}'), 71.32 (C-3^{Glc}), 71.29 (C-5^{Glc}), 71.2 (d, J = 7.5 Hz, C-2^{Glc}), 70.4 (C-2^{Gal}'), 70.0 (C-2^{Gal}), 68.5 (-OCH₂CH₂OPh), 68.44 (C-4^{Gal}), 68.38 (C-4^{Gal}'), 67.8 (C-4^{GlcNAc}), 66.9 (d, J = 6.0 Hz, -OCH₂CH₂-), 60.8 (2C, C-6^{Gal}, C-6^{GlcNAc}), 60.7 (C-6^{Gal}'), 59.4 (C-6^{Glc}), 51.2 (C-2^{GlcNAc}), 29.64 (d, J = 7.5 Hz, -OCH₂CH₂-), 28.5 (3C), 28.3, 28.2, 28.1, 24.9, 24.7 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 22.1 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.78 (d, J = 19.4 Hz), -13.18 (d, J = 19.4 Hz); ESI-(-)-TOF HRMS *m/z*: Calcd for C₄₃H₇₂NO₂₈P₂ 1112.3722 [M - H]⁻; Found 1112.3728.

4d: ¹H NMR (600 MHz, D₂O): δ 7.34 (t, J = 7.8 Hz, 2H, Ph), 7.03–6.96 (m, 3H, Ph), 5.55 (dd, J = 6.6, 3.0 Hz, 1H, H-1^{Glc}), 4.65 (d, J = 7.8 Hz, 1H, H-1^{Glc}'), 4.46 (d, J = 7.8 Hz, 1H, H-1^{Gal}), 4.40 (d, J = 7.8 Hz, 1H, H-1^{Gal}'), 4.13 (d, J = 2.4 Hz, 1H, H-4^{Gal}), 4.06 (t, J = 6.6 Hz, 2H, -CH₂CH₂OPh), 3.99 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.94–3.81 (m, 7H, H-3^{Glc}, H-6a,b^{Glc}, H-6a^{Glc}', H-4^{Gal}', -OCH₂CH₂-), 3.80–3.74 (m, 4H, H-3,6a^{Gal}, H-6b^{Glc}', H-6a^{Gal}'), 3.74–3.59 (m, 9H, H-4^{Glc}, H-2,5,6b^{Gal}, H-3,4^{Glc}', H-3,5,6b^{Gal}'), 3.56–3.47 (m, 3H, H-2^{Glc}, H-5^{Glc}', H-2^{Gal}'), 3.38 (t, J = 8.4 Hz, 1H, H-2^{Glc}'), 1.77–1.69 (m, 2H, -CH₂CH₂-), 1.64–1.57 (m, 2H, -CH₂CH₂-), 1.44–1.37 (m, 2H, -

*CH₂CH₂-), 1.35–1.23 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 158.0, 129.8 (2C), 121.3, 114.8 (2C), 103.5 (C-1^{Glc'}), 102.8 (C-1^{Gal'}), 102.4 (C-1^{Gal}), 95.1 (d, *J* = 6.0 Hz, C-1^{Glc}), 81.9 (C-3^{Gal}), 77.9 (C-4^{Glc'}), 77.3 (C-4^{Glc}), 75.2 (C-5^{Gal'}), 74.8 (C-5^{Gal}), 74.5 (C-5^{Glc'}), 74.0 (C-3^{Glc'}), 72.9 (C-2^{Glc'}), 72.4 (C-3^{Gal'}), 71.4 (C-3^{Glc}), 71.31 (C-5^{Glc}), 71.27 (d, *J* = 9.0 Hz, C-2^{Glc}), 70.8 (C-2^{Gal'}), 70.0 (C-2^{Gal}), 68.6 (-OCH₂CH₂OPh), 68.4 (C-4^{Gal'}), 68.2 (C-4^{Gal}), 66.9 (d, *J* = 6.0 Hz, -OCH₂CH₂-), 60.9 (2C, C-6^{Gal}, C-6^{Gal'}), 59.7 (C-6^{Glc'}), 59.5 (C-6^{Glc}), 29.70 (d, *J* = 7.5 Hz, -OCH₂CH₂-), 28.6, 28.5 (2C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh); ³¹P NMR (243 MHz, D₂O): -10.74 (d, *J* = 21.8 Hz), -13.18 (d, *J* = 21.8 Hz); ESI-(-)-TOF HRMS *m/z*: Calcd for C₄₁H₆₉O₂₈P₂ 1071.3456 [M - H]⁻; Found 1071.3450.*

4e: ¹H NMR (600 MHz, D₂O): δ 7.33 (t, *J* = 7.8 Hz, 2H, Ph), 7.02–6.96 (m, 3H, Ph), 5.53 (dd, *J* = 6.6, 3.6 Hz, 1H, H-1^{Glc}), 4.65 (d, *J* = 7.8 Hz, 1H, H-1^{Glc'}), 4.62 (d, *J* = 7.8 Hz, 1H, H-1^{Gal'}), 4.45 (d, *J* = 7.8 Hz, 1H, H-1^{Gal}), 4.11 (d, *J* = 2.4 Hz, 1H, H-4^{Gal}), 4.04 (t, *J* = 6.4 Hz, 2H, -CH₂CH₂OPh), 3.97 (br d, *J* = 10.2 Hz, 1H, H-5^{Glc}), 3.92–3.80 (m, 7H, H-3^{Glc}, H-6a,b^{Glc}, H-6a^{Glc'}, H-4^{Gal'}, -OCH₂CH₂-), 3.79–3.63 (m, 11H, H-4^{Glc}, H-2,3,5^{Gal}, H-6a,6b^{Gal}, H-3,6b^{Glc'}, H-5^{Gal'}, H-6a,6b^{Gal'}), 3.61 (dd, *J* = 9.6, 3.0 Hz, 1H, H-3^{Gal'}), 3.55–3.45 (m, 4H, H-2^{Glc}, H-2,4^{Glc'}, H-2^{Gal'}), 3.44–3.39 (m, 1H, H-5^{Glc'}), 1.75–1.68 (m, 2H, -CH₂CH₂-), 1.62–1.55 (m, 2H, -CH₂CH₂-), 1.43–1.35 (m, 2H, -CH₂CH₂-), 1.34–1.21 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 157.9, 129.7 (2C), 121.2, 114.7 (2C), 103.4 (C-1^{Glc'}), 103.1 (C-1^{Gal'}), 102.3 (C-1^{Gal}), 95.0 (d, *J* = 6.0 Hz, C-1^{Glc}), 83.8 (C-3^{Glc'}), 81.8 (C-3^{Gal}), 77.1 (C-4^{Glc}), 75.1 (2C, C-5^{Glc'}, C-5^{Gal'}), 74.8 (C-5^{Gal}), 72.9 (C-2^{Glc'}), 72.4 (C-3^{Gal'}), 71.3 (C-3^{Glc}), 71.3 (C-5^{Glc}), 71.2 (d, *J* = 9.0 Hz, C-2^{Glc}), 71.0 (C-2^{Gal'}), 69.9 (C-2^{Gal}), 68.5 (-OCH₂CH₂OPh), 68.4 (C-4^{Gal'}), 68.1 (C-4^{Gal}), 67.8 (C-4^{Glc'}), 66.8 (d, *J* = 6.0 Hz, -OCH₂CH₂-), 60.9 (C-6^{Gal}), 60.8 (C-6^{Gal'}), 60.2 (C-6^{Glc'}), 59.4 (C-6^{Glc}), 29.63 (d, *J* = 6.0 Hz, -OCH₂CH₂-), 28.50, 28.49 (2C), 28.3, 28.2, 28.1, 24.9, 24.7 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh); ³¹P NMR (243 MHz, D₂O): δ -10.74 (d, *J* = 21.8 Hz), -13.17 (d, *J* = 21.8 Hz); ESI-(-)-TOF HRMS *m/z*: Calcd for C₄₁H₆₉O₂₈P₂ 1071.3456 [M - H]⁻; Found 1071.3446.

Enzymatic synthesis of pentasaccharides **5a**, **5b**, **5c**, **5d**, and **5e**

Procedure A. Milligram-scale production systems of NeuNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5a** and NeuNAc α -2,3-Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5d** were 2 mL reaction mixture of 50 mmol/L Tris-HCl buffer (pH 7.0) containing 3 mmol/L CMP-NeuNAc, 2 mmol/L **4a** or **4d** and 40 μ g/mL Cps1aK. Milligram-scale production systems of NeuNAc α -2,3-Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5b**, NeuNAc α -2,3-Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **1c**, and NeuNAc α -2,3-Gal β -1,3-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **1e** were 2 mL reaction mixture of 50 mmol/L Tris-HCl buffer (pH 7.0) containing 3 mmol/L CMP-NeuNAc, 2 mmol/L **4b**, **4c**, or **4e** and 400 μ g/mL Cps1bJ. All of the reaction systems were incubated at 25 °C for 2 h, at which time TLC indicated the complete consumption of acceptor substrate, and then quenched by boiling for 30 s. After purification protocol as mentioned above, the fractions containing desired product were collected, then concentrated and lyophilized to afford **5a** (4.8 mg, 86%), **5d** (4.5 mg, 81%), **5b** (4.6 mg, 82%), **5c** (4.5 mg, 80%) and **5e** (4.6 mg, 84%) as white powders.

Procedure B. The one-pot three-enzyme synthesis of **5a**, **5b**, **5c**, **5d** and **5e** was carried out with Cps1aI, Cps1aJ/Cps1bJ and Cps1aK/Cps1bK enzymes using Lac α -PP-(CH₂)₁₁-OPh **2** as the starting acceptor substrate. A 5 mL reaction mixture of 50 mmol/L Tris-HCl buffer (pH 7.5) containing 2.4 mmol/L UDP-GlcNAc or UDP-Glc, 2.4 mmol/L UDP-Gal, 3 mM CMP-NeuNAc, 2 mmol/L **2**, 5 mmol/L MgCl₂, 50 μ g/mL Cps1aI, 200 mg/mL Cps1aJ and 40 μ g/mL Cps1aK to produce **5a**/**5d**. A 5 mL reaction mixture of 50 mmol/L

Tris-HCl buffer (pH 7.5) containing 2.4 mmol/L UDP-GlcNAc, UDP-GalNAc, or UDP-Glc, 2.4 mmol/L UDP-Gal, 3 mM CMP-NeuNAc, 2 mmol/L **2**, 5 mmol/L MgCl₂, 50 µg/mL Cps1aI, 200 mg/mL Cps1bJ and 400 µg/mL Cps1bK to producing **5b/5c/5e**. All of the reaction systems were incubated at 25 °C for 2 h, at which time TLC indicated the complete consumption of acceptor substrate, and then quenched by boiling for 30 s. After purification protocol as mentioned above, the fractions containing desired product were collected, then concentrated and lyophilized to afford **5a** (11.6 mg, 83%), **5d** (10.9 mg, 80%), **5b** (11.2 mg, 80%), **5c** (11.0 mg, 78%), and **5e** (10.5 mg, 77%) as white powders.

5a: ¹H NMR (600 MHz, D₂O): δ 7.36 (t, *J* = 7.8 Hz, 2H, Ph), 7.05–7.00 (m, 3H, Ph), 5.56 (dd, *J* = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.68 (d, *J* = 8.4 Hz, 1H, H-1^{GlcNAc}), 4.53 (d, *J* = 7.8 Hz, 1H, H-1^{Gal}'), 4.41 (d, *J* = 7.8 Hz, 1H, H-1^{Gal}), 4.12 (d, *J* = 3.2 Hz, 1H, H-4^{Gal}), 4.10 (dd, *J* = 9.6, 3.0 Hz, 1H, H-3^{Gal}'), 4.07 (t, *J* = 6.6 Hz, 2H, -CH₂CH₂OPh), 4.00 (br d, *J* = 10.2 Hz, 1H, H-5^{Glc}), 3.96–3.81 (m, 11H, H-3^{Glc}, H-6a,b^{Glc}, H-6a,b^{GlcNAc}, H-4^{Gal}'), H-5,8,9a^{NeuNAc}, -OCH₂CH₂-), 3.80–3.65 (m, 12H, H-4^{Glc}, H-3,5^{Gal}, H-6a,b^{Gal}, H-2,3,4^{GlcNAc}, H-5^{Gal}'), H-6a,b^{Gal}', H-4^{NeuNAc}), 3.65–3.53 (m, 7H, H-2^{Glc}, H-2^{Gal}, H-5^{GlcNAc}, H-2^{Gal}'), H-6,7,9b^{NeuNAc}), 2.74 (dd, *J* = 12.0, 4.2 Hz, 1H, H-3_{eq}^{NeuNAc}), 2.01 (s, 6H, CH₃CO), 1.81–1.72 (m, 3H, H-3_{ax}^{NeuNAc}, -CH₂CH₂-), 1.65–1.59 (m, 2H, -CH₂CH₂-), 1.46–1.39 (m, 2H, -CH₂CH₂-), 1.37–1.24 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 174.9 (CH₃CO), 174.8 (CH₃CO), 173.8 (C-1^{NeuNAc}), 158.0, 129.8 (2C), 121.3, 114.9 (2C), 102.8 (C-1^{Gal}'), 102.7 (C-1^{GlcNAc}), 102.5 (C-1^{Gal}'), 99.7 (C-2^{NeuNAc}), 95.1 (d, *J* = 6.0 Hz, C-1^{Glc}), 81.8 (C-3^{Gal}), 77.9 (C-4^{GlcNAc}), 77.5 (C-4^{Glc}), 75.4 (C-3^{Gal}'), 75.1 (C-5^{Gal}'), 74.8 (C-5^{Glc}), 74.5 (C-5^{GlcNAc}), 72.8 (C-6^{NeuNAc}), 72.08 (C-3^{GlcNAc}), 71.7 (C-8^{NeuNAc}), 71.44 (C-3^{Glc}), 71.37 (C-5^{Glc}), 71.26 (d, *J* = 9.0 Hz, C-2^{Glc}), 69.9 (C-2^{Gal}), 69.3 (C-2^{Gal}'), 68.7 (-OCH₂CH₂OPh), 68.4 (C-4^{Gal}), 68.3 (C-4^{NeuNAc}), 68 (C-7^{NeuNAc}), 67.4 (C-4^{Gal}'), 66.9 (d, *J* = 6.0 Hz, -OCH₂CH₂-), 62.5 (C-9^{NeuNAc}), 60.93 (C-6^{Gal}), 60.91 (C-6^{Gal}'), 59.7 (C-6^{GlcNAc}), 59.5 (C-6^{Glc}), 55.1 (C-2^{GlcNAc}), 51.6 (C-5^{NeuNAc}), 39.5 (C-3^{NeuNAc}), 29.72 (d, *J* = 6.0 Hz, -OCH₂CH₂-), 28.6, 28.5 (2C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 22.1 (CH₃CO), 21.9 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.68 (d, *J* = 17.0 Hz), -13.09 (d, *J* = 17.0 Hz); ESI(-)-TOF HRMS *m/z*: [M - 2H]²⁻ Calcd for C₅₄H₈₈N₂O₃₆P₂ 701.2301; Found 701.2308.

5b: ¹H NMR (600 MHz, D₂O): δ 7.36 (t, *J* = 7.8 Hz, 2H, Ph), 7.05–7.00 (m, 3H, Ph), 5.56 (dd, *J* = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.72 (d, *J* = 8.4 Hz, 1H, H-1^{GlcNAc}), 4.48 (d, *J* = 7.8 Hz, 1H, H-1^{Gal}'), 4.41 (d, *J* = 7.8 Hz, 1H, H-1^{Gal}), 4.11 (d, *J* = 3.2 Hz, 1H, H-4^{Gal}), 4.09–4.05 (m, 3H, H-3^{Gal}', -CH₂CH₂OPh), 4.00 (br d, *J* = 10.2 Hz, 1H, H-5^{Glc}), 3.95–3.90 (m, 3H, H-4^{Gal}', -OCH₂CH₂-), 3.90–3.80 (m, 8H, H-3^{Glc}, H-6a,b^{Glc}, H-2,6a^{GlcNAc}, H-5,8,9a^{NeuNAc}), 3.80–3.73 (m, 3H, H-6a^{Gal}, H-3,6b^{GlcNAc}), 3.73–3.64 (m, 8H, H-4^{Glc}, H-3,5,6b^{Gal}, H-5^{Gal}'), H-6a,b^{Gal}', H-4^{NeuNAc}), 3.64–3.49 (m, 7H, H-2^{Glc}, H-2^{Gal}, H-4^{GlcNAc}, H-2^{Gal}'), H-6,7,9b^{NeuNAc}), 3.47–3.43 (m, 1H, H-5^{GlcNAc}), 2.73 (dd, *J* = 12.6, 4.8 Hz, 1H, H-3_{eq}^{NeuNAc}), 2.01 (s, 3H, CH₃CO), 2.00 (s, 3H, CH₃CO), 1.79–1.72 (m, 3H, H-3_{ax}^{NeuNAc}, -CH₂CH₂-), 1.65–1.59 (m, 2H, -CH₂CH₂-), 1.46–1.40 (m, 2H, -CH₂CH₂-), 1.36–1.26 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 174.8 (2C, CH₃CO), 173.8 (C-1^{NeuNAc}), 158.0, 129.8 (2C), 121.3, 114.9 (2C), 103.3 (C-1^{Gal}'), 102.8 (C-1^{Gal}), 102.4 (C-1^{GlcNAc}), 99.5 (C-2^{NeuNAc}), 95.0 (d, *J* = 4.5 Hz, C-1^{Glc}), 82.1 (C-3^{GlcNAc}), 81.6 (C-3^{Gal}), 77.5 (C-4^{Glc}), 75.5 (C-3^{Gal}'), 75.1 (C-5^{GlcNAc}), 74.9 (C-5^{Gal}'), 74.8 (C-5^{Gal}), 72.7 (C-6^{NeuNAc}), 71.7 (C-8^{NeuNAc}), 71.4 (C-3^{Glc}), 71.3 (C-5^{Glc}), 71.24 (d, *J* = 7.5 Hz, C-2^{Glc}), 69.97 (C-2^{Gal}), 68.98 (C-2^{Gal}'), 68.6 (-OCH₂CH₂OPh), 68.4 (C-4^{GlcNAc}), 68.33 (C-4^{Gal}), 68.31 (C-4^{NeuNAc}), 67.9 (C-7^{NeuNAc}), 67.1 (C-4^{Gal}'), 66.9 (d, *J* = 4.5 Hz, -OCH₂CH₂-), 62.3 (C-9^{NeuNAc}), 60.90 (C-6^{Gal}), 60.87 (C-6^{Gal}'), 60.4 (C-6^{GlcNAc}), 59.5 (C-6^{Glc}), 54.4 (C-2^{GlcNAc}), 51.6 (C-5^{NeuNAc}), 39.6 (C-3^{NeuNAc}), 29.7 (d, *J* = 7.5 Hz, -OCH₂CH₂-), 28.5, 28.5 (2C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 22.2 (CH₃CO), 21.9 (CH₃CO); ³¹P NMR

(243 MHz, D₂O): δ -10.67 (d, J = 21.8 Hz), -13.10 (d, J = 21.8 Hz); ESI-(-)-TOF HRMS m/z : [M - 2H]²⁻ Calcd for C₅₄H₈₈N₂O₃₆P₂ 701.2301; Found 701.2295.

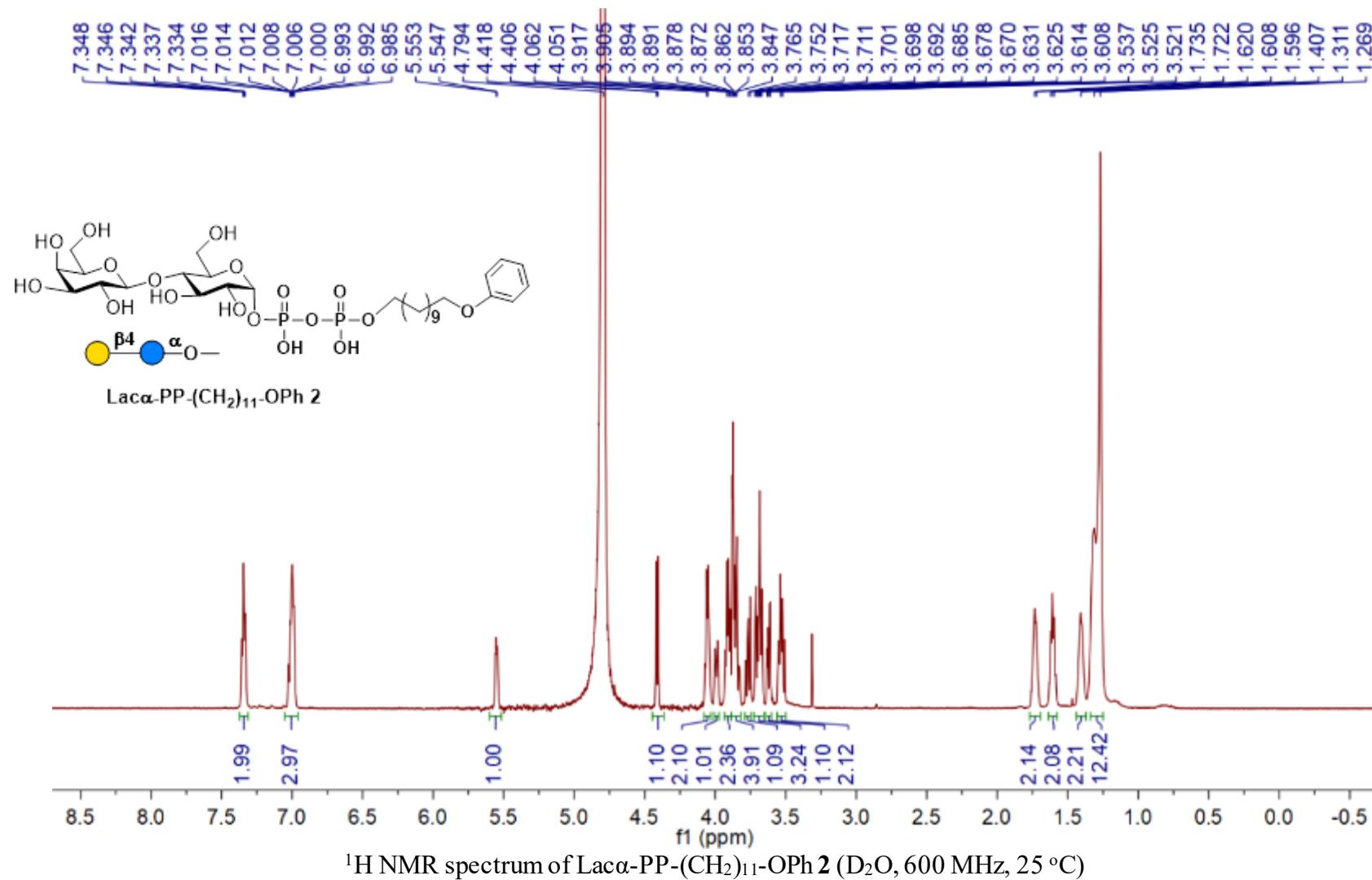
5c: ¹H NMR (600 MHz, D₂O): δ 7.35 (t, J = 7.8 Hz, 2H, Ph), 7.03–6.98 (m, 3H, Ph), 5.55 (dd, J = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.65 (d, J = 8.4 Hz, 1H, H-1^{GalNAc}), 4.48 (d, J = 7.8 Hz, 1H, H-1^{Gal}'), 4.40 (d, J = 7.8 Hz, 1H, H-1^{Gal}), 4.14 (d, J = 1.8 Hz, 1H, H-4^{GalNAc}), 4.09 (d, J = 2.4 Hz, 1H, H-4^{Gal}), 4.08–4.01 (m, 4H, H-2^{GlcNAc}, H-3^{Gal}', -CH₂CH₂OPh), 3.99 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.94–3.78 (m, 10H, H-3^{Glc}, H-6a,b^{Glc}, H-3^{GalNAc}, H-4^{Gal}', H-5,8,9a^{NeuNAc}, -OCH₂CH₂-), 3.78–3.61 (m, 11H, H-4^{Glc}, H-3,5^{Gal}, H-6a,b^{Gal}, H-6a,b^{GalNAc}, H-5^{Gal}', H-6a,b^{Gal}', H-4^{NeuNAc}), 3.61–3.47 (m, 7H, H-2^{Glc}, H-2^{Gal}, H-5^{GalNAc}, H-2^{Gal}', H-6,7,9b^{NeuNAc}), 2.71 (dd, J = 12.0, 4.2 Hz, 1H, H-3_{eq}^{NeuNAc}), 1.99 (s, 6H, CH₃CO), 1.78–1.70 (m, 3H, H-3_{ax}^{NeuNAc}, -CH₂CH₂-), 1.64–1.57 (m, 2H, -CH₂CH₂-), 1.44–1.37 (m, 2H, -CH₂CH₂-), 1.35–1.23 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 175.0 (CH₃CO), 174.8 (CH₃CO), 173.9 (C-1^{NeuNAc}), 158.0, 129.8 (2C), 121.3, 114.8 (2C), 104.46 (C-1^{Gal}'), 102.9 (C-1^{GlcNAc}), 102.8 (C-1^{Gal}), 99.5 (C-2^{NeuNAc}), 95.1 (d, J = 7.5 Hz, C-1^{Glc}), 81.3 (C-3^{Gal}), 79.6 (C-3^{GalNAc}), 77.4 (C-4^{Glc}), 75.4 (C-3^{Gal}'), 74.8 (C-5^{GalNAc}), 74.6 (C-5^{Gal}), 74.5 (C-5^{Gal}'), 72.7 (C-6^{NeuNAc}), 71.7 (C-8^{NeuNAc}), 71.4 (C-3^{Glc}), 71.3 (C-5^{Glc}), 71.2 (d, J = 7.5 Hz, C-2^{Glc}), 70.0 (C-2^{Gal}), 68.9 (C-2^{Gal}'), 68.6 (-OCH₂CH₂OPh), 68.5 (C-4^{Gal}), 68.3 (C-4^{NeuNAc}), 67.9 (C-7^{NeuNAc}), 67.7 (C-4^{GalNAc}), 67.2 (C-4^{Gal}'), 66.9 (d, J = 6.0 Hz, -OCH₂CH₂-), 62.3 (C-9^{NeuNAc}), 60.9 (2C, C-6^{Gal}, C-6^{GalNAc}), 60.8 (C-6^{Gal}'), 59.5 (C-6^{Glc}), 51.5 (C-5^{NeuNAc}), 51.2 (C-2^{GalNAc}), 39.6 (C-3^{NeuNAc}), 29.7 (d, J = 7.5 Hz, -OCH₂CH₂-), 28.54, 28.51, 28.50, 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 22.2 (CH₃CO), 21.9 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.73 (d, J = 19.4 Hz), -13.16 (d, J = 19.4 Hz); ESI-(-)-TOF HRMS m/z : [M - 2H]²⁻ Calcd for C₅₄H₈₈N₂O₃₆P₂ 701.2301; Found 701.2291.

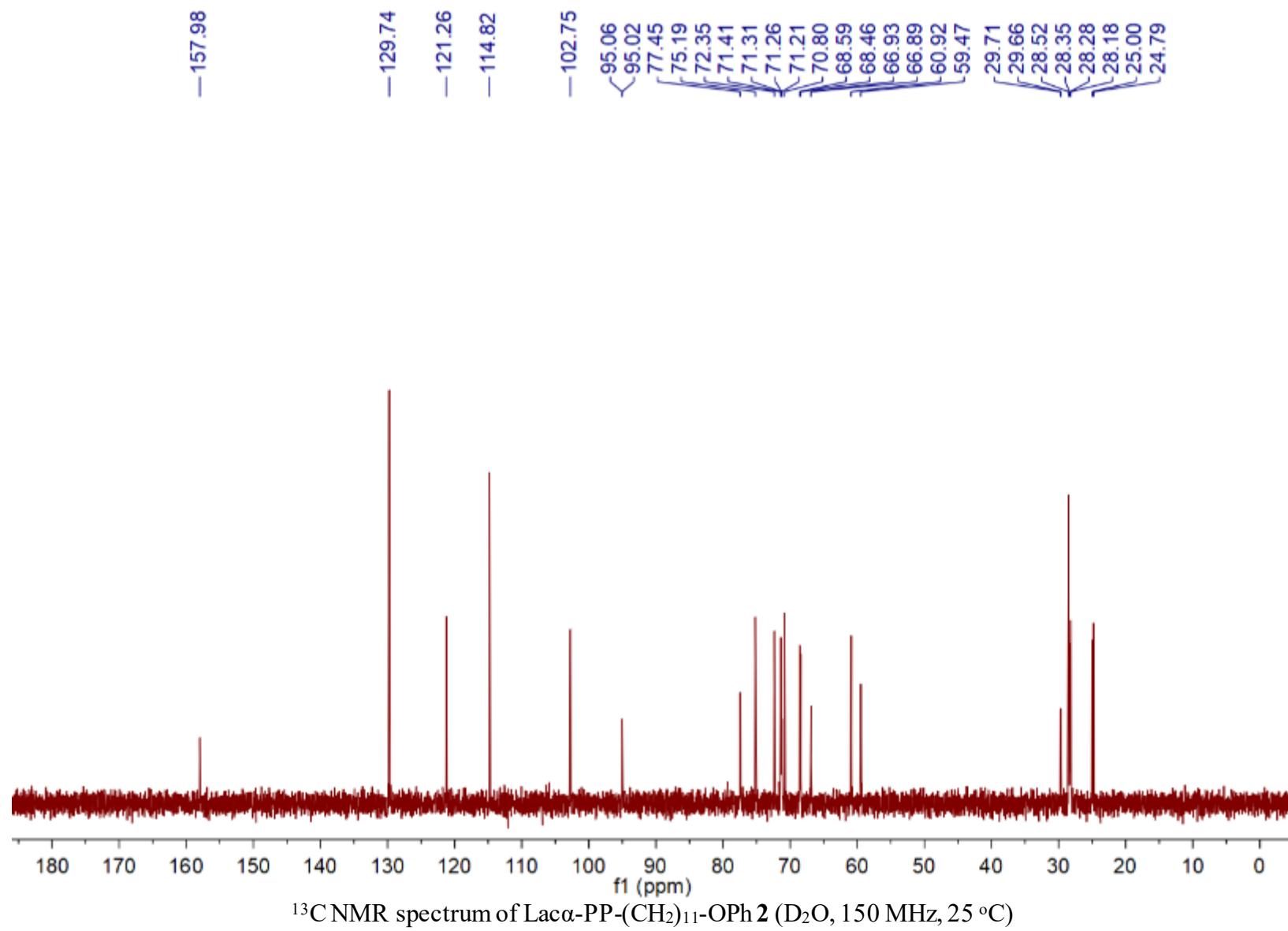
5d: ¹H NMR (600 MHz, D₂O): δ 7.32 (t, J = 7.8 Hz, 2H, Ph), 7.01–6.95 (m, 3H, Ph), 5.52 (dd, J = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.62 (d, J = 8.4 Hz, 1H, H-1^{Glc}'), 4.45 (d, J = 7.2 Hz, 1H, H-1^{Gal}'), 4.45 (d, J = 7.2 Hz, 1H, H-1^{Gal}), 4.12 (d, J = 3.0 Hz, 1H, H-4^{Gal}), 4.05 (dd, J = 9.0, 2.4 Hz, 1H, H-3^{Gal}'), 4.03 (t, J = 6.6 Hz, 2H, -CH₂CH₂OPh), 3.96 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.92–3.71 (m, 13H, H-3^{Glc}, H-6a,b^{Glc}, H-3,6a^{Gal}, H-6a,b^{Glc}, H-4^{Gal}', H-5,8,9a^{NeuNAc}, -OCH₂CH₂-), 3.70–3.54 (m, 9H, H-4^{Glc}, H-2,5,6b^{Gal}, H-3^{Glc}', H-5^{Gal}', H-6a,b^{Gal}', H-4^{NeuNAc}), 3.54–3.48 (m, 7H, H-2^{Glc}, H-4,5^{Glc}', H-2^{Gal}'), H-6,7,9b^{NeuNAc}), 3.34 (t, J = 9.0 Hz, 1H, H-2^{Glc}'), 2.69 (dd, J = 12.0, 4.2 Hz, 1H, H-3_{eq}^{NeuNAc}), 1.96 (s, 3H, CH₃CO), 1.76–1.68 (m, 3H, H-3_{ax}^{NeuNAc}, -CH₂CH₂-), 1.61–1.55 (m, 2H, -CH₂CH₂-), 1.41–1.35 (m, 2H, -CH₂CH₂-), 1.33–1.20 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 174.9 (CH₃CO), 173.8 (C-1^{NeuNAc}), 158.0, 129.8 (2C), 121.3, 114.8 (2C), 103.5 (C-1^{Glc}'), 102.5 (C-1^{Gal}'), 102.4 (C-1^{Gal}), 99.7 (C-2^{NeuNAc}), 95.1 (d, J = 4.5 Hz, C-1^{Glc}), 82.0 (C-3^{Gal}), 77.8 (C-4^{Glc}'), 77.3 (C-4^{Glc}), 75.4 (C-3^{Gal}'), 75.0 (C-5^{Gal}'), 74.8 (C-5^{Gal}), 74.5 (C-5^{Glc}'), 73.9 (C-3^{Glc}'), 72.8 (C-2^{Glc}'), 72.7 (C-6^{NeuNAc}), 71.6 (C-8^{NeuNAc}), 71.4 (C-3^{Glc}), 71.3 (C-5^{Glc}), 71.2 (d, J = 9.0 Hz, C-2^{Glc}), 69.9 (C-2^{Gal}), 69.2 (C-2^{Gal}'), 68.6 (-OCH₂CH₂OPh), 68.3 (C-4^{NeuNAc}), 68.1 (C-4^{Gal}), 67.9 (C-7^{NeuNAc}), 67.3 (C-4^{Gal}'), 66.9 (d, J = 6.0 Hz, -OCH₂CH₂-), 62.4 (C-9^{NeuNAc}), 60.9 (2C, C-6^{Gal}, C-6^{Gal}'), 59.7 (C-6^{Glc}'), 59.5 (C-6^{Glc}), 51.6 (C-5^{NeuNAc}), 39.5 (C-3^{NeuNAc}), 29.7 (d, J = 7.5 Hz, -OCH₂CH₂-), 28.56, 28.53 (2C), 28.4, 28.3, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 21.9 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.70 (d, J = 19.4 Hz), -13.14 (d, J = 19.4 Hz); ESI-(-)-TOF HRMS m/z : [M - 2H]²⁻ Calcd for C₅₂H₈₅NO₃₆P₂ 680.7169; Found 680.7170.

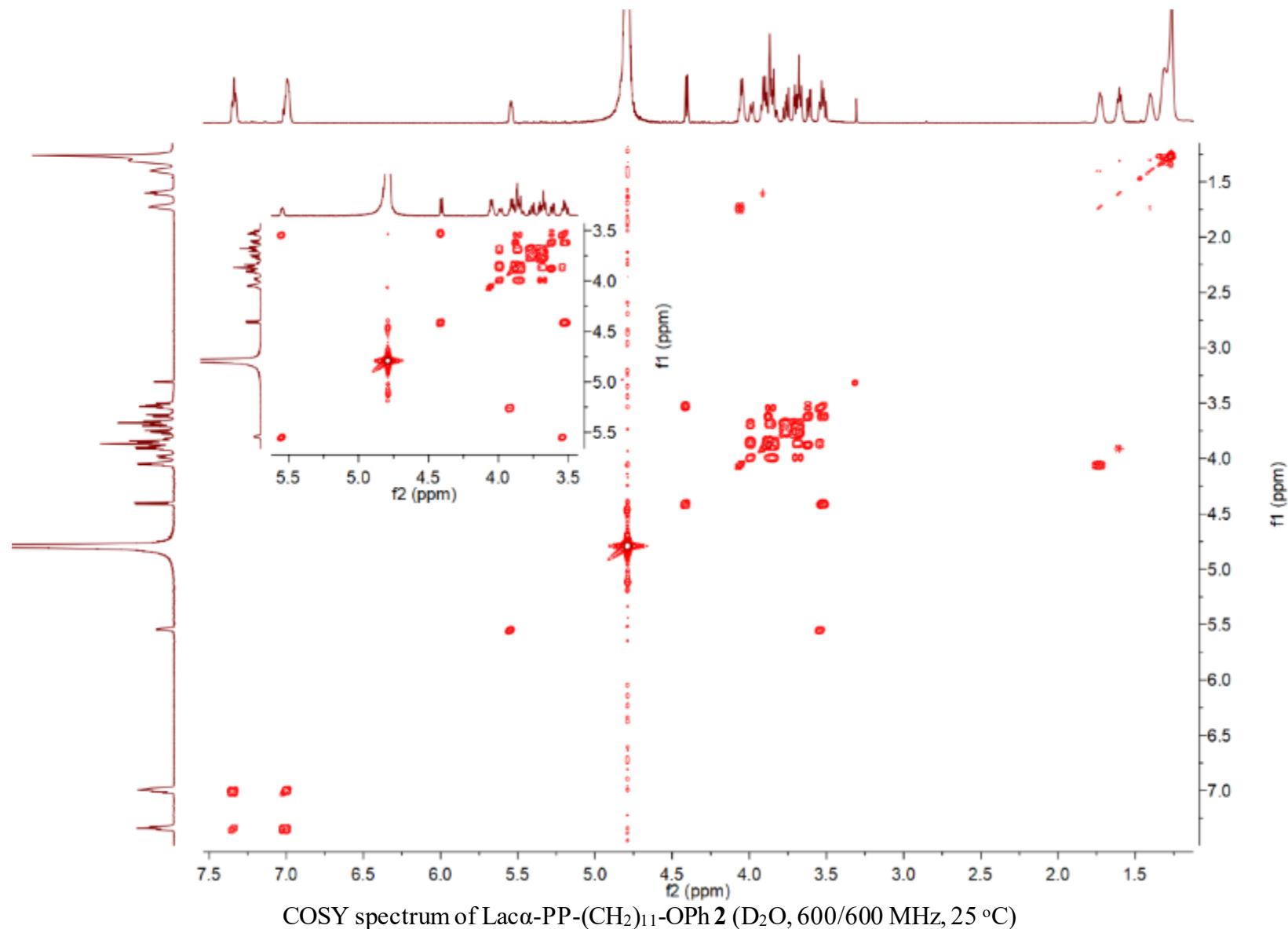
5e: ¹H NMR (600 MHz, D₂O): δ 7.31 (t, J = 7.8 Hz, 2H, Ph), 7.00–6.95 (m, 3H, Ph), 5.52 (dd, J = 7.2, 3.6 Hz, 1H, H-1^{Glc}), 4.68 (d, J = 7.8 Hz, 1H, H-1^{Gal}'), 4.63 (d, J = 7.8 Hz, 1H, H-1^{Glc}'), 4.43 (t, J = 7.8 Hz, 1H, H-1^{Gal}), 4.10 (d, J = 3.2 Hz, 1H, H-4^{Gal}), 4.06 (dd, J = 10.2, 3.0 Hz, 1H, H-3^{Gal}'), 4.03 (t, J = 6.6 Hz, 2H, -CH₂CH₂OPh), 3.96 (br d, J = 10.2 Hz, 1H, H-5^{Glc}), 3.91–3.77 (m, 10H, H-3^{Glc}, H-6a,b^{Glc}, H-6a^{Glc}', H-4^{Gal}', H-5,8,9a^{NeuNAc}, -OCH₂CH₂-), 3.76–3.59 (m, 12H, H-4^{Glc}, H-2,3,5^{Gal}, H-6a,b^{Gal}, H-3,6b^{Glc}', H-5^{Gal}', H-6a,b^{Gal}',

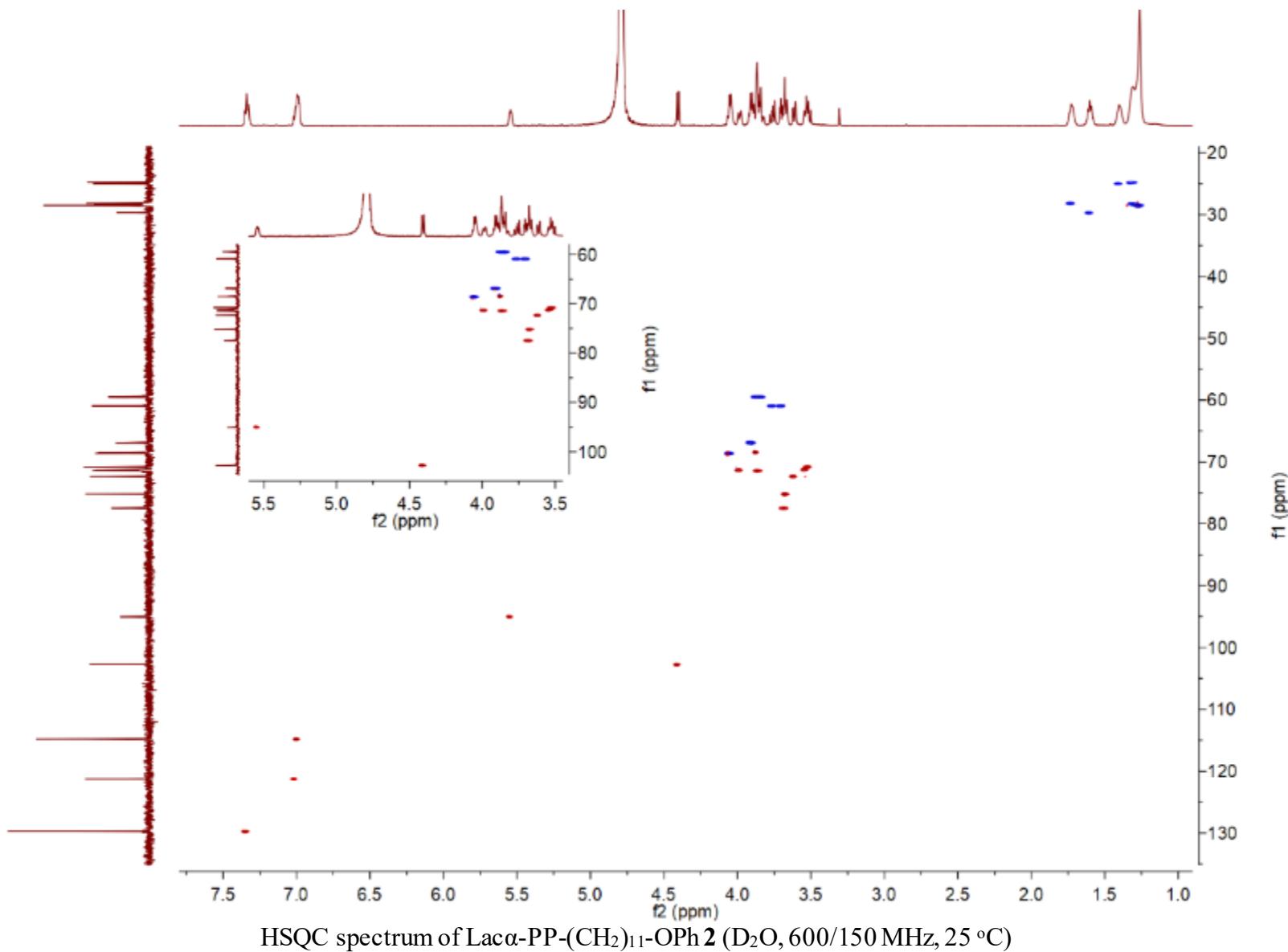
H-4^{NEuNAc}), 3.59–3.44 (m, 7H, H-2^{Glc}, H-2,4^{Glc'}, H-2^{Gal'}, H-6,7,9b^{NeuNAc}), 3.42–3.33 (m, 1H, H-5^{Glc'}), 2.69 (dd, J = 12.0, 4.2 Hz, 1H, H-3_{eq}^{NeuNAc}), 1.96 (s, 3H, CH₃CO), 1.77–1.67 (m, 3H, H-3_{ax}^{NeuNAc}, -CH₂CH₂-), 1.61–1.54 (m, 2H, -CH₂CH₂-), 1.41–1.34 (m, 2H, -CH₂CH₂-), 1.33–1.21 (m, 12H, -CH₂CH₂-); ¹³C NMR (150 MHz, D₂O): δ 174.8 (CH₃CO), 173.8 (C-1^{NeuNAc}), 157.9, 129.7 (2C), 121.2, 114.8 (2C), 103.3 (C-1^{Glc'}), 102.7 (C-1^{Gal'}), 102.3 (C-1^{Gal}), 99.6 (C-2^{NeuNAc}), 95.0 (d, J = 6.0 Hz, C-1^{Glc}), 83.8 (C-3^{Glc'}), 81.8 (C-3^{Gal}), 77.2 (C-4^{Glc}), 75.4 (C-3^{Gal'}), 75.1 (C-5^{Glc'}), 74.9 (C-5^{Gal'}), 74.8 (C-5^{Gal}), 72.8 (C-2^{Glc'}), 72.7 (C-6^{NeuNAc}), 71.7 (C-8^{NeuNAc}), 71.4 (C-3^{Glc}), 71.3 (C-5^{Glc}), 71.2 (d, J = 9.0 Hz, C-2^{Glc}), 69.9 (C-2^{Gal}), 69.5 (C-2^{Gal'}), 68.6 (-OCH₂CH₂OPh), 68.25 (C-4^{NeuNAc}), 68.16 (C-4^{Gal}), 67.9 (C-7^{NeuNAc}), 67.8 (C-4^{Glc'}), 67.3 (C-4^{Gal'}), 66.9 (d, J = 6.0 Hz, -OCH₂CH₂-), 62.4 (C-9^{NeuNAc}), 60.88 (C-6^{Gal}), 60.85 (C-6^{Gal'}), 60.3 (C-6^{Glc'}), 59.4 (C-6^{Glc}), 51.5 (C-5^{NeuNAc}), 39.4 (C-3^{NeuNAc}), 29.6 (d, J = 7.5 Hz, -OCH₂CH₂-), 28.52, 28.50, 28.48, 28.32, 28.26, 28.2, 25.0, 24.8 (8C, -OCH₂CH₂(CH₂)₈CH₂OPh), 21.9 (CH₃CO); ³¹P NMR (243 MHz, D₂O): δ -10.71 (d, J = 21.8 Hz), -13.14 (d, J = 21.8 Hz); ESI(-)-TOF HRMS *m/z*: [M - 2H]²⁻ Calcd for C₅₂H₈₅NO₃₆P₂ 680.7169; Found 680.7163.

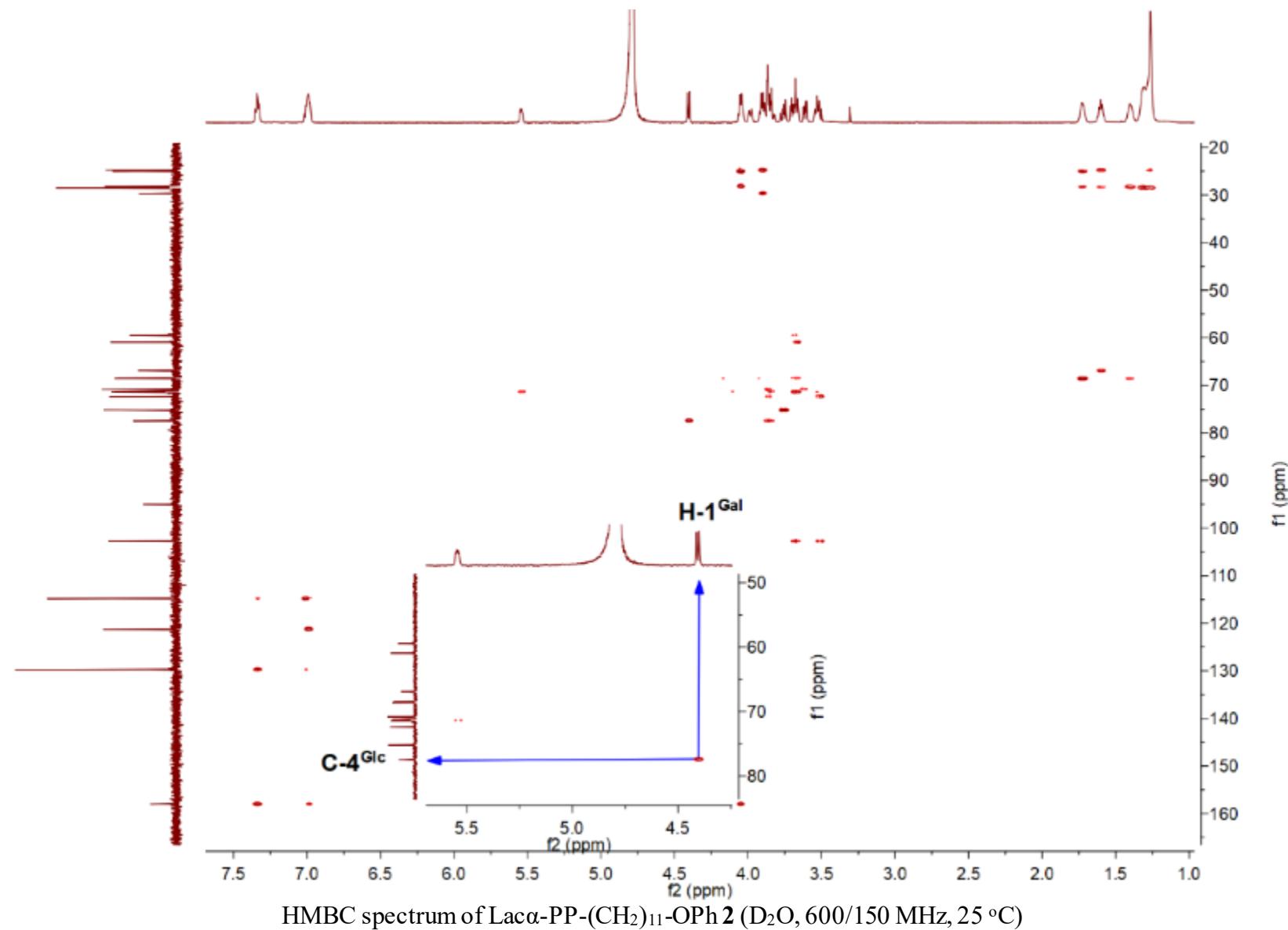
II. NMR and ESI-HRMS spectra

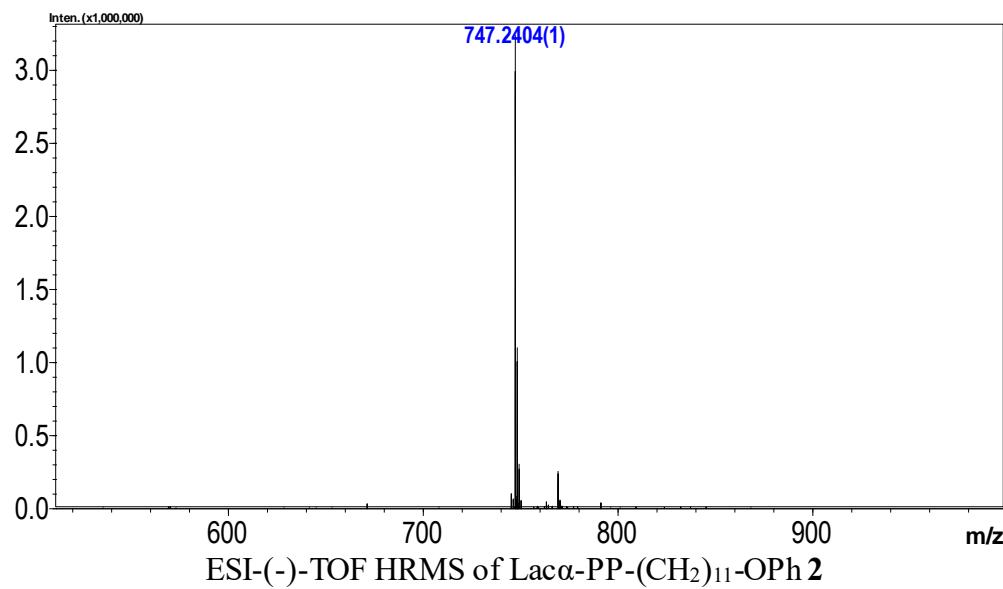
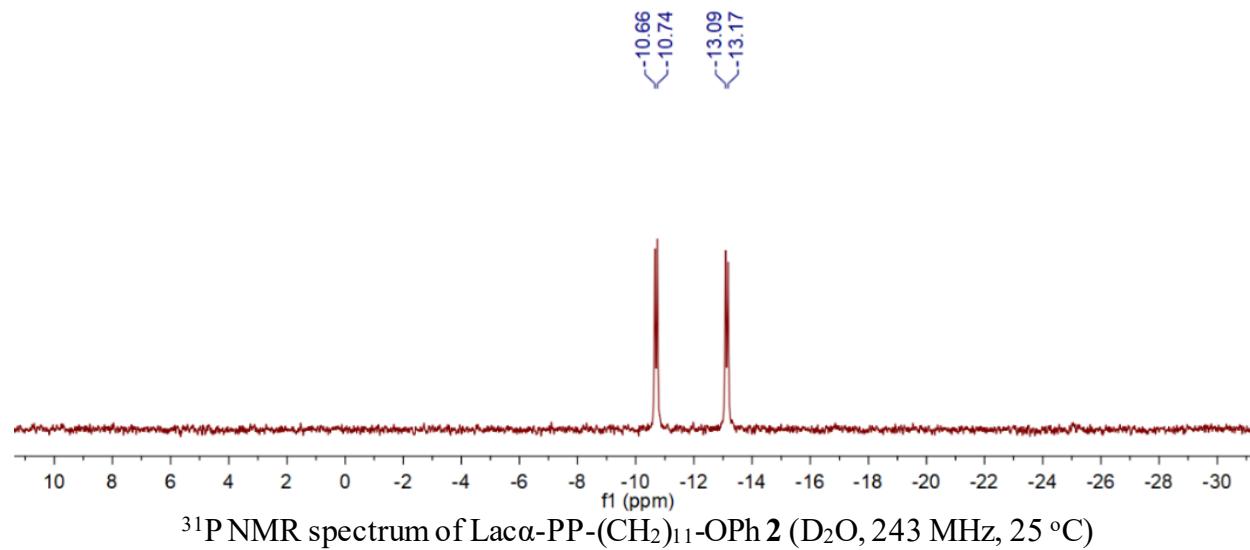


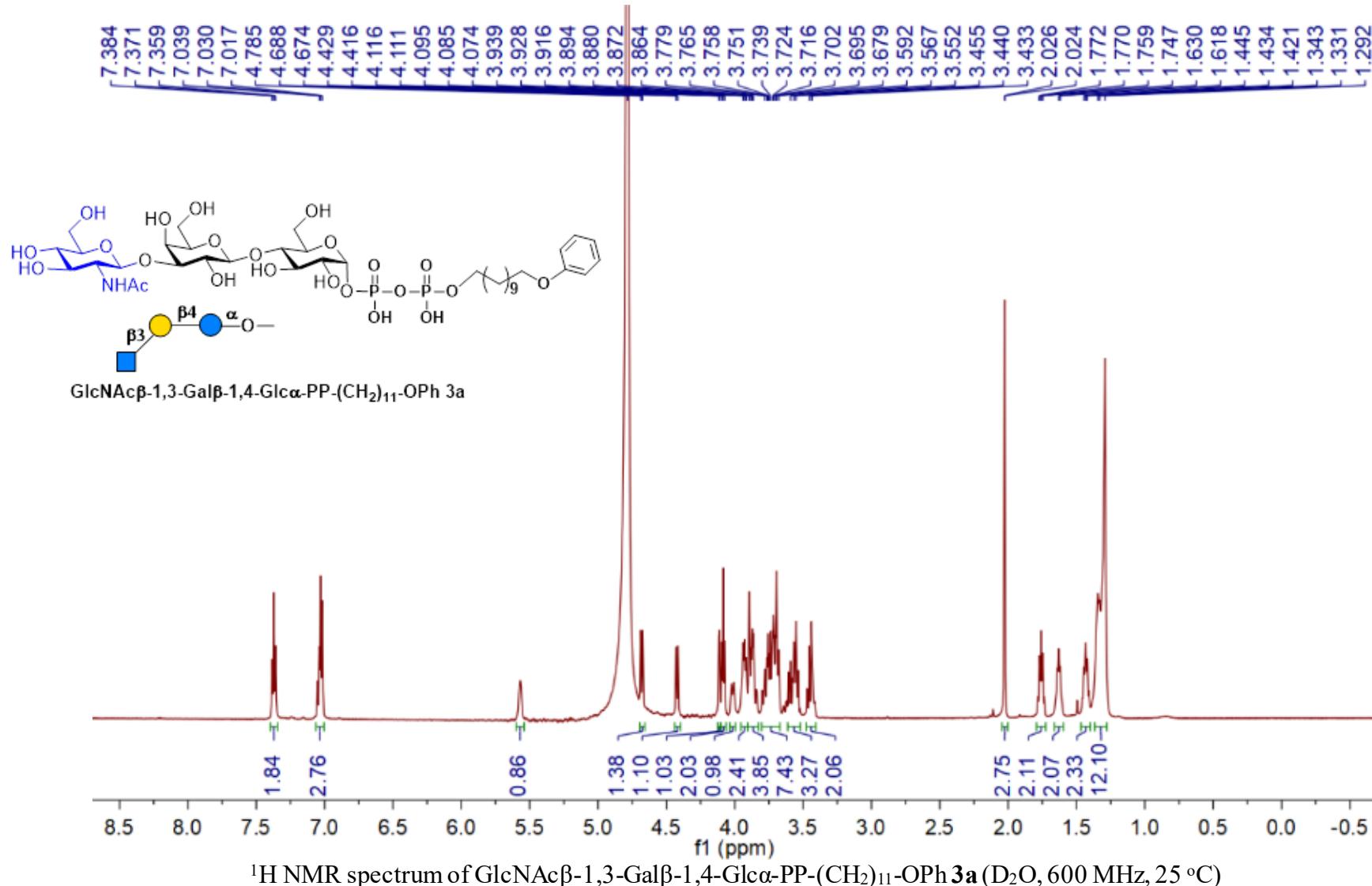


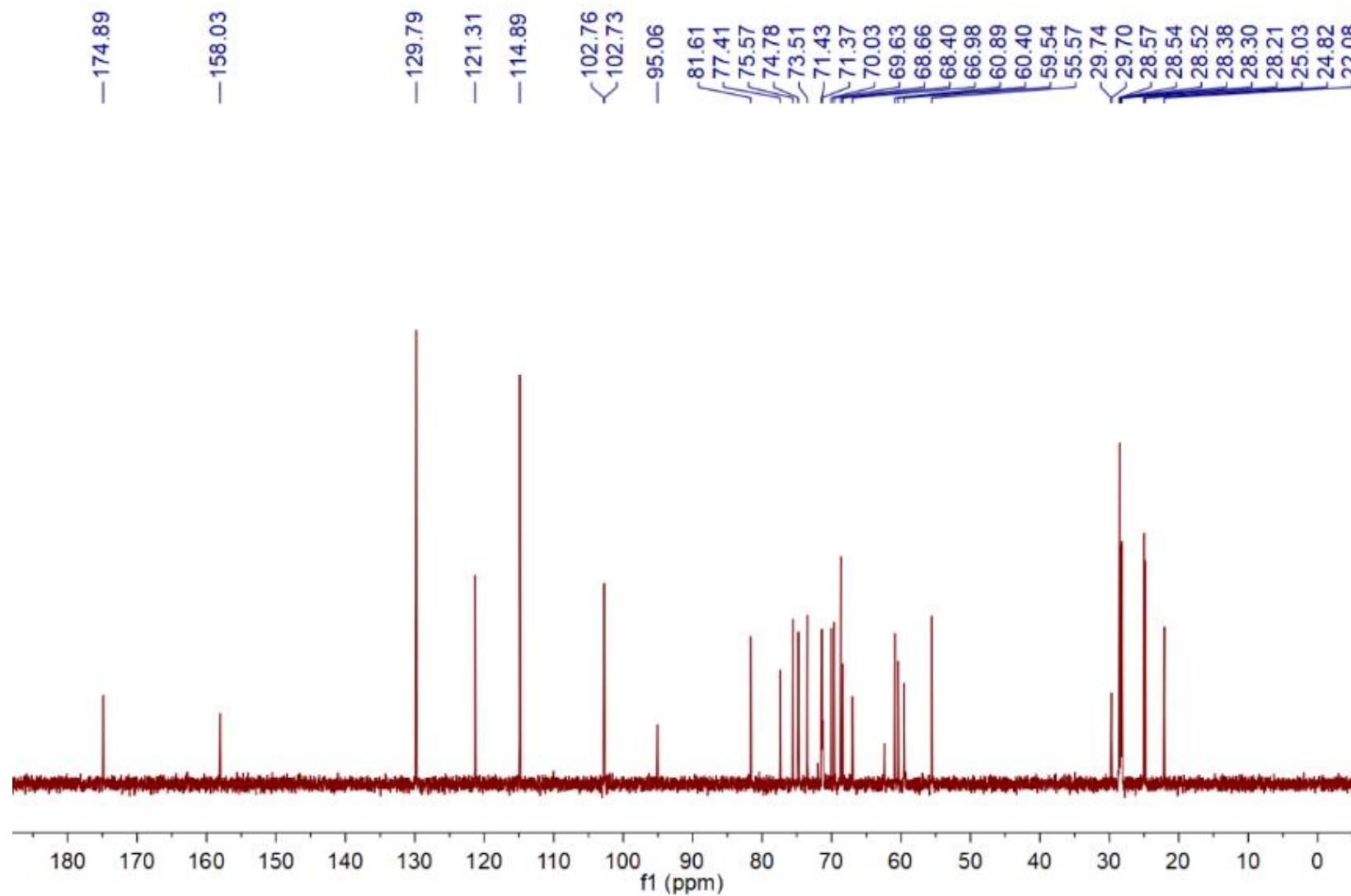


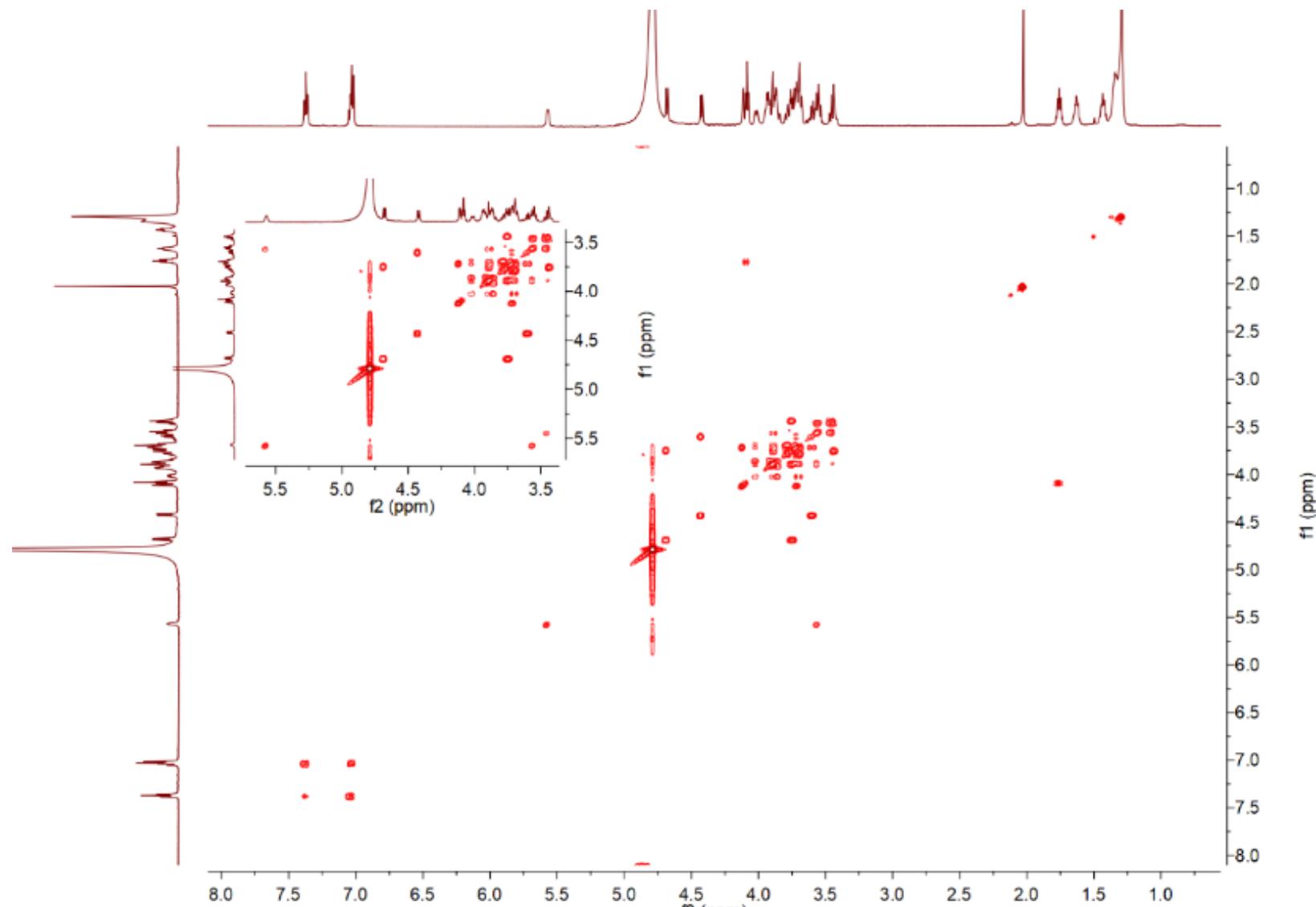




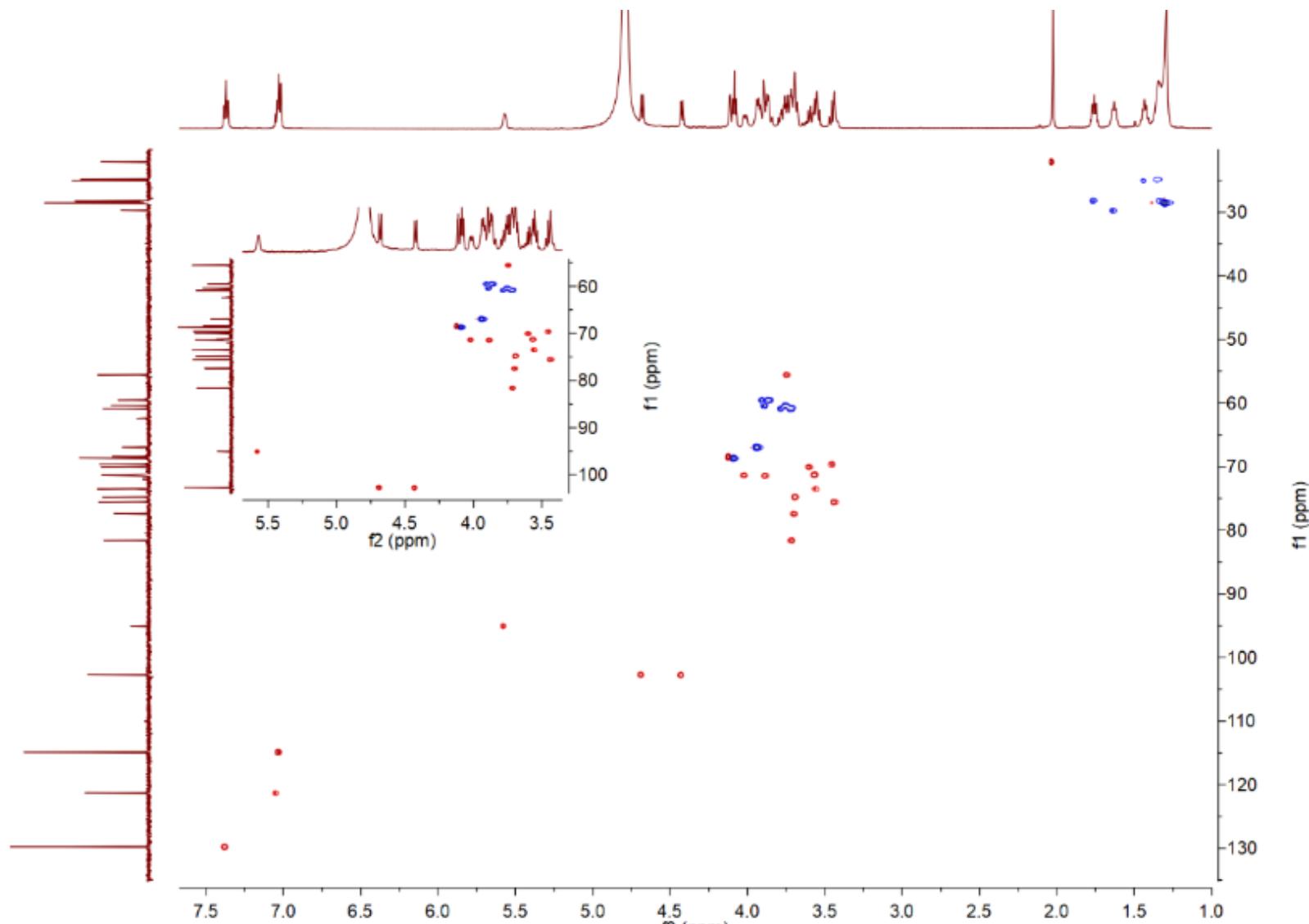


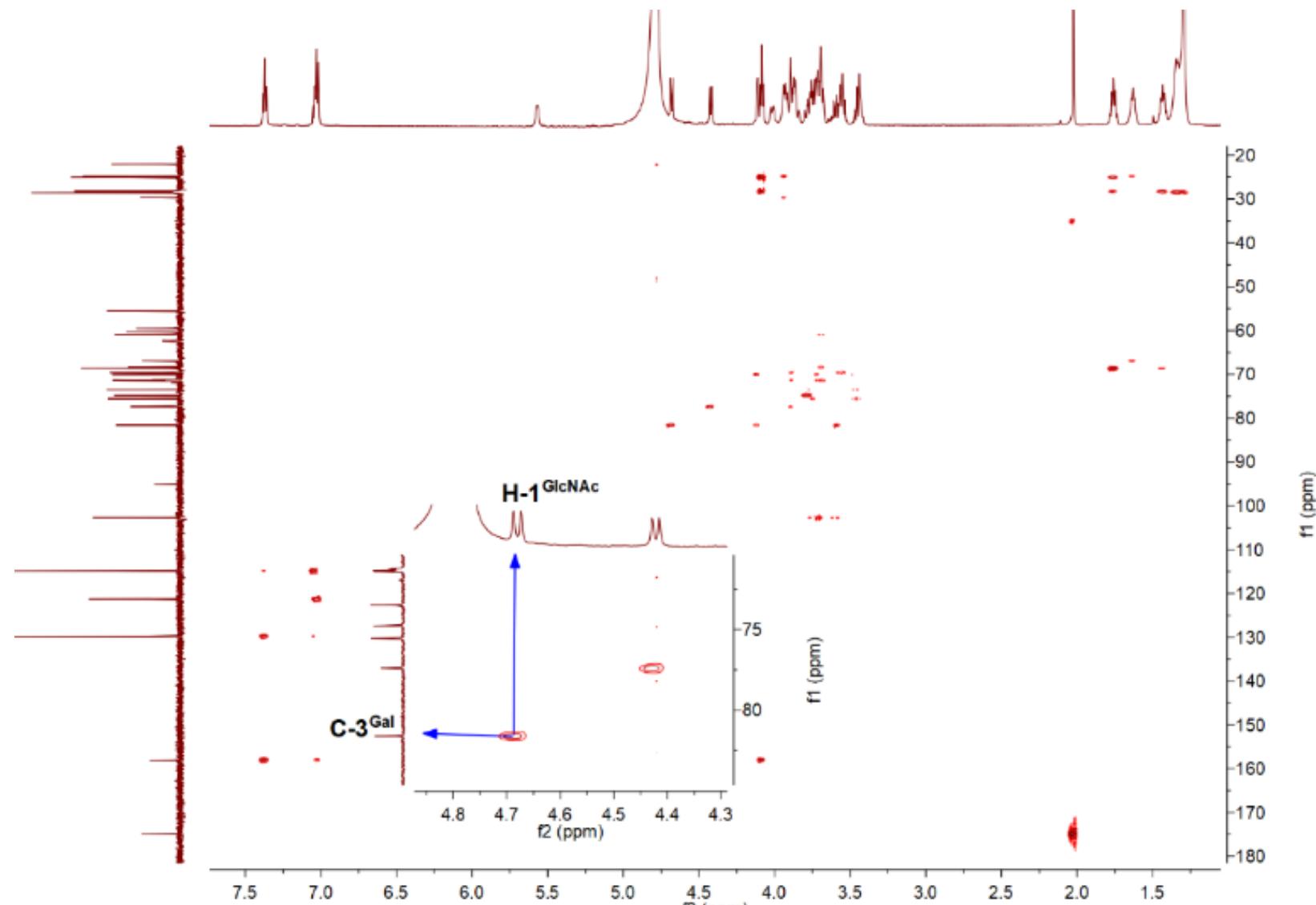




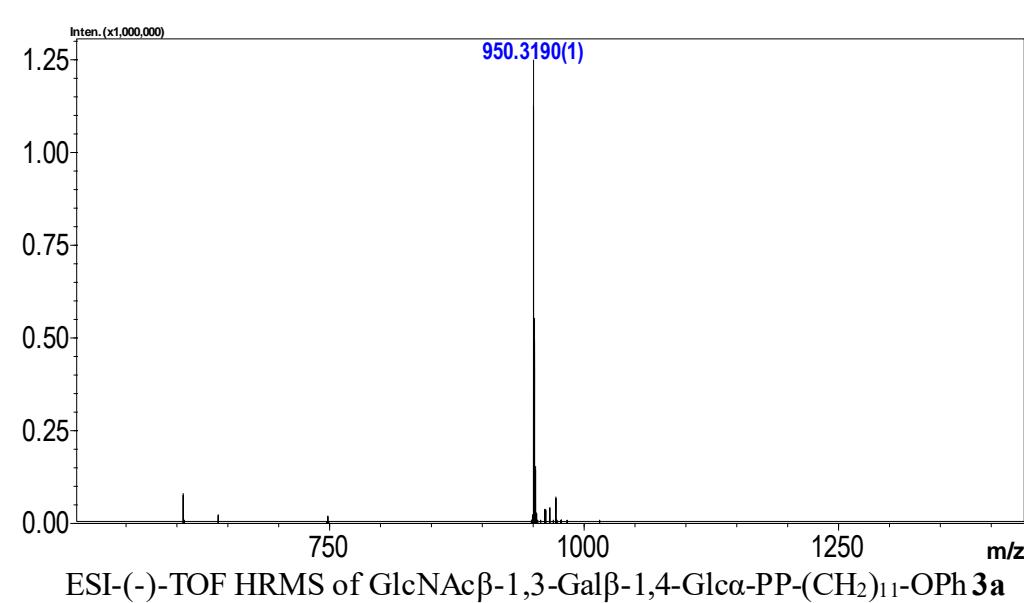
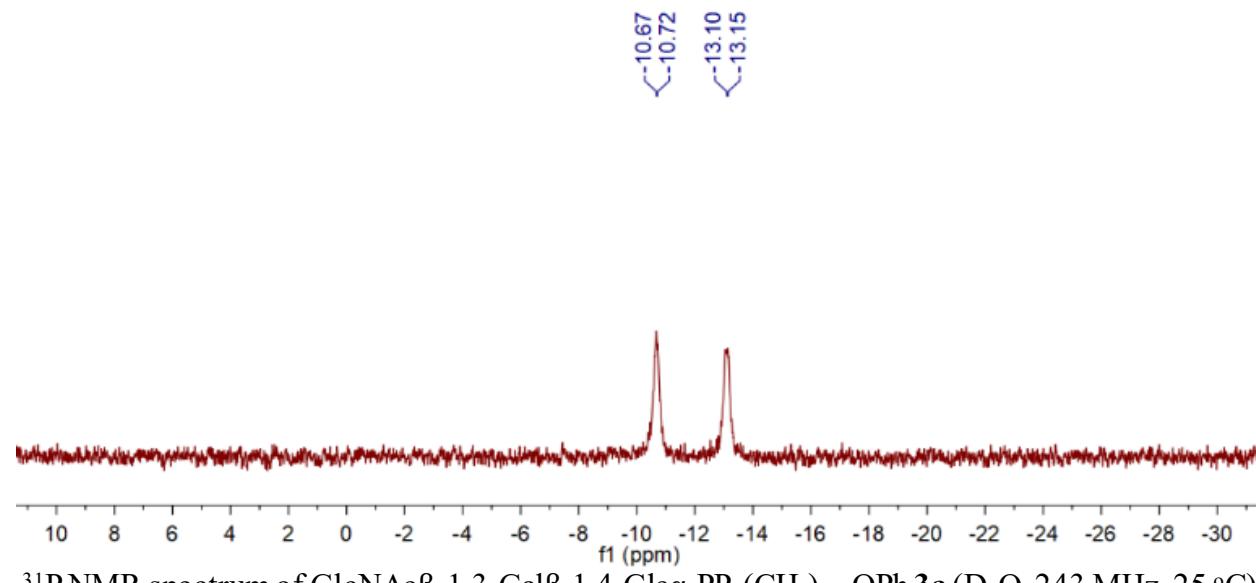


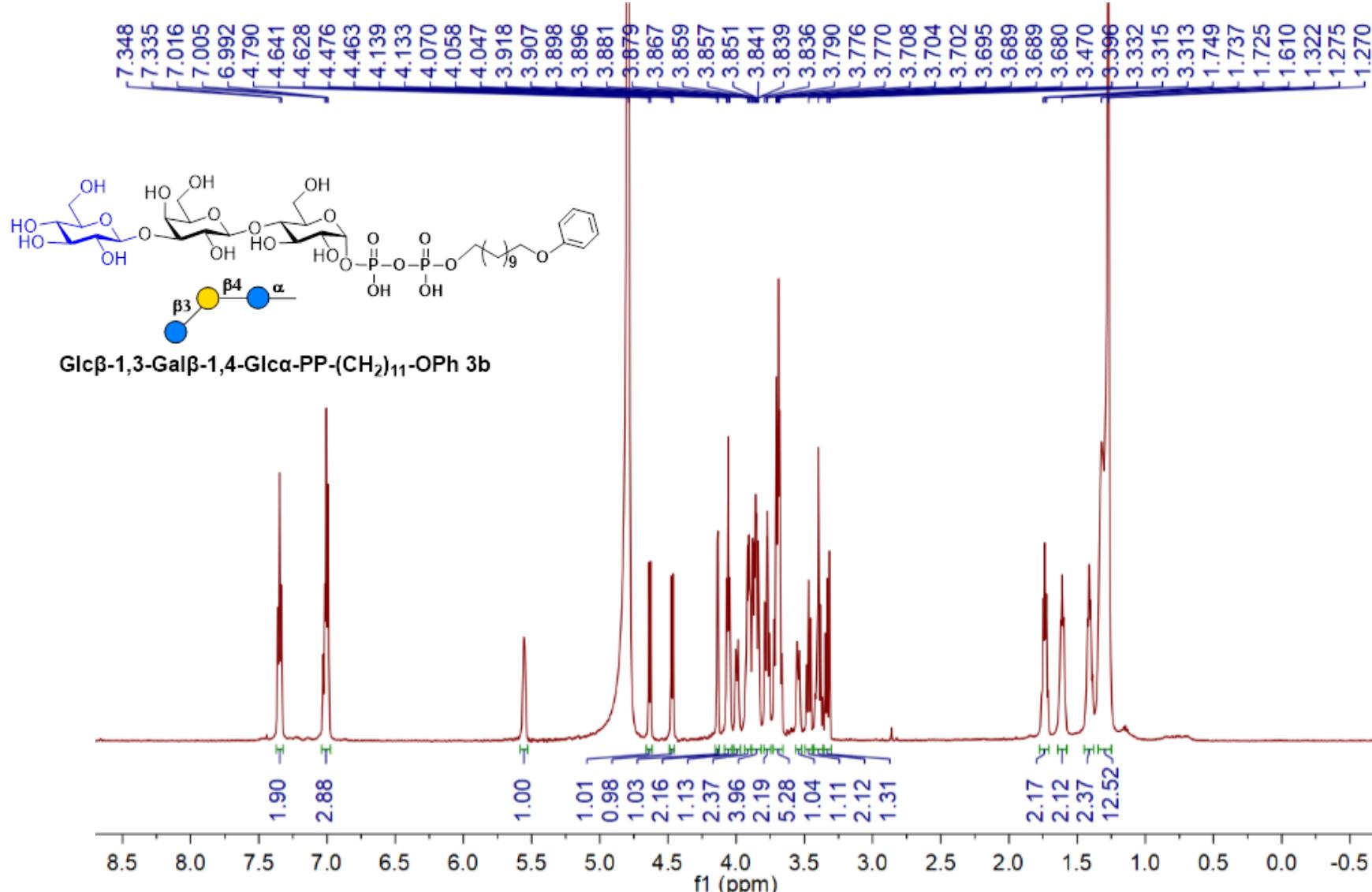
COSY spectrum of GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3a** (D₂O, 600/600 MHz, 25 °C)



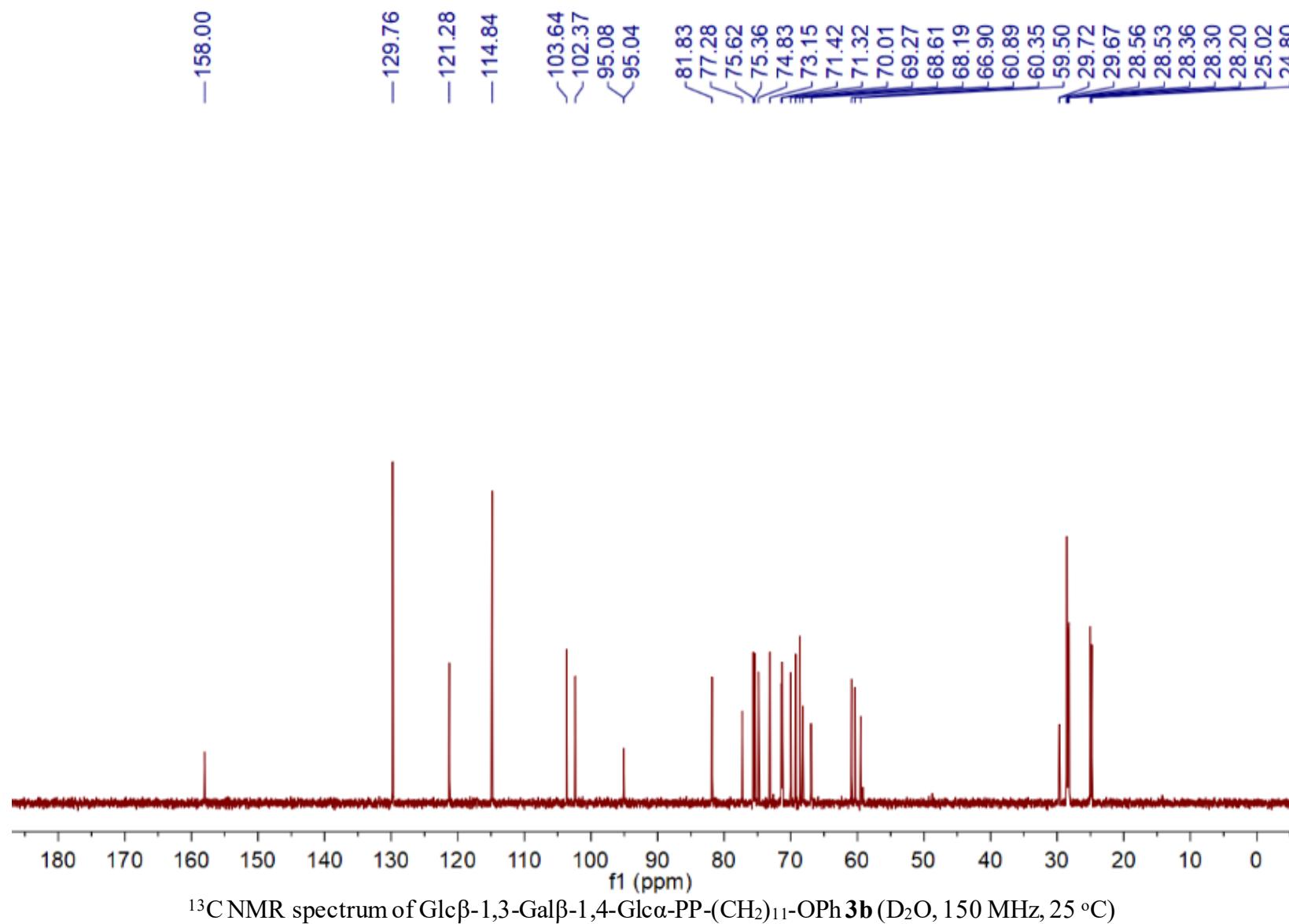


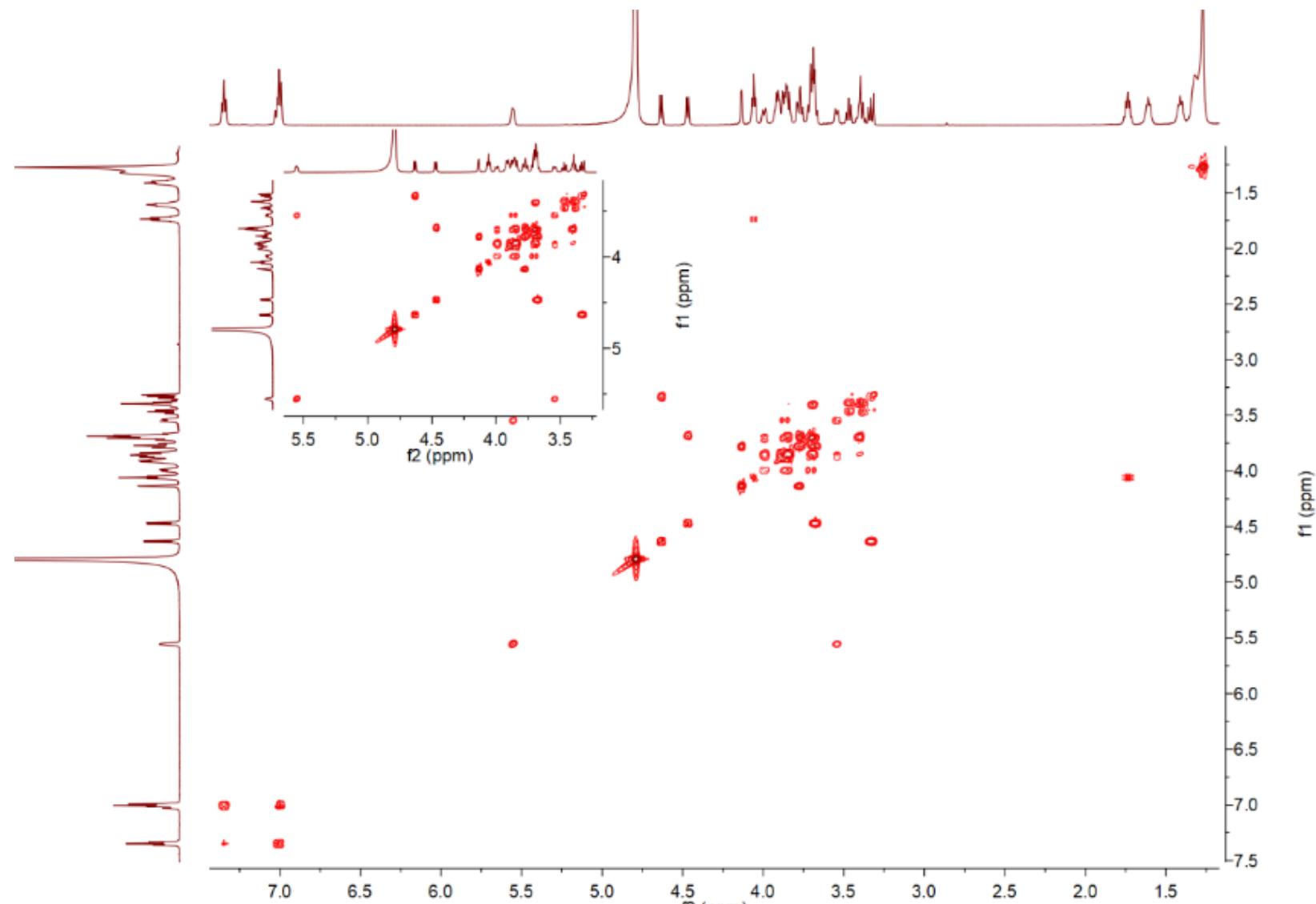
HMBC spectrum of GlcNAc β -1,3-Gal β -1,4-Glca-PP-(CH₂)₁₁-OPh **3a** (D₂O, 600/150 MHz, 25 °C)



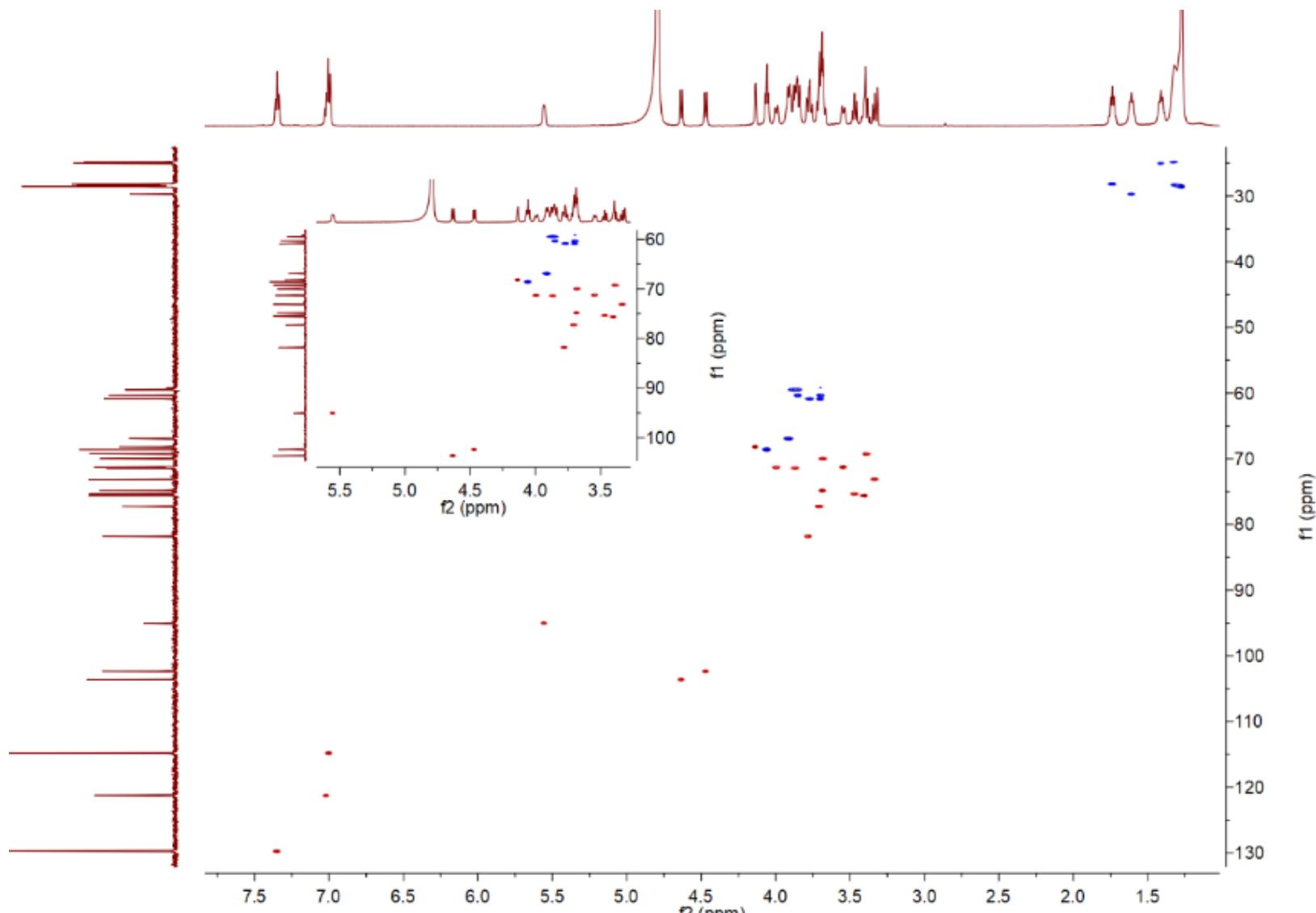


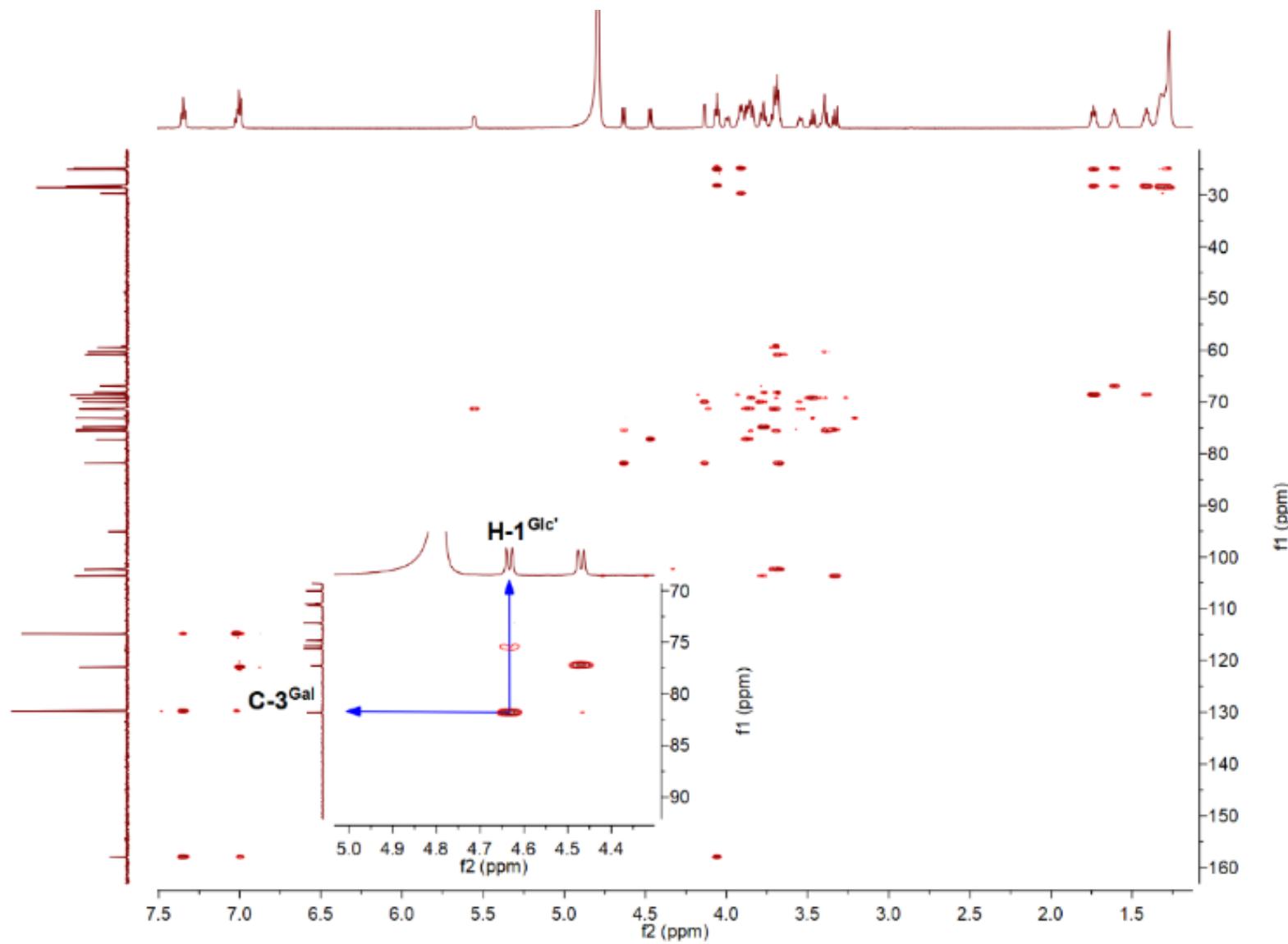
¹H NMR spectrum of Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3b** (D₂O, 600 MHz, 25 °C)



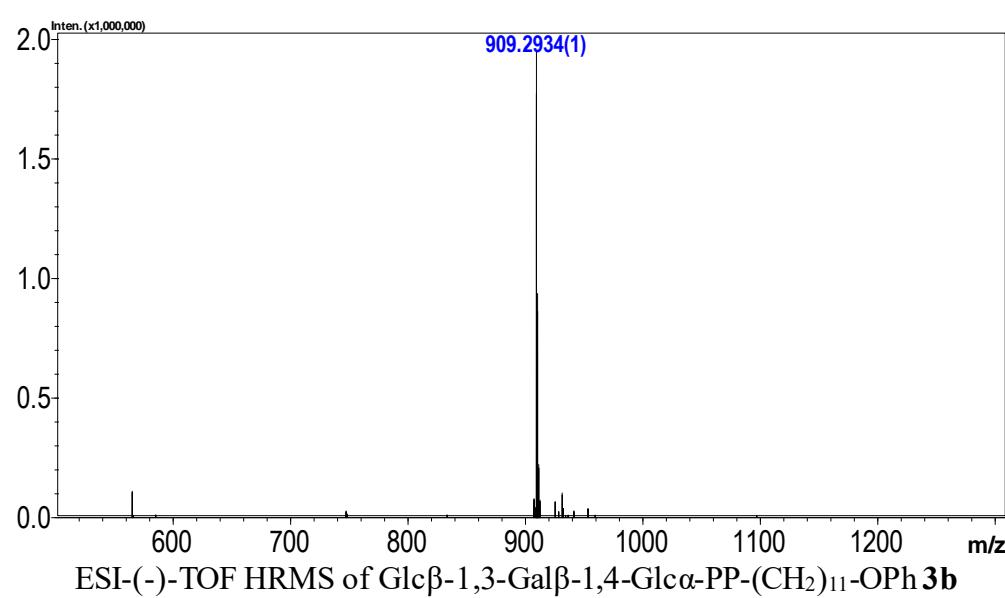
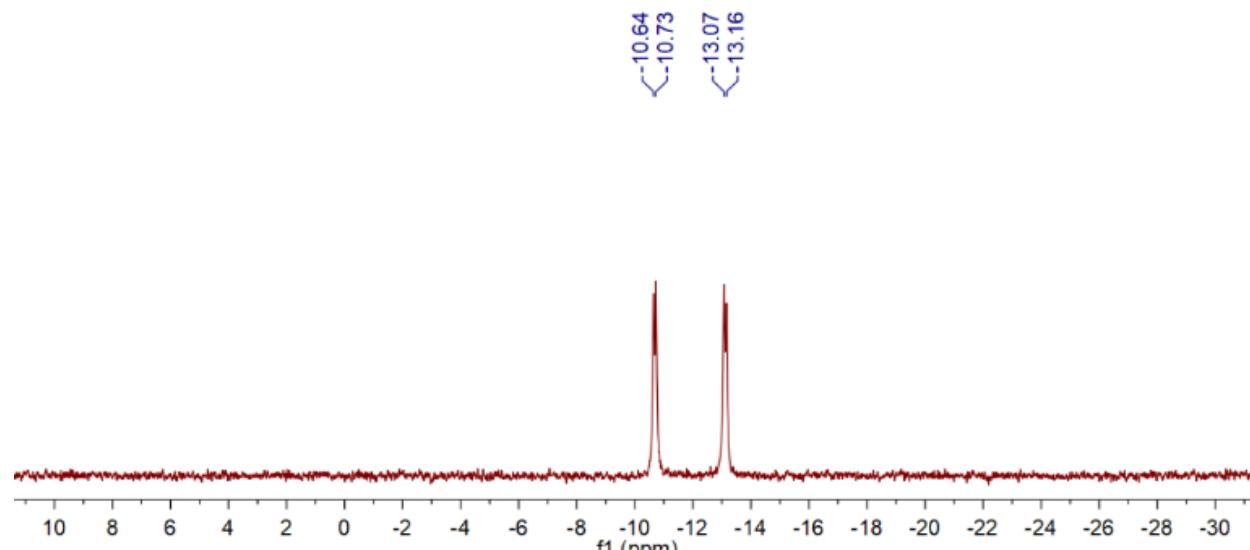


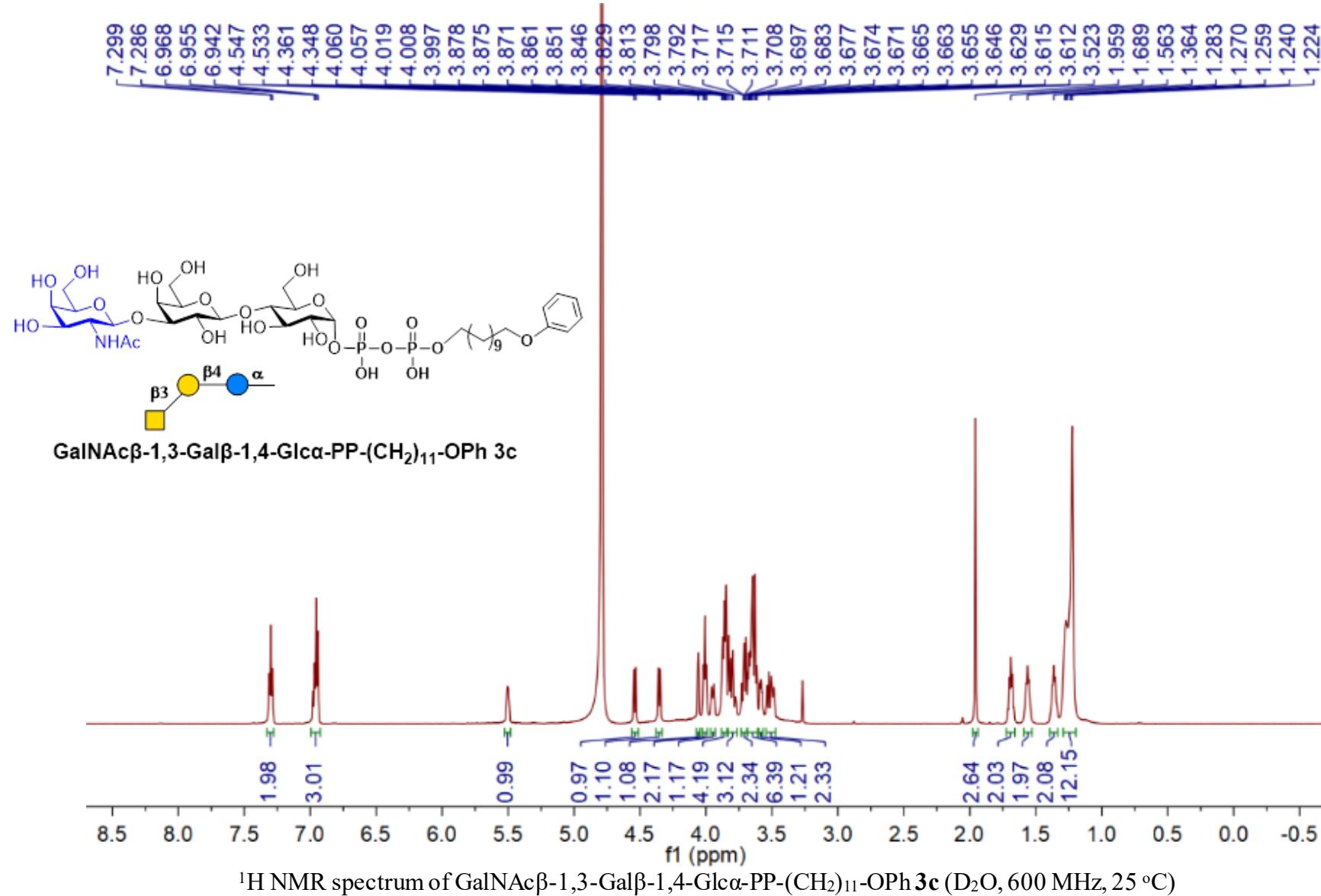
COSY spectrum of $\text{Glc}\beta\text{-}1,3\text{-Gal}\beta\text{-}1,4\text{-Glc}\alpha\text{-PP-(CH}_2\text{)}_{11}\text{-OPh}$ **3b** (D_2O , 600/600 MHz, 25 °C)

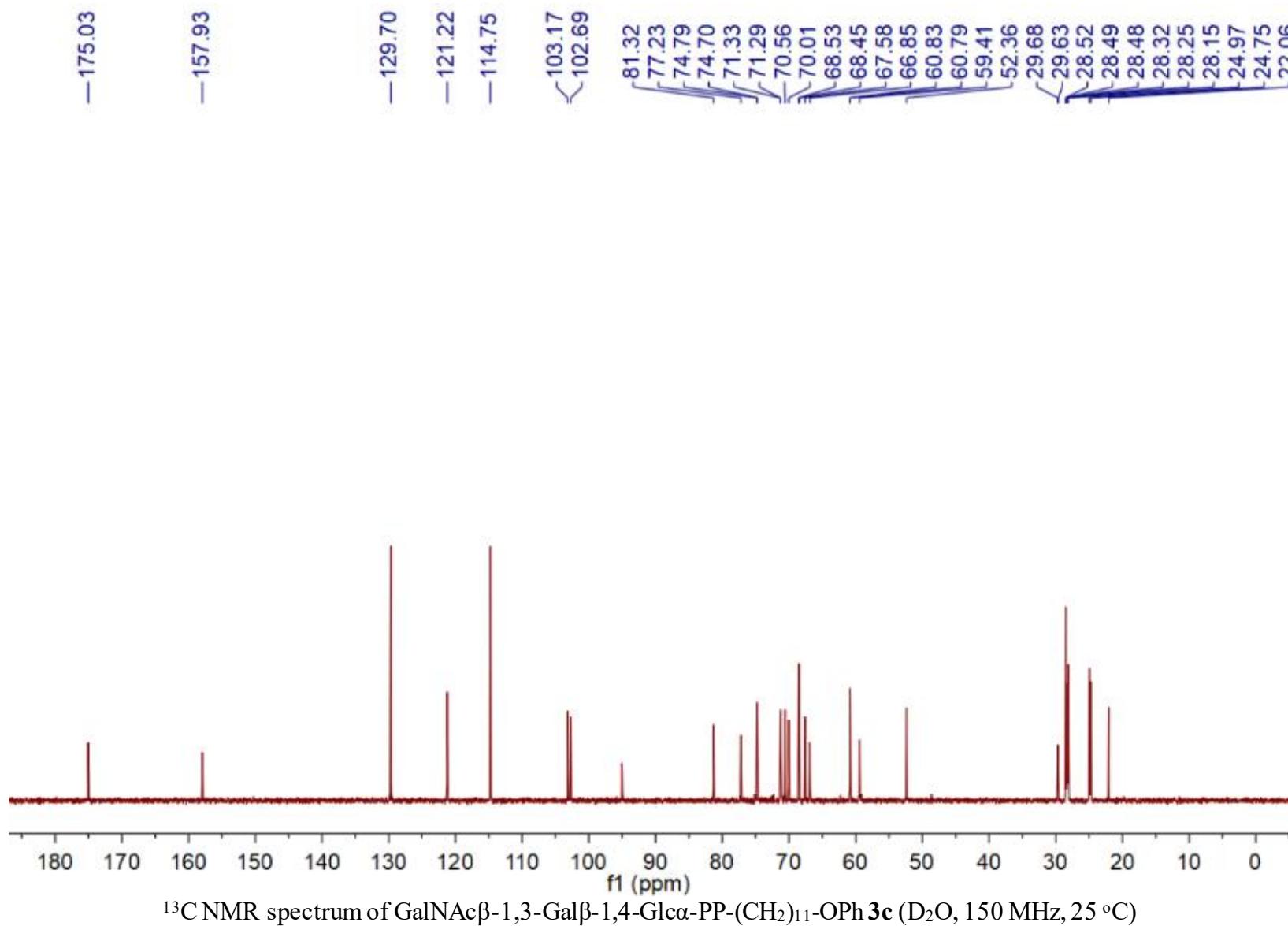


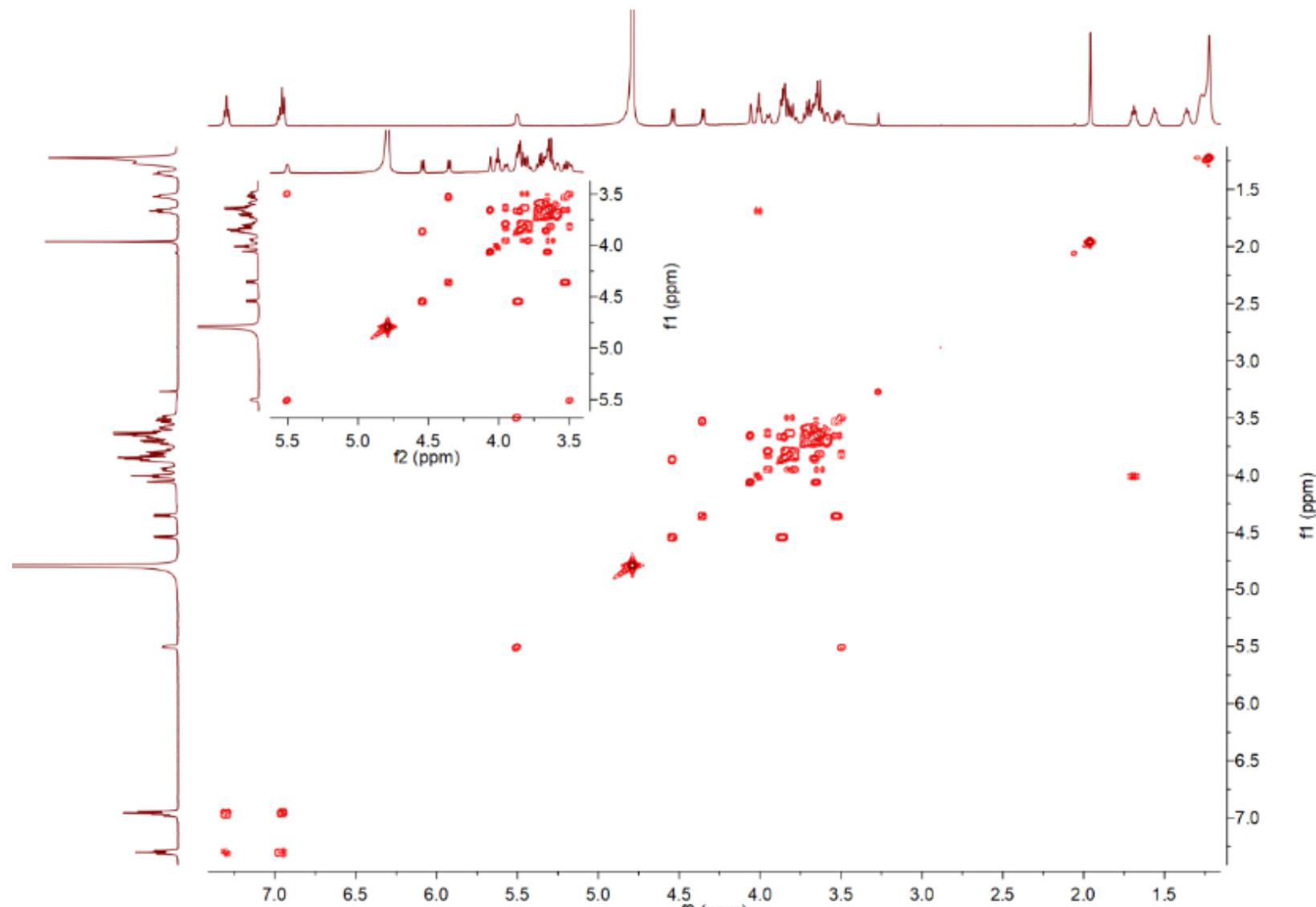


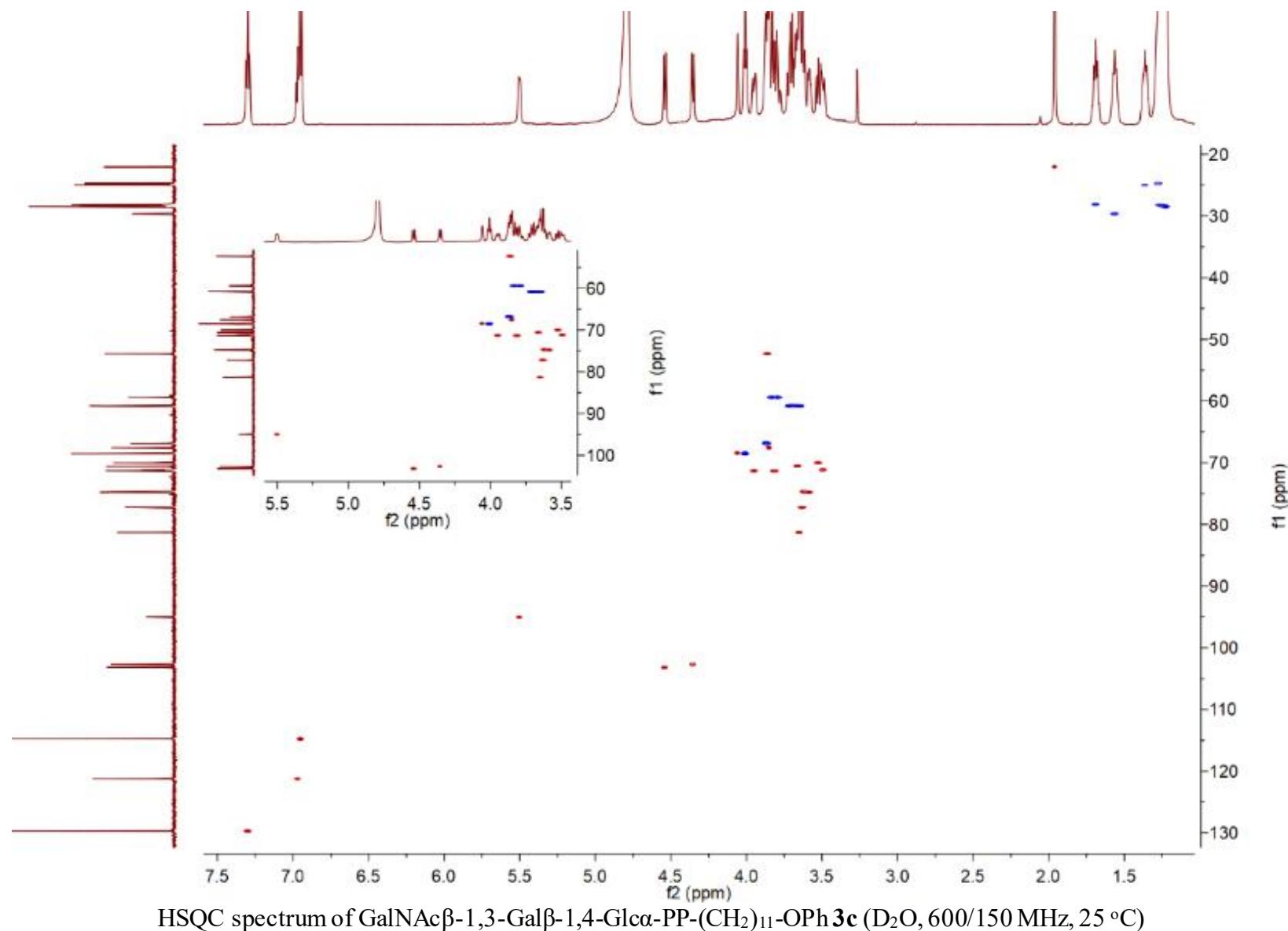
HMBC spectrum of Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **3b** (D₂O, 600/150 MHz, 25 °C)

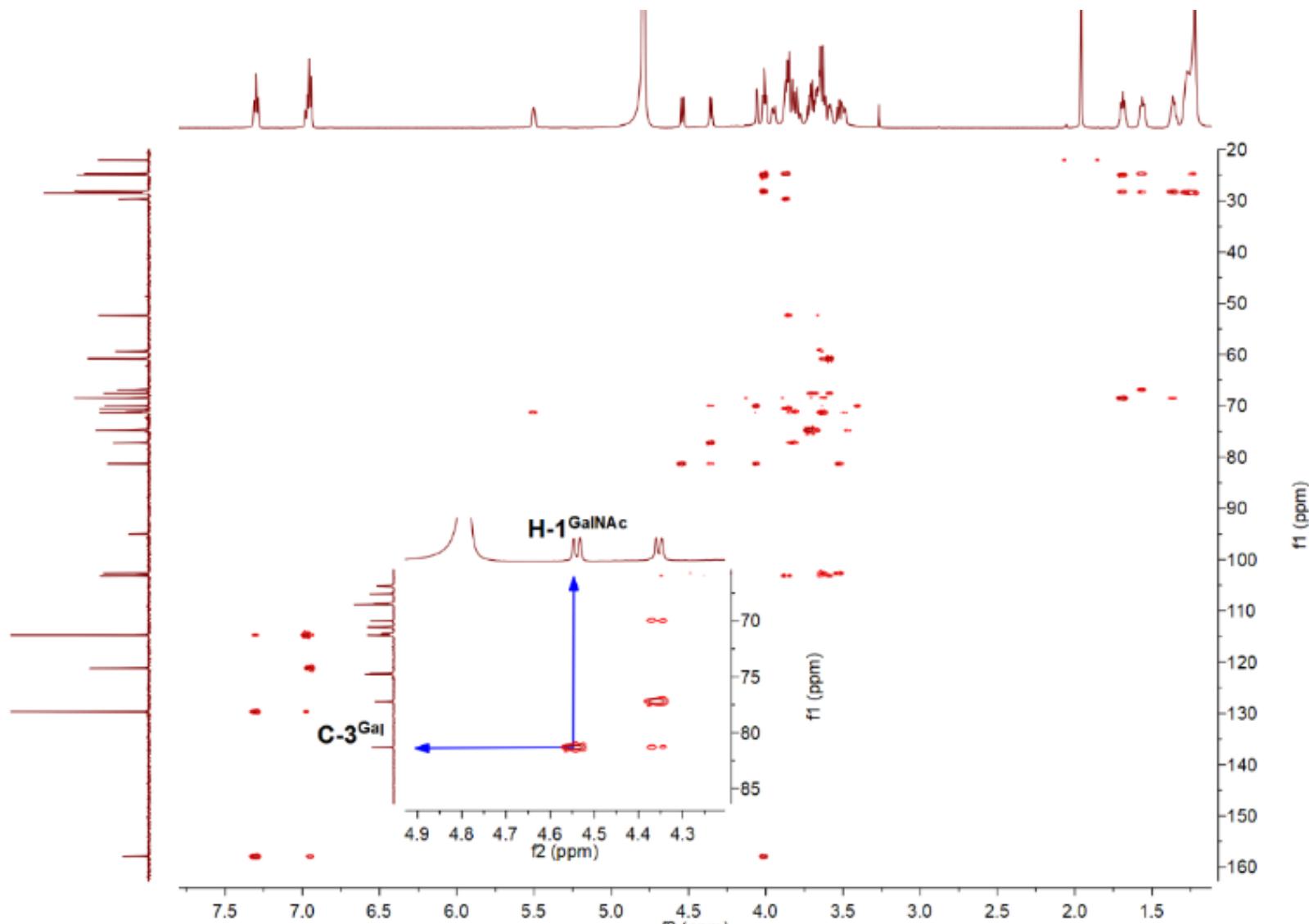


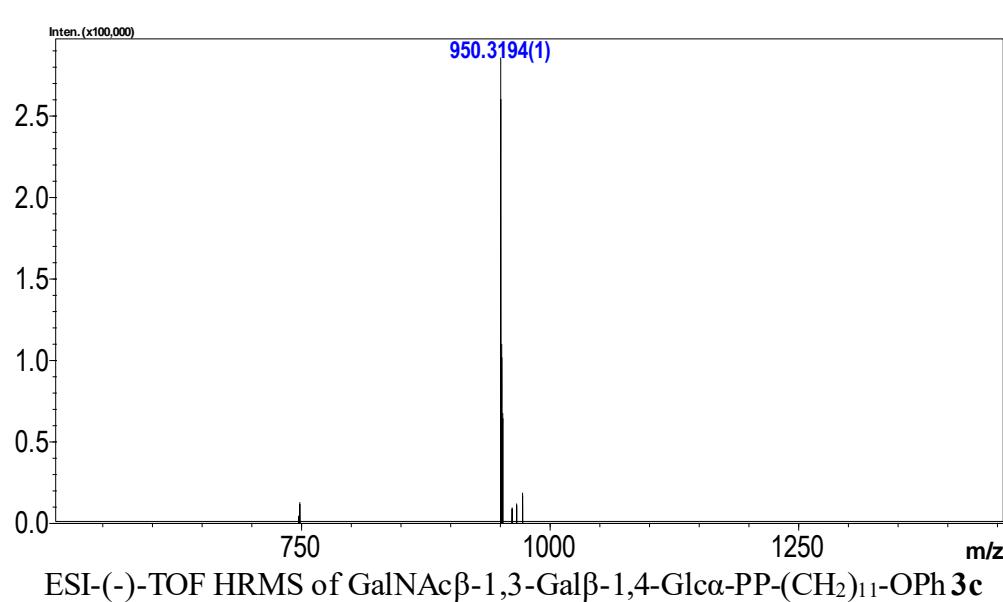
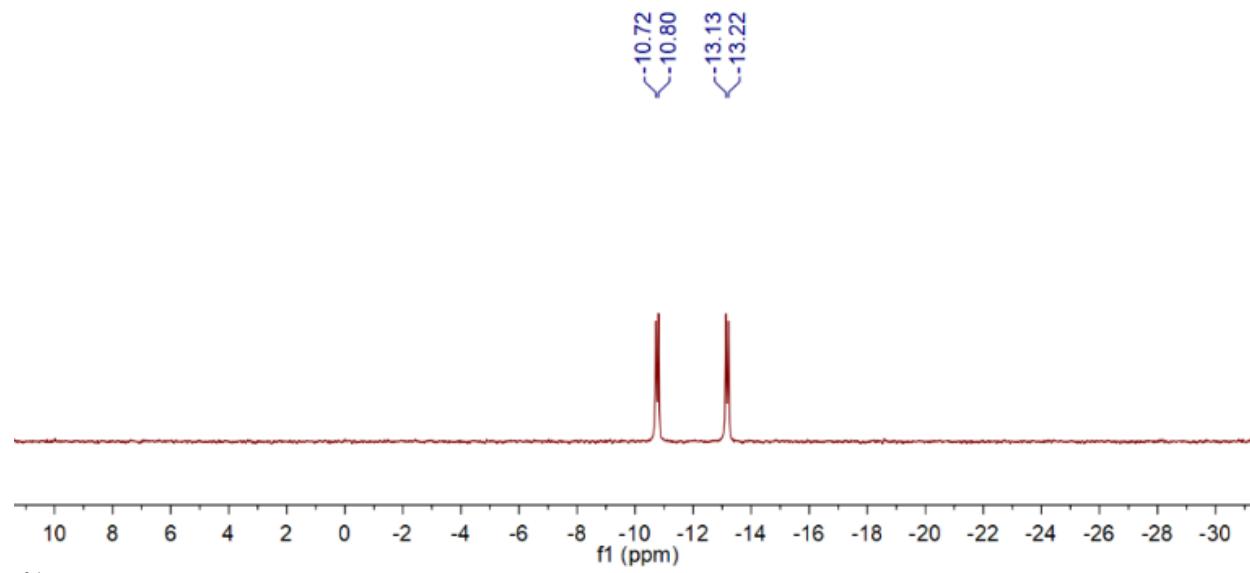


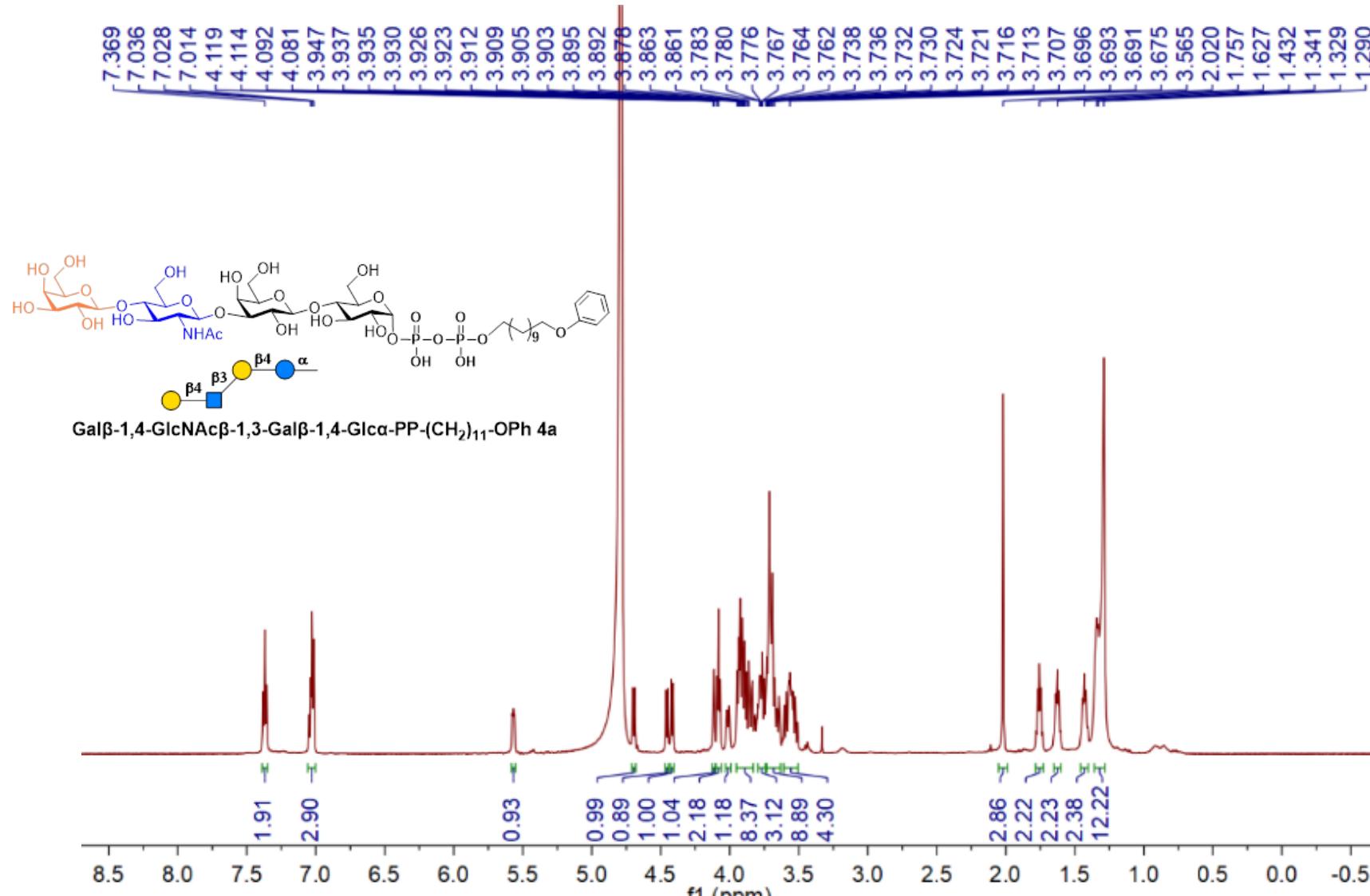




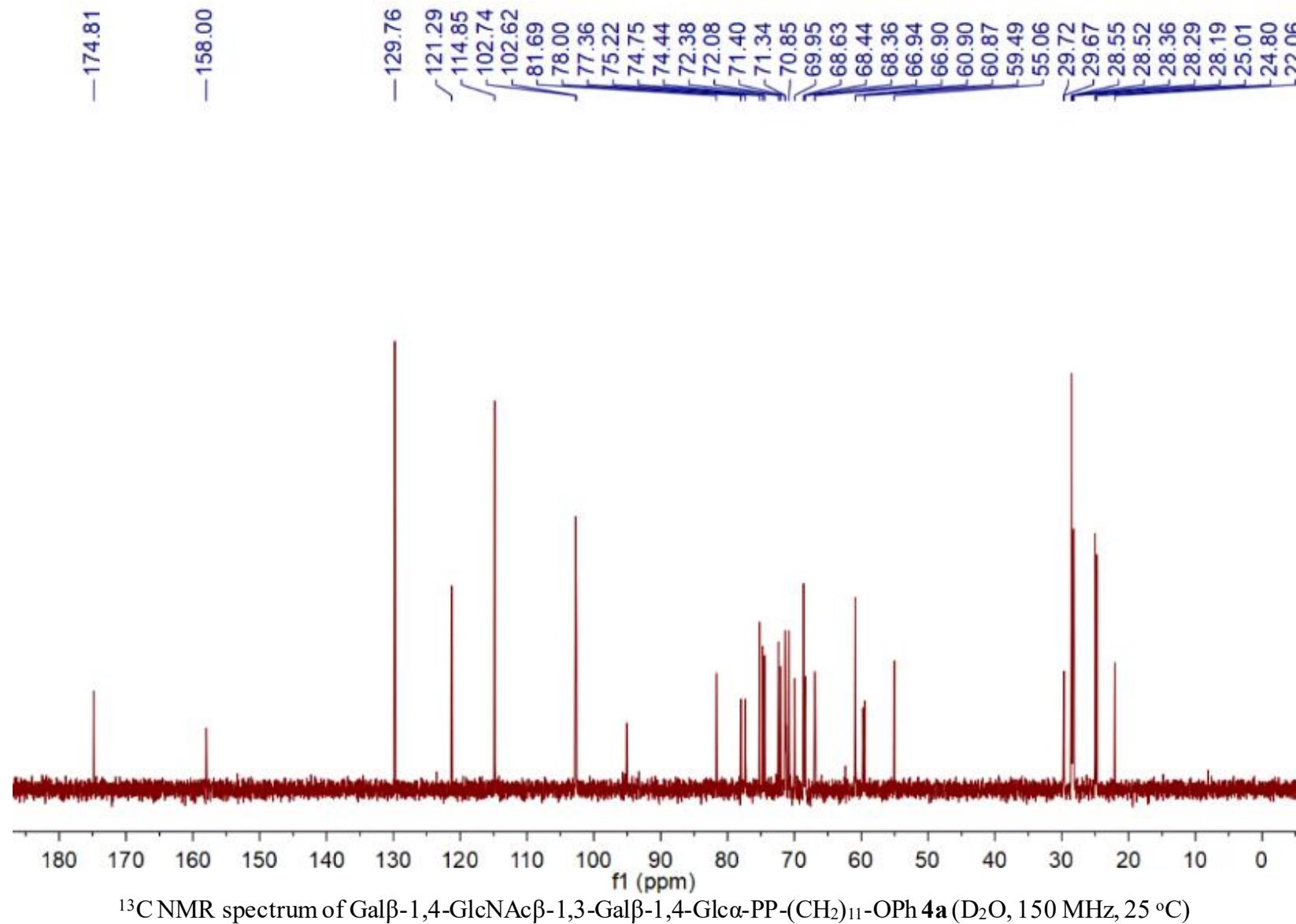


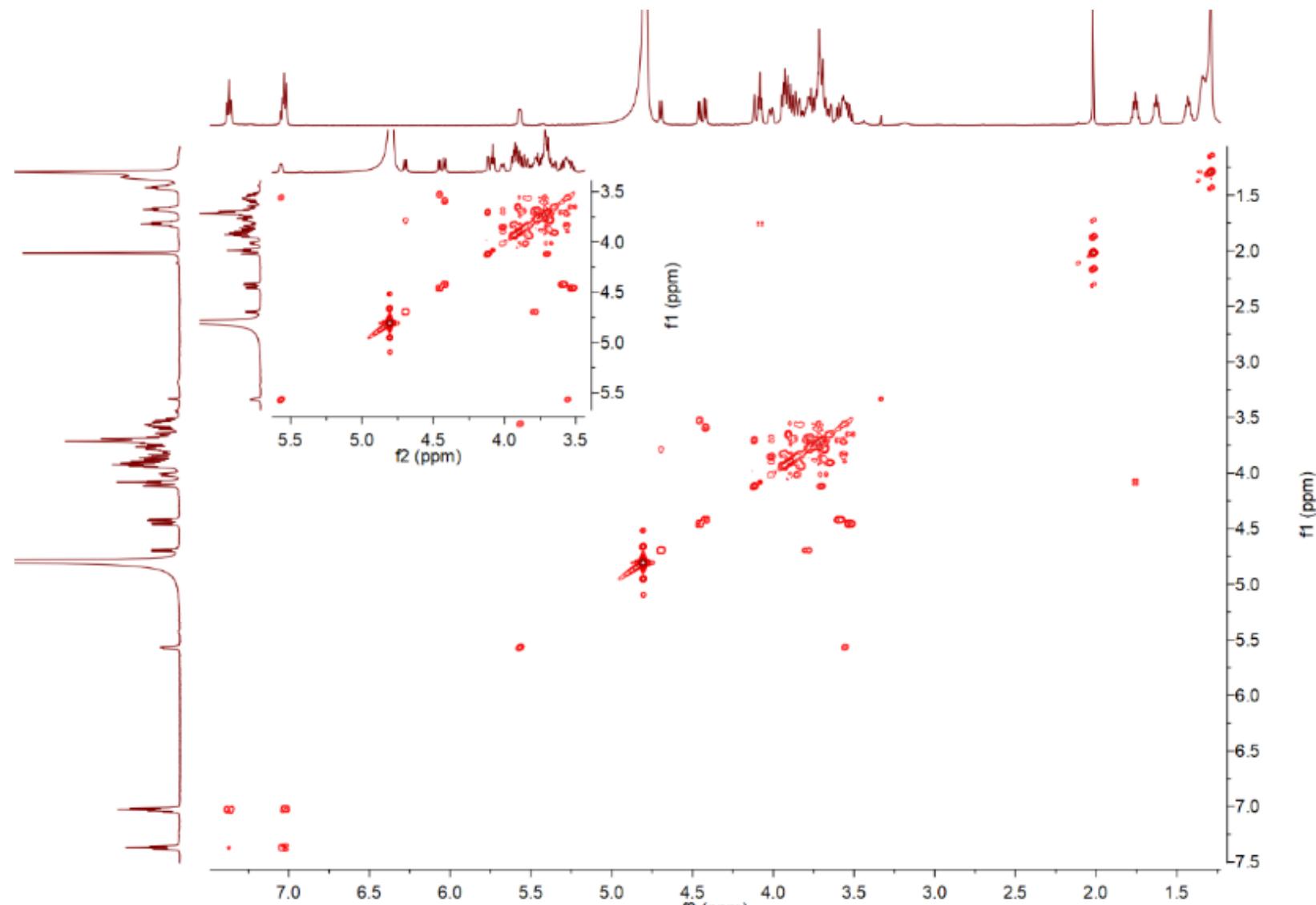




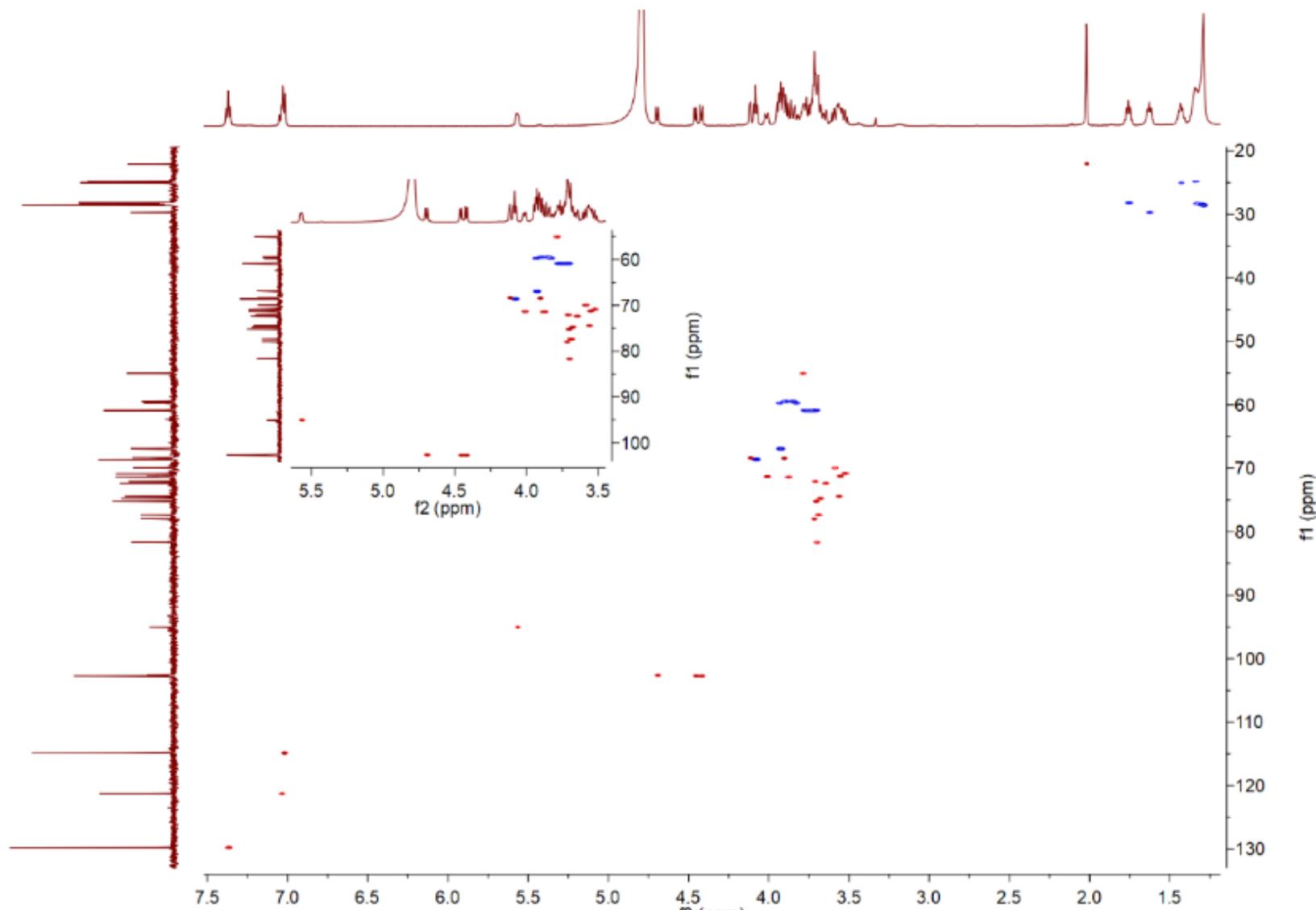


¹H NMR spectrum of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a** (D₂O, 600 MHz, 25 °C)

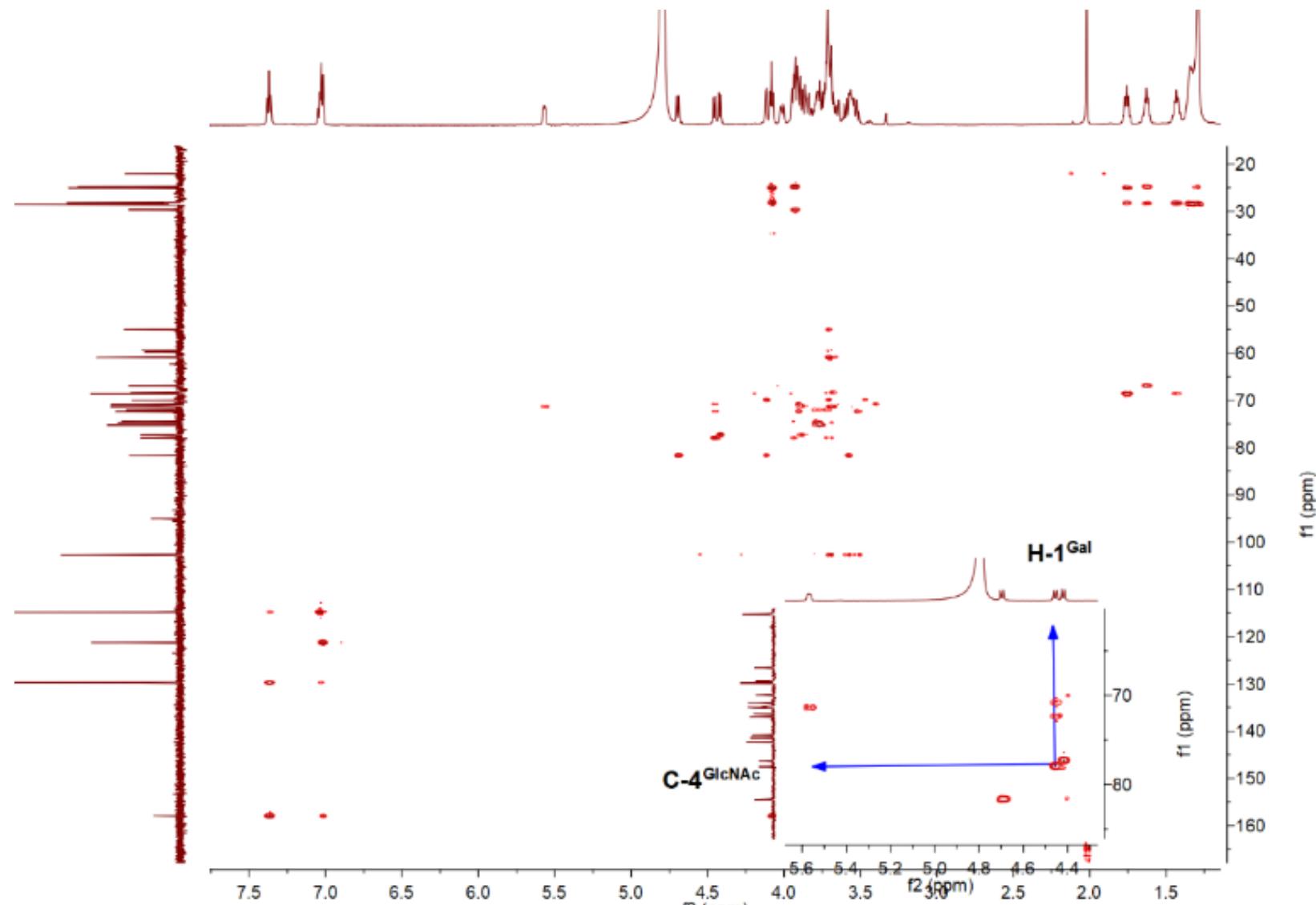




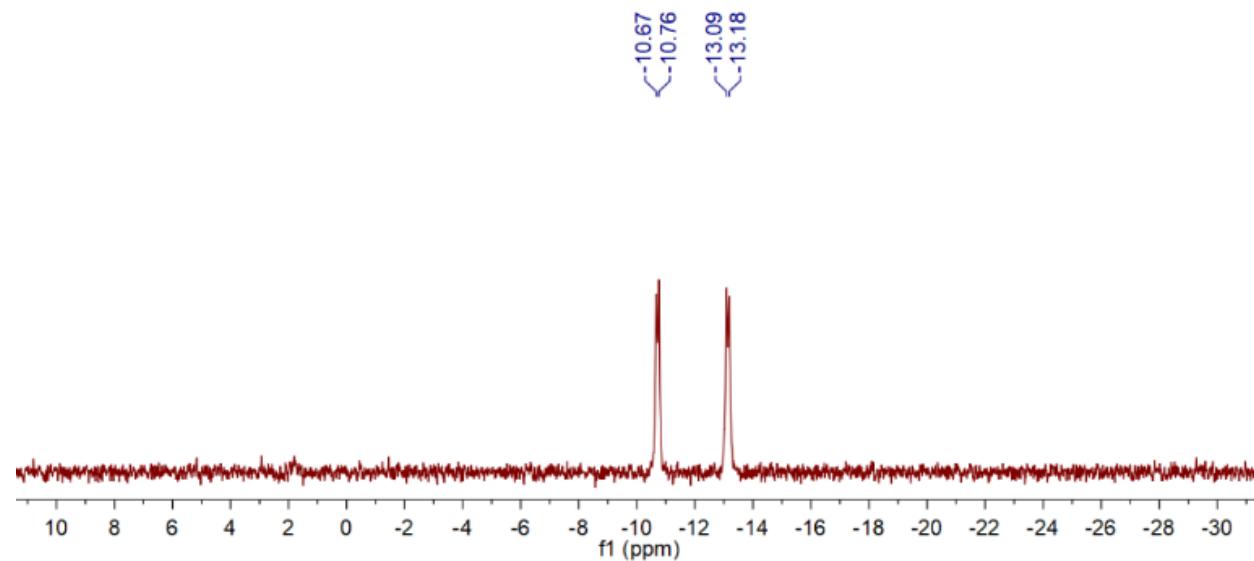
COSY spectrum of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glca-PP-(CH₂)₁₁-OPh **4a** (D₂O, 600/600 MHz, 25 °C)



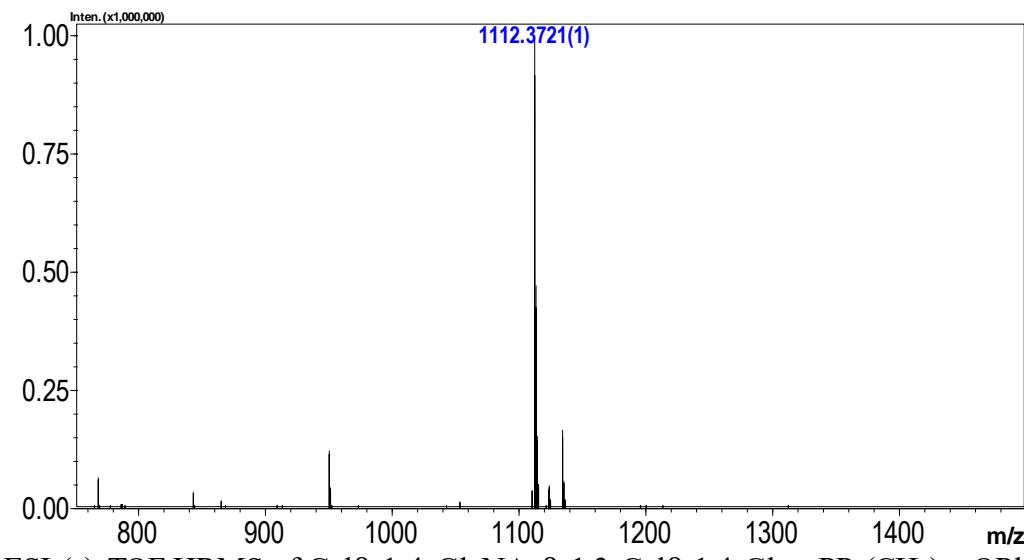
HSQC spectrum of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a** (D₂O, 600/150 MHz, 25 °C)



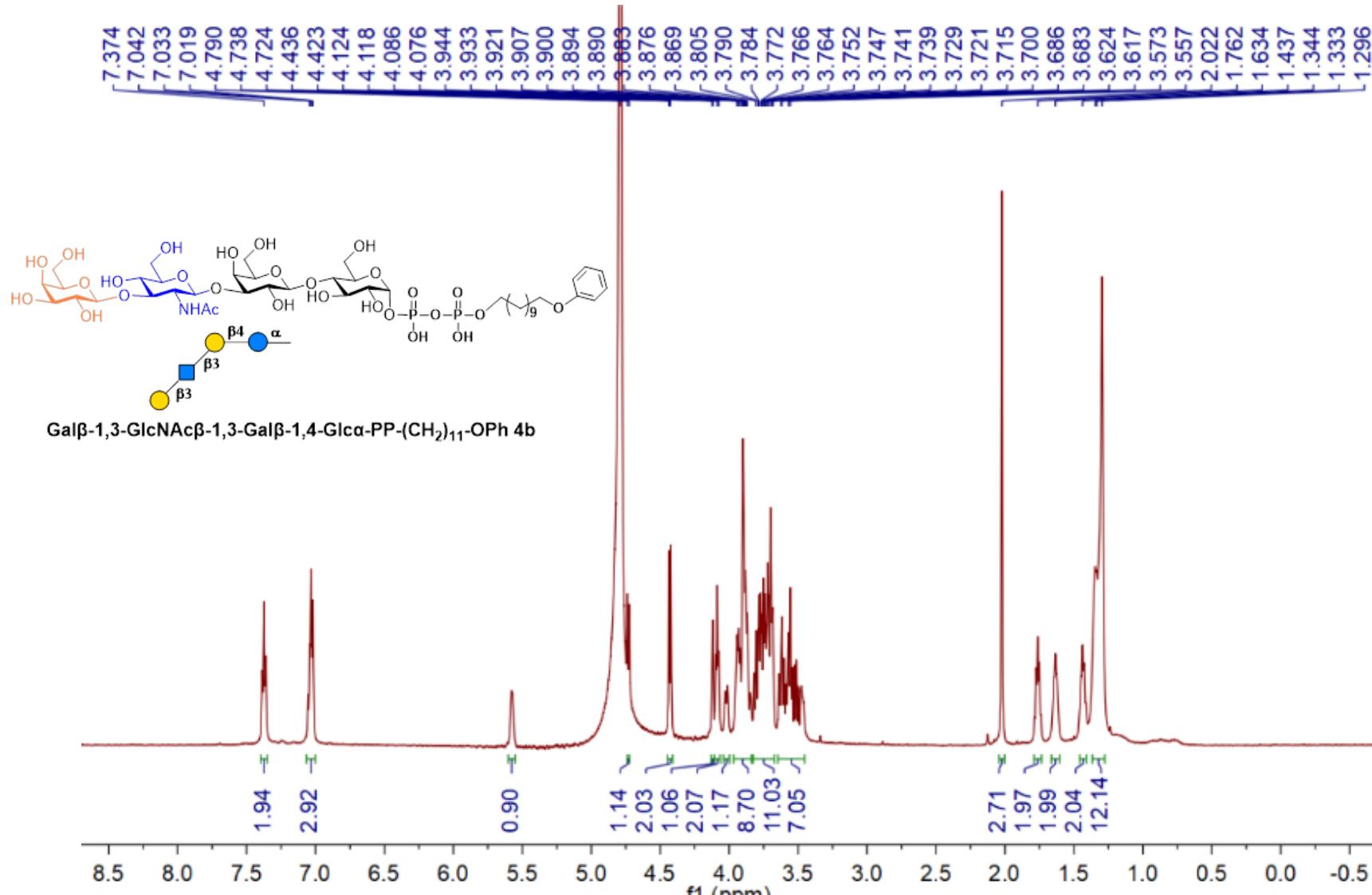
HMBC spectrum of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a** (D₂O, 600/150 MHz, 25 °C)



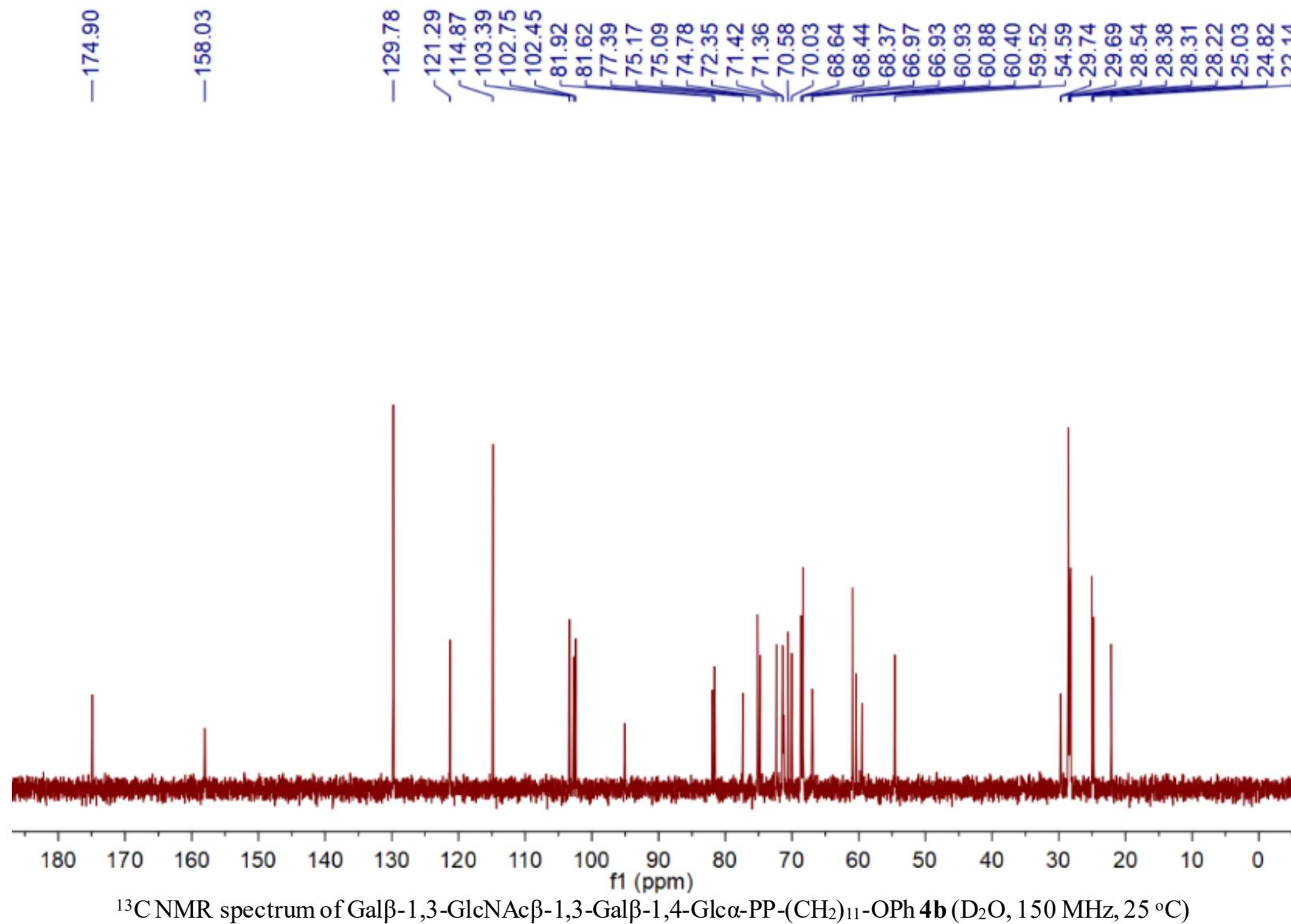
^{31}P NMR spectrum of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a** (D₂O, 243 MHz, 25 °C)

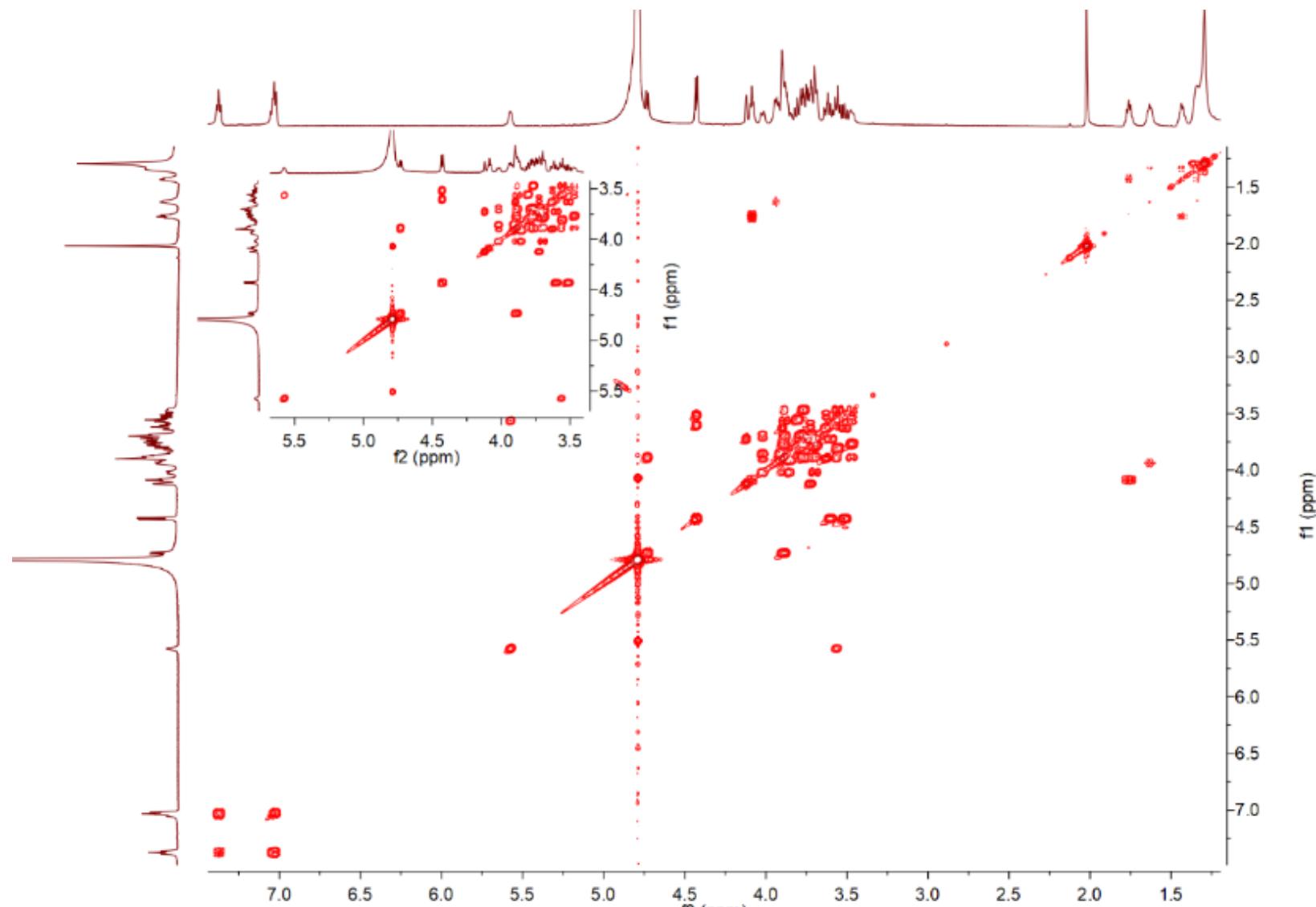


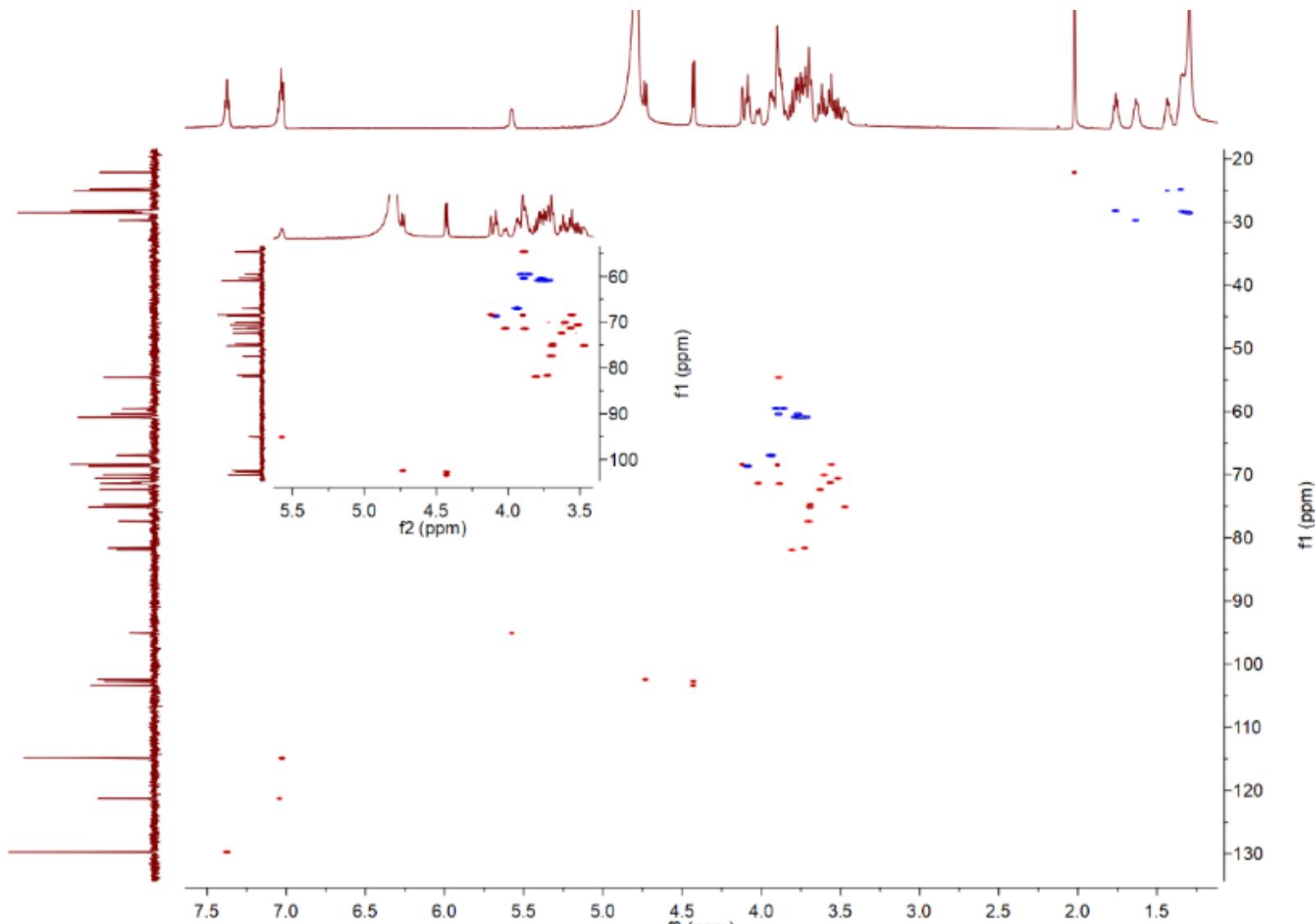
ESI(-)-TOF HRMS of Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4a**



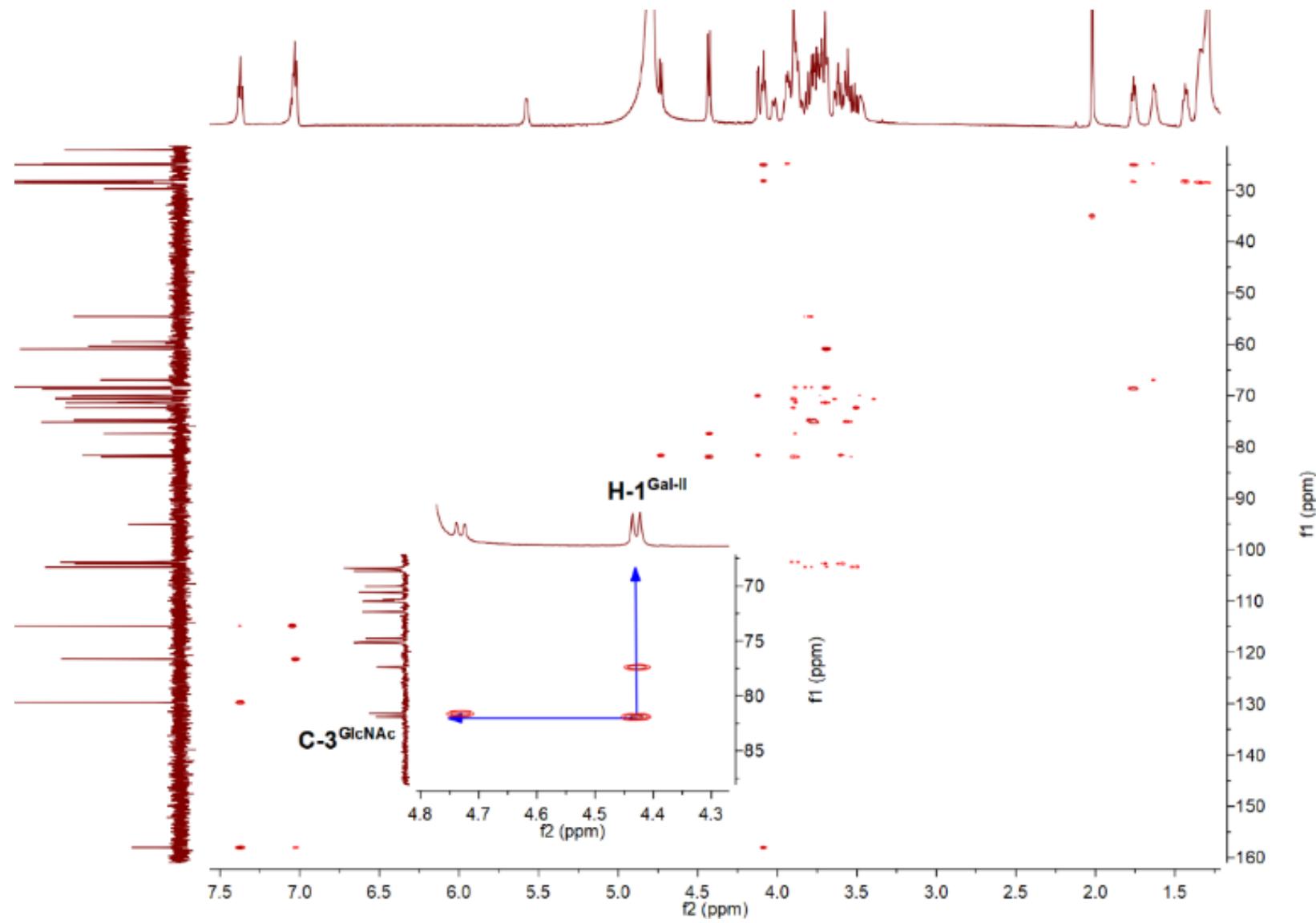
¹H NMR spectrum of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b** (D₂O, 600 MHz, 25 °C)



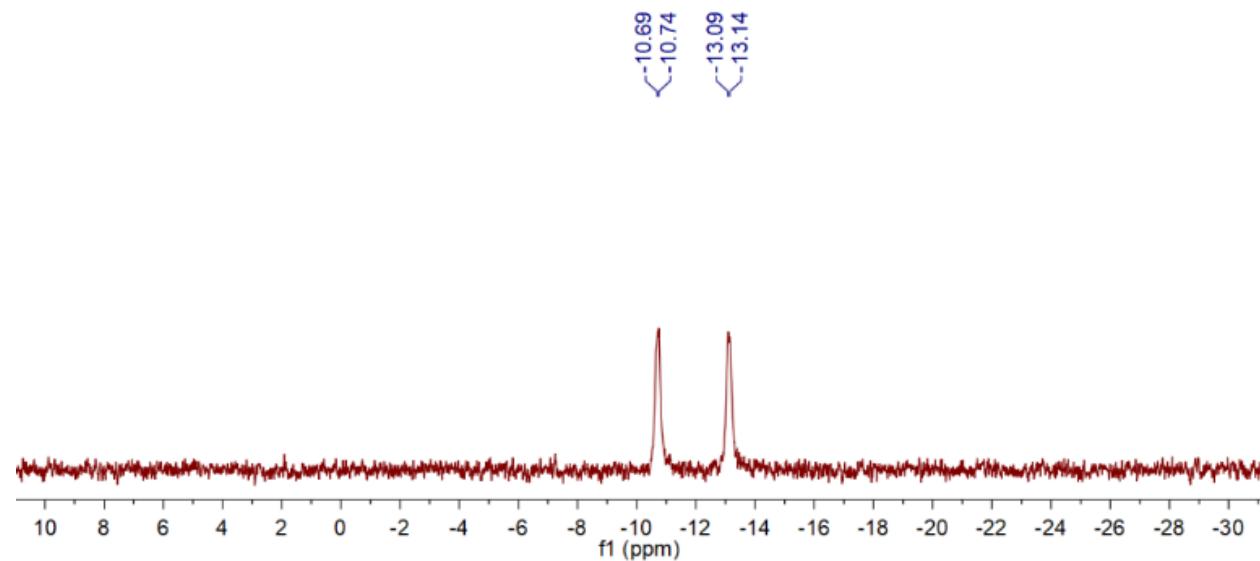




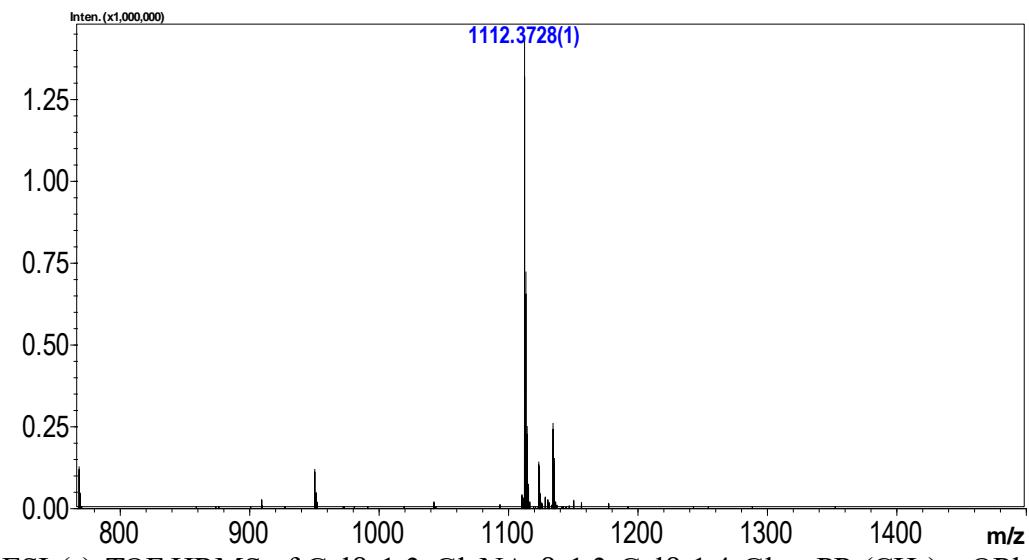
HSQC spectrum of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glca-PP-(CH₂)₁₁-OPh **4b** (D₂O, 600/150 MHz, 25 °C)



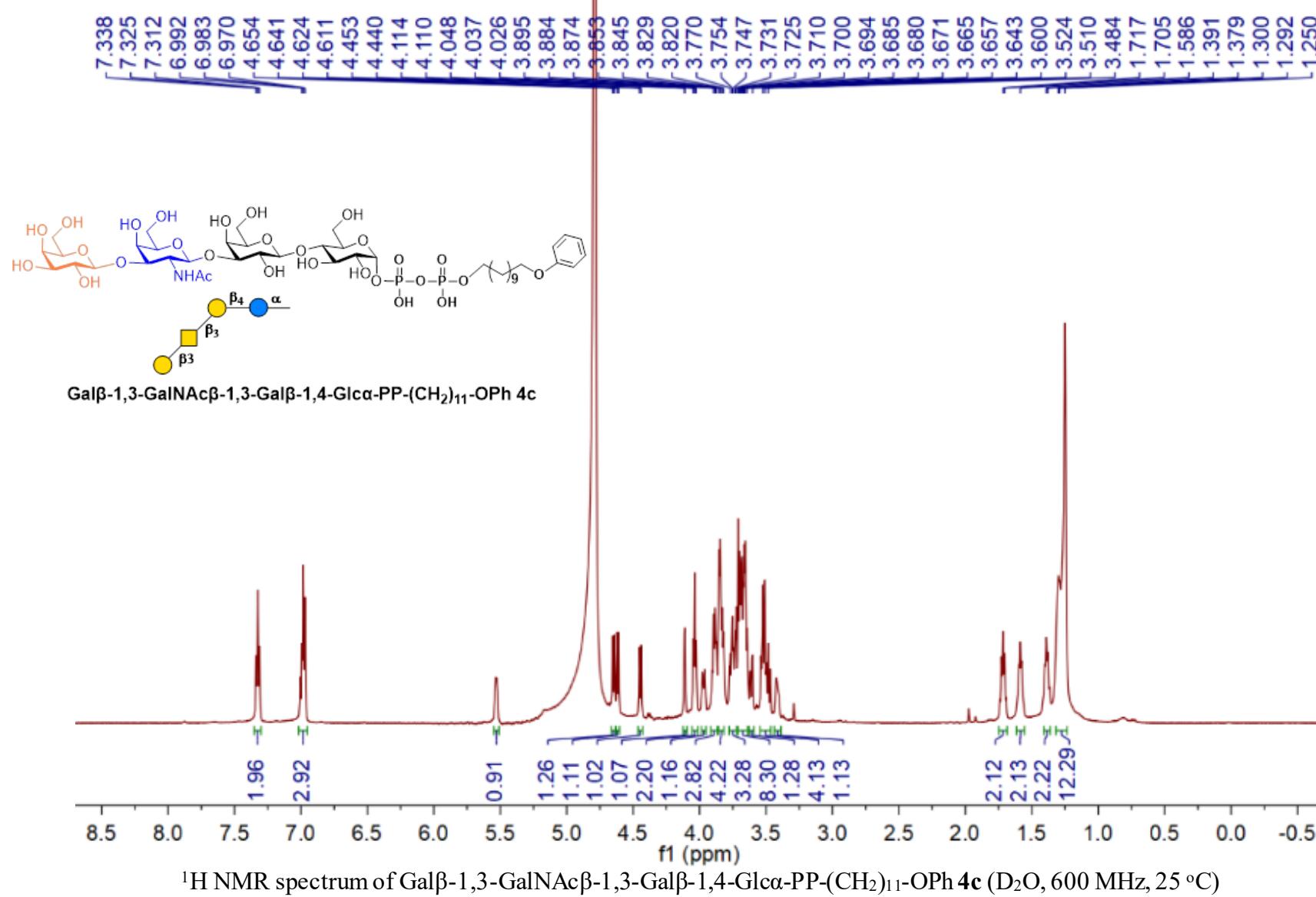
HMBC spectrum of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b** (D₂O, 600/150 MHz, 25 °C)



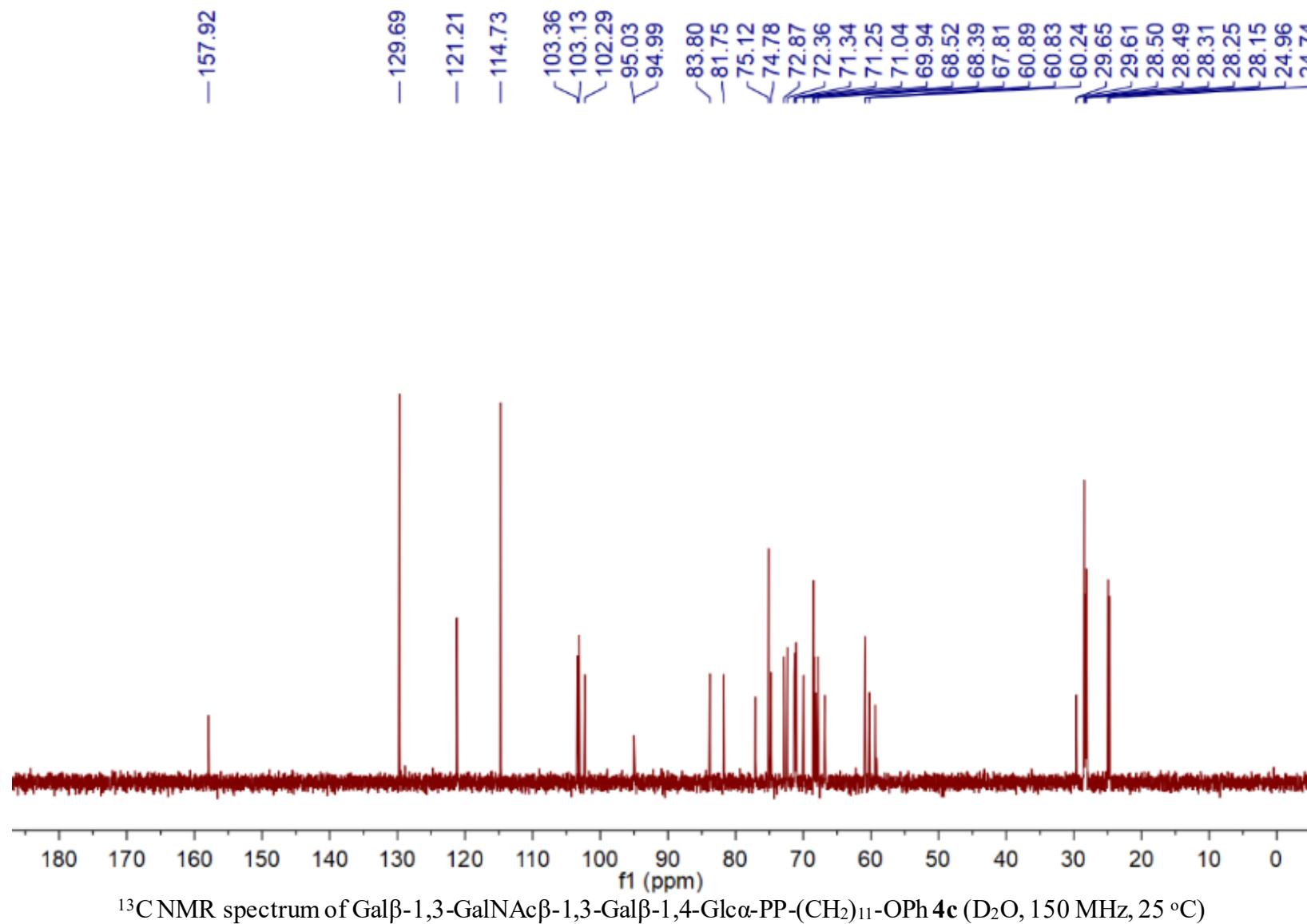
^{31}P NMR spectrum of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b** (D₂O, 243 MHz, 25 °C)

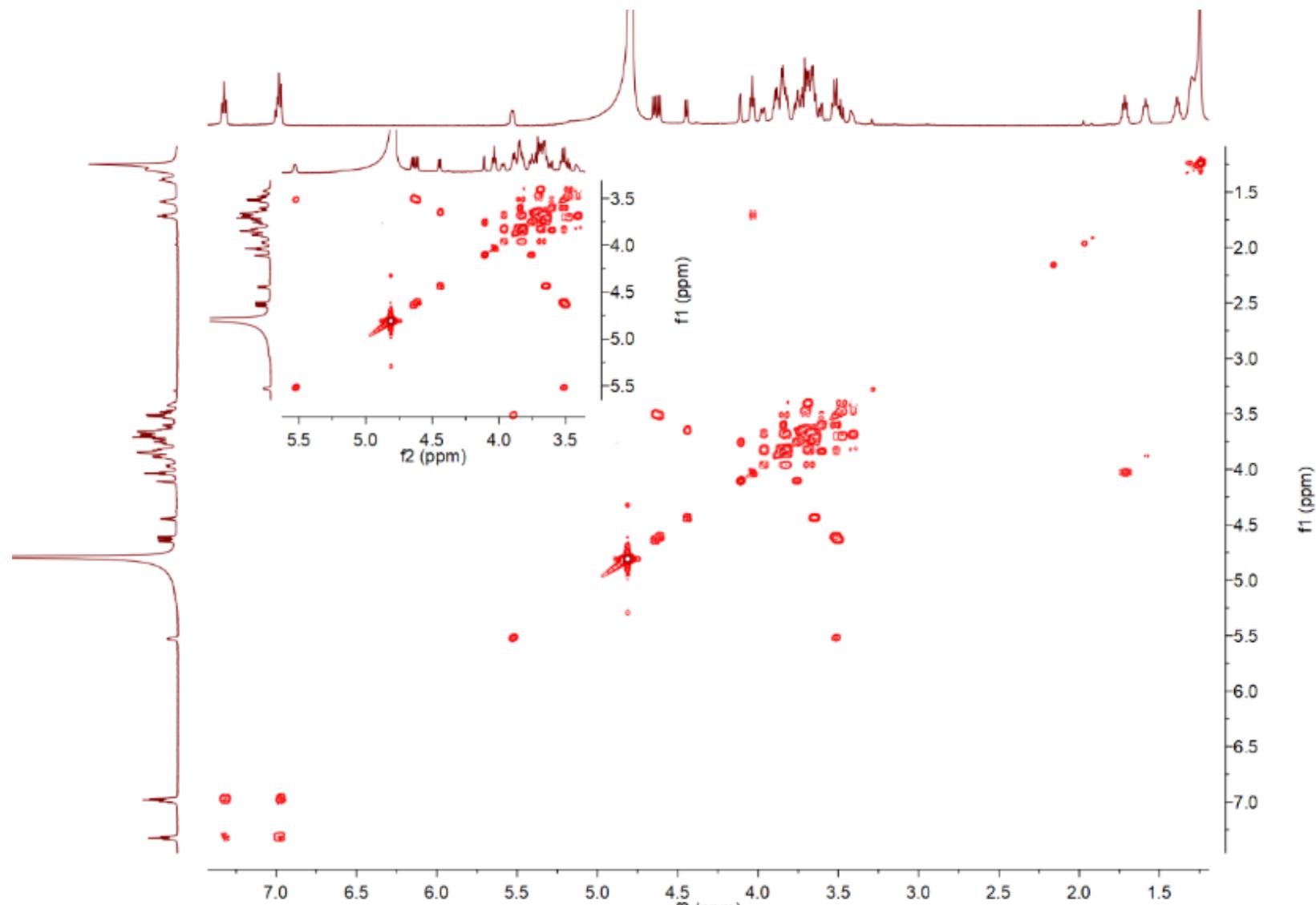


ESI(-)-TOF HRMS of Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4b**

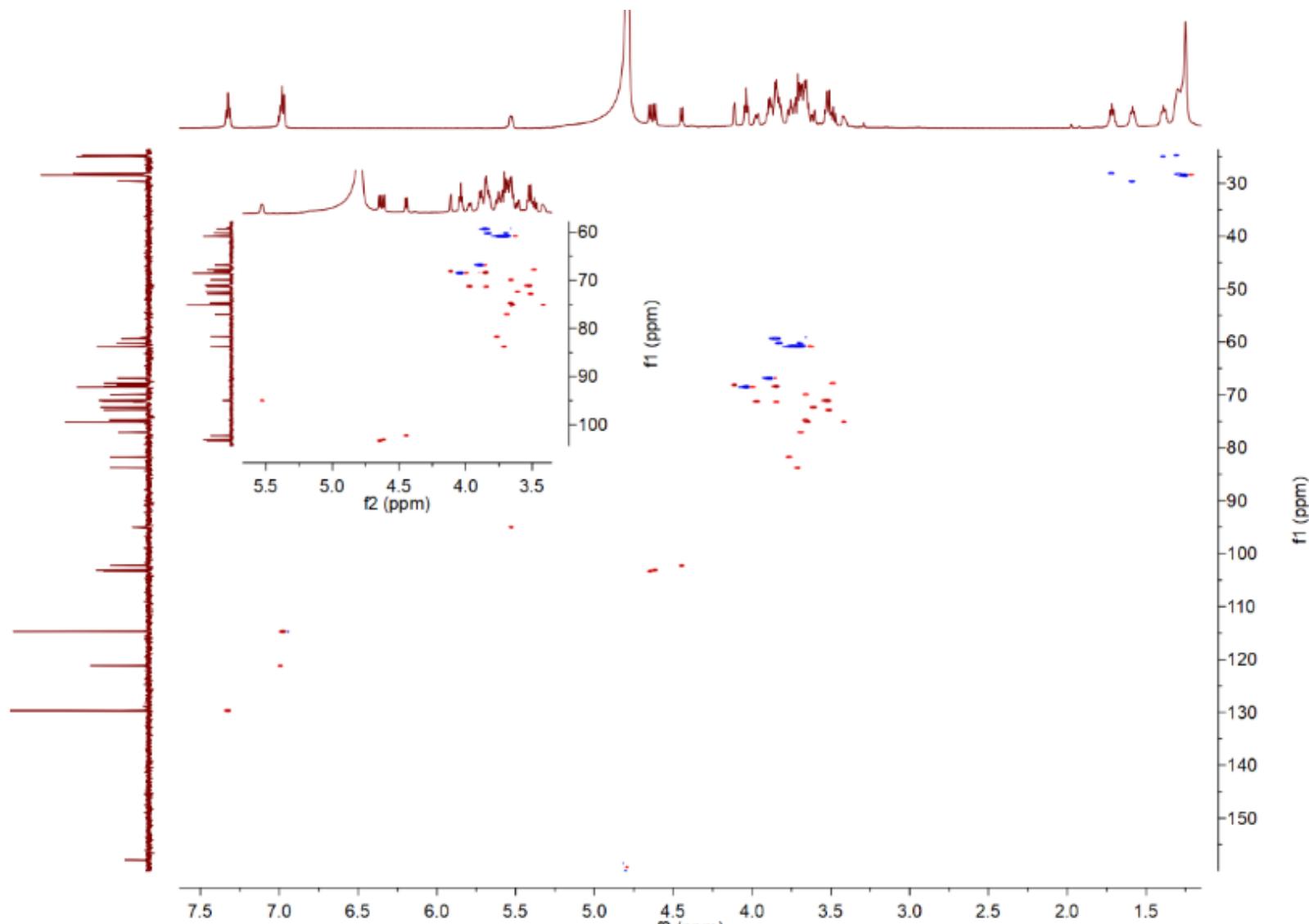


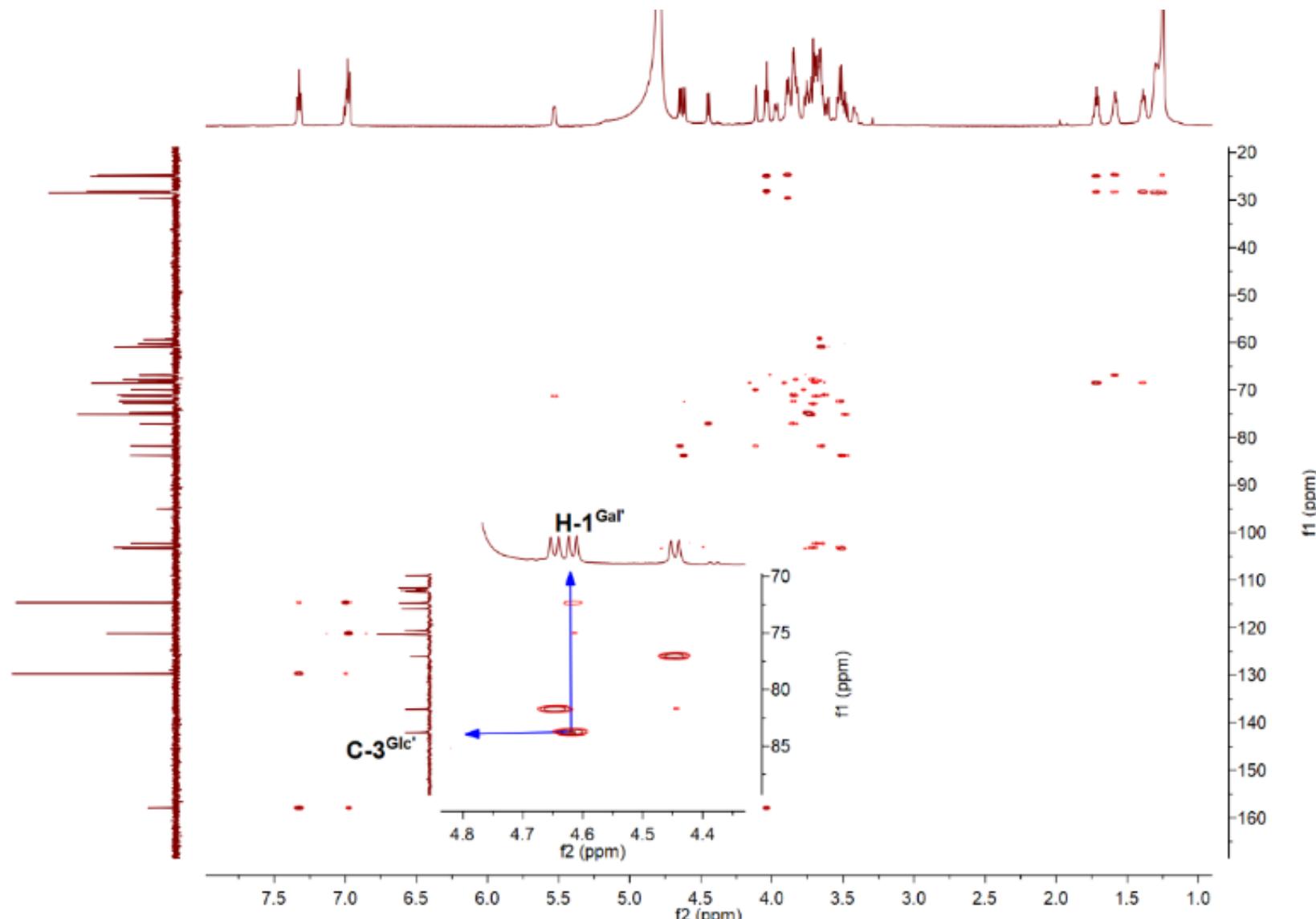
¹H NMR spectrum of Galβ-1,3-GalNAcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **4c** (D₂O, 600 MHz, 25 °C)



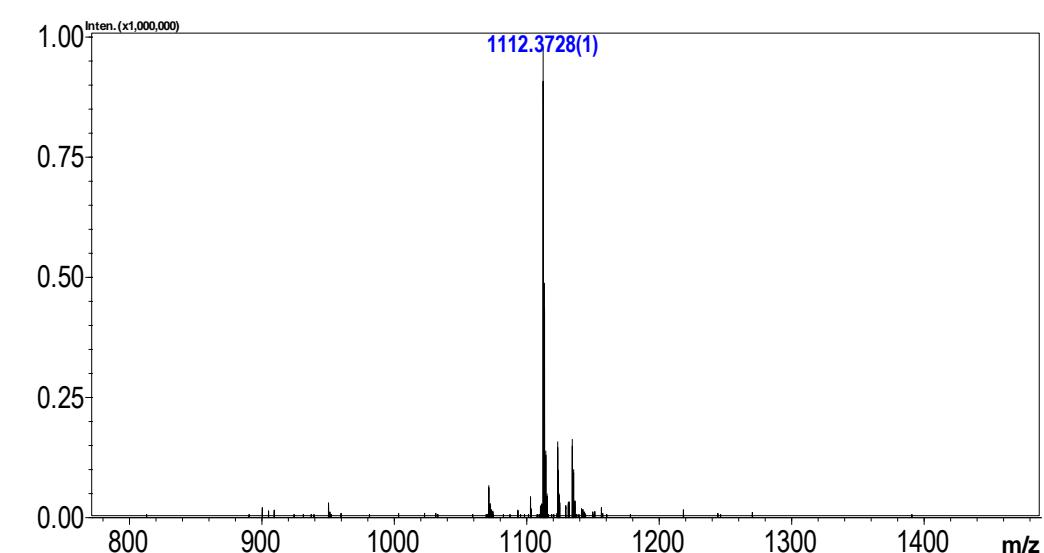
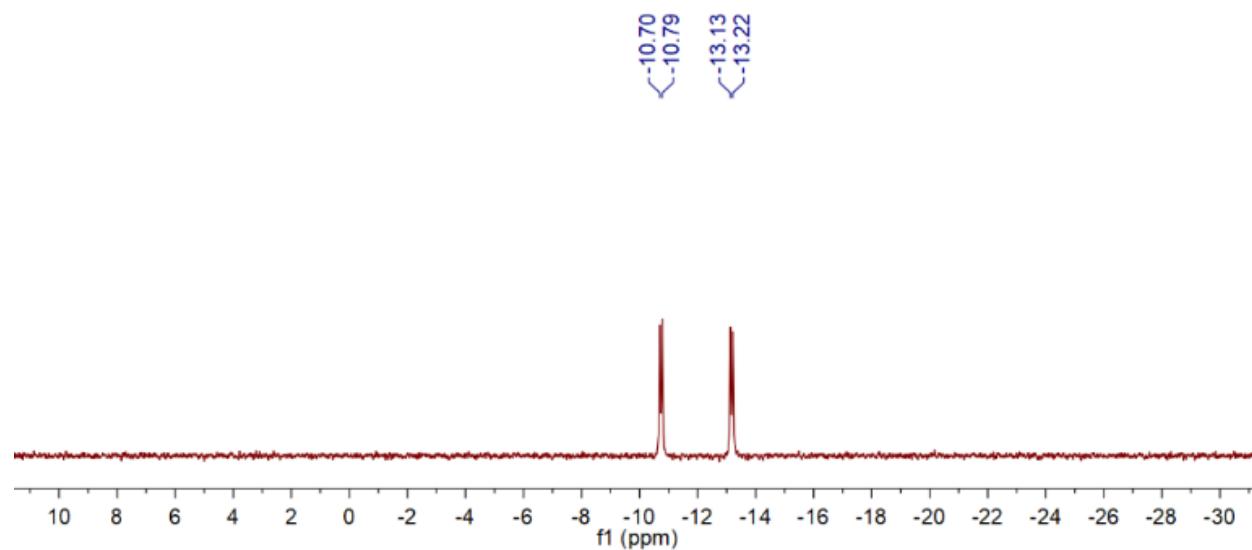


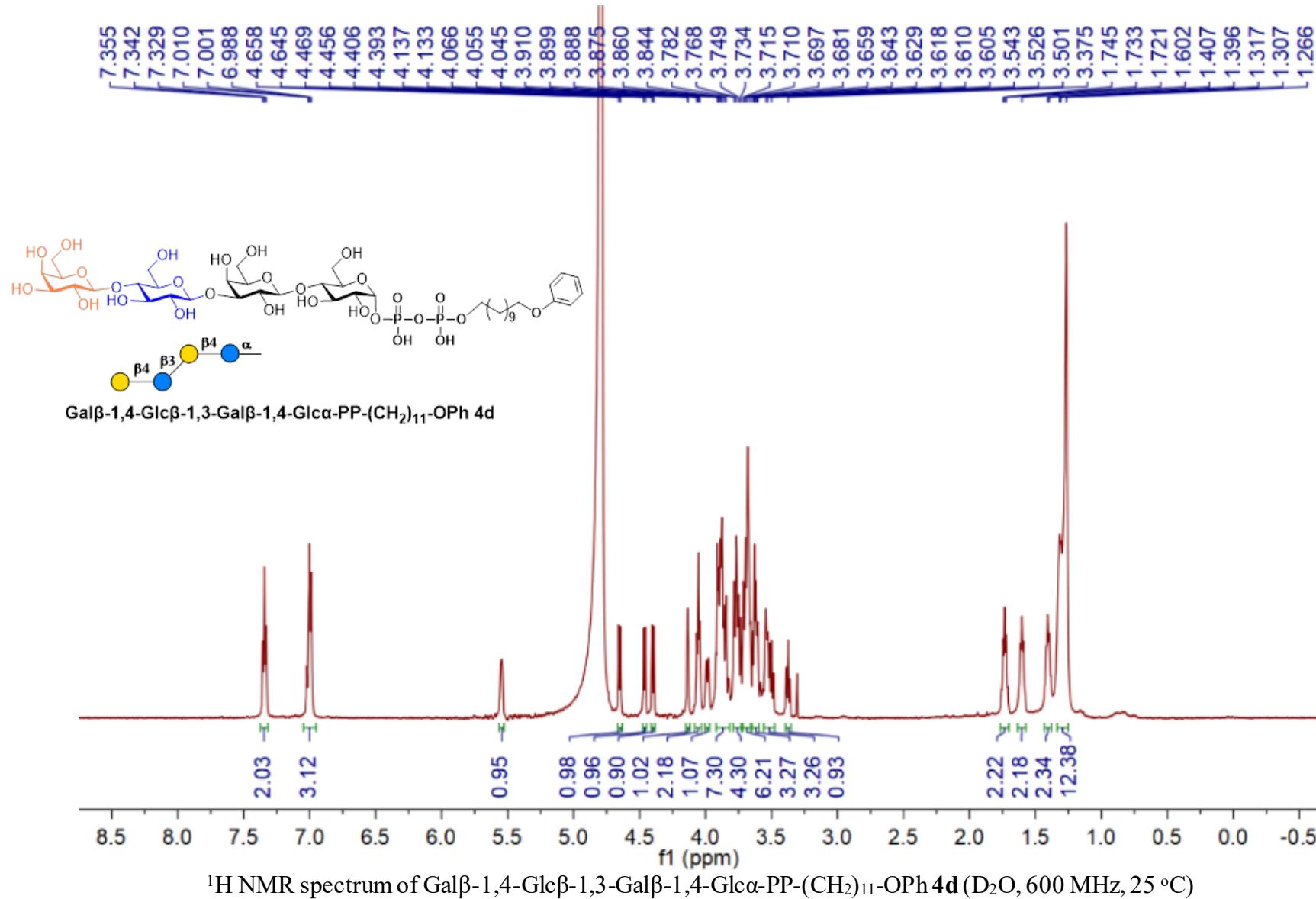
COSY spectrum of Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glca-PP-(CH₂)₁₁-OPh **4c** (D₂O, 600/600 MHz, 25 °C)

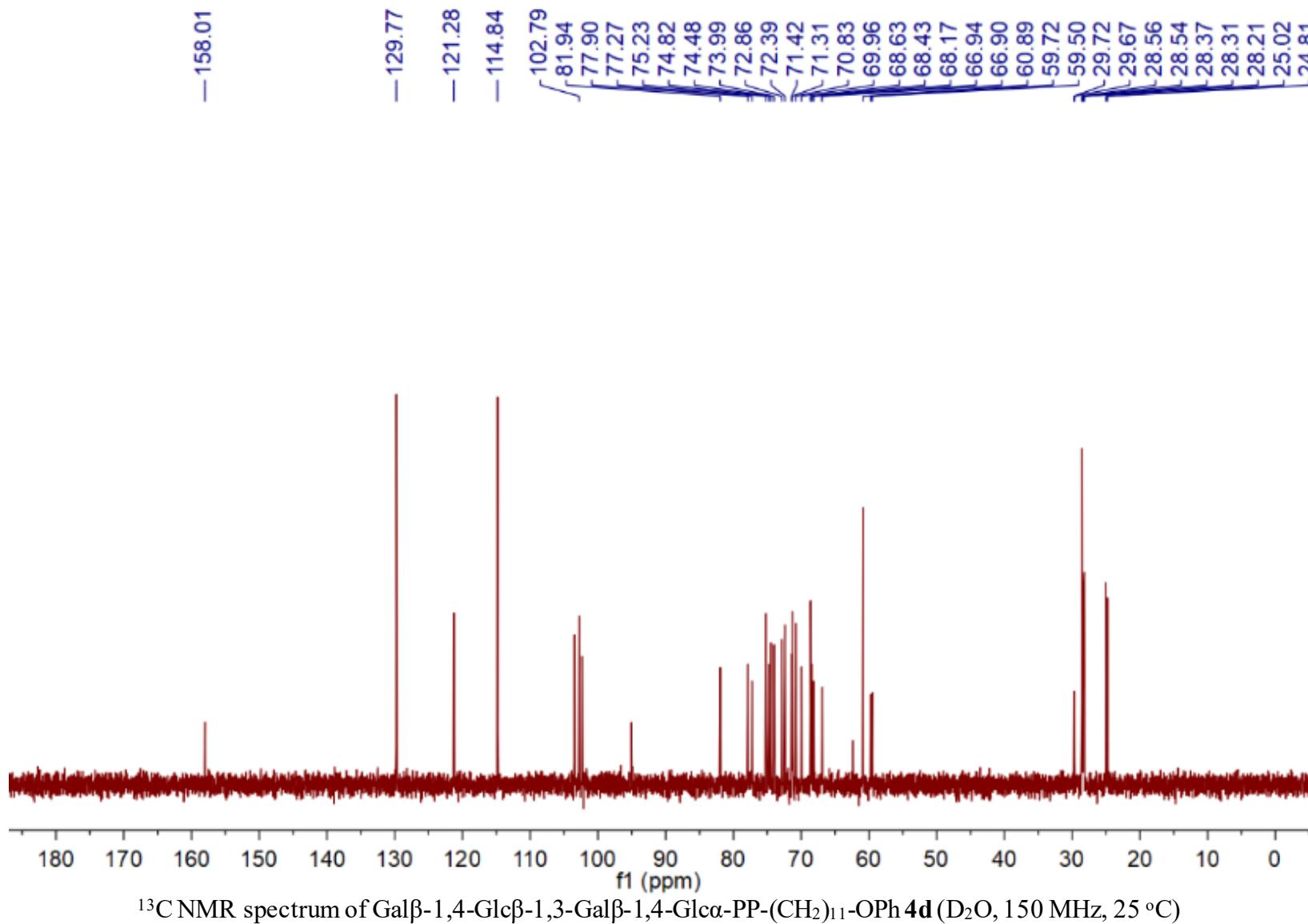


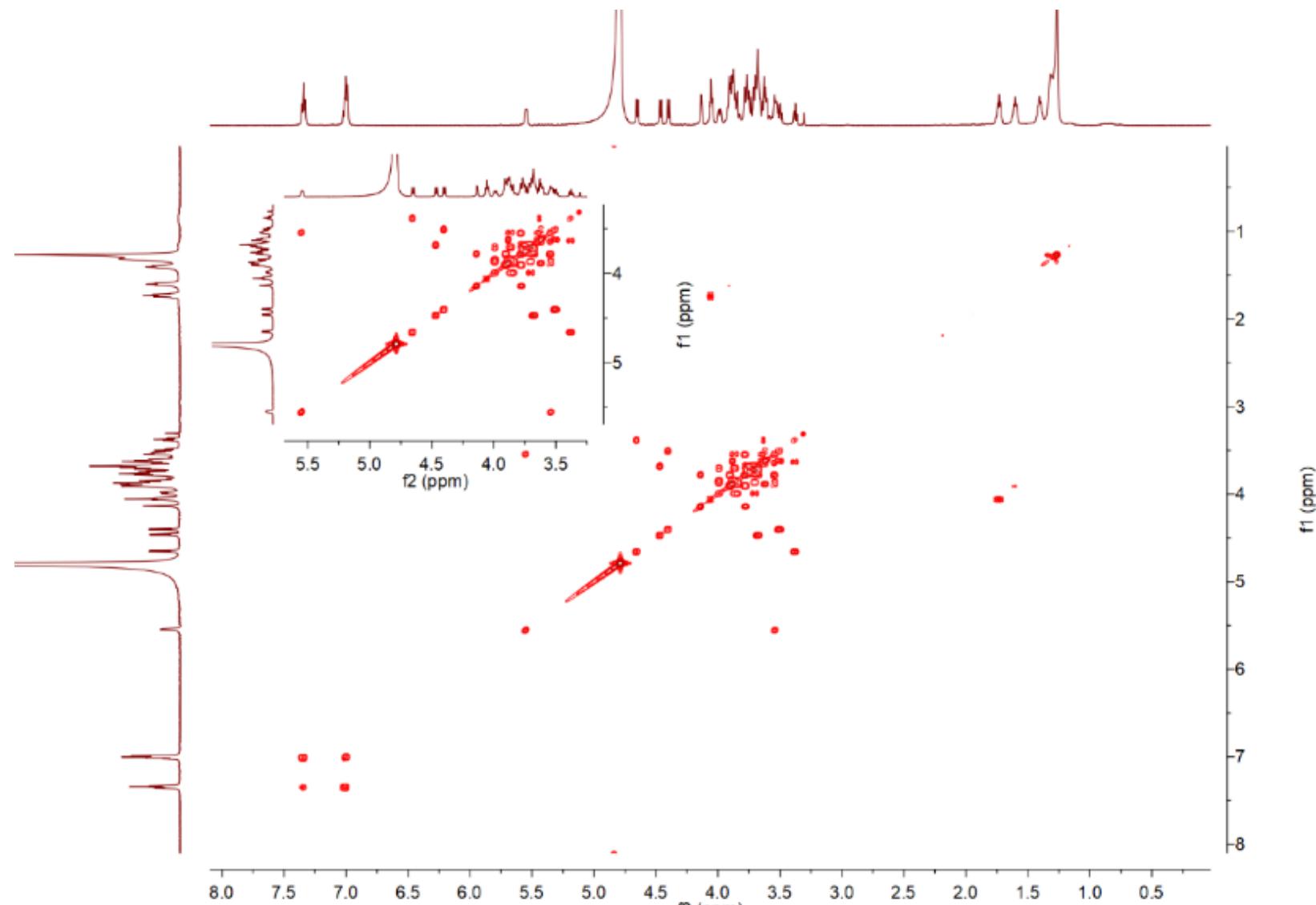


HMBC spectrum of Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glca-PP-(CH₂)₁₁-OPh **4c** (D₂O, 600/150 MHz, 25 °C)

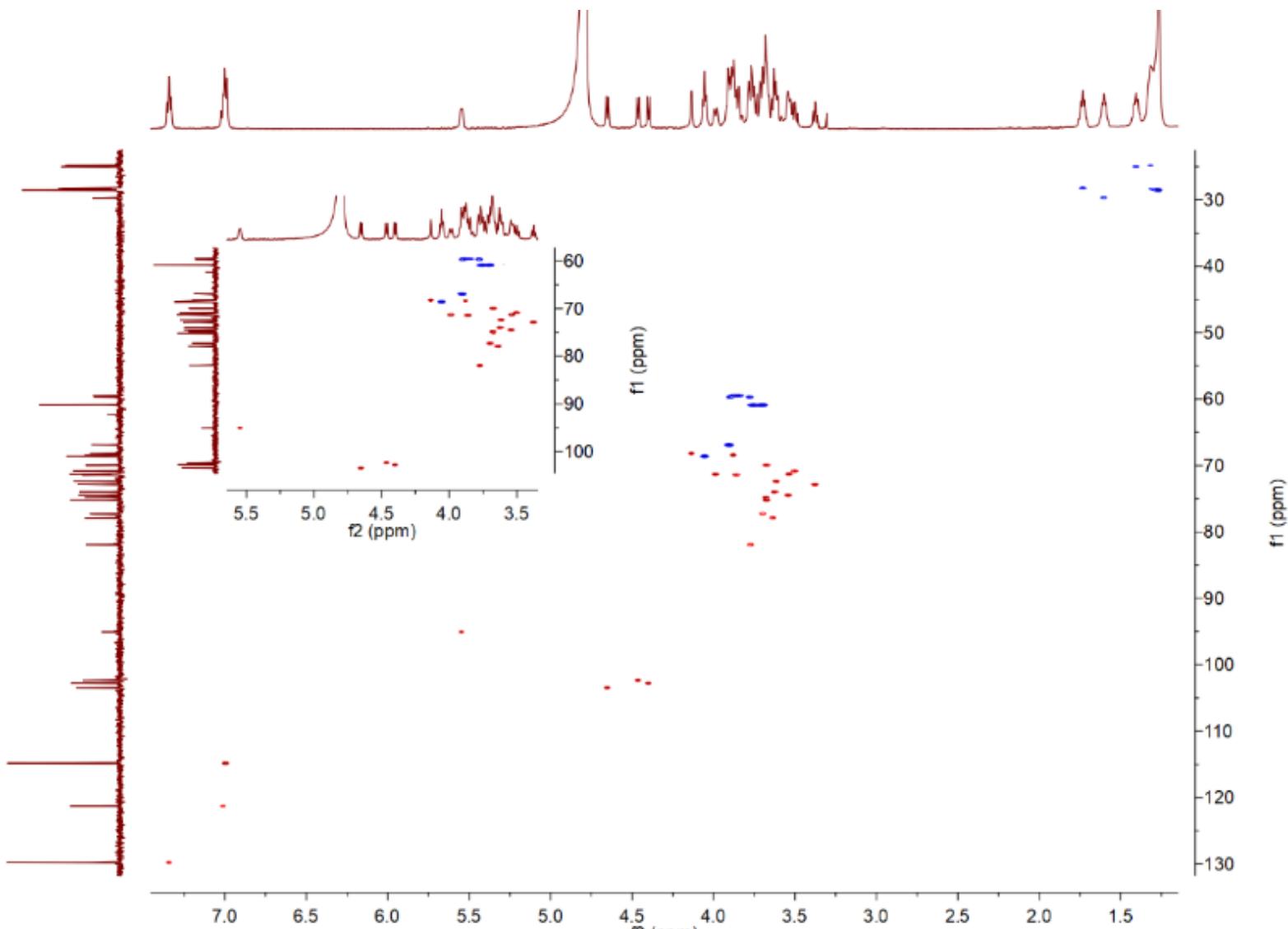


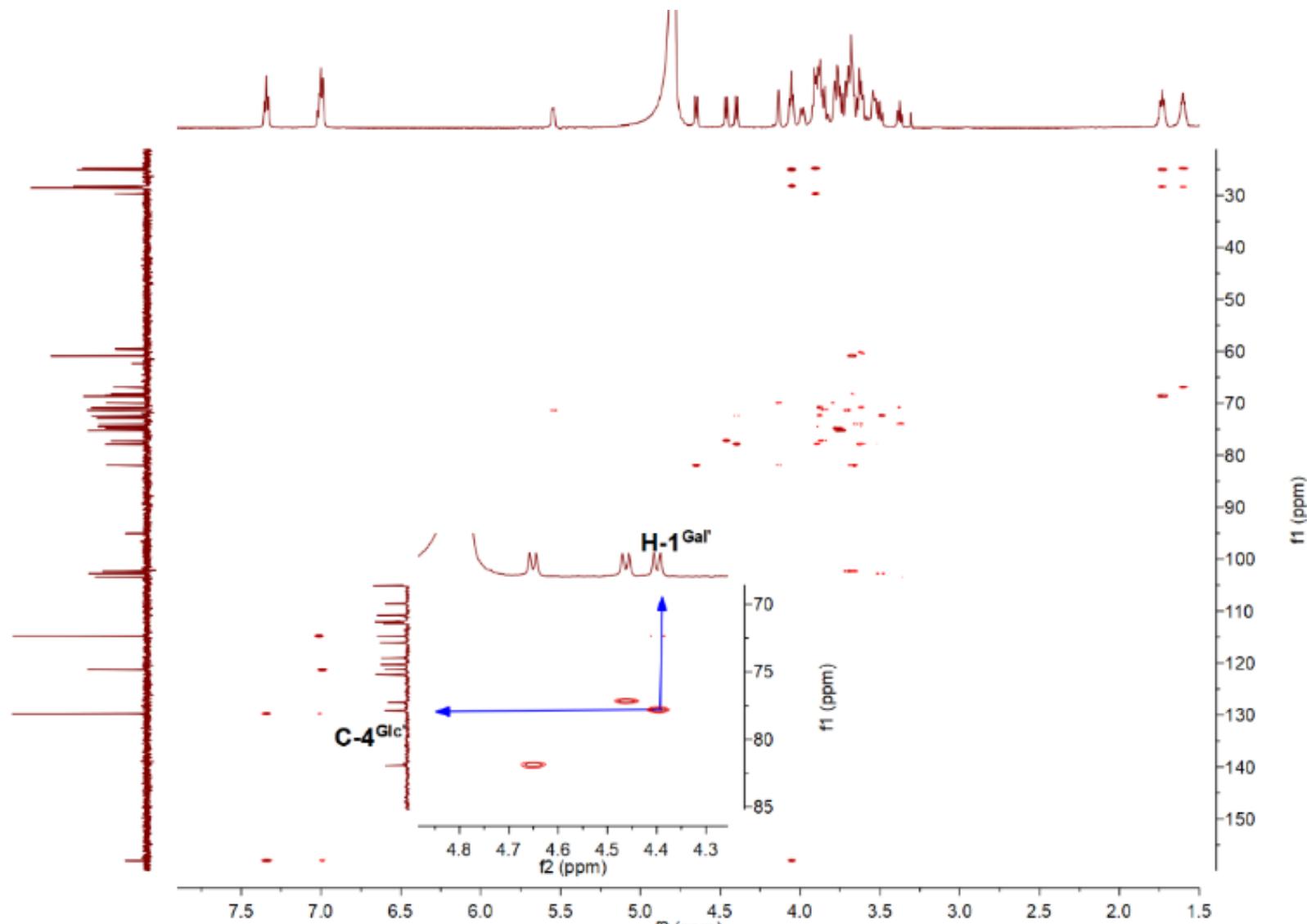


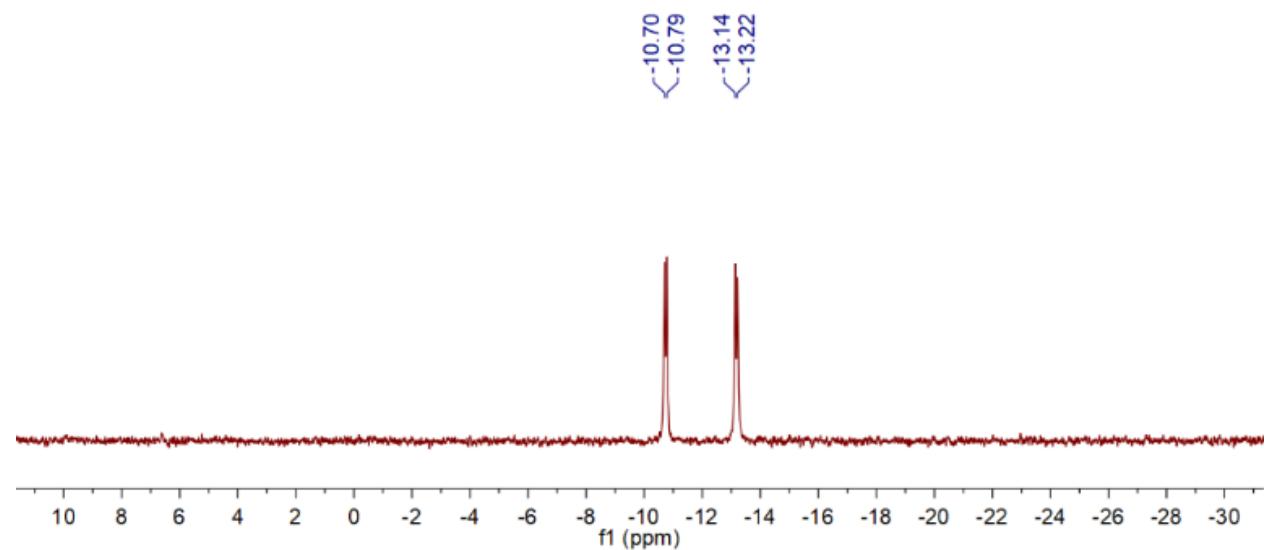




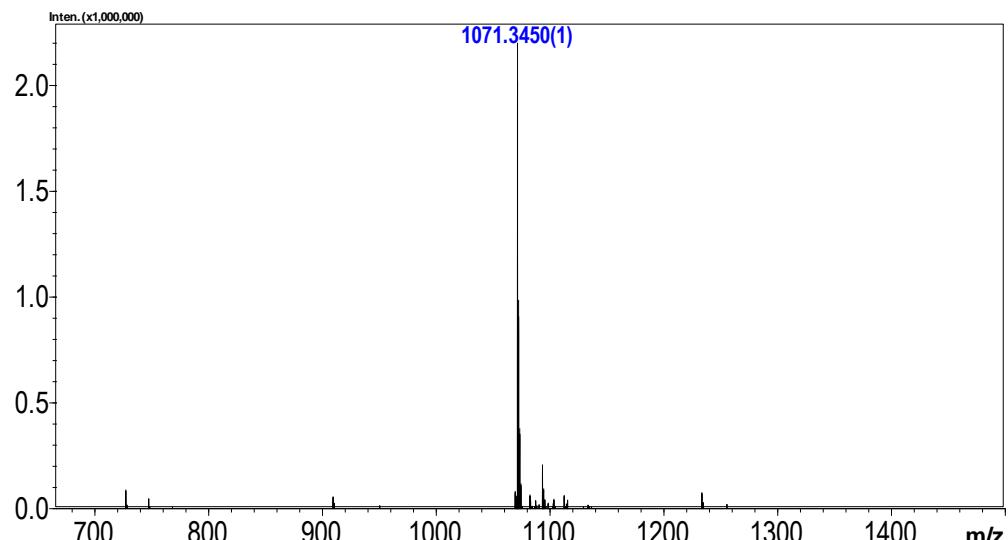
COSY spectrum of Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4d** (D₂O, 600/600 MHz, 25 °C)



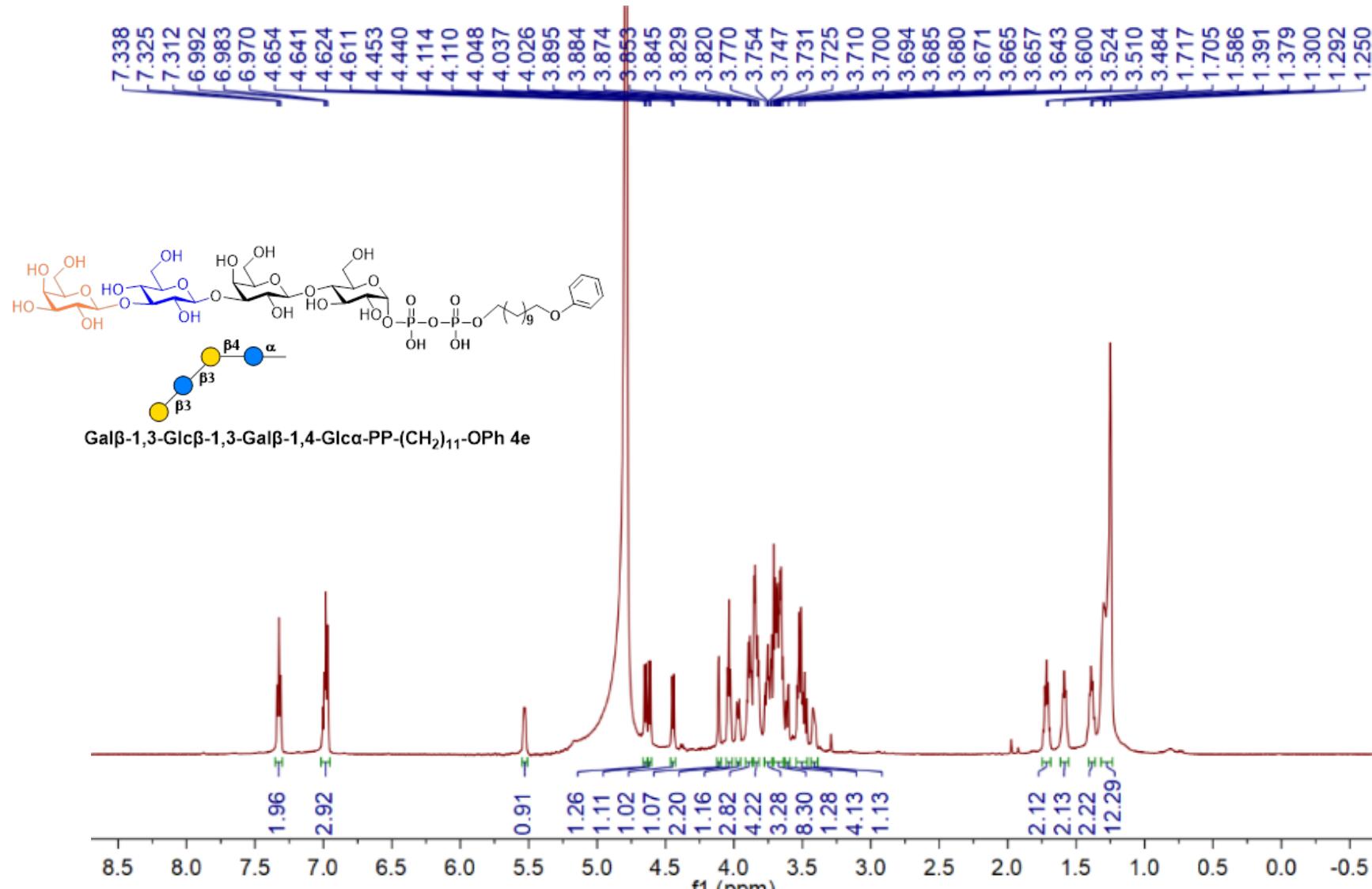




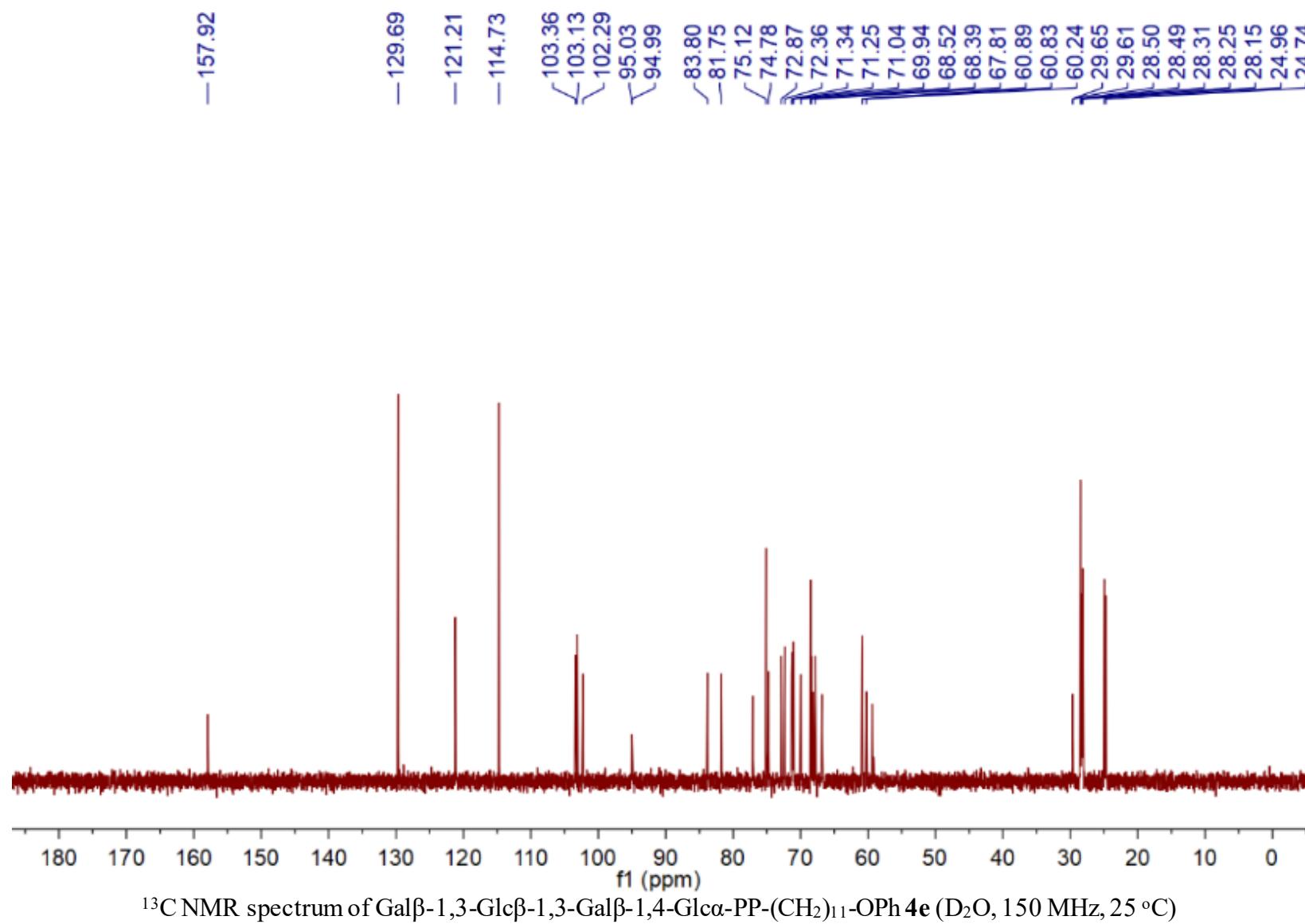
^{31}P NMR spectrum of Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4d** (D₂O, 243 MHz, 25 °C)

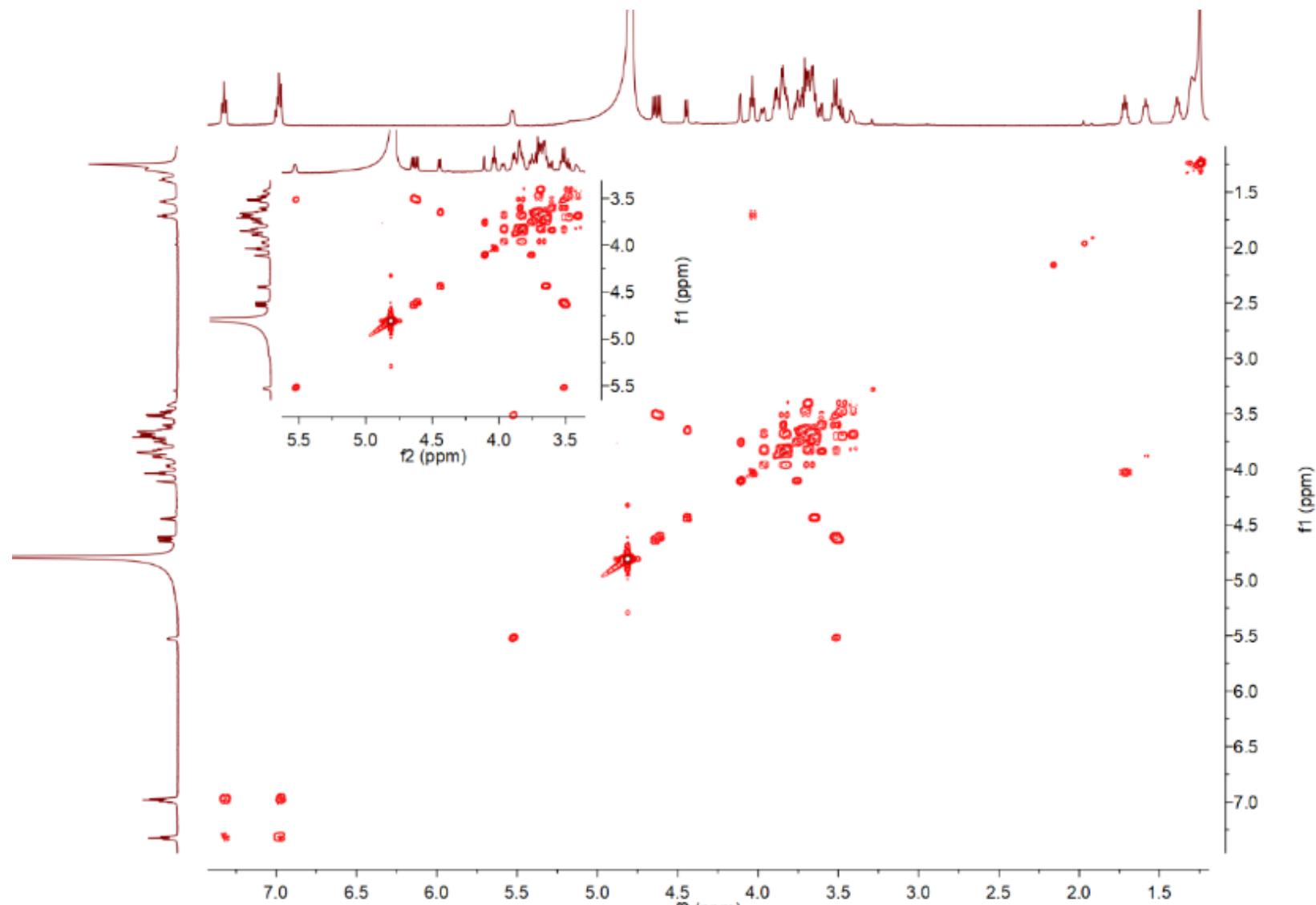


ESI(-)-TOF HRMS of Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4d**

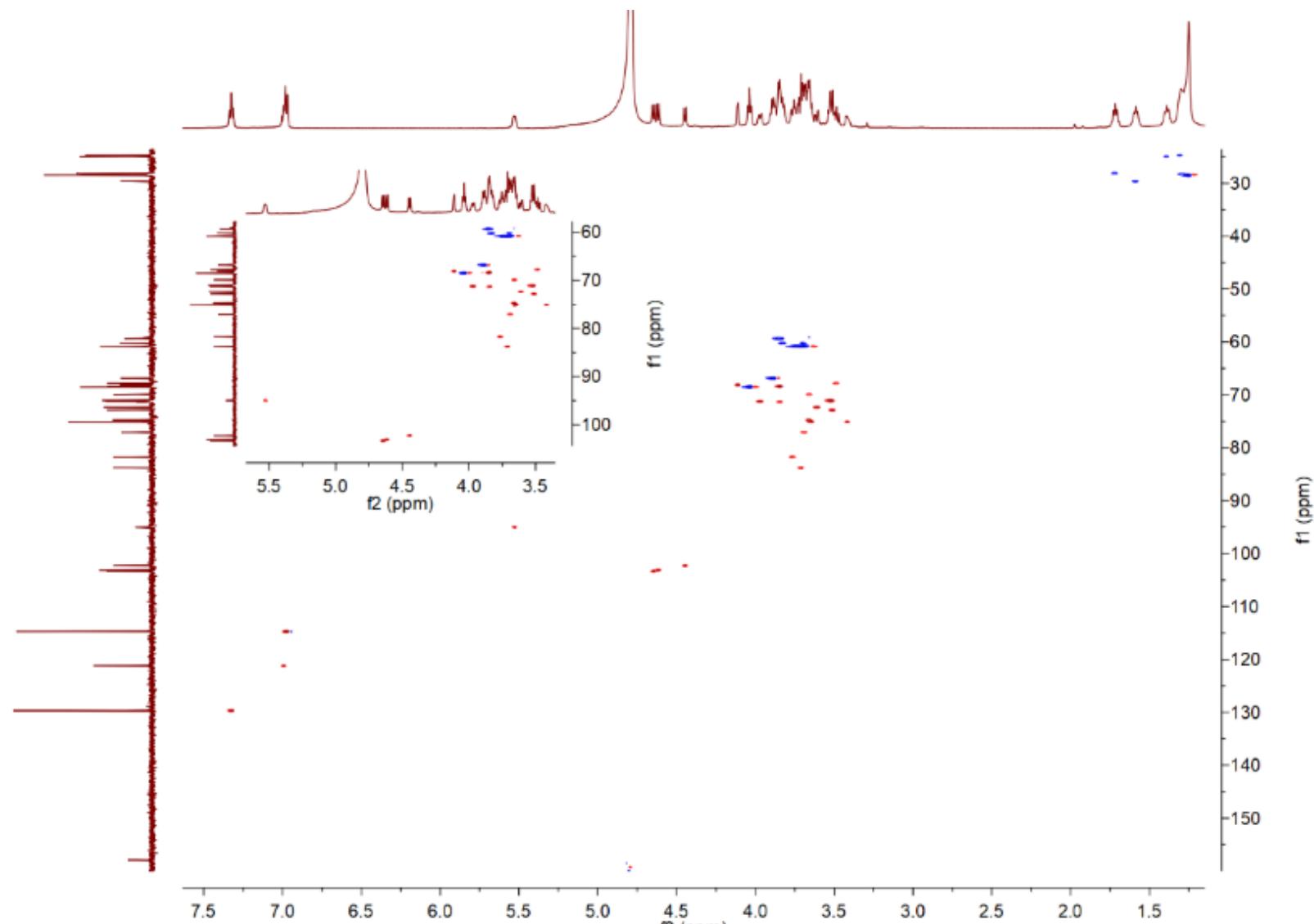


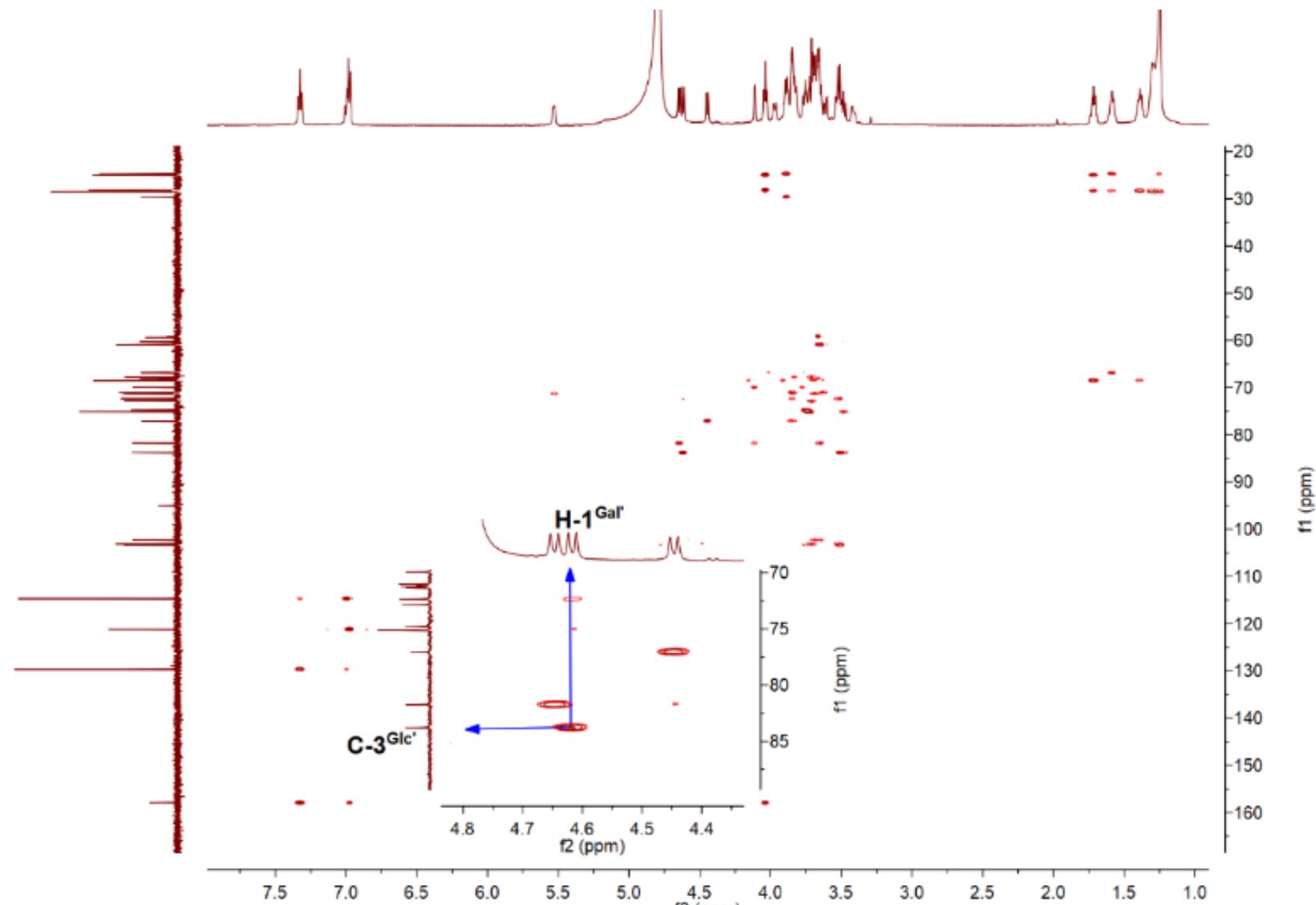
¹H NMR spectrum of Galβ-1,3-Glcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh **4e** (D₂O, 600 MHz, 25 °C)



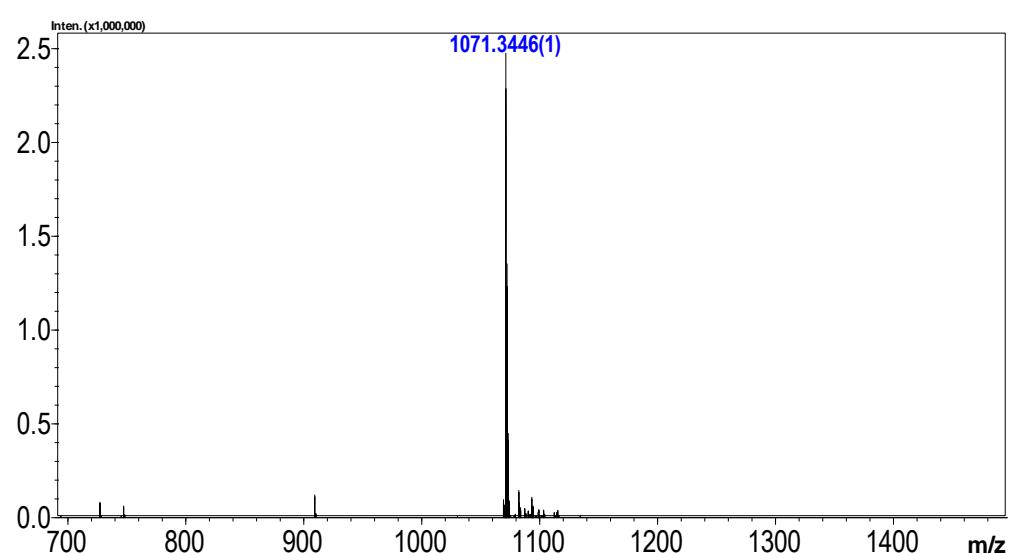
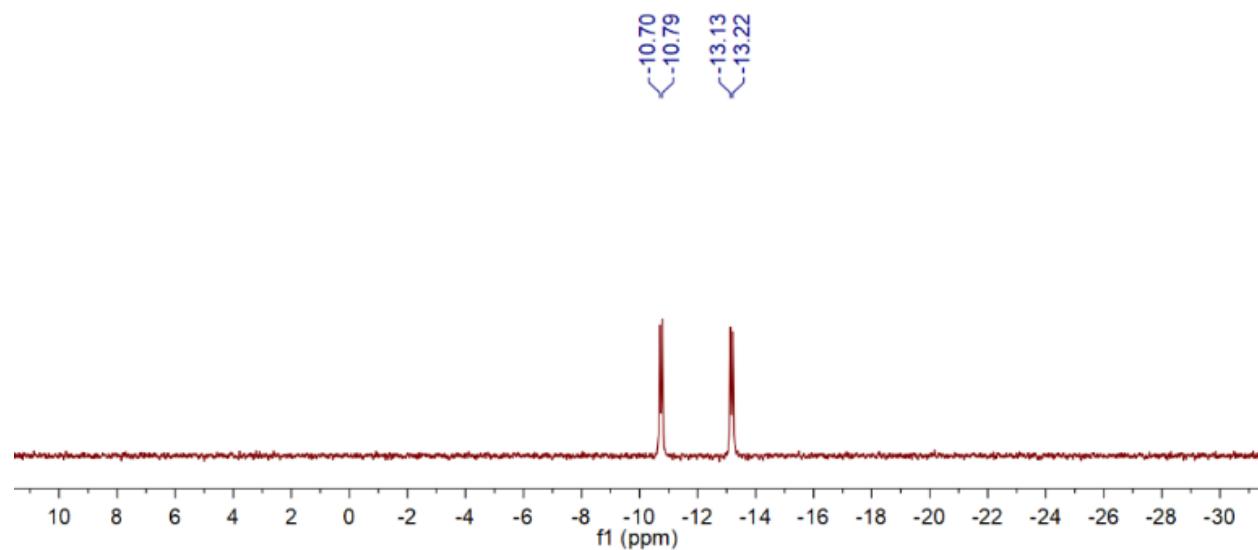


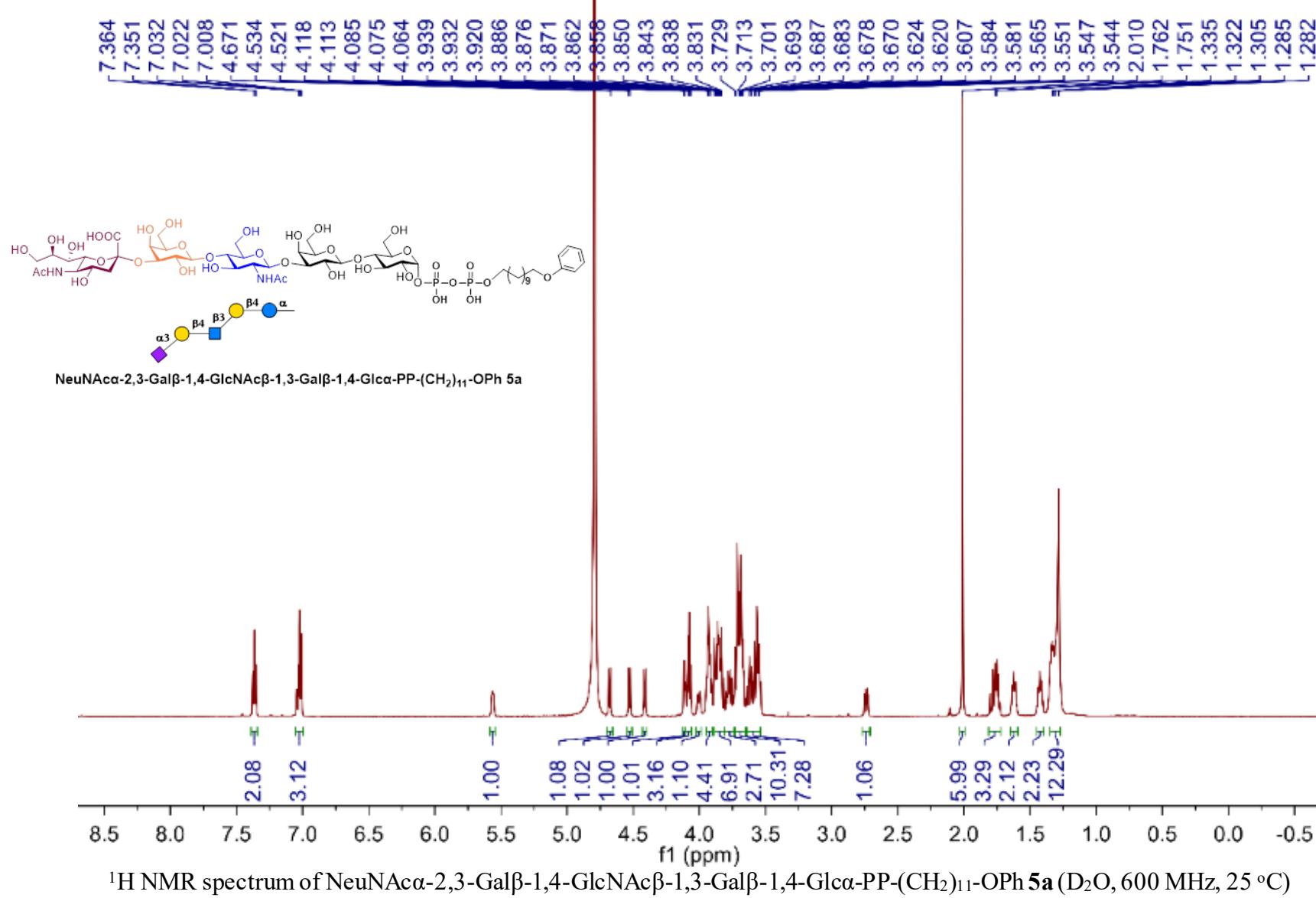
COSY spectrum of Gal β -1,3-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **4e** (D₂O, 600/600 MHz, 25 °C)

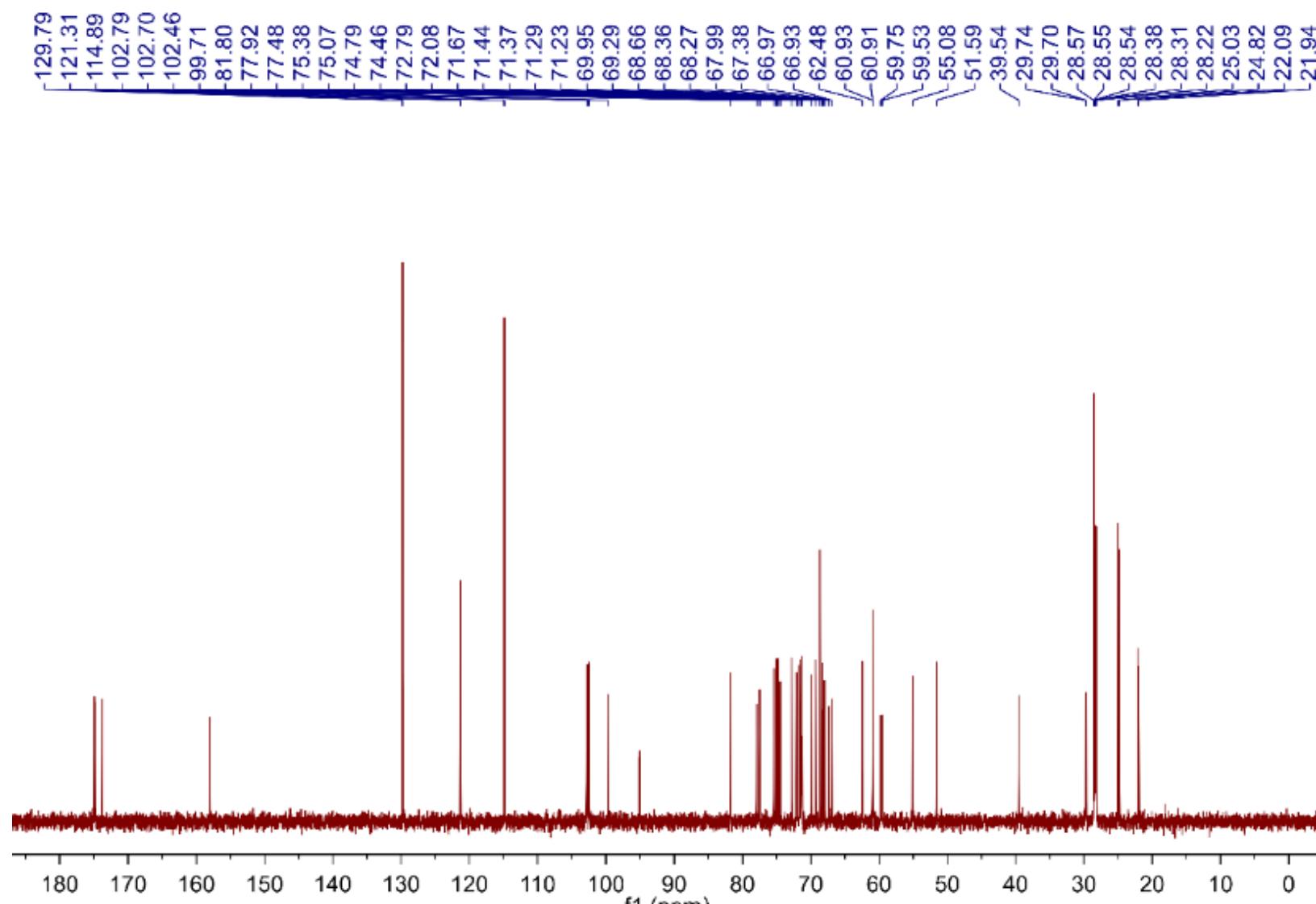


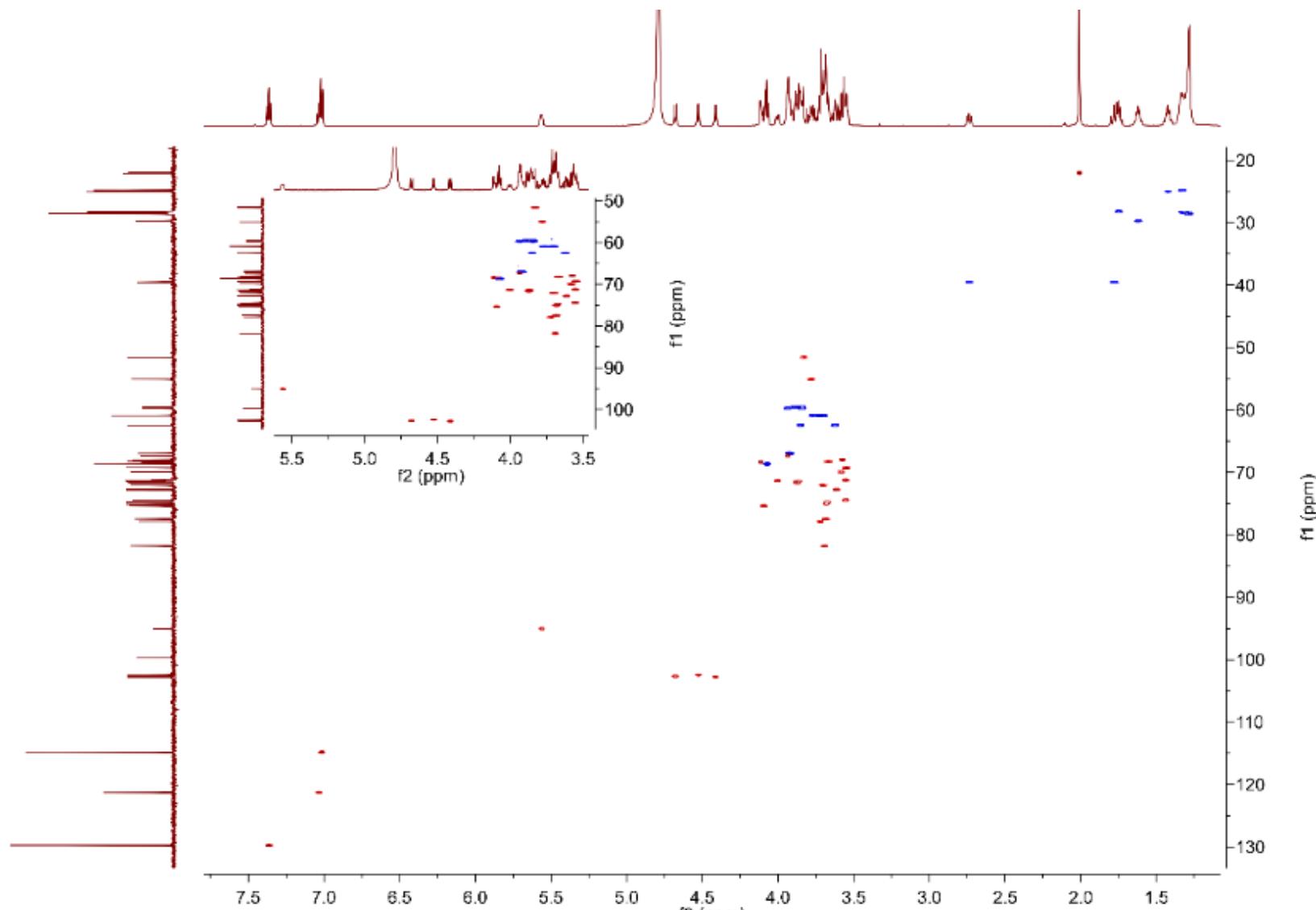


HMBC spectrum of Gal β -1,3-Glc β -1,3-Gal β -1,4-Glca-PP-(CH₂)₁₁-OPh **4e** (D₂O, 600/150 MHz, 25 °C)

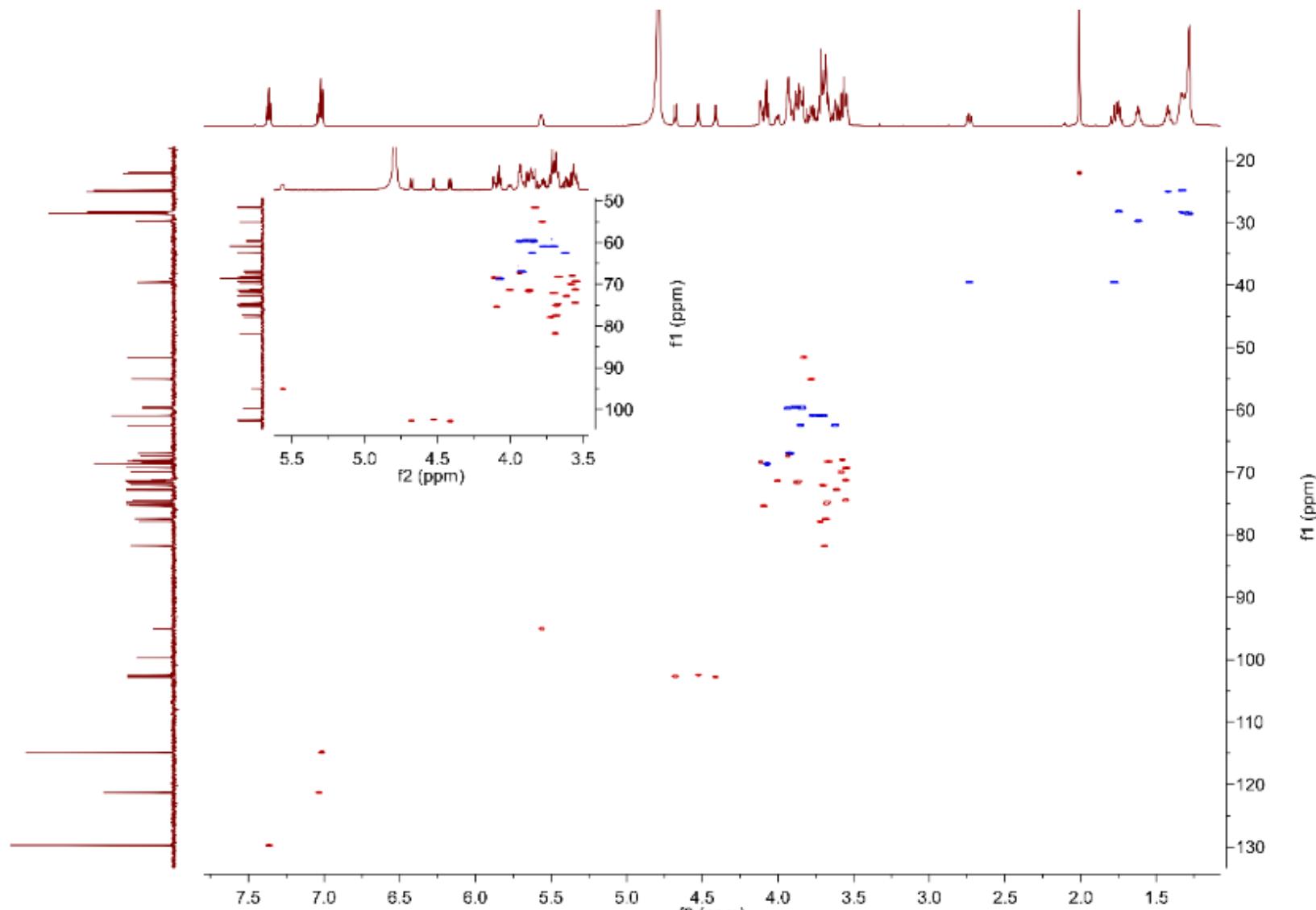


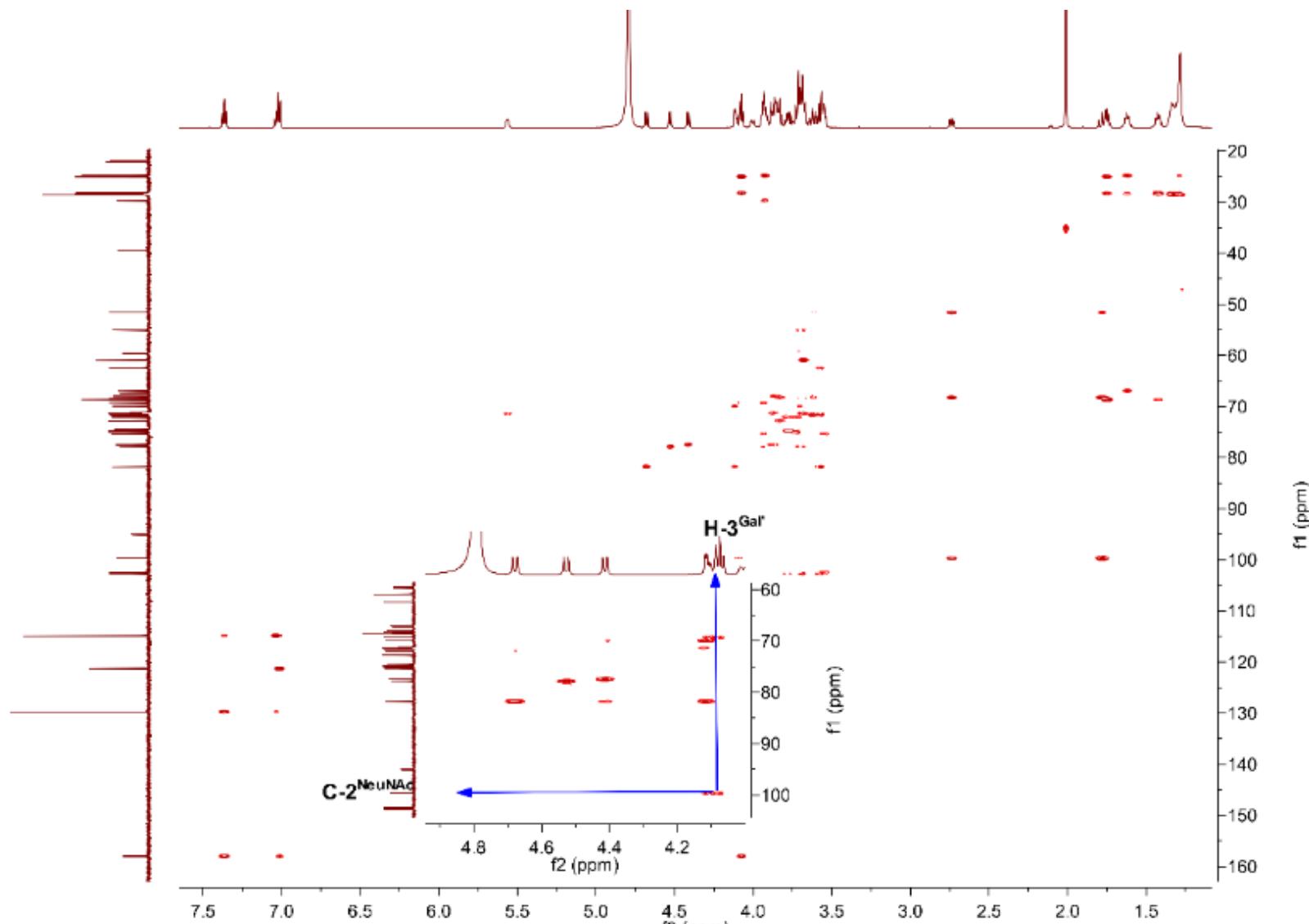




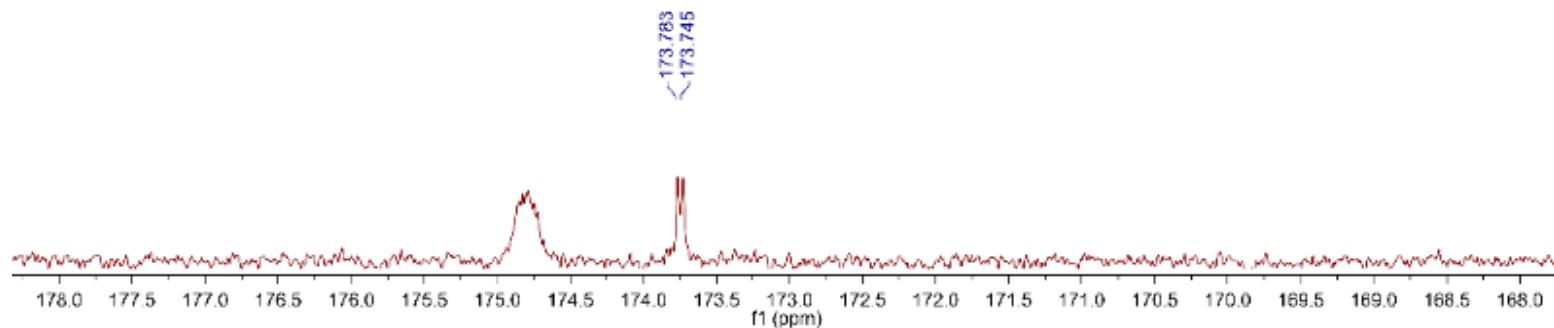


COSY spectrum of NeuNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5a** (D₂O, 600/600 MHz, 25 °C)

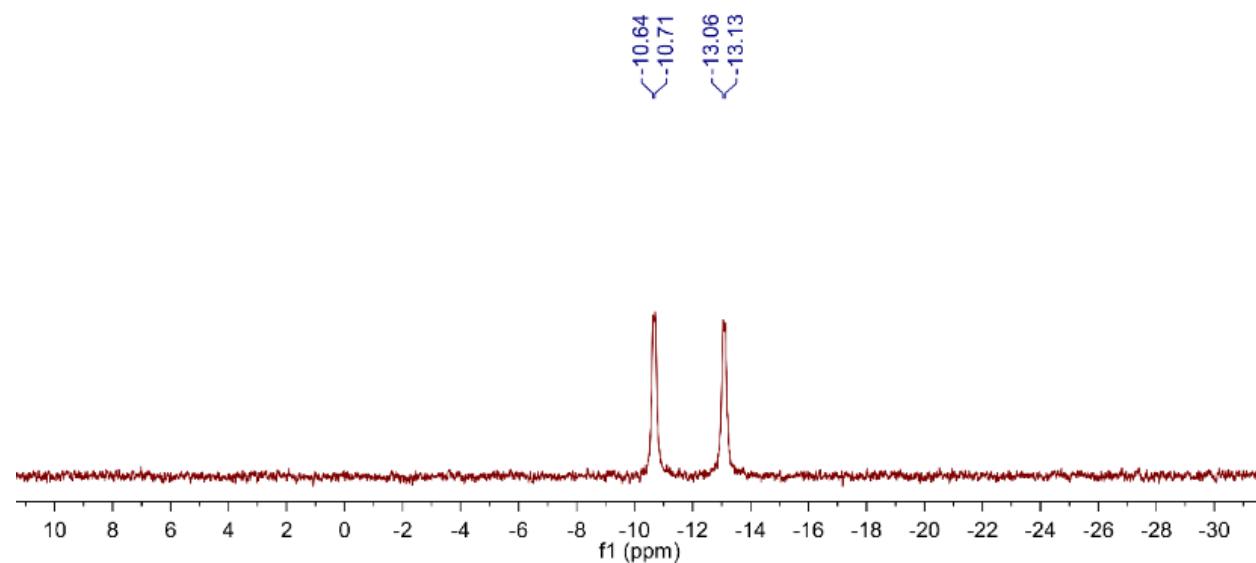




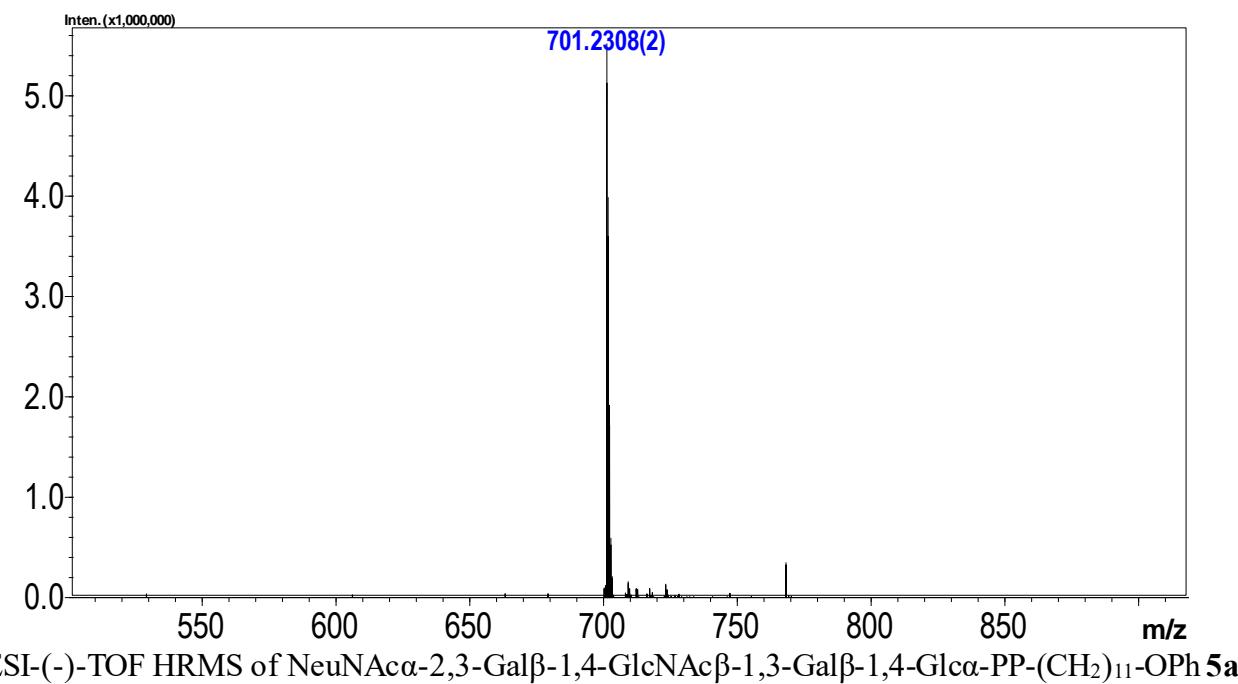
HMBC spectrum of NeuNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5a** (D₂O, 600/150 MHz, 25 °C)

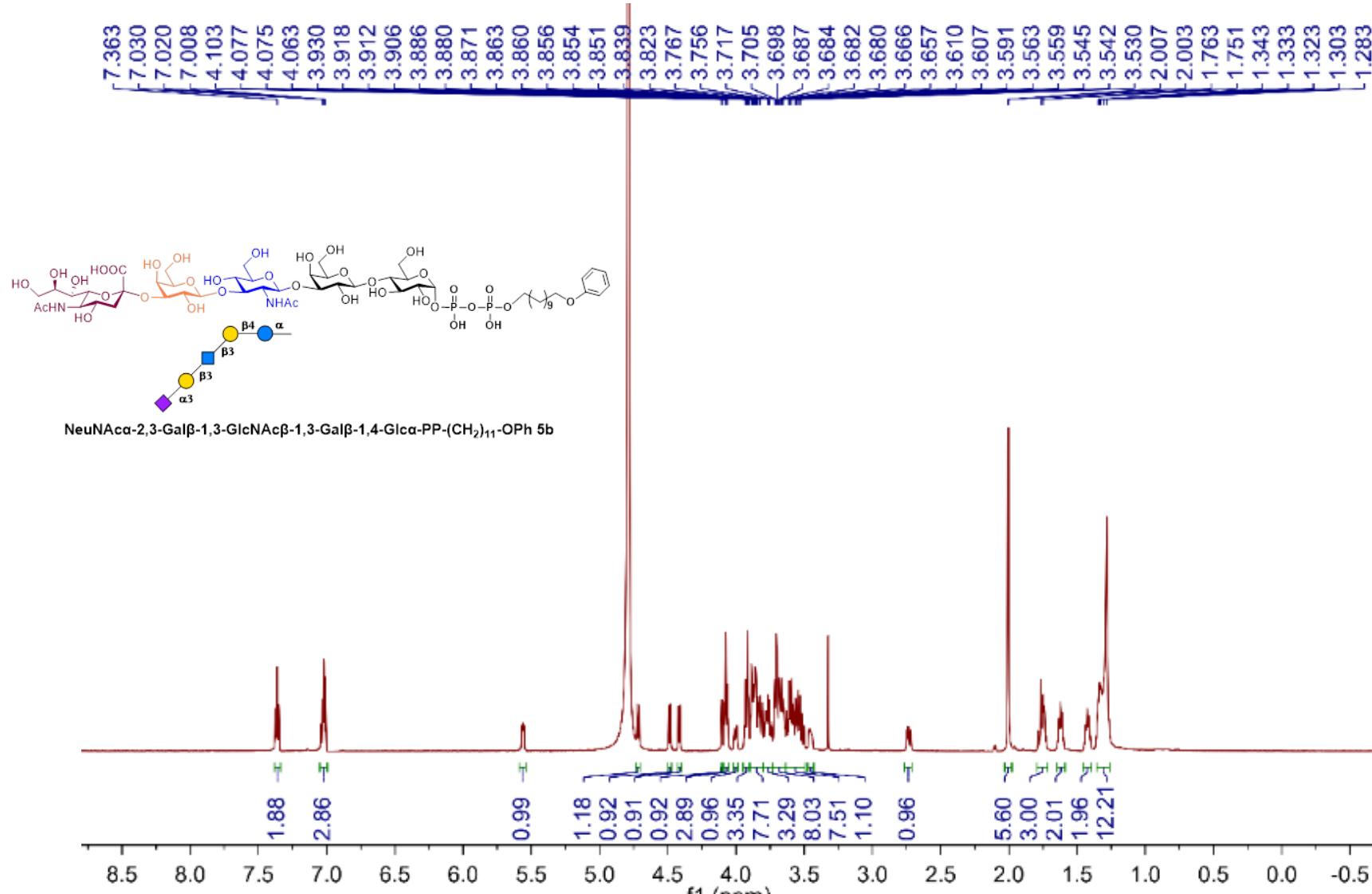


Selected non-decoupling ^{13}C NMR spectrum of NeuNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH $_2$) $_{11}$ -OPh **5a** (D $_2$ O, 150 MHz, 25 °C)

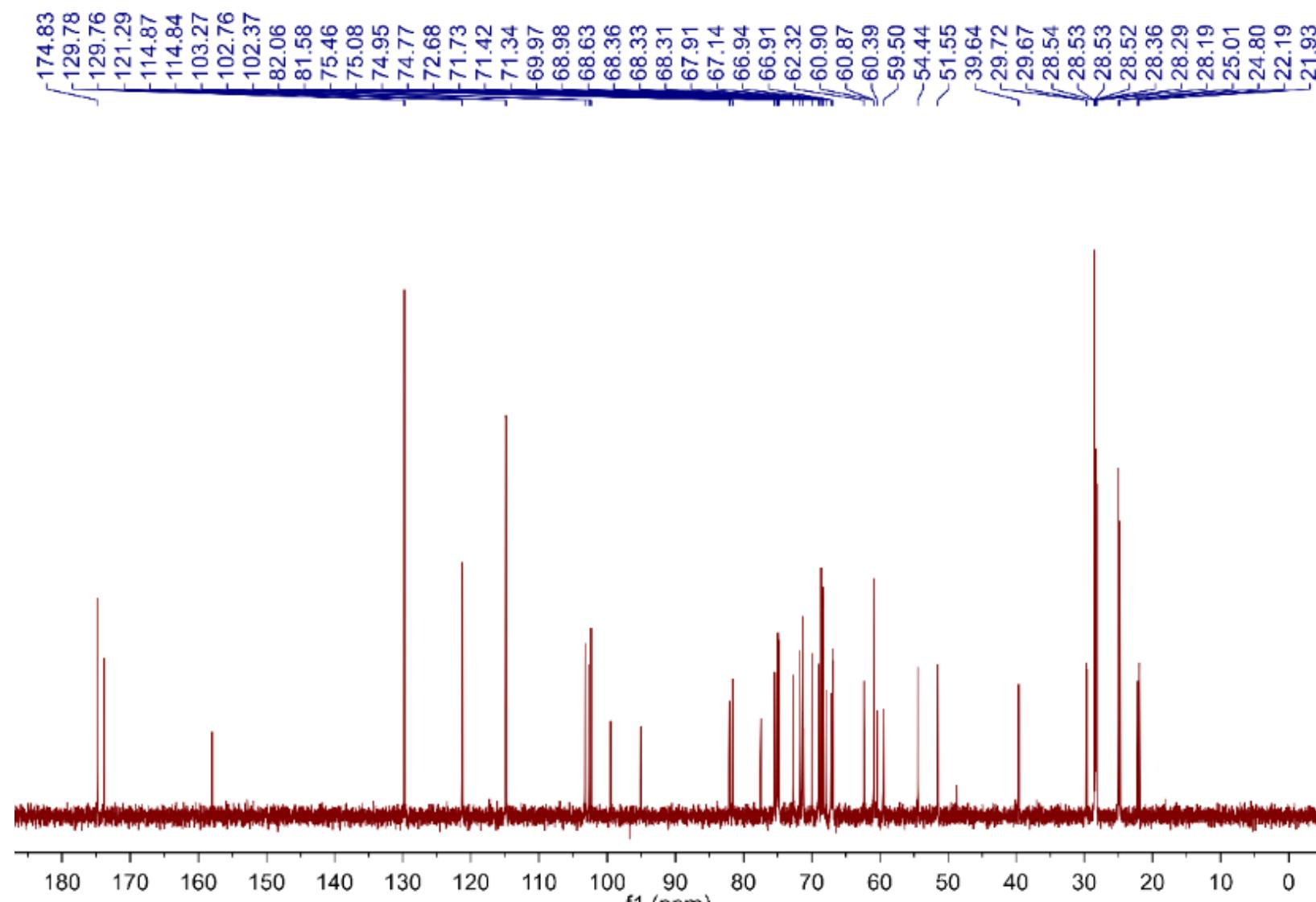


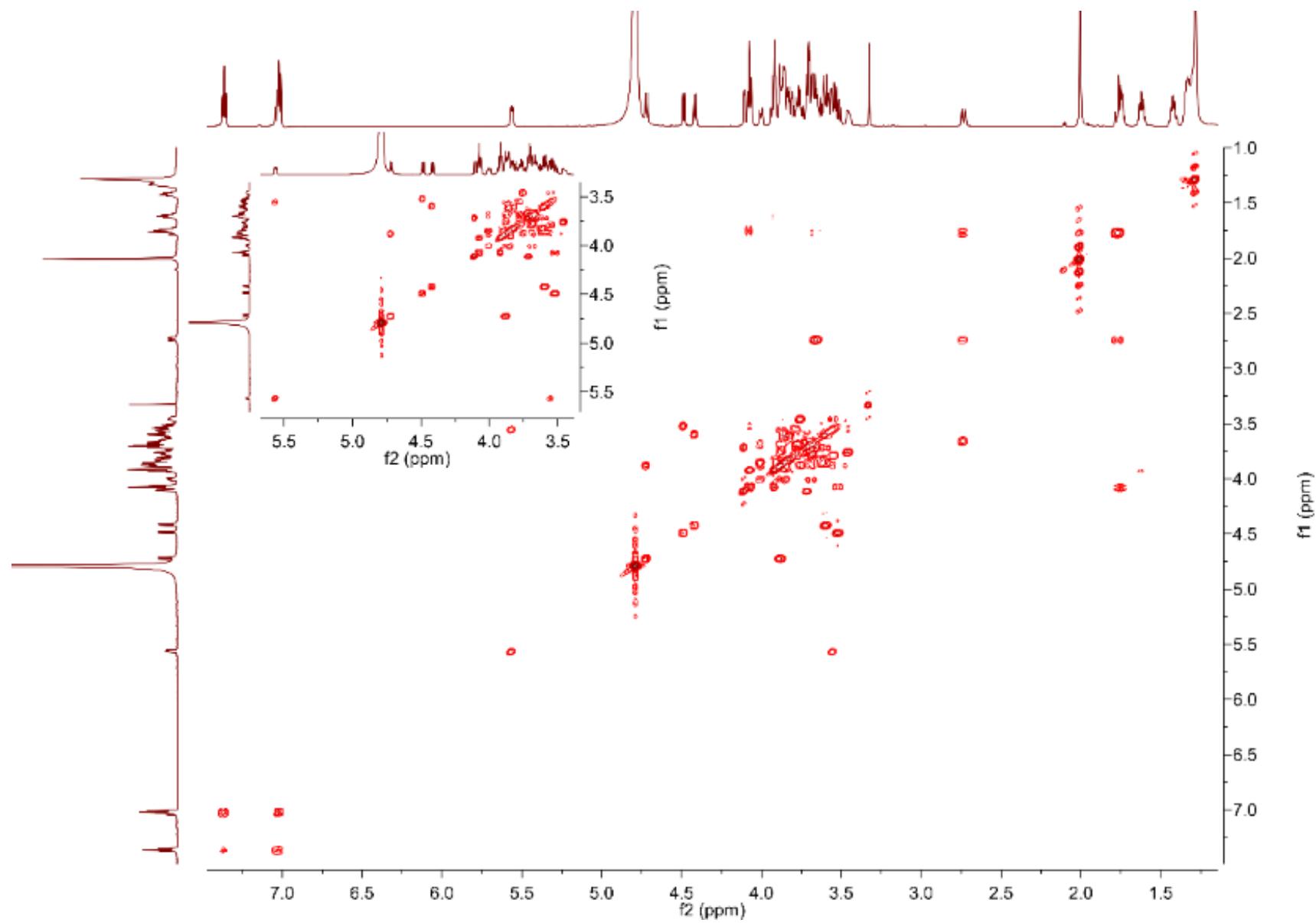
^{31}P NMR spectrum of NeuNAc α -2,3-Gal β -1,4-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH $_2$) $_{11}$ -OPh **5a** (D $_2$ O, 243 MHz, 25 °C)



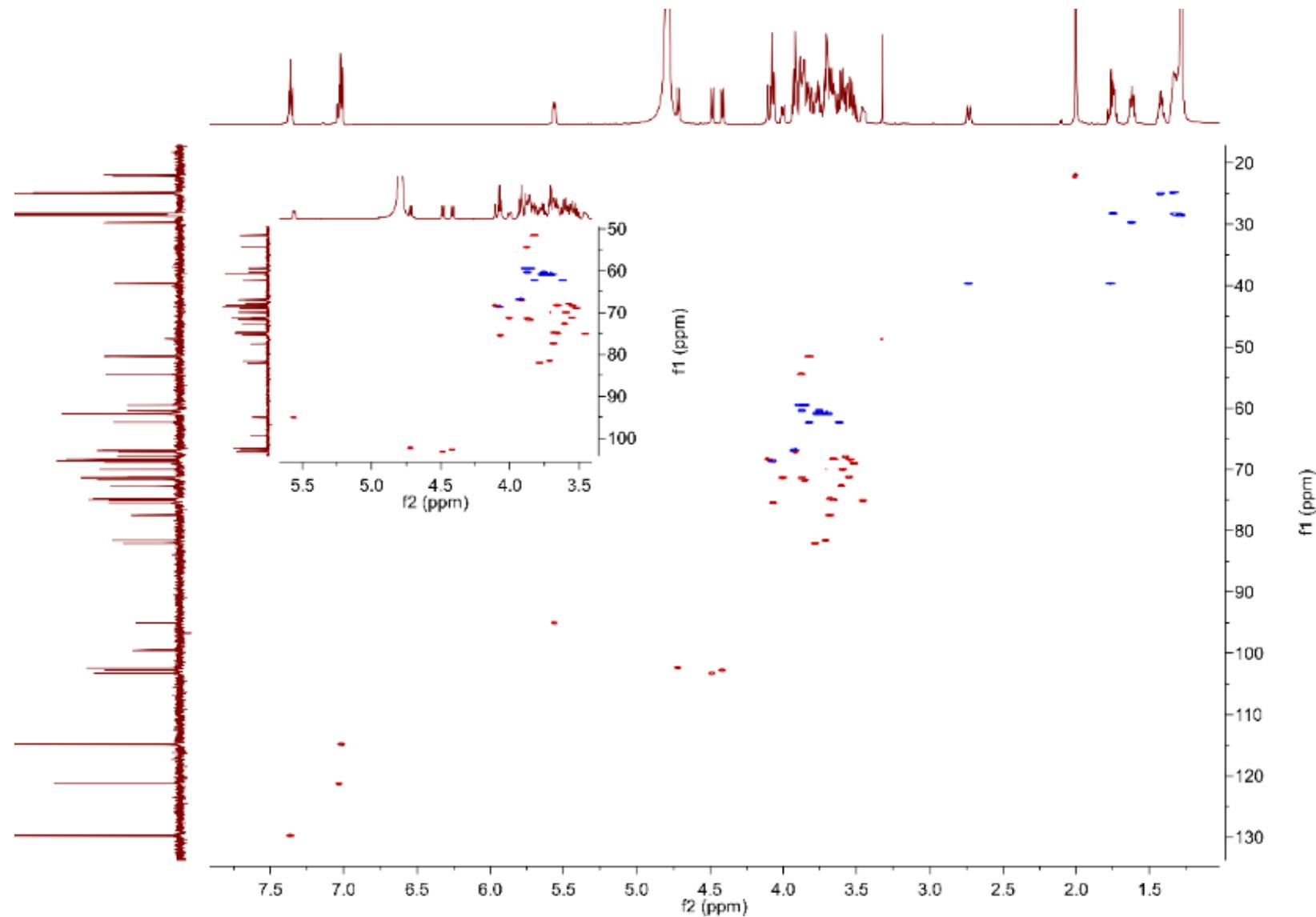


¹H NMR spectrum of NeuNAc-2,3-Gal-1,3-GlcNAc-1,3-Gal-1,4-Glc-PP-(CH₂)₁₁-OPh **5b** (D₂O, 600 MHz, 25 °C)

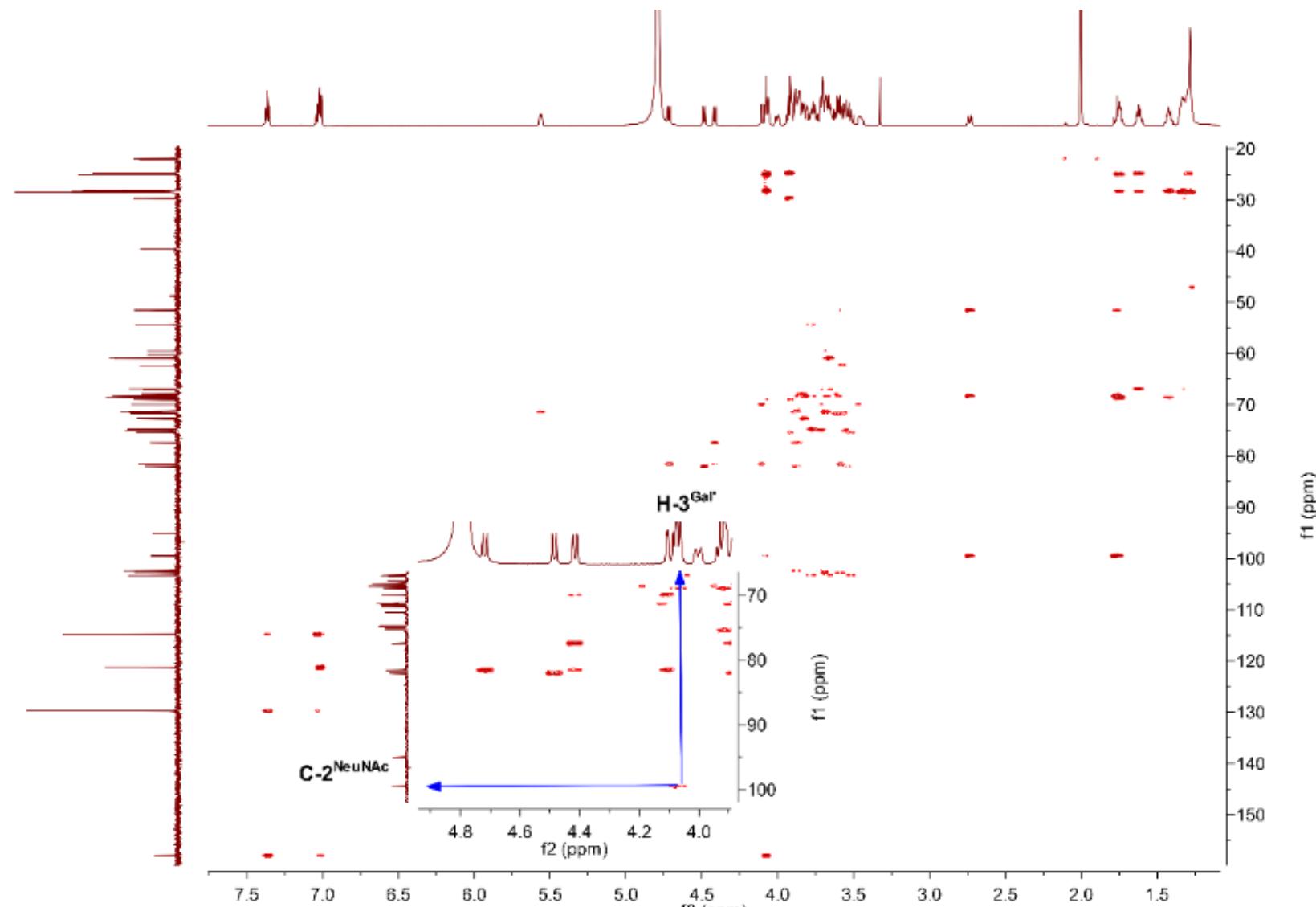




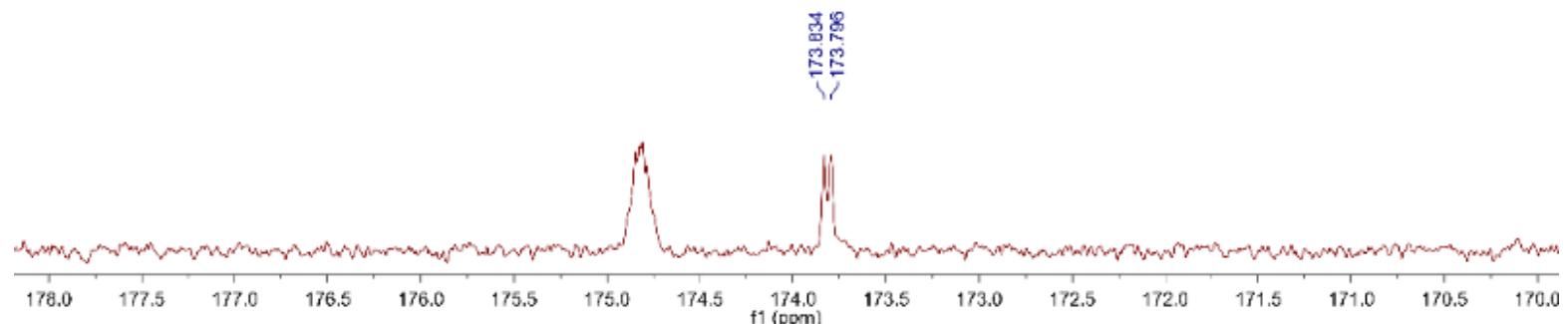
COSY spectrum of NeuNAc α -2,3-Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5b** (D₂O, 600/600 MHz, 25 °C)



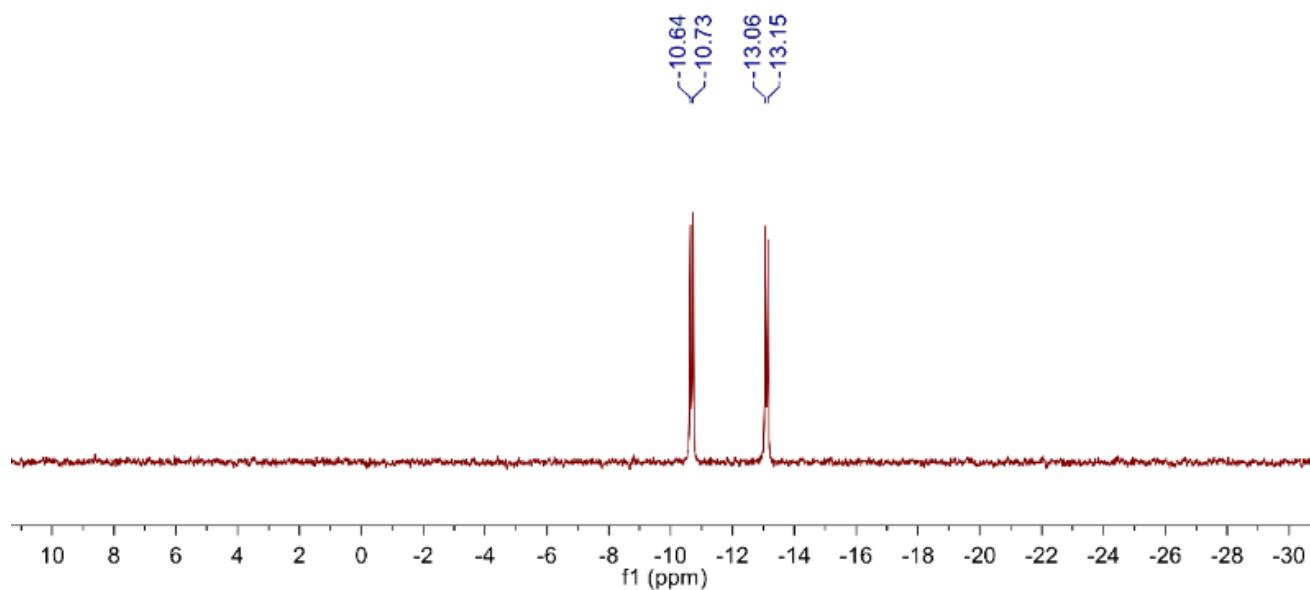
HSQC spectrum of NeuNAc-2,3-Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5b** (D₂O, 600/150 MHz, 25 °C)



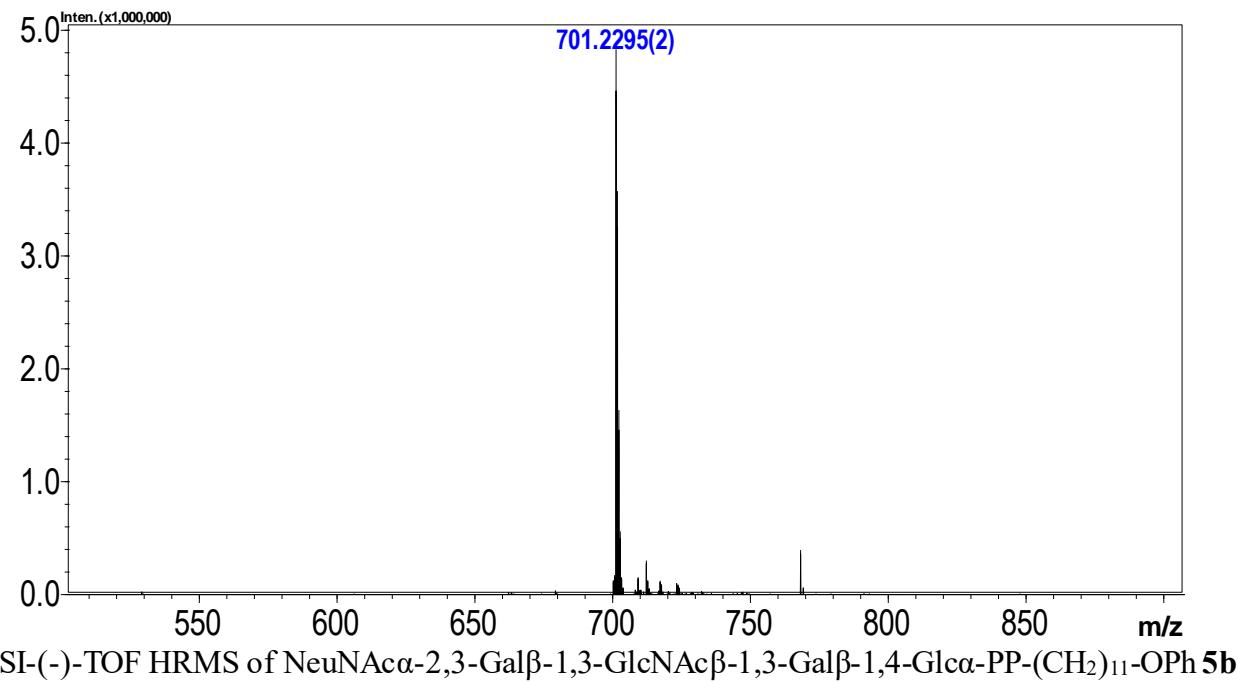
HMBC spectrum of NeuNAc-2,3-Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5b** (D₂O, 600/150 MHz, 25 °C)

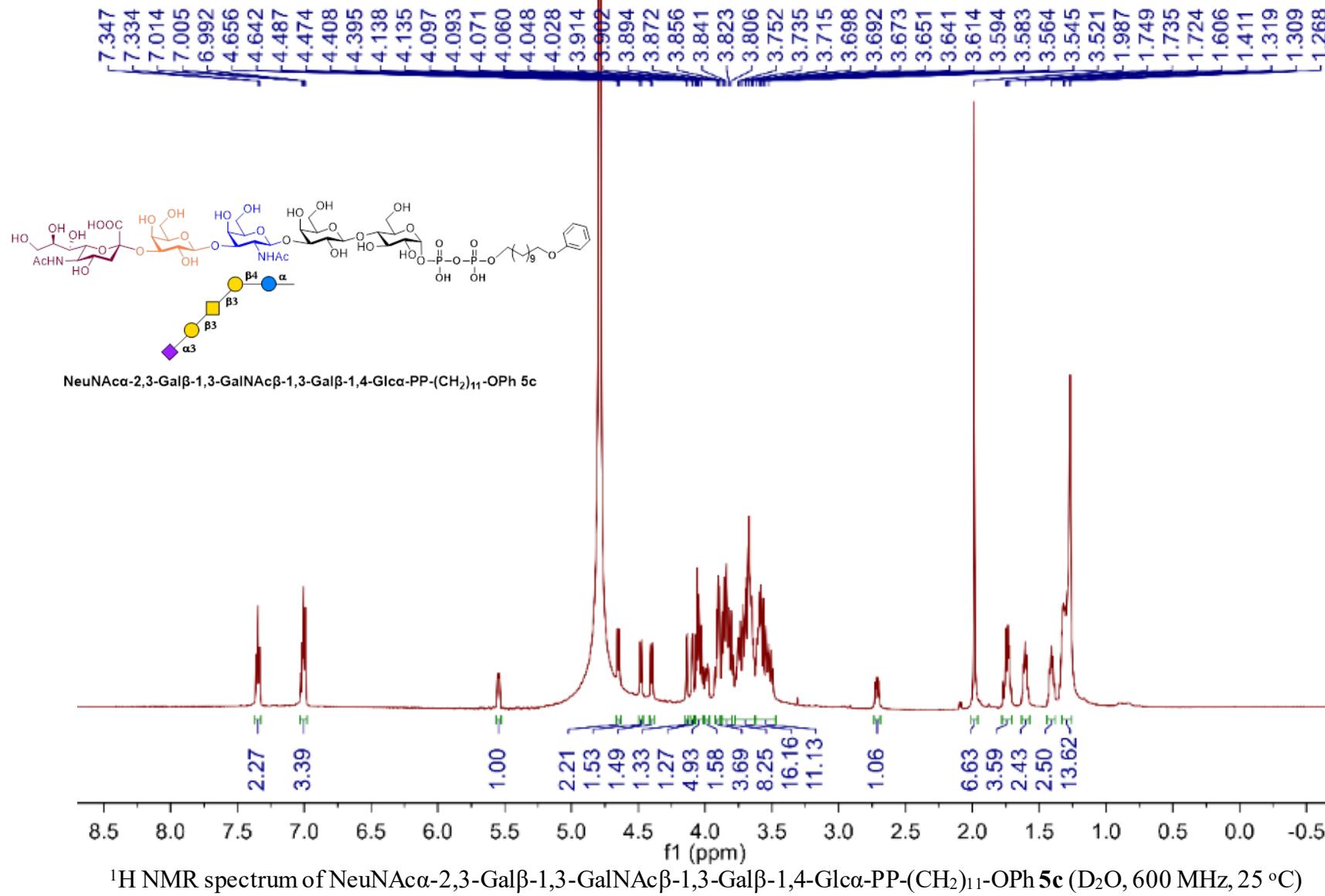


Selected non-decoupling ^{13}C NMR spectrum of NeuNAc α -2,3-Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH $_2$) $_{11}$ -OPh **5b** (D $_2$ O, 150 MHz, 25 °C)

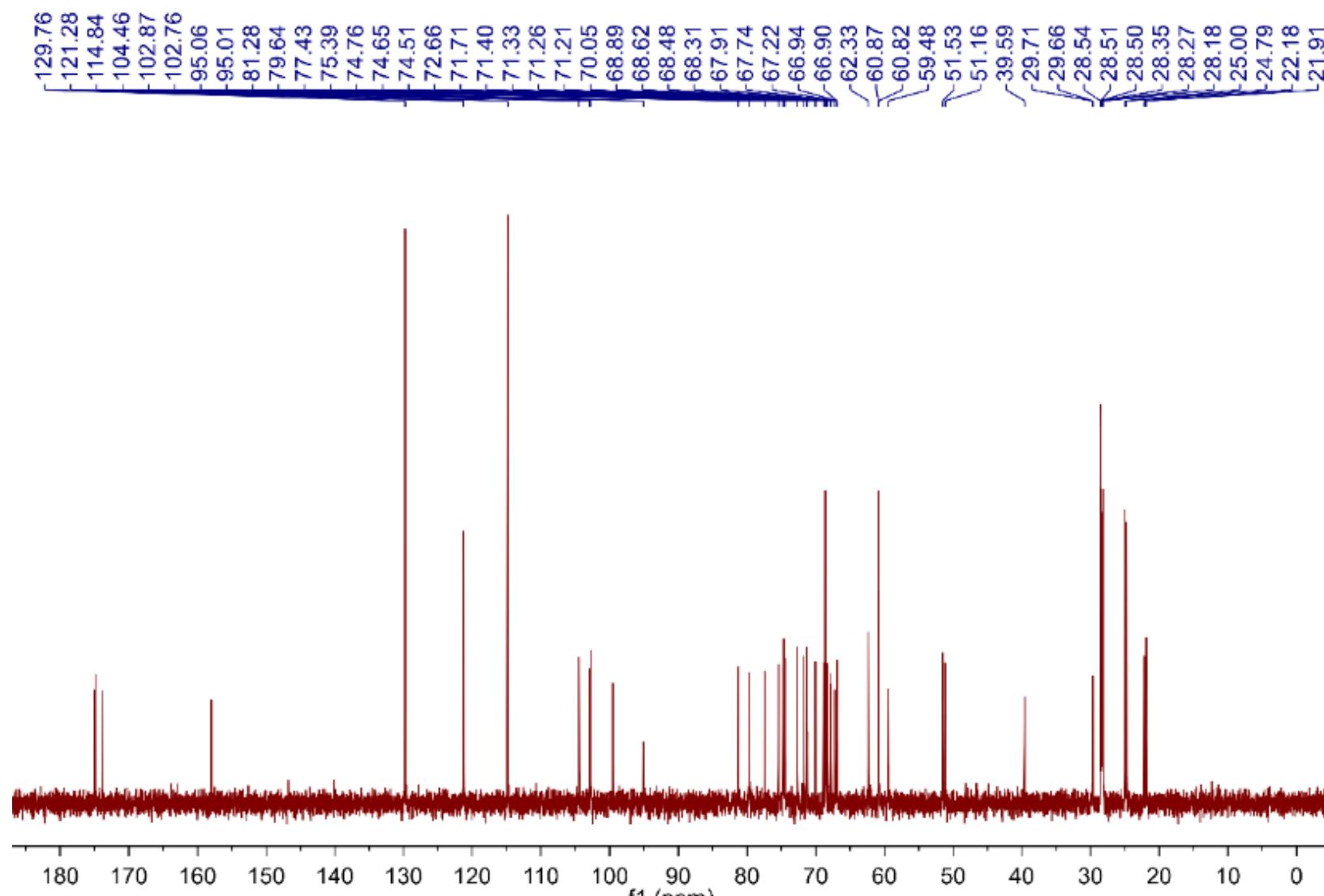


^{31}P NMR spectrum of NeuNAc α -2,3-Gal β -1,3-GlcNAc β -1,3-Gal β -1,4-Glc α -PP-(CH $_2$) $_{11}$ -OPh **5b** (D $_2$ O, 243 MHz, 25 °C)

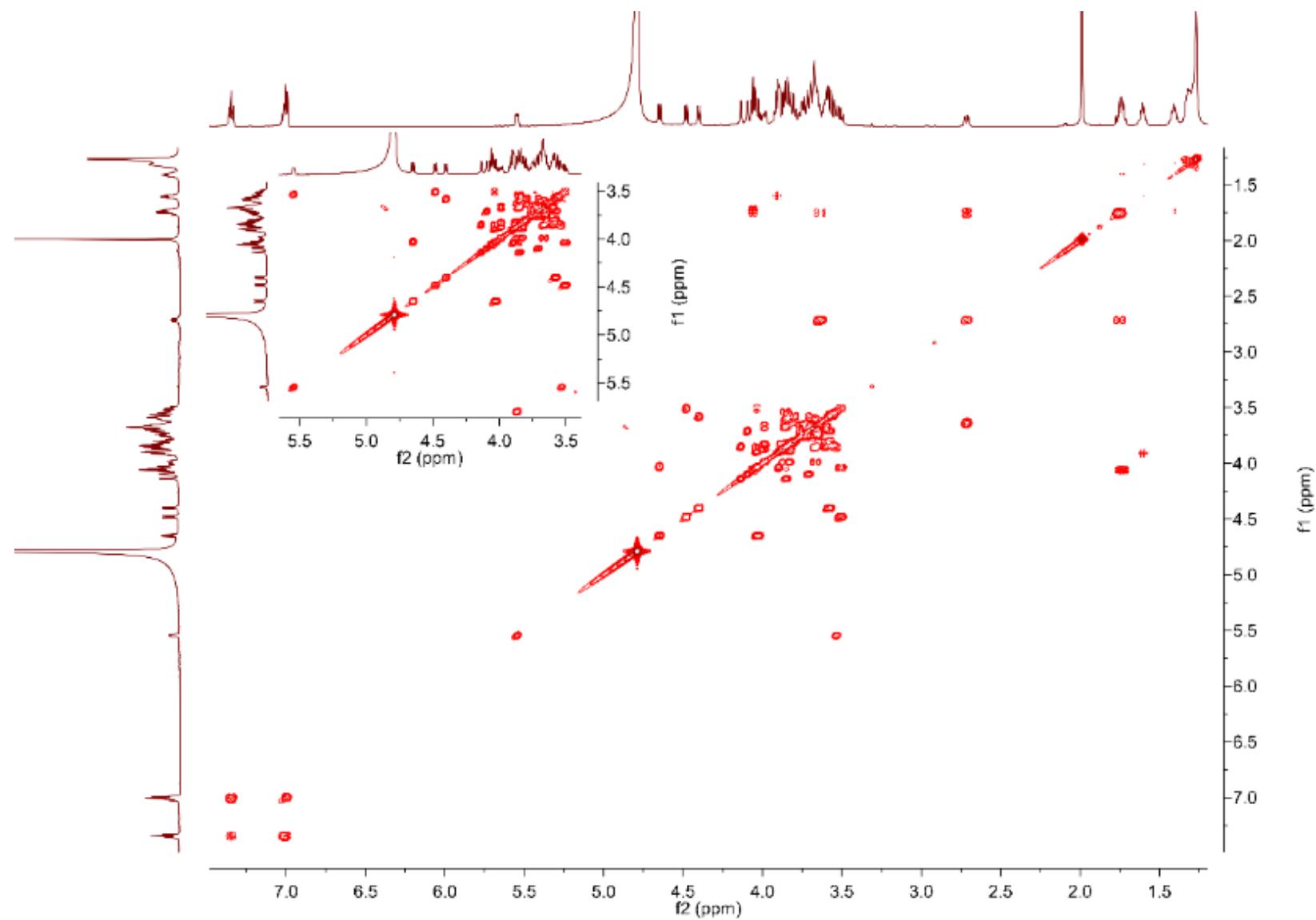




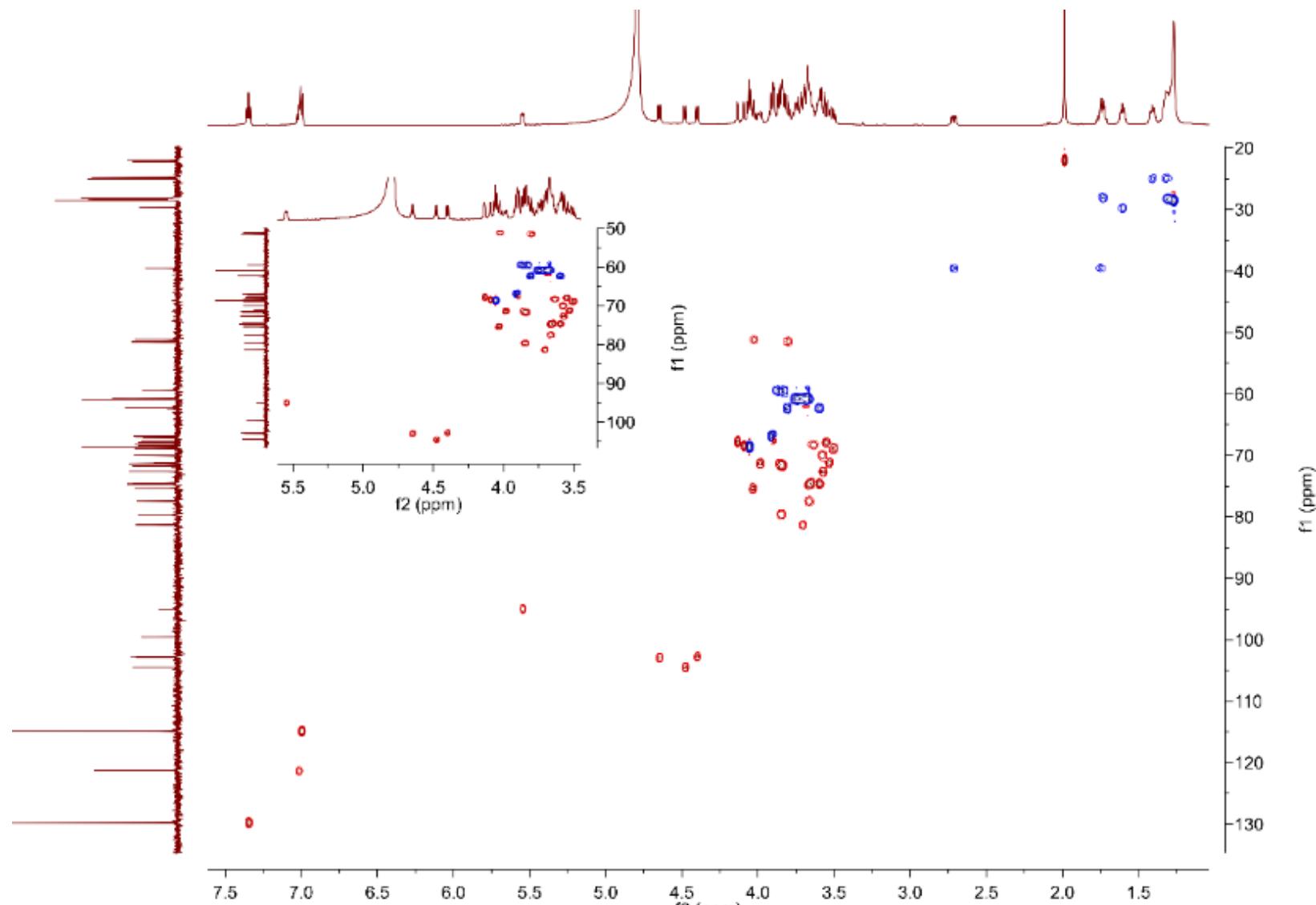
¹H NMR spectrum of NeuNAc α -2,3-Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5c** (D₂O, 600 MHz, 25 °C)

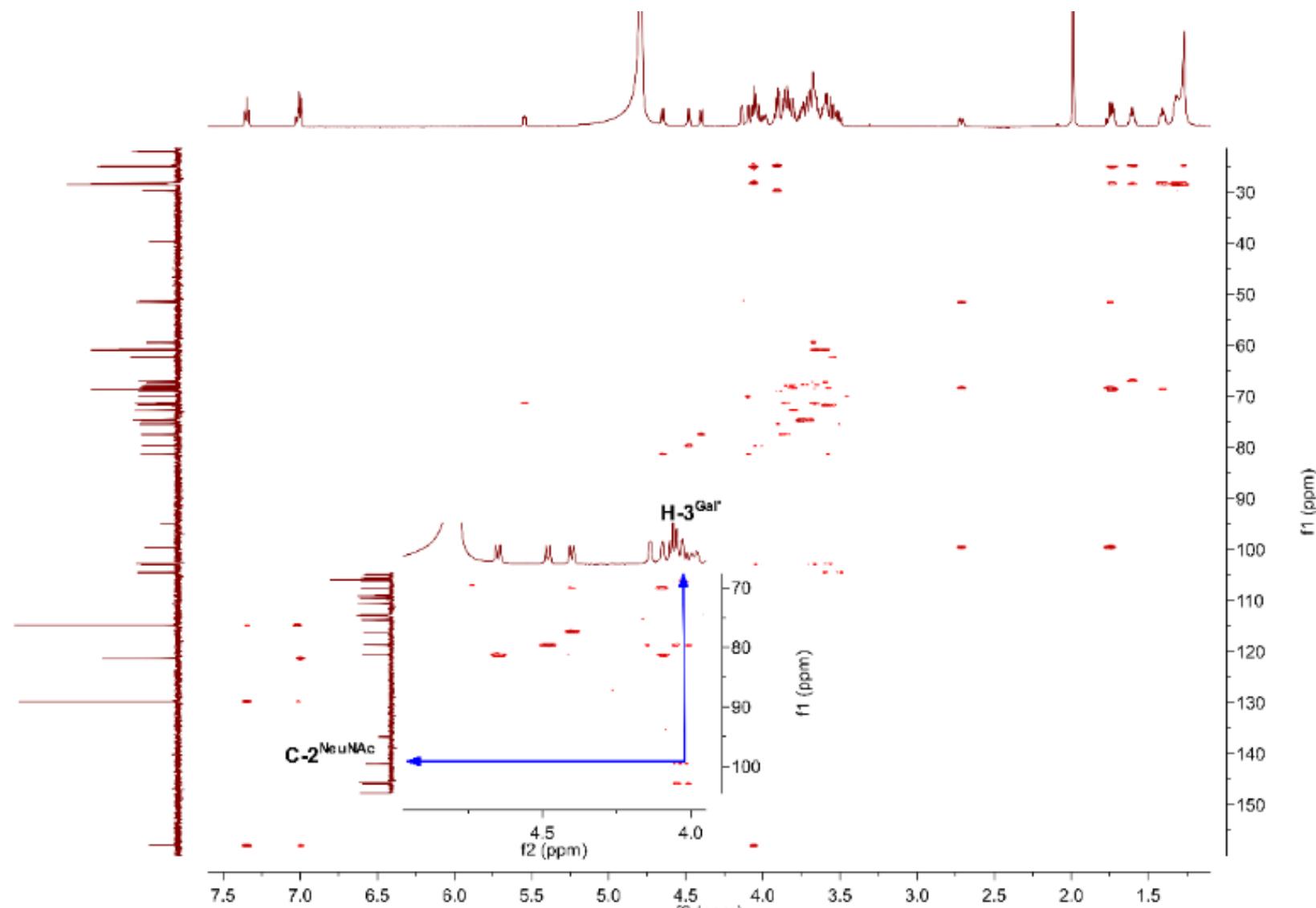


^{13}C NMR spectrum of NeuNAc α -2,3-Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5c** (D₂O, 150 MHz, 25 °C)

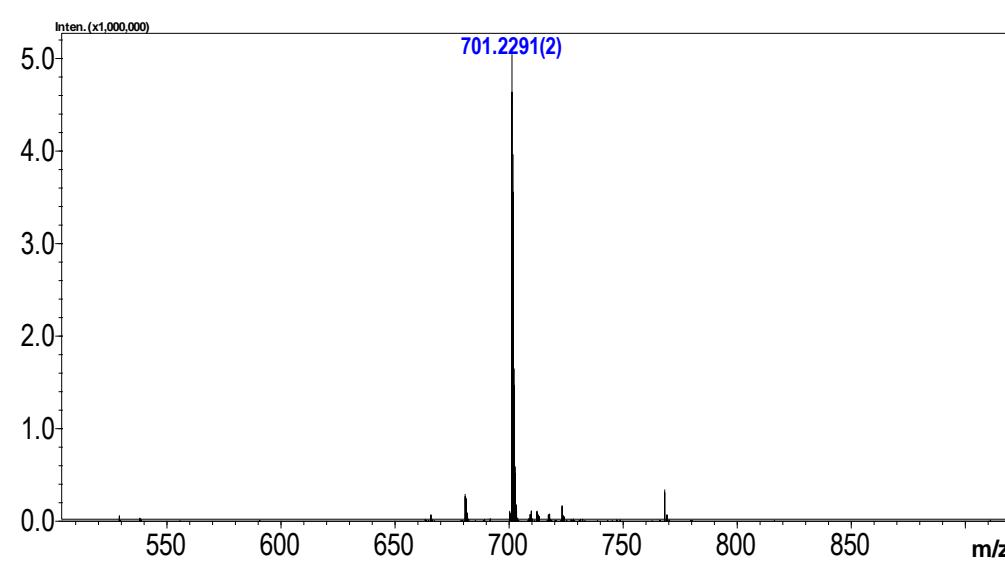
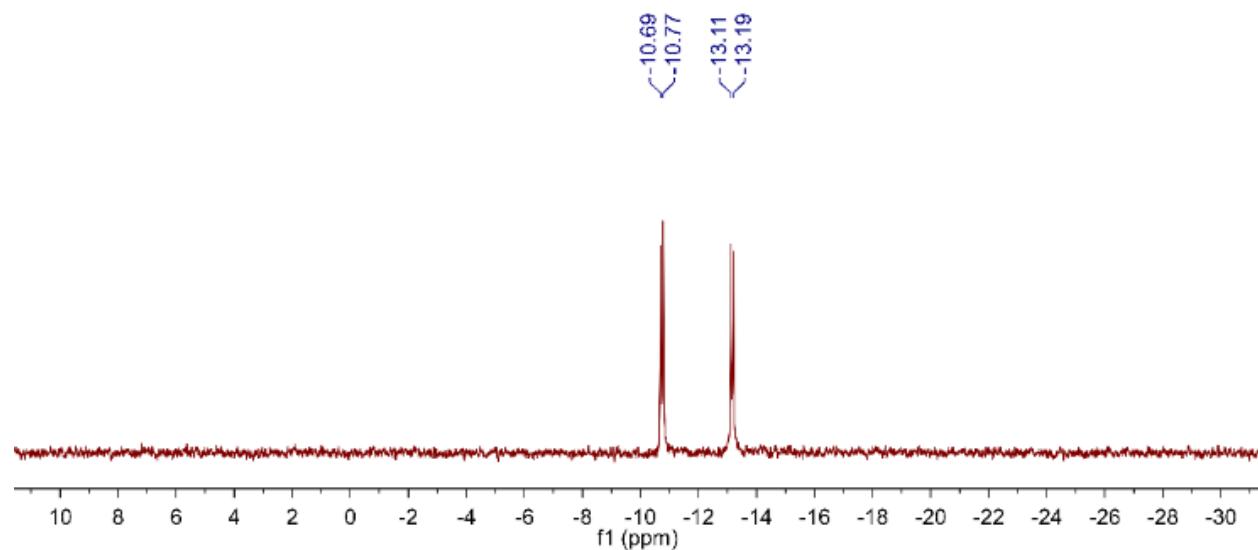


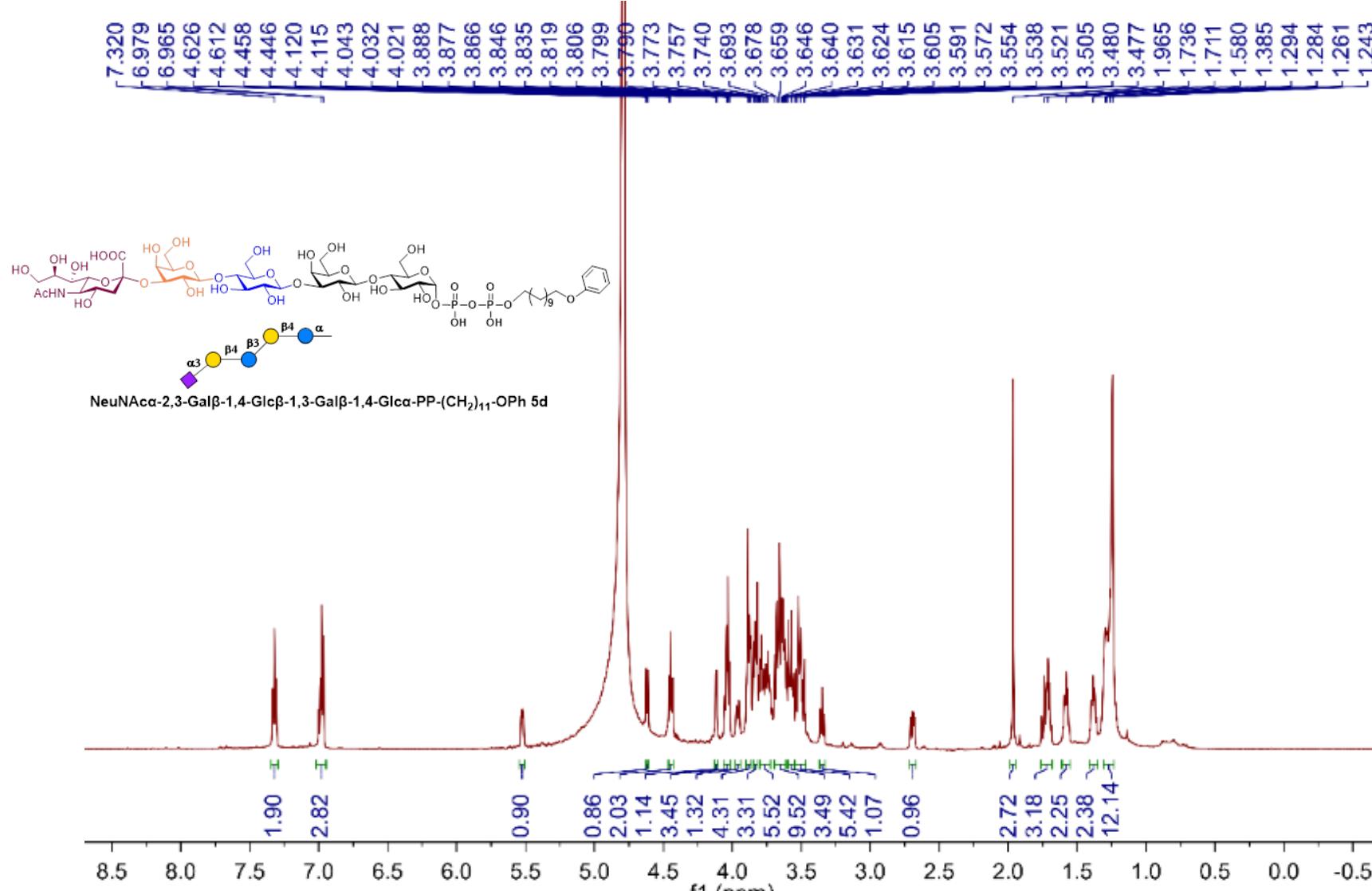
COSY spectrum of NeuNAc α -2,3-Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5c** (D₂O, 600/600 MHz, 25 °C)

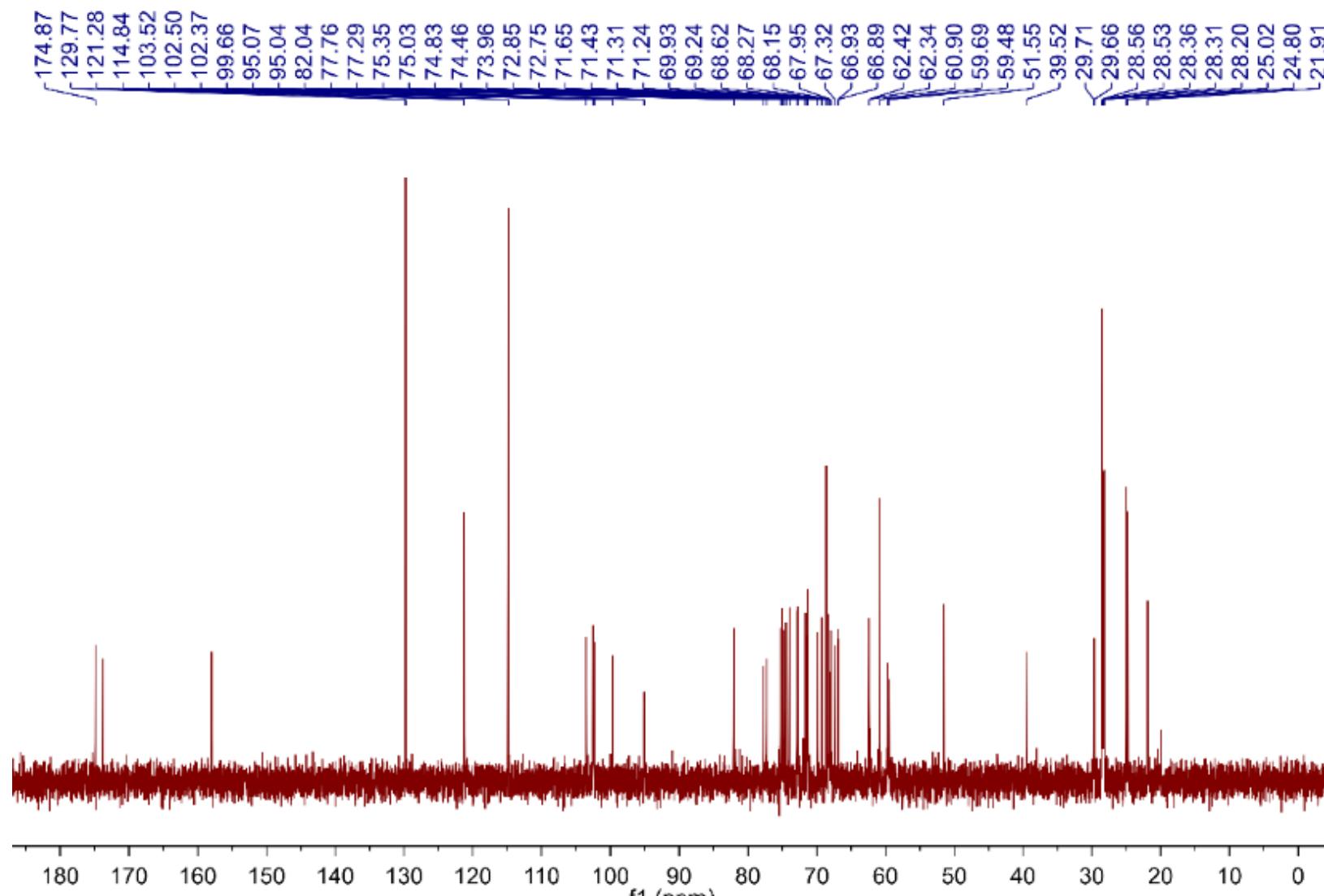




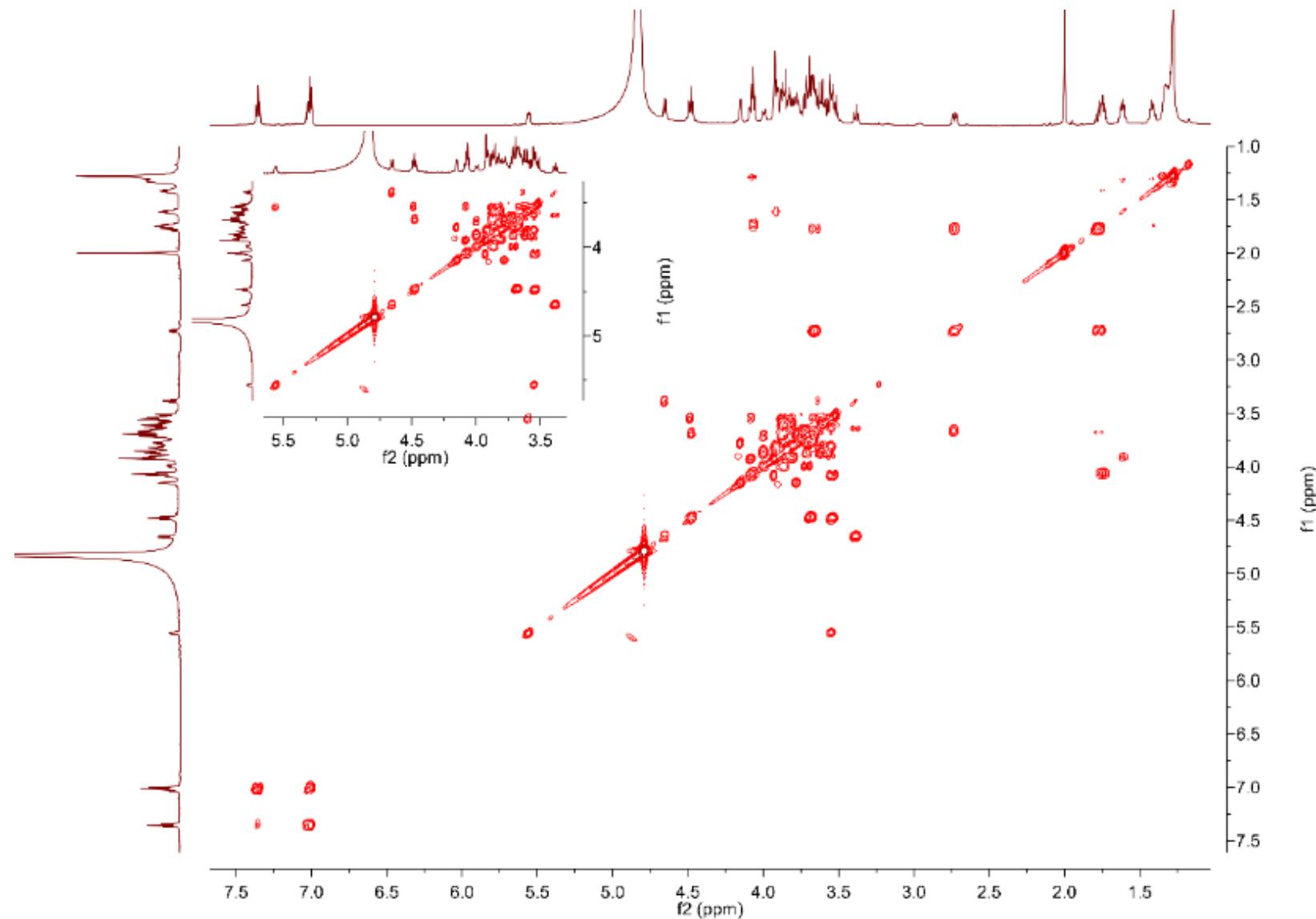
HMBC spectrum of NeuNAc-2,3-Gal β -1,3-GalNAc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5c** (D₂O, 600/150 MHz, 25 °C)



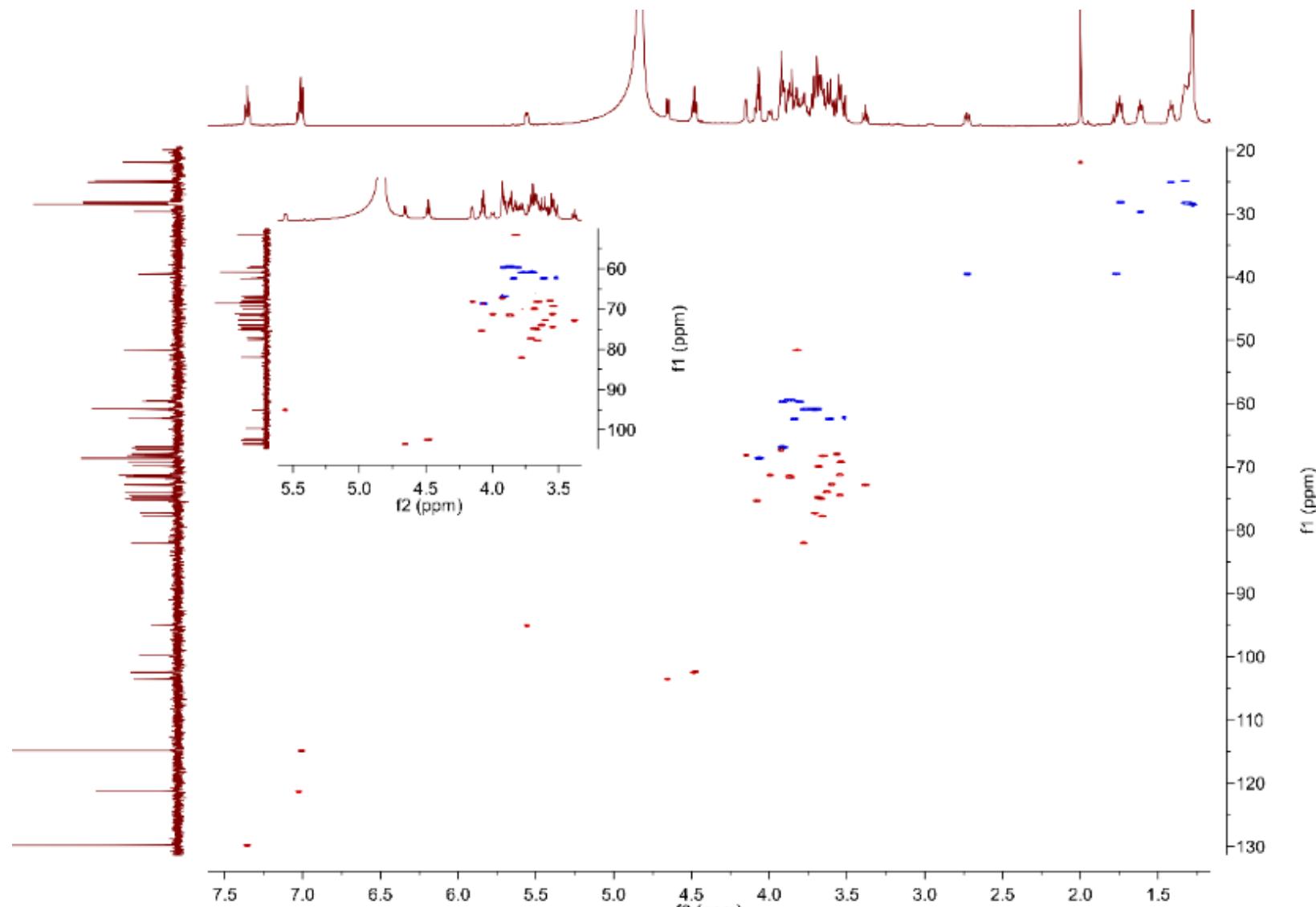




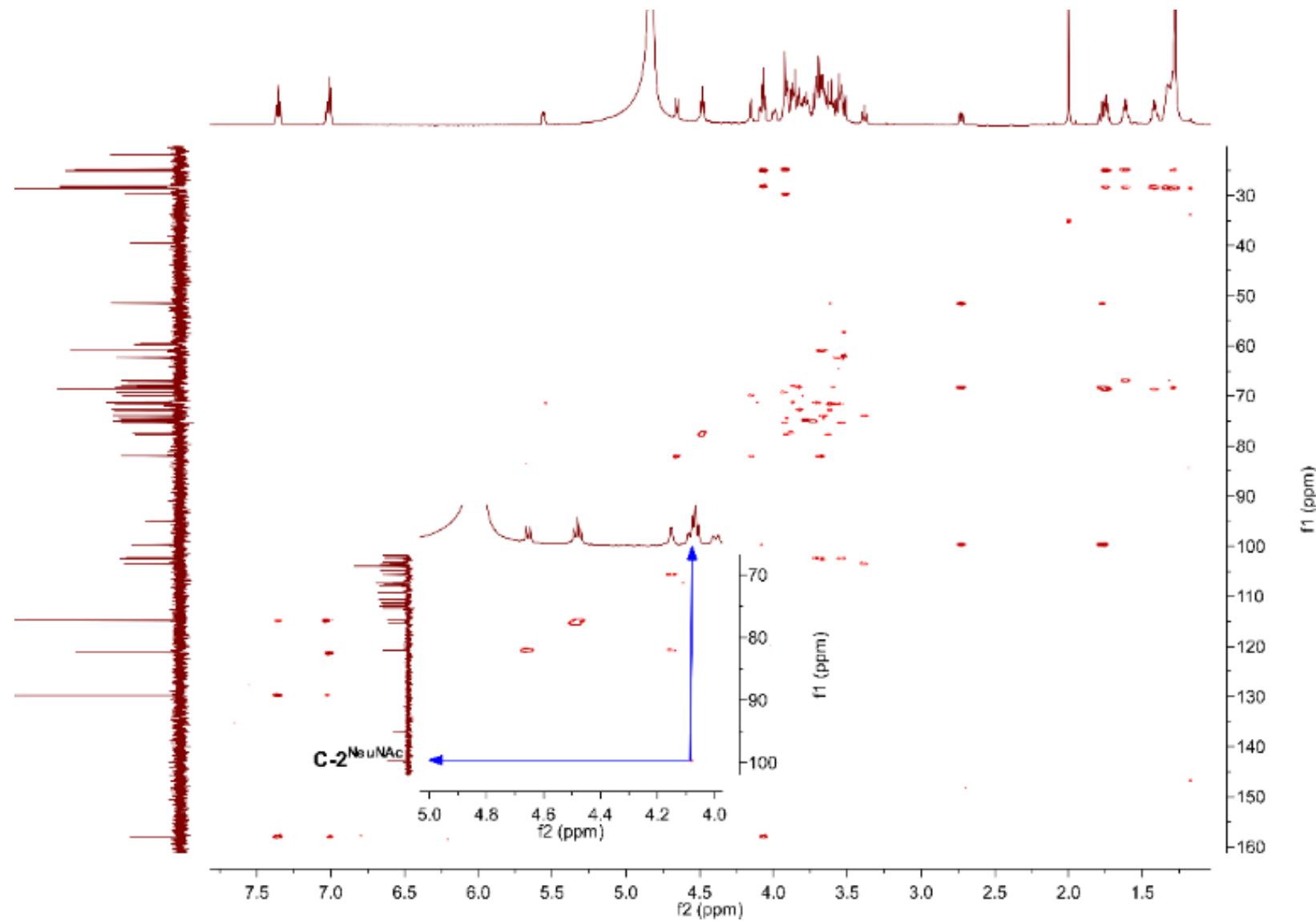
^{13}C NMR spectrum of NeuNAc-2,3-Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5d** (D₂O, 150 MHz, 25 °C)



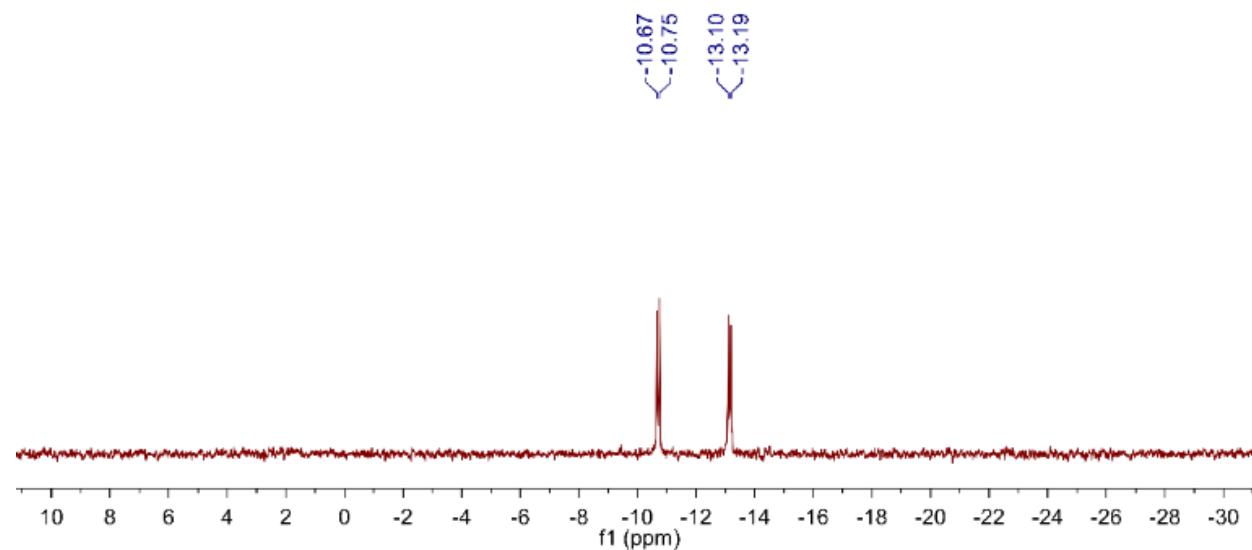
COSY spectrum of NeuNAc α -2,3-Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5d** (D₂O, 600/600 MHz, 25 °C)



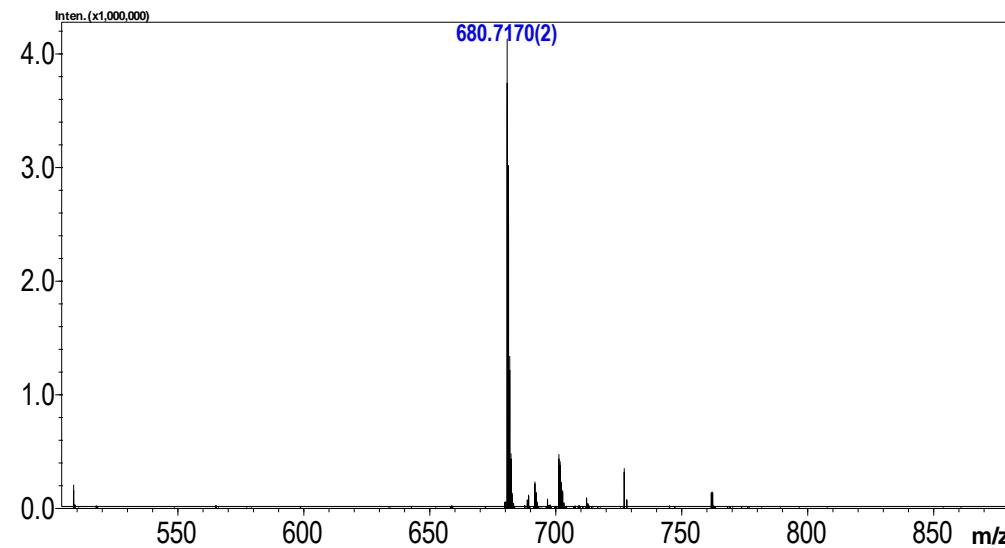
HSQC spectrum of NeuNAc α -2,3-Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5d** (D₂O, 600/150 MHz, 25 °C)



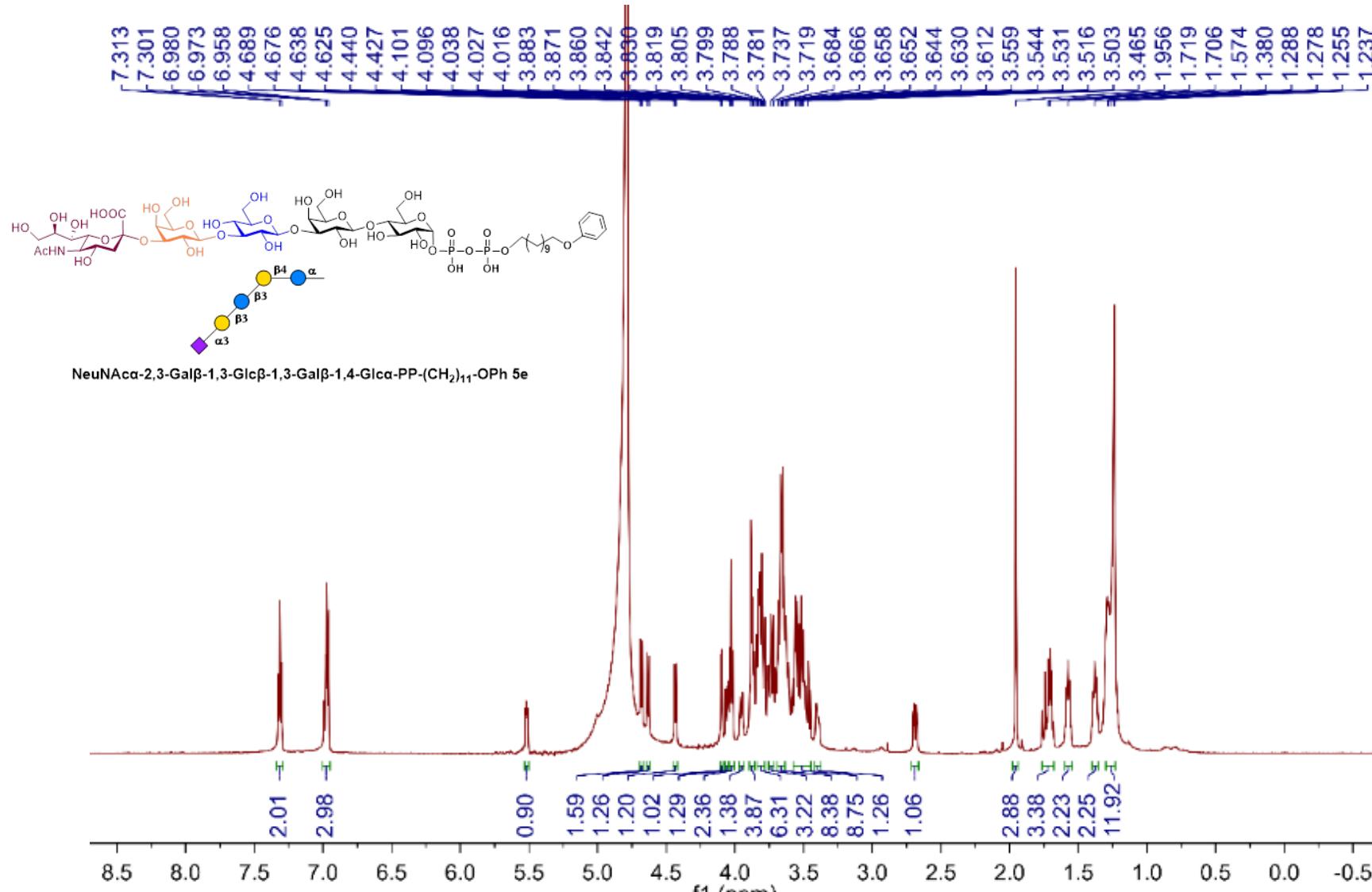
HMBC spectrum of NeuNAc-2,3-Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5d** (D₂O, 600/150 MHz, 25 °C)



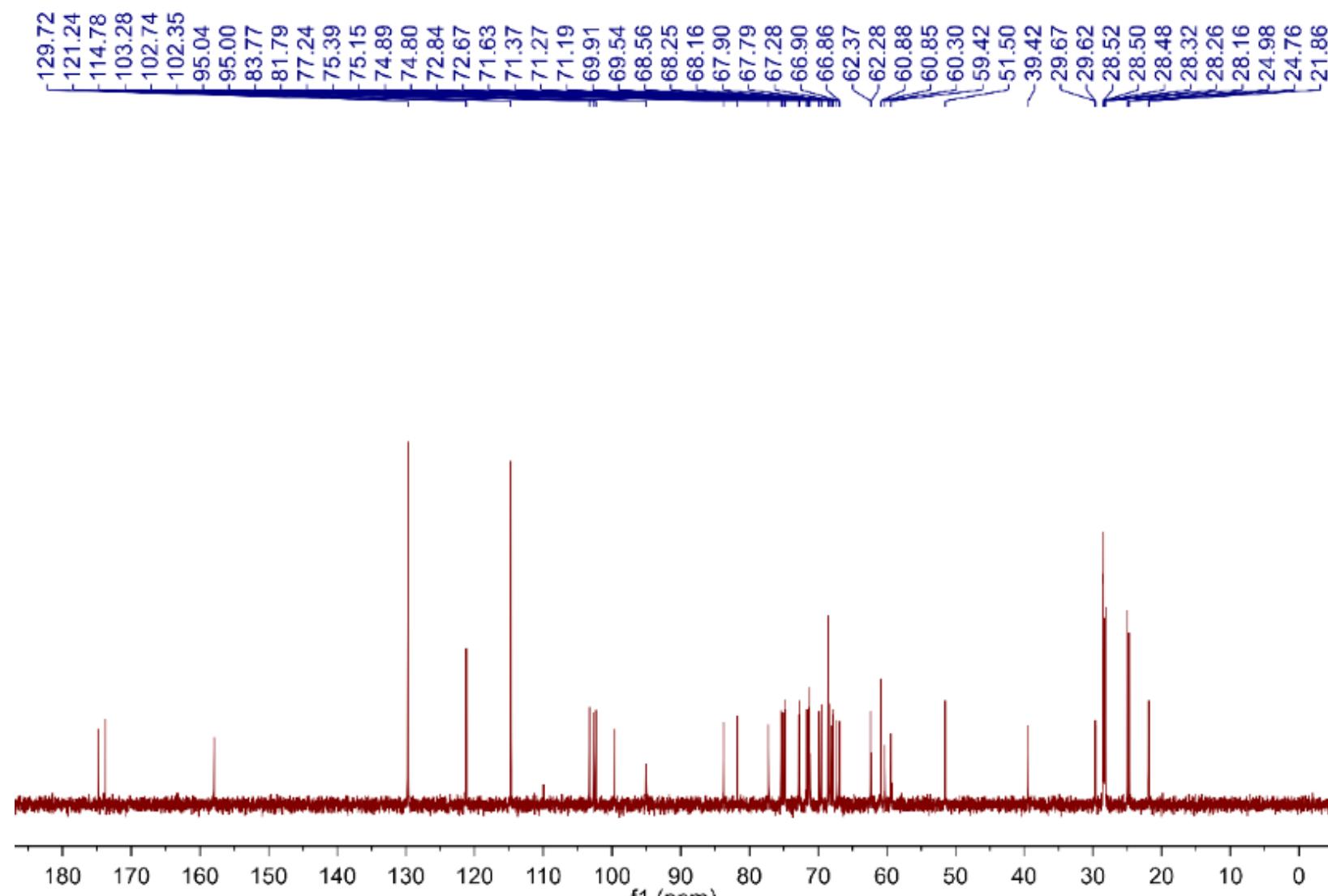
^{31}P NMR spectrum of NeuNAc α -2,3-Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5d** (D₂O, 243 MHz, 25 °C)

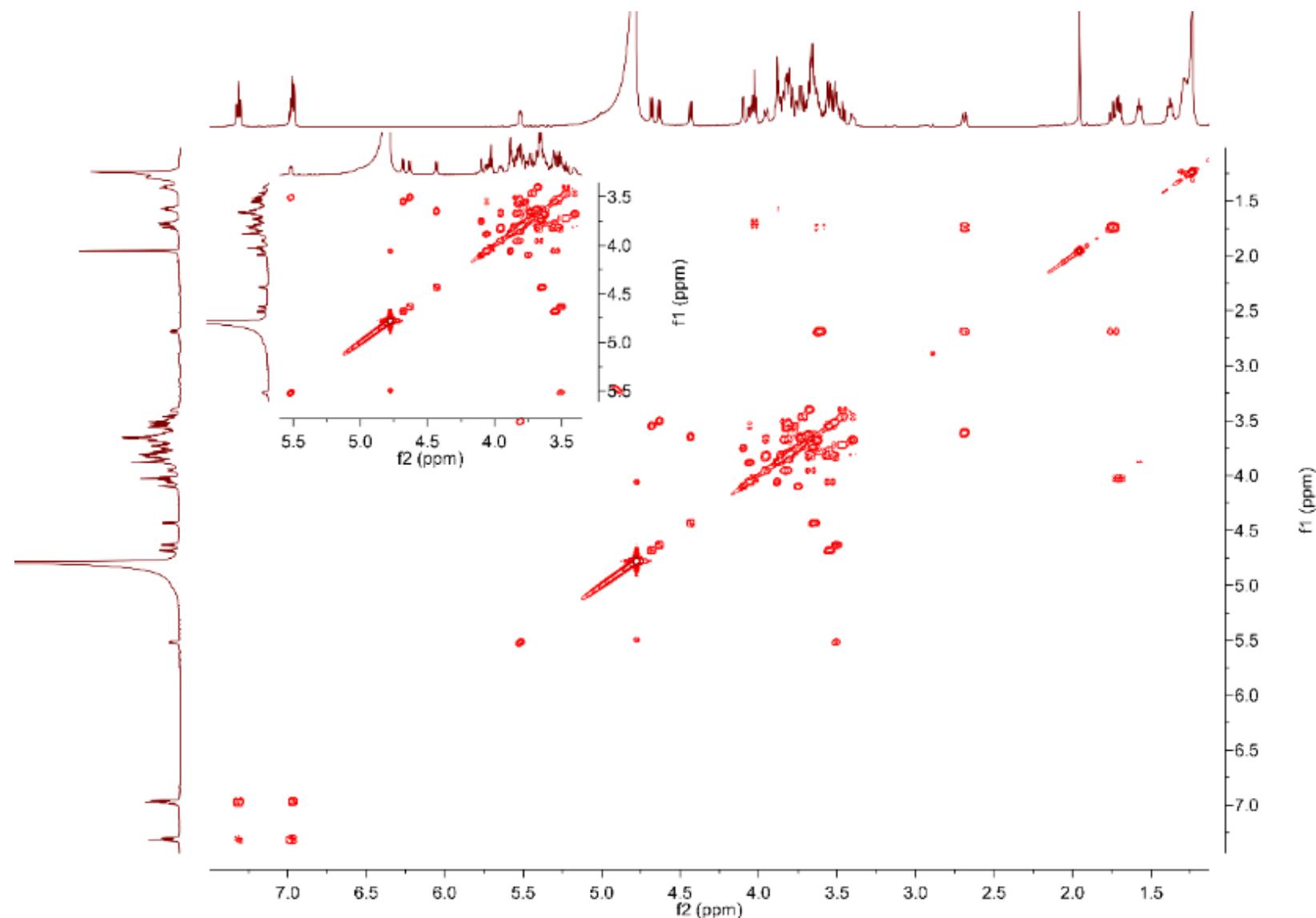


ESI-(-)-TOF HRMS of NeuNAc α -2,3-Gal β -1,4-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5d**

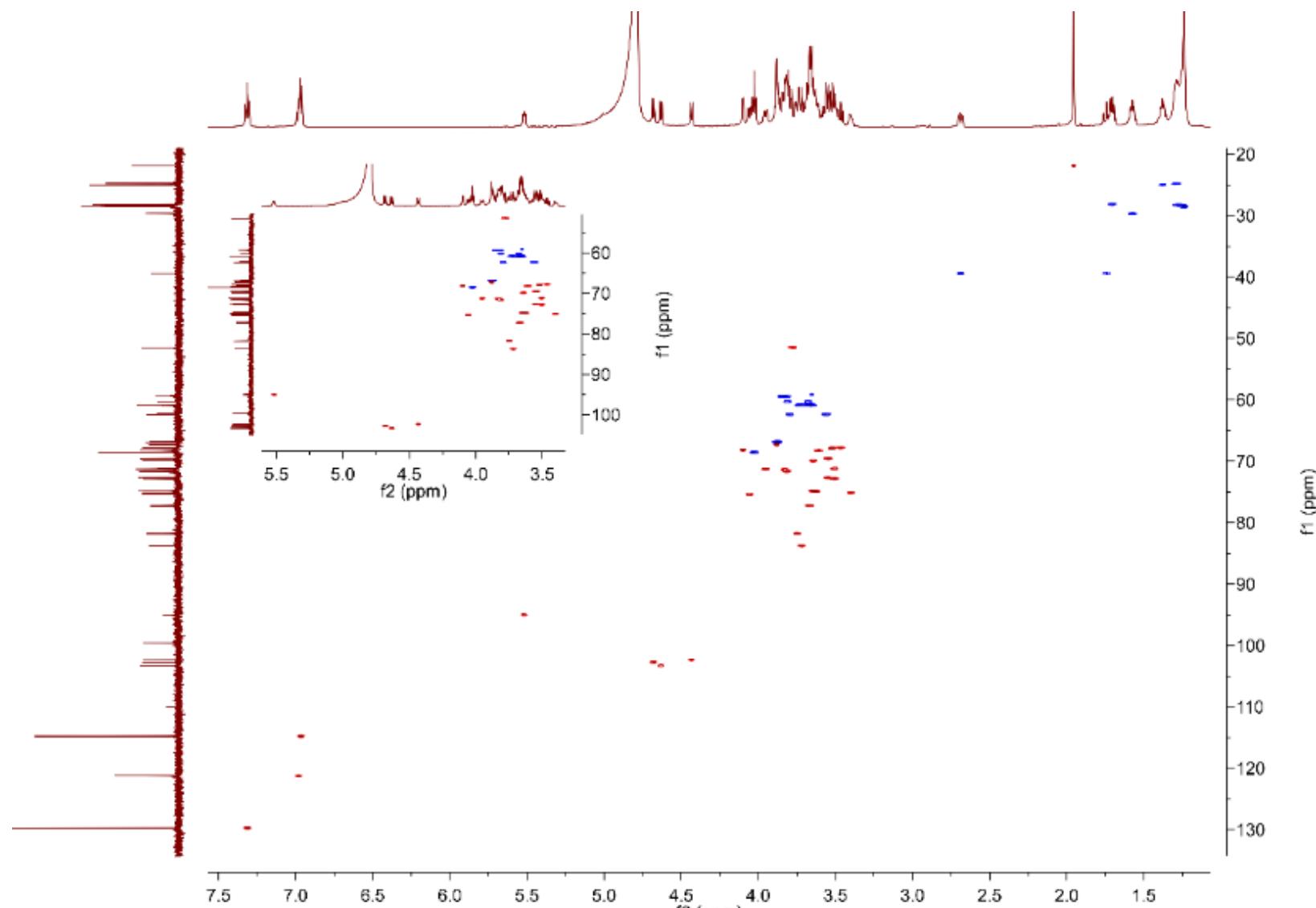


¹H NMR spectrum of NeuNAc-2,3-Galβ-1,3-Glcβ-1,3-Galβ-1,4-Glcα-PP-(CH₂)₁₁-OPh 5e (D₂O, 600 MHz, 25 °C)

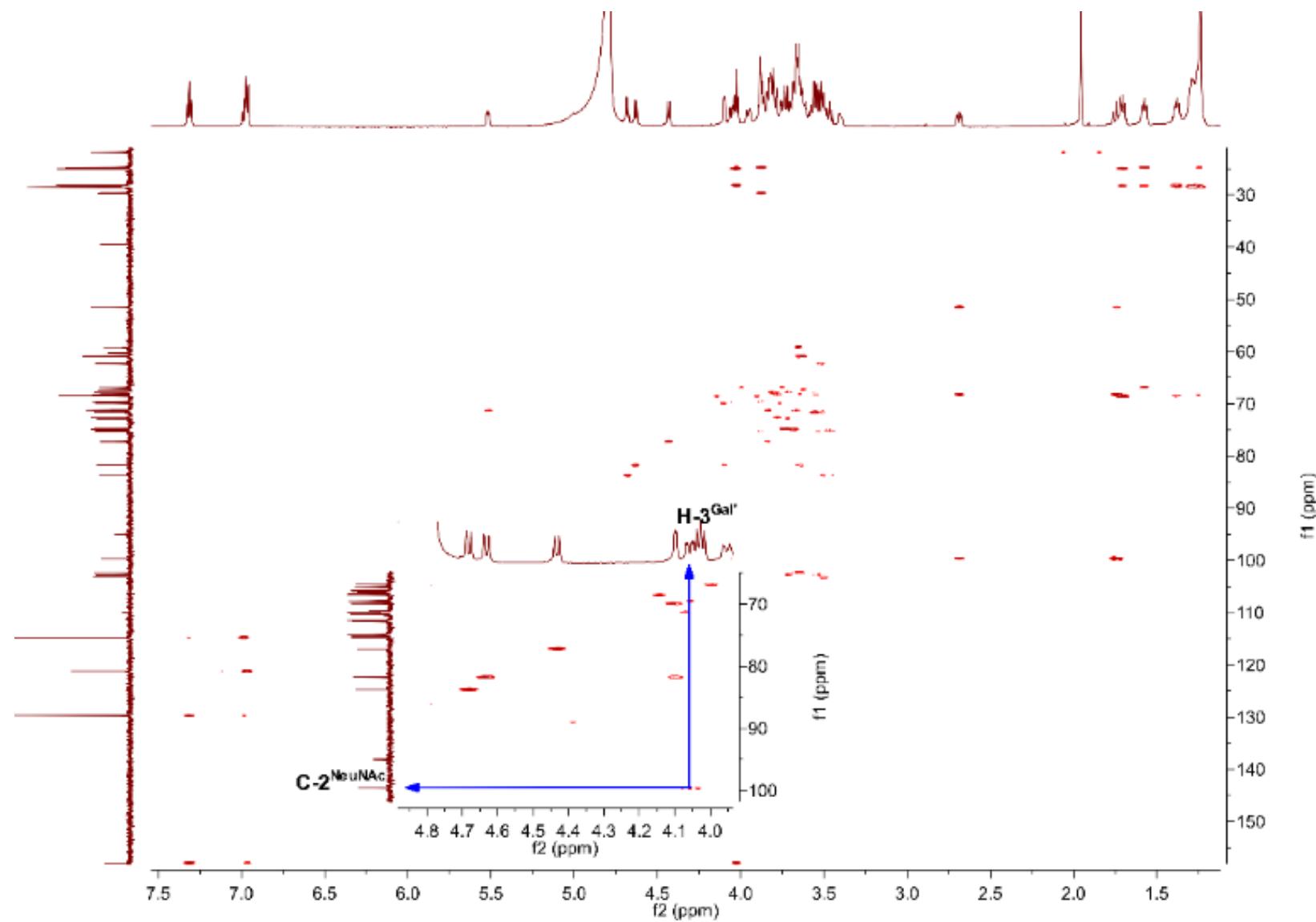


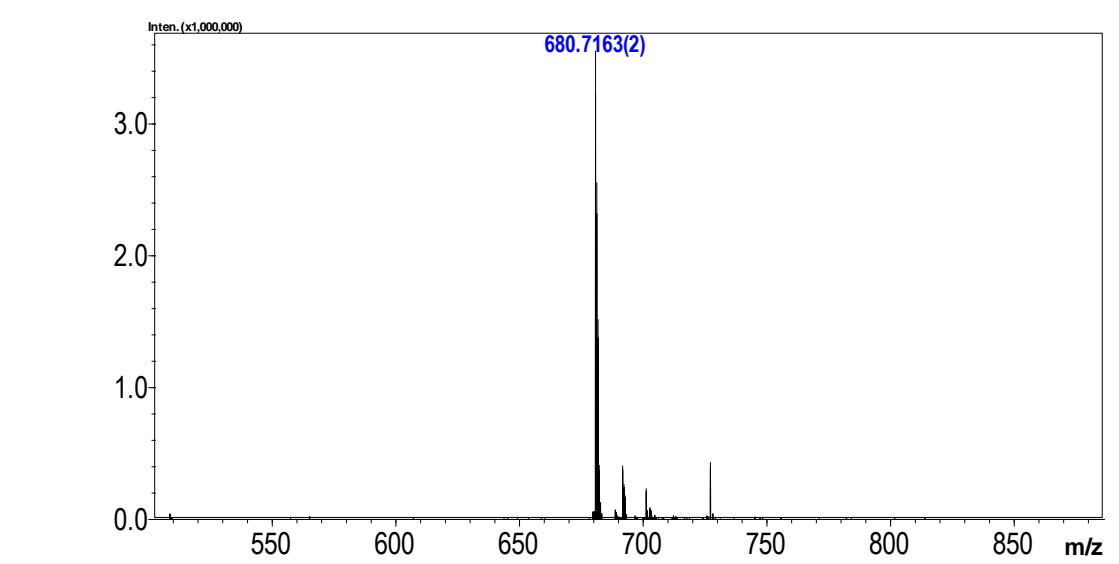
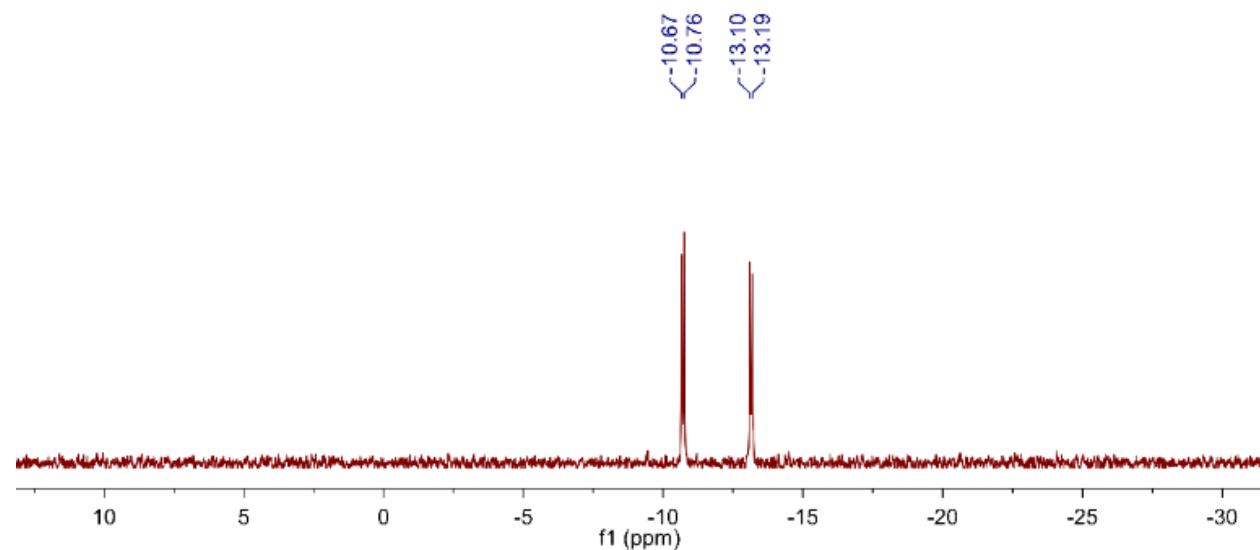


COSY spectrum of NeuNAc α -2,3-Gal β -1,3-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5e** (D₂O, 600/600 MHz, 25 °C)



HSQC spectrum of NeuNAc α -2,3-Gal β -1,3-Glc β -1,3-Gal β -1,4-Glc α -PP-(CH₂)₁₁-OPh **5e** (D₂O, 600/150 MHz, 25 °C)





IV. References

- [1] S. Li, H. Wang, G. Jin, Z. Chen, G. Gu. Exploring the broad nucleotide triphosphate and sugar-1-phosphate specificity of thymidyltransferase Cps23FL from *Streptococcus pneumonia* serotype 23F. *RSC adv.*, **2020**, 10(50): 30110-30114.
- [2] X. Wu, J. Liu, X. Yin, D. Ma, S. Zhang, X. Liu. Protein fusion of biosynthetic enzymes and a thermo-responsive polypeptide expedites facile access to biocatalysts for nucleotide sugars. *ChemBioChem*, **2025**, 26(8), e202401005.
- [3] M. Liang, G. Gong, C. Sun, J. Zhao, H. Wang, Z. Chen, M. Xiao, G. Gu. Sequential one-pot three-enzyme synthesis of the tetrasaccharide repeating unit of Group B *Streptococcus* serotype VIII capsular polysaccharide. *Chin. J. Chem.*, **2022**, 40(9): 1039-1044.
- [4] W. Gong, M. Liang, J. Zhao, H. Wang, Z. Chen, F. Wang, G. Gu. Biochemical characterization and synthetic application of WciN and its mutants from *Streptococcus pneumoniae* serotype 6B. *Front. Chem.*, **2022**, 10: 914698.
- [5] S. Masuko, S. Bera, D. E. Green, M. Wēwer, J. Liu, P. L. DeAngelis, R. J. Linhardt. Chemoenzymatic synthesis of uridine diphosphate-GlcNAc and uridine diphosphate-GalNAc analogs for the preparation of unnatural glycosaminoglycans. *J. Org. Chem.*, **2012**, 77(3): 1449-1456.
- [6] D. L. Macdonald. Preparation of glycosyl phosphates. β -fructopyranose 2-phosphate. *J. Org. Chem.*, **1966**, 31(2): 513-516.
- [7] P. Dabrowski-Tumanski, J. Kowalska, L. Jemielity. Efficient and rapid synthesis of nucleoside diphosphate sugars from nucleoside phosphorimidazolides. *Eur. J. Org. Chem.*, **2013**, 2013(11): 2147-2154.