

Supporting Information for:

Probing the Aglycon Specificity of an Engineered Glycosyltransferase

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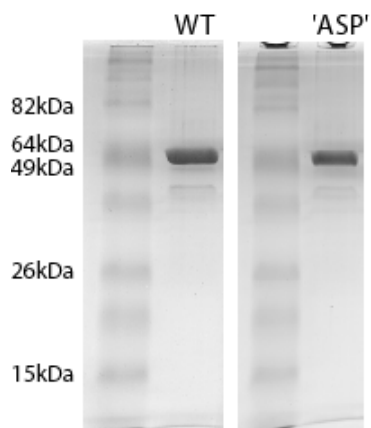
1. Full citations of references [13], [22], [30], and [35].

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- [30] C. Zhang, C. Albermann, X. Fu, N. R. Peters, J. D. Chisholm, G. Zhang, E. J. Gilbert, P. G. Wang, D. L. Van Vranken, J. S. Thorson, *ChemBioChem* **2006**, *7*, 795-804.
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2. General methods. All chemicals and reagents were purchased from Sigma-Aldrich, Fluka, or New England Biolabs unless otherwise stated. In addition to the previously reported putative aglycons [(**11**, **118-119**)^[S1] (**14**, **32**, and **38**)^[S2] (**45**, **70** and **139**)^[S3] **55**,^[S4] **73**^[S5] and **132-136**^[S6]], **29**, **50**, and **56** were a gift from Prof. Joe Langenhan, Seattle Univ.; **39** was a gift from Prof. Ben Shen, Univ. of Wisconsin-Madison; **74** and **127** were a gift from Prof. Dr. Sarah O'Connor, Massachusetts Inst. of Tech., USA; **85** was a gift from Prof. Dr. Michael Thomas, Univ. of Wisconsin-Madison; **105-112** were a gift from Prof. Dr. David L. Jakeman, Dalhousie University, Nova Scotia; **41** and **64** were isolated from fermentation; and the synthesis of **16** and **136** (Timmons, S. unpublished) will be reported elsewhere. Plasmid pET28/OleD was obtained from Prof. Hung-Wen Liu (Univ. of Texas-Austin, Austin, USA). LC/ESI-MS mass spectra were obtained using electrospray ionization on an Agilent 1100 HPLC-MSD SL quadrupole mass spectrometer connected to a UV/Vis diode array detector. High resolution mass spectra were determined utilizing electrospray ionization on a Waters (Micromass) LCT instrument (Beverly, MA, USA) with a time-of-flight analyzer and all HRMS samples contained an aliquot of a known compound (lock mass). Routine TLC analyses were accomplished on aluminum TLC plates coated with 0.2 mm silica gel from Sigma-Aldrich and monitoring at 254 nm. Flash column chromatography was achieved on 40 – 63 μm , 60 \AA silica gel (Silicycle, Quebec, Canada). Unless otherwise noted, compounds were characterized by NMR with a UNITY INOVA 400 MHz instrument (Varian, Palo Alto, CA, USA) in conjunction with a QN Switchable BB probe (Varian, Palo Alto, CA, USA). ^1H and ^{13}C chemical shifts were referenced to internal solvent resonances and reported relative to TMS. Multiplicities are indicated by s (singlet), d (doublet), t (triplet), q (quartet), qn (quintet), m (multiplet) and br (broad). Italicized elements or groups are those that are responsible for the shifts. Chemical shifts are reported in parts per million (ppm) and coupling constants J are given in Hz. NMR assignments were performed with the aid of COSY, TOCSY, HSQC, and HMBC experiments.

3. Protein expression and purification. Single colonies of *E. coli* BL21(DE3)pLysS (Stratagene, La Jolla, CA, USA) transformed with either pET28a/OleD or pET28a/OleD[A242V/S132F/P67T] vector^[S7] were used to inoculate 3 mL LB medium supplemented with 50 $\mu\text{g mL}^{-1}$ kanamycin and cultured overnight at 37°C. The entire starter culture was then transferred to 1 L LB medium supplemented with 50 $\mu\text{g mL}^{-1}$ kanamycin and grown at 37°C until the OD600 reached ~0.7. Isopropyl β -D-thiogalactoside (IPTG) was subsequently added to a final concentration of 0.4 mM and the culture was incubated at 28°C for approximately 18 hours. Cell pellets were collected by centrifugation at 10,000 g at 4°C for 20 min and the supernatant discarded. Pellets were resuspended in 10 mL lysis buffer (20 mM phosphate buffer, pH 7.4, 0.5 M NaCl, 10 mM imidazole) and were lysed by sonication. Cell debris was removed by centrifugation at 10,000 g at 4°C for 20 min and the cleared supernatant immediately applied to 2 mL of nickel nitrilotriacetic acid (Ni-NTA) resin (QIAGEN)

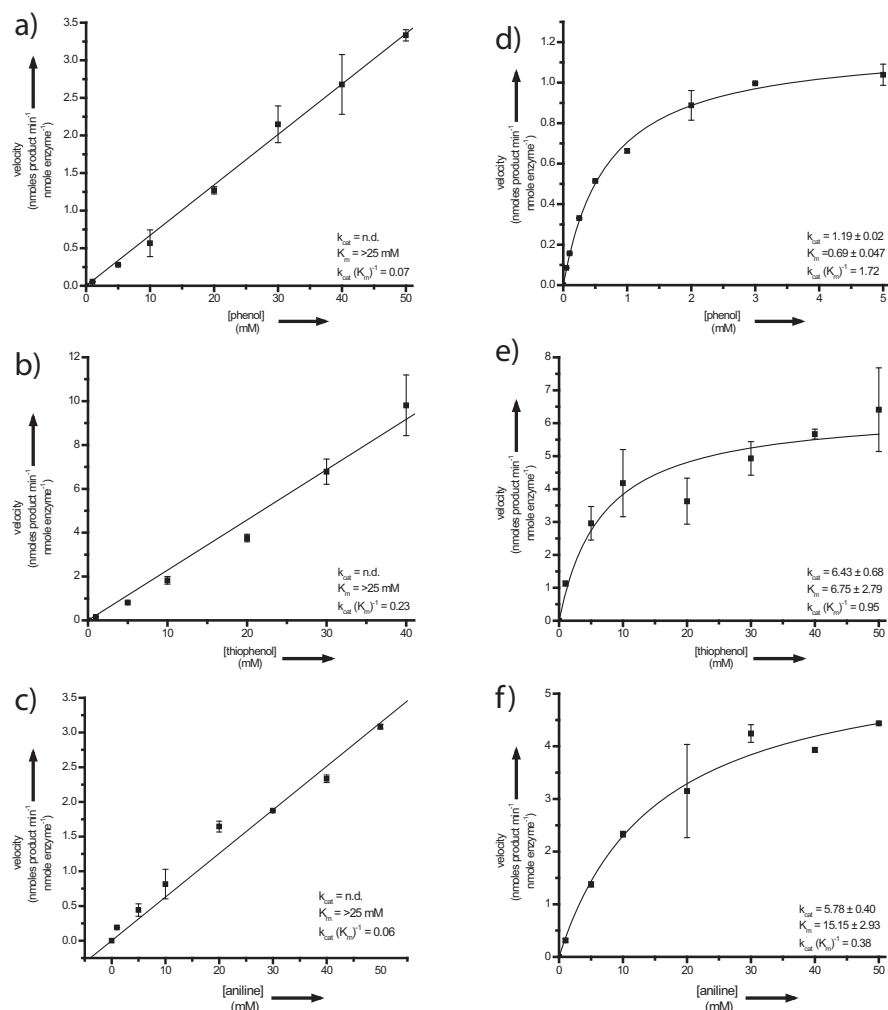
Valencia, CA, USA) pre-equilibrated with lysis buffer. Protein was allowed to bind for 30 min at 4°C with gentle agitation, and the resin washed with 50 mL lysis buffer (x 4). Finally, the enzyme was eluted by incubation of the resin with 2 mL lysis buffer containing 250 mM imidazole for 10 min at 4°C with gentle agitation. The purified protein was applied to a PD-10 desalting column (Amersham Biosciences, Piscataway, NJ, USA), equilibrated with 50 mM Tris-HCl (pH 8.0), and eluted as described by the manufacturer. Protein aliquots were immediately flash frozen in liquid nitrogen and stored at -80°C. Protein purity was confirmed by SDS-PAGE to be >95% and protein concentration for all studies was determined using the Bradford Protein Assay Kit from Bio-Rad (Hercules, CA, USA).



Supplementary Figure 1. SDS-PAGE gel of purified WT and 'ASP' OleD. Molecular weights for the standard ladder are labeled at left.

4. Acceptor library screening. Reactions were conducted in a final volume of 100 μ L and contained 50 μ g of purified enzyme, 2.5 mM UDP-glucose, 50 mM Tris-HCl (pH 8.0), 5 mM $MgCl_2$, and 1 mM of aglycon unless otherwise noted. Two separate control reactions for each aglycon that withheld either enzyme or UDP-glucose were performed in parallel. Reactions were allowed to proceed at 25 °C for ~16 hr, quenched with an equal volume of MeOH, centrifuged at 10,000 g for 10 min and the supernatant removed for analysis. The clarified reaction mixtures were analyzed by analytical reverse-phase HPLC with a 250 mm x 4.6 mm Gemini 5 μ C18 column (Phenomenex, Torrance, CA, USA) using various methods (See **Supp. Info. 8.2.**). HPLC peak areas were integrated with Star Chromatography Workstation Software (Varian, Palo Alto, CA, USA) and the total percent conversion calculated as a percent of the total peak area of substrate and product(s). Reactions which displayed potential new product(s) via HPLC were further analyzed by tandem LC/ESI-MS using a 250 mm x 4.6 mm Gemini 5 μ C18 analytical column. If both WT and 'ASP' reactions resulted in chromatographically identical product(s), only the reaction with the highest percent conversion was analyzed via LC-MS. Chromatographic methods, reactant and product retention times, and MS determinations for all positive substrates are summarized in **Supp. Info. 8.1-8.2.**

5. Determination of Kinetic Parameters. Assays were performed in a final volume of 200 μ L 50 mM Tris-HCl (pH 8.0), and contained constant concentrations of enzyme (40 μ g for 'ASP' reactions or 120 μ g for WT reactions) and saturating UDP-glucose (2.5 mM) while varying the phenol (**8**), thiophenol (**6**), or aniline (**34**) concentration (0-50 mM). Aliquots (50 μ L) were removed at 10 min (where the rate of product formation was determined to be linear), mixed with an equal volume of ice cold MeOH, and centrifuged at 10,000 g for 10 min. Supernatants were analyzed by analytical reverse-phase HPLC as described in the preceding section. HPLC peak areas were integrated with Star Chromatography Workstation Software (Varian, Palo Alto, CA, USA) and the total percent conversion calculated as a percent of the total peak area of substrate and product(s). All experiments were performed in triplicate. Initial velocities were fitted to the Michaelis-Menten equation using Origin Pro 7.0 software. OleD wild-type enzyme could not be saturated with acceptors **8**, **6**, and **34**. Consequentially, k_{cat}/K_m for wild-type was determined by linear regression.



Supplementary Figure 2. Determination of kinetic parameters for WT and 'ASP' OleD; experiments were performed in triplicate with standard deviation noted. WT with varied acceptors (a) **8**, (b) **6**, and (c) **34**. 'ASP' with varied acceptor (d) **8**, (e) **6**, and (f) **34**.

5.1. Supplementary Table 1. Kinetic parameters determined for WT and 'ASP' OleD with phenol (**8**), thiophenol (**6**), or aniline (**34**) and saturating [UDP-Glc].

Substrate	Enzyme	k_{cat} (min^{-1})	K_m (mM)	k_{cat} / K_m ($\text{mM}^{-1}\text{min}^{-1}$)	$(k_{cat} / K_m)_{ASP} / (k_{cat} / K_m)_{WT}$
phenol (8)	WT	n.a. ^[a]	n.a.	0.07	---
	'ASP'	1.19 ± 0.02	0.69 ± 0.05	1.72	24.6
thiophenol (6)	WT	n.a.	n.a.	0.23	---
	'ASP'	6.43 ± 0.68	6.75 ± 2.79	0.95	4.1
aniline (34)	WT	n.a.	n.a.	0.08	---
	'ASP'	5.78 ± 0.40	15.15 ± 2.93	0.38	4.8

^[a] n.a., not available

6. Scale-Up and Characterization of Representative Glucosides.

6.1. General Reaction Procedure. Reactions were conducted in 10 mL Tris-HCl (50 mM, pH 8.0) containing substrate **6**, **8**, or **24** (20 mM), UDP-glucose (100 mM), and MgCl_2 (5 mM) at 25 °C with agitation. Aliquots of 'ASP' enzyme (4.4 mg) were added to each reaction at 0, 7,

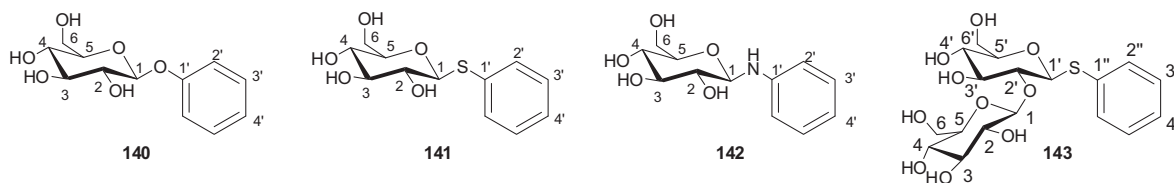
and 19 hours. The reactions allowed to proceed for a total of 42 hours and were subsequently frozen (-80°C), lyophilized, resuspended in 2 mL of ice cold MeOH, and filtered. Glucosides were isolated by collecting fractions from analytical reverse-phase HPLC with a 250 mm x 4.6 mm Gemini 5 μ C18 column (Phenomenex, Torrance, CA, USA) using the appropriate method (**Supp. Info. 8.2**). The product-containing fractions were subsequently collected and lyophilized. Products were confirmed by HRMS as previously described and via ^1H and ^{13}C NMR using a Varian ^{UNITY}INOVA 500 MHz instrument (Palo Alto, CA, USA) with a Protasis/MRM CapNMR capillary probe (Savoy, IL, USA). ^{13}C assignments were attained through gHSQC and gHMBC methods.

6.2. Phenyl β -D-glucopyranoside (140). ^1H NMR (CD_3OD , 500 MHz) δ 7.32 (m, 2 H, $\text{H}_{3'}$), 7.14 (d, $J = 8.2$ Hz, 2 H, $\text{H}_{2'}$), 7.05 (m, 1 H, $\text{H}_{4'}$), 4.96 (d, $J = 6.7$ Hz, 1 H, $\text{H}_{1'}$), 3.94 (d, $J = 12.0$ Hz, 1 H, $\text{H}_{6\text{A}}$), 3.75 (dd, $J = 12.0, 5.0$ Hz, 1 H, $\text{H}_{6\text{B}}$), 3.55-3.45 (m, 4 H, H_{2-5}); ^{13}C NMR (CD_3OD , 500 MHz) δ 158.0 ($\text{C}_{1'}$), 129.3, ($\text{C}_{3'}$), 122.3 ($\text{C}_{4'}$), 116.6 ($\text{C}_{2'}$), 101.2 (C_1), 76.9 (C_5), 73.8 (C_2), 70.3 (C_4), 63.4 (C_3), 61.4 (C_6); HRMS-ESI (m/z): $[\text{M}+\text{Na}]^+$ calcd for $\text{C}_{12}\text{H}_{16}\text{NaO}_6$, 279.0840; found 279.0833.

6.3. Phenyl 1-thio- β -D-glucopyranoside (141). ^1H NMR (CD_3OD , 500 MHz) δ 7.64 (d, $J = 7.3$ Hz, 2 H, $\text{H}_{2'}$), 7.38 (m, 2 H, $\text{H}_{3'}$), 7.35-7.30 (m, 1 H, $\text{H}_{4'}$), 4.68 (d, $J = 9.6$ Hz, 1 H, $\text{H}_{1'}$), 3.95 (d, $J = 12.0$ Hz, 1 H, $\text{H}_{6\text{A}}$), 3.74 (dd, $J = 12.0, 5.2$ Hz, 1 H, $\text{H}_{6\text{B}}$), 3.46 (m, 1 H, H_3), 3.43-3.24 (m, 2 H, $\text{H}_{4,5}$), 3.29 (m, 1 H, H_2); ^{13}C NMR (CD_3OD , 500 MHz) δ 134.2 ($\text{C}_{1'}$), 131.5 (C_2), 128.8 (C_3), 127.2 (C_4), 88.2 (C_1), 80.9 (C_4), 78.6 (C_3), 72.6 (C_2), 70.2 (C_5), 61.7 (C_6); HRMS-ESI (m/z): $[\text{M}+\text{Na}]^+$ calcd for $\text{C}_{12}\text{H}_{16}\text{NaO}_5\text{S}$, 295.0611; found 295.0624.

6.4. Phenyl 1-amino- β -D-glucopyranoside (142). ^1H NMR (CD_3OD , 500 MHz) δ 7.16 (m, 2 H, $\text{H}_{3'}$), 6.81 (d, $J = 7.6$ Hz, 2 H, $\text{H}_{2'}$), 6.73 (m, 1 H, $\text{H}_{4'}$), 4.57 (d, $J = 7.8$ Hz, 1 H, $\text{H}_{1'}$), 3.87 (d, $J = 11.7$ Hz, 1 H, $\text{H}_{6\text{A}}$), 3.73-3.66 (m, 1 H, $\text{H}_{6\text{B}}$), 3.49 (t, $J = 7.9$ Hz, 1 H, H_3), 3.44-3.33 (m, 3 H, $\text{H}_{2,4,5}$); ^{13}C NMR (CD_3OD , 500 MHz) δ 147.0 ($\text{C}_{1'}$), 128.8, ($\text{C}_{3'}$), 118.4 ($\text{C}_{4'}$), 113.9 ($\text{C}_{2'}$), 85.8 (C_1), 77.9 (C_3), 77.2 (C_5), 73.5 (C_4), 70.7 (C_2), 61.6 (C_6); HRMS-ESI (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{12}\text{H}_{16}\text{NNaO}_5$, 278.0999; found 278.1006.

6.5. Phenyl β -D-glucopyranose-(1 \rightarrow 2)-1-thio- β -D-glucopyranoside (143). ^1H NMR (acetone- d_6 , 500 MHz) δ 7.61 (m, 2 H, $\text{H}_{2'}$), 7.33 (m, 2 H, $\text{H}_{3'}$), 7.28 (m, 1 H, $\text{H}_{4'}$), 4.77 (d, $J = 9.8$ Hz, 1 H, $\text{H}_{1'}$), 4.65 (d, $J = 7.8$ Hz, 1 H, H_1), 3.91-3.85 (m, 2 H, $\text{H}_{6\text{A},6\text{A}'}$), 3.73-3.66 (m, 3 H, $\text{H}_{3',6\text{B},6\text{B}'}$), 3.51-3.47 (m, 1 H, H_2), 3.46-3.37 (m, 5 H, $\text{H}_{3-5,4',5'}$), 3.31-3.27 (m, 1 H, H_2); ^{13}C NMR (acetone- d_6 , 500 MHz) δ 131.5 ($\text{C}_{2'}$), 128.9 ($\text{C}_{3'}$), 127.1 ($\text{C}_{4'}$), 104.6 (C_1), 85.7 ($\text{C}_{1'}$), 81.8 (C_2), 80.7, 78.5 (C_3), 77.3, 76.9, 75.3 (C_2), 71.0, 70.3, 62.4 (C_6/C_6'), 62.3 (C_6/C_6'); MS-ESI (m/z): $[\text{M}+\text{Cl}]^-$ calcd for $\text{C}_{18}\text{H}_{26}\text{ClO}_{10}\text{S}$, 469.1; 469.0 observed.¹



¹ The disaccharide linkage of **143** was determined to be 1 \rightarrow 2 through comparison of COSY and TOCSY NMR experiments of the peracetylated analogs of **143** and its monosaccharide, **141**. The H2 proton of peracetylated **141** appears at 4.97 ppm (CDCl_3 , 500 MHz), while the H2 proton of the proximal sugar of peracetylated **143** is at 3.86 ppm, indicating the position of glycosylation.

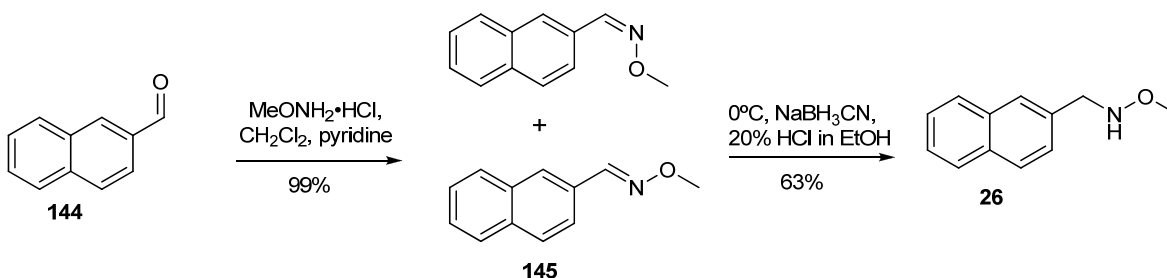
7. Aglycon syntheses and spectral data.

7.1. General reductive amination procedure. Aldehyde (12.5 mmol) was dissolved in CH_2Cl_2 to a final concentration of 0.45 M. To this was added 1.4 equivalents of $\text{MeONH}_2\cdot\text{HCl}$ and 2.2 equivalents of pyridine and the mixture was stirred for 2 hours at RT. TLC analysis at this stage revealed the substrate to be completely consumed with two products being formed. The reaction mixture was subsequently washed with 5% aqueous HCl (3 x 50 mL) and saturated NaCl (2 x 50 mL). The resulting organic layer was dried over Na_2SO_4 and concentrated under reduced pressure to provide the crude oxime (>80%) which was used in subsequent reactions without further purification.

Crude oxime was dissolved in EtOH to a final concentration of 1.5 M. The reaction mixture was cooled to 0 °C, 3 equivalents of NaBH_3CN were added, and the solution was stirred for 15 min. An equal volume of 20% HCl in EtOH chilled to 0 °C was subsequently added in a drop-wise fashion over 10 min. The reaction was then allowed to warm to RT and stirred overnight. TLC analysis revealed complete consumption of substrate and a single new product. The reaction was neutralized with the addition of Na_2CO_3 until the evolution of gas halted, concentrated under reduced pressure, and CH_2Cl_2 (20 mL) was added. The resulting mixture was washed with saturated NaHCO_3 (2 x 50 mL), dried over Na_2SO_4 , and the collected organics concentrated under reduced pressure. The concentrate was purified by flash chromatography (1:1 hexanes: CH_2Cl_2) to yield the desired methoxyamine product in > 50% yield.

7.2. N-methoxy-2-naphthalenemethanamine (26). 2-Naphthaldehyde (**144**; 2.0 g, 12.5 mmol) provided of the desired oxime **145** (2.3 g, 99% crude yield) as a white solid. TLC R_f = 0.48, 0.60 (1:8 EtOAc:hexanes); ^1H NMR (400 MHz, CDCl_3) δ 8.24 (s, 1 H, NCH), 7.92-7.80 (m, 5 H, Ph), 7.56-7.48 (m, 2 H, Ph), 4.058 (s, 3 H, OCH_3),² 4.055 (s, 3 H, OCH_3); ^{13}C NMR (100 MHz, CDCl_3) δ 148.0, 134.4, 133.5, 130.2, 128.9, 128.6, 128.6, 128.2, 127.2, 126.9, 62.4; HRMS-ESI (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{12}\text{H}_{12}\text{NO}$, 186.0841; found 186.0920.

Oxime **145** (2.3 gm, 12.4 mmol) yielded **26** (1.5 g, 63% yield) as an orange oil. TLC R_f = 0.36 (1:4 EtOAc:hexanes); ^1H NMR (400 MHz, CDCl_3) δ 7.84-7.74 (m, 4 H, Ph), 7.52-7.39 (m, 3 H, Ph), 5.80 (br s, 1 H, NH), 4.18 (s, 2 H, NHCH_2), 3.50 (t, J = 0.4 Hz, 3 H, OCH_3); ^{13}C NMR (100 MHz, CDCl_3) δ 135.5, 133.7, 133.1, 128.4, 128.1, 128.0, 127.9, 127.2, 126.3, 126.1, 62.2, 56.6; HRMS-ESI (m/z): $[\text{M}+\text{H}]^+$ calcd for $\text{C}_{12}\text{H}_{14}\text{NO}$, 188.1070; found 188.1069.

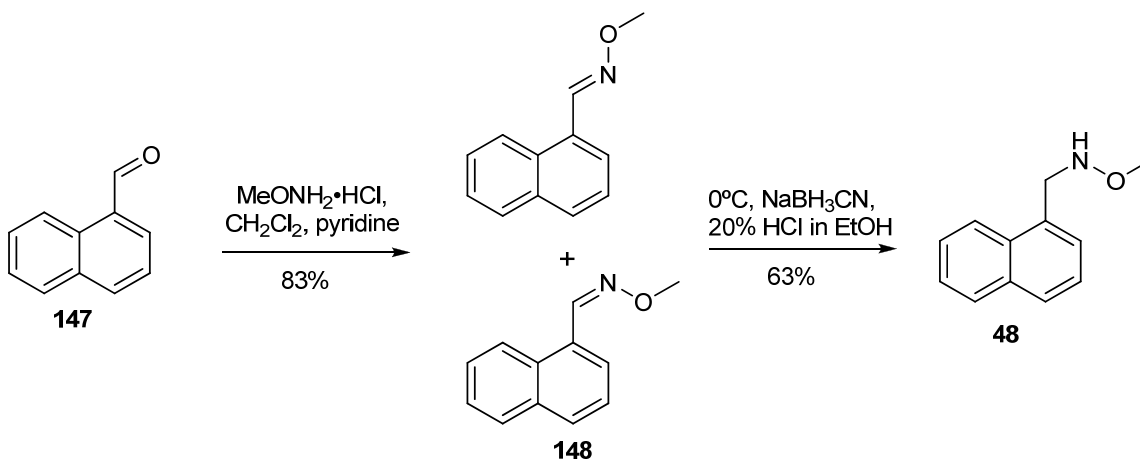


7.3. N-methoxy-1-naphthalenemethanamine (48). 1-Naphthaldehyde (**147**; 2.0 g, 12.7 mmol) gave oxime **148** (2.0 g, 83% crude yield) as a yellow oil. TLC R_f = 0.56, 0.65 (1:4 EtOAc:hexanes.); ^1H NMR (400 MHz, CDCl_3) δ 8.71 (s, 1 H, NCH), 8.52 (m, 1 H, Ph), 7.84

² Signals observed in ^1H spectrum at 4.058 and 4.055 for OCH_3 are from the presence of (*E*) and (*Z*)-isomers.

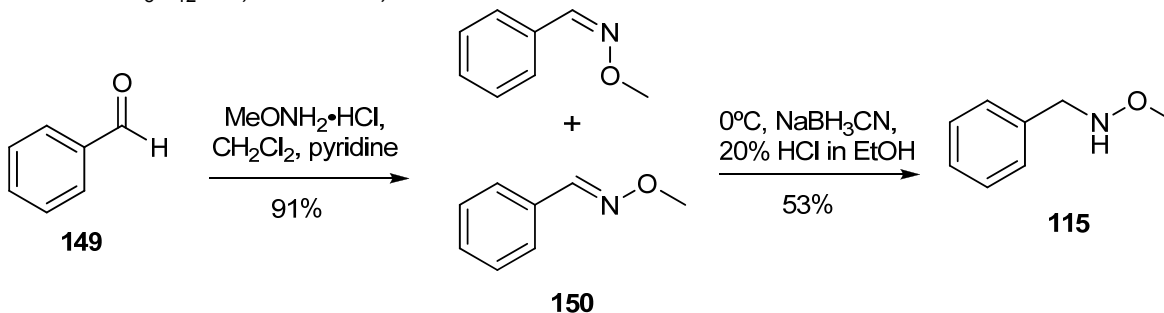
(m, 2 H, Ph), 7.74 (m, 1 H, Ph), 7.58-7.42 (m, 3 H, Ph), 4.06 (s, 3 H, OCH₃); ¹³C NMR (100 MHz, CDCl₃) δ 148.5, 133.9, 130.8, 130.5, 128.8, 128.1, 127.4, 127.1, 126.2, 125.4, 124.6, 62.2; HRMS-ESI (*m/z*): [M]⁺⁺ calcd for C₁₂H₁₁NO, 185.0836; found 185.0842.

Oxime **148** (2.0 gm, 10.5 mmol) yielded **48** (1.2 g, 63% yield) as a yellow oil. TLC R_f = 0.41 (1:4::EtOAc:hexanes); ¹H NMR (400 MHz, CDCl₃) δ 8.15 (d, *J* = 8.8 Hz, 1 H, Ph), 7.84 (d, *J* = 8.4 Hz, 1 H, Ph), 7.77 (d, *J* = 8.4 Hz, 1 H, Ph), 7.56-7.36 (m, 4 H, Ph), 5.77 (br s, 1 H, NH), 4.51 (s, 2 H, NHCH₂), 3.53 (d, *J* = 0.4 Hz, 3 H, OCH₃); ¹³C NMR (100 MHz, CDCl₃) δ 134.1, 133.0, 132.3, 129.0, 128.7, 127.8, 126.5, 126.0, 125.7, 124.0, 62.1, 54.1; HRMS-ESI (*m/z*): [M+H]⁺ calcd for C₁₂H₁₄NO, 188.1070; found 188.1070.



7.4. N-methoxybenzylamine (115). Benzaldehyde (**149**; 4.8 g, 49.2 mmol) afforded oxime **150** (6.1 g, 91% crude yield) as a colorless oil. TLC R_f = 0.82 (1:8 EtOAc:hexanes); ¹H NMR (400 MHz, CDCl₃) δ 8.03 (s, 1 H, NCH), 7.58-7.51 (m, 2 H, Ph), 7.37-7.32 (m, 3 H, Ph), 3.95 (s, 3 H, OCH₃); ¹³C NMR (100 MHz, CDCl₃) δ 148.8, 132.5, 130.0, 129.0, 127.3, 62.2³; HRMS-ESI (*m/z*): [M]⁺⁺ calcd for C₈H₉NO, 135.0679; found 135.0684.

Oxime **150** (6.1 g, 44.9 mmol) provided desired methoxyamine **115** (3.3 g, 53% yield) as a colorless oil. TLC R_f = 0.31 (1:8 EtOAc:hexanes); ¹H NMR (400 MHz, CDCl₃) δ 7.38-7.22 (m, 5 H, Ph), 5.71 (br s, 1 H, NH), 4.04 (s, 2 H, CH₂NH), 3.50 (d, *J* = 0.4 Hz, 3 H, OCH₃); ¹³C NMR (100 MHz, CDCl₃) δ 137.9, 129.1, 128.7, 127.7, 62.1, 56.5⁴; HRMS-ESI (*m/z*): [M+H]⁺ calcd for C₈H₁₂NO, 138.0914; found 138.0916.



³ Not all phenyl ¹³C resonances were observed, presumably due to spectral overlap in the 132.5-127.3 ppm region.

⁴ Not all phenyl ¹³C resonances were observed, presumably due to spectral overlap in the 129.1-127.1 ppm region.

8. Acceptor library screening summary and methods.

8.1. Data summary.

Compound Number	Compound Name	WT % conversion ^(A)	'ASP' % conversion ^(A)	substrate retention time	calculated substrate mass (m/z)	observed substrate mass (m/z)	product retention time	expected product mass (m/z)	observed product mass (m/z)	product designation ^(B)	HPLC method ^(C)	comments		
3	benzhydrol	>99.9	>99.9	19.7	184.2	n.f.	12.9	508.2	531.0 [M+Na] ⁺	DI	A			
							13.3	508.2	531.0 [M+Na] ⁺	DI				
							14.0	346.4	361.2 [M+Cl] ⁻	MONO				
4	β -zearalenol	91.8	>99.9	18.1	320.2	321.2 [M+H] ⁺	8.1	482.2	483.2 [M+H] ⁺	MONO	A			
							9.8	482.2	483.2 [M+H] ⁺	MONO				
							9.7	644.3	679.2 [M+Cl] ⁻	DI				
							10.5	644.3	667.7 [M+Na] ⁺	DI				
							11.7	482.2	483.2 [M+H] ⁺	MONO				
							12.7	644.3	643.2 [M-H] ⁻	DI				
							12.8	644.3	643.2 [M-H] ⁻	DI				
							13.9	482.2	483.2 [M+H] ⁺	MONO				
							11.7	382.0	409.4 [M+Na] ⁺	MONO			A	
							8.4	434.1	469.0 [M+Cl] ⁻	DI			A	
9.4	272.1	307.2 [M+Cl] ⁻	MONO	A										
8.7	338.3	373.2 [M+Cl] ⁻ ; 339.0 [M+H] ⁺	MONO	A	0.2 mM final concentration									
7.9	256.3	291.2 [M+Cl] ⁻	MONO	A										
9	kaempferol	71.3	88.0	17.4	286.0	285.0 [M+H] ⁺	7.1	448.1	483.0 [M+Cl] ⁻	MONO	A			
							8.2	610.2	609.0 [M-H] ⁻	DI				
							8.6	610.2	609.0 [M-H] ⁻	DI				
							8.8	610.2	609.0 [M-H] ⁻	DI				
							9.5	610.2	639.9 [M+H] ⁺ ; 645.0 [M+Cl] ⁻	DI				
							10.0	610.2	609.0 [M-H] ⁻ ; 645.0 [M+Cl] ⁻	DI				
							10.9	610.2	609.0 [M-H] ⁻	DI				
							11.2	610.2	645.0 [M+Cl] ⁻	DI				
							11.8	610.2	645.0 [M+Cl] ⁻	DI				
							12.3	448.1	447.0 [M-H] ⁻	MONO				
							12.4	448.1	447.0 [M-H] ⁻	MONO				
							12.8	448.1	447.0 [M-H] ⁻	MONO				
13.3	402.1	403.0 [M+H] ⁺	MONO	A										
12.8	565.3	566.0 [M+H] ⁺	MONO	A	0.25 mM final concentration									
9.6	301.3	336.4 [M+Cl] ⁻	MONO	A										
11.3	270.1	270.8 [M+H] ⁺ ; 292.8 [M+Na] ⁺ ; 268.8 [M+H] ⁺ ; 304.8 [M+Cl] ⁻	MONO	C										
11.89	746.1	781.0 [M+Cl] ⁻	MONO	B	0.1 final concentration									
13.8	299.1	300.2 [M+H] ⁺	MONO	J										
13.7	501.2	524.2 [M+Na] ⁺	MONO	D										
13.9	501.2	524.0 [M+Na] ⁺	MONO											
14.2	501.2	524.0 [M+Na] ⁺	MONO											
15.6	606.3	607.0 [M+H] ⁺	MONO	E	0.25 mM final concentration									
11.8	596.3	619.2 [M+Na] ⁺	DI	A										
13.1	596.3	619.4 [M+Na] ⁺	DI											
14.7	434.2	435.4 [M+H] ⁺	MONO											
13.3	493.2	494.2 [M+H] ⁺ ; 516.0 [M+Na] ⁺ ; 491.8 [M-H] ⁻ ; 528.2 [M+Cl] ⁻	DI	G										
14.0	331.1	331.6 [M+H] ⁺ ; 353.6 [M+Na] ⁺	MONO											
8.2	478.2	513.0 [M+Cl] ⁻	DI	A										
11.7	316.1	351.0 [M+Cl] ⁻	MONO											
16.2	482.2	505.0 [M+Na] ⁺	DI	E										
17.1	482.2	504.8 [M+Na] ⁺	DI											
17.5	320.1	342.9 [M+Na] ⁺	MONO											
10.3	286.3	321.2 [M+Cl] ⁻	MONO	A										
9.4	445.2	446.2 [M+H] ⁺ ; 468.2 [M+Na] ⁺	DI	G										
10.7	283.1	283.9 [M+H] ⁺ ; 305.9 [M+Na] ⁺	MONO											
12.2	536.3	537.0 [M+H] ⁺	MONO											
13.5	536.3	537.0 [M+H] ⁺	MONO	A										
13.0	323.0	328.8 [M+H] ⁺	MONO	D	0.2 mM final concentration									
11.6	511.2	510.2 [M+H] ⁺	DI	G										
12.1	511.2	534.2 [M+Na] ⁺	DI											
13.6	349.1	372.2 [M+Na] ⁺	MONO											
11.1	350.1	348.6 [M+H] ⁺ ; 384.8 [M+Cl] ⁻	MONO	A										
11.5	313.2	314.1 [M+H] ⁺	MONO	E										
15.9	613.3	613.8 [M+H] ⁺ ; 635.6 [M+Na] ⁺	DI	D										
17.2	451.1	452.2 [M+H] ⁺ ; 474.0 [M+Na] ⁺	MONO											

Compound Number	Compound Name	WT % Conversion	'ASP' % Conversion	substrate retention time	calculated substrate mass (m/z)	observed substrate mass (m/z)		product retention time	expected product mass (m/z)	observed product mass (m/z)	product designation	HPLC method	comments
30	alazarin	27	31	19.6	240.0	238.6 [M+H] ⁺		13.4	402.1	425.0 [M+Na] ⁺ ; 400.8 [M+H] ⁺	MONO	A	
31	cannabidiol	49	29	24.8	314.2	415.2 [M+H] ⁺		24.5	476.3	477.3 [M+H] ⁺	MONO	E	0.1 mM final concentration
32	pseudo-aglycon - calicheamicin fragment	43	27	19.4	1050.1	1049.1 [M+H] ⁺		9.9	1212.1	1247.1 [M+Cl] ⁺	MONO	B	0.1 mM final concentration
33	<i>N</i> -methylaniline	4.4	27.2	18.1	107.1	n.f.		8.1	431.2	432.2 [M+H] ⁺ ; 454.0 [M+Na] ⁺	DI		
								9.1	269.1	269.8 [M+H] ⁺ ; 291.8 [M+Na] ⁺	MONO	G	0.2 mM final concentration
34	aniline	3	27	12.8	93.1	94.2 [M+H] ⁺		8.0	255.1	256.0 [M+H] ⁺	MONO	G	
35	1-pyrenemethanol	2.6	26.7	22.5	232.1	255.0 [M+Na] ⁺		14.0	556.2	591.2 [M+Cl] ⁺	DI		
								15.2	556.2	591.2 [M+Cl] ⁺	DI	A	
								16.0	394.1	429.0 [M+Cl] ⁺	MONO		
36	ampicillin	3	27	9.8	349.1	350.2 [M+H] ⁺		15.4	511.2	511.8 [M+H] ⁺ ; 533.8 [M+Na] ⁺	MONO		
								16.1	511.2	511.8 [M+H] ⁺ ; 533.8 [M+Na] ⁺	MONO	A	
37	benzhydrilamine	34	27	19.1	183.1	208.0 [M+Na] ⁺		18.1	345.2	345.8 [M+H] ⁺ ; 367.8 [M+Na] ⁺	MONO	E	
38	α31 - calicheamicin fragment	21	25	17.4	1210.2	1245.1 [M+Cl] ⁺		15.2	1372.2	1407.1 [M+Cl] ⁺	MONO	B	0.1 mM final concentration
39	fredericamycin A	17	25	23.6		540.0 [M+H] ⁺		10.0	701.2	700.8 [M+H] ⁺	MONO	A	0.25 mM final concentration
40	dynemicin A	45	25	24.9	537.1	537.8 [M+H] ⁺		14.8	699.6	722.2 [M+Na] ⁺	MONO	M	0.1 mM final concentration
41	hydramycin	7	24	19.1	408.1	407.0 [M+H] ⁺		15.7	570.1	569.0 [M+H] ⁺	MONO	L	
42	2-phenylethanethiol	31	23	22.8	138.2	172.2 [M+Cl] ⁺		11.8	300.1	335.0 [M+Cl] ⁺	MONO	A	
43	pyrimethamine	36	21	20.2	248.1	249.2 [M+H] ⁺		17.8	410.1	411.2 [M+H] ⁺	MONO	J	
44	7-amino-4-methylcoumarin	1	18	12.3	175.1	176.2 [M+H] ⁺		9.2	337.3	338.0 [M+H] ⁺	MONO	A	0.2 mM final concentration
45	staurosporine aglycon	5	15	19.5	311.1	312.2 [M+H] ⁺		14.6	473.2	474.8 [M+H] ⁺	MONO	A	0.05 mM final concentration
46	benzylalcohol	8	14	11.4	108.1	107.0 [M+H] ⁺		8.75	270.1	305.0 [M+Cl] ⁺	MONO	A	
47	phenylacetic hydrazide	0	13	12.8	150.1	151.2 [M+H] ⁺		15.2	312.1	313.2 [M+H] ⁺	MONO	J	
48	<i>N</i> -methoxy-1-naphthalenemethanamine	18	11	19.8	187.1	188.2 [M+H] ⁺		13.5	349.1	350.2 [M+H] ⁺	MONO	H	
49	brefeldin A	1	10	16.0	280.2	281.2 [M+H] ⁺			442.2	477.2 [M+Cl] ⁺	MONO	A	
50	<i>N</i> -ethoxybenzhydrilamine	41	10	24.8	227.1	250.0 [M+Na] ⁺		14.2	389.2	412.2 [M+Na] ⁺	MONO	D	
51	benzaldehyde oxime	3	9	14.8	121.1	122.2 [M+H] ⁺		9.9	283.1	284.2 [M+H] ⁺	MONO	A	
52	benzoic hydrazide	3	7	5.7	136.1	137.0 [M+H] ⁺		7.2	298.1	299.2 [M+H] ⁺	MONO	A	
53	sulfanilamide	0	6.7	10.6	172.0	173.1 [M+H] ⁺		3.7	334.1	335.1 [M+H] ⁺	MONO		
								5.9	334.1	335.1 [M+H] ⁺ ; 369.0 [M+Cl] ⁺	MONO	E	
54	4-biphenylacetic acid	0.8	5.9	20.3	212.1	213.2 [M+H] ⁺		15.9	374.1	397.0 [M+Na] ⁺	MONO		
								16.0	374.1	397.0 [M+Na] ⁺	MONO	A	
								16.2	374.1	397.0 [M+Na] ⁺	MONO		
55	deacetyl-colchicine	21	3	11.0	357.2	358.0 [M+H] ⁺		11.4	519.0	520.0 [M+H] ⁺	MONO		0.5 mM final concentration
56	<i>N</i> -methoxybenzhydrilamine	9	2	23.6	213.2	248.6 [M+Cl] ⁺		13.4	375.2	398.0 [M+Na] ⁺	MONO	D	
57	nystatin A1	41.6	0.0	15.2	925.5	926.0 [M+H] ⁺		12.7	1087.6	1088.4 [M+H] ⁺	MONO		
								13.9	1087.6	1087.8 [M+H] ⁺	MONO	F	0.25 mM final concentration
								14.1	1087.6	1088.2 [M+H] ⁺	MONO		
58	2-naphthoic acid	0.2	4.6	17.4	172.1	173.2 [M+H] ⁺		12.0	334.1	357.2 [M+Na] ⁺	MONO		
								12.3	334.1	357.2 [M+Na] ⁺	MONO	A	
								12.6	334.1	357.2 [M+Na] ⁺	MONO		
59	1-hydroxyanthraquinone	3	4	24.1	224.1	224.8 [M+H] ⁺		13.1	386.1	386.8 [M+H] ⁺	MONO	A	
60	3-iodobenzoic acid	0.6	3.2	18.7	248.0	249.0 [M+H] ⁺		13.4	410.0	445.0 [M+Cl] ⁺	MONO		
								13.6	410.0	445.0 [M+Cl] ⁺	MONO	A	
								13.9	410.0	445.0 [M+Cl] ⁺	MONO		
								15.0	410.0	445.0 [M+Cl] ⁺	MONO		
61	sulfapyridine	1	3	15.4	249.1	250.1 [M+H] ⁺		12.6	411.1	412.2 [M+H] ⁺	MONO	E	
62	daunorubicin	0.4	2.8	14.5	527.2	528.2 [M+H] ⁺		11.1	689.2	690.2 [M+H] ⁺	MONO	A	
63	trimethoprim	2	2	15.4	290.1	291.3 [M+H] ⁺		13.3	689.2	690.2 [M+H] ⁺	MONO		
								12.0	452.2	453.2 [M+H] ⁺	MONO	E	
64	calicheamicin γ	3	1	15.6	1367.3	1402.2 [M+Cl] ⁺		9.9	1529.3	1584.3 [M+Cl] ⁺	MONO	B	0.1 mM final concentration
65	3-chlorobenzoic acid	0.1	1.8	17.2	156.0	157.0 [M+H] ⁺		11.9	318.1	317.0 [M+H] ⁺	MONO		
								12.4	318.1	317.0 [M+H] ⁺	MONO	A	
								12.8	318.1	317.0 [M+H] ⁺	MONO		

Compound Number	Compound Name	WT % Conversion	'ASP' % Conversion	substrate retention time	calculated substrate mass (m/z)	observed substrate mass (m/z)		product retention time	expected product mass (m/z)	observed product mass (m/z)	product designation	HPLC method	comments
66	ivermectin	0	2	33.2	874.5	897.2 [M+Na] ⁺		23.4	1036.6	1059.2 [M+Na] ⁺	MONO	K	0.1 mM final concentration
67	phenylacetic acid	0	2	14.1	136.1	137.2 [M+H] ⁺		9.9	298.1	333.2 [M+Cl] ⁻	MONO	A	
68	2-benzoylbenzoic acid	1	1	18.2	226.1	227.2 [M+H] ⁺		13.2	388.1	411.2 [M+Na] ⁺	MONO	A	
								13.4	388.1	411.0 [M+Na] ⁺	MONO		
								13.6	388.1	411.0 [M+Na] ⁺	MONO		
								13.7	388.1	411.2 [M+Na] ⁺	MONO		
								13.9	388.1	411.2 [M+Na] ⁺	MONO		
69	chloramphenicol	1	1	14.4	322.0	321.0 [M-H] ⁻		12.4	484.1	483.0 [M-H] ⁻	MONO	A	
70	staurosporine glucoside	0	1	14.6	473.2	472.2 [M-H] ⁻		7.6	635.2	670.0 [M+Cl] ⁻	MONO	A	0.05 mM final concentration
71	avermectin	0	1	26.3	872.5	895.2 [M+Na] ⁺		18.2	1034.5	1058.0 [M+Na] ⁺	MONO	K	0.1 mM final concentration
72	indole	0	0.4	18.4	117.1	118.2 [M+H] ⁺		10.6	279.1	280.0 [M+H] ⁺	MONO	A	
73	ivermectin pseudo-aglycon	0	0.2	27.8	730.4	753.0 [M+Na] ⁺		18.65	892.5	915.6 [M+Na] ⁺	MONO	K	0.01 mM final concentration

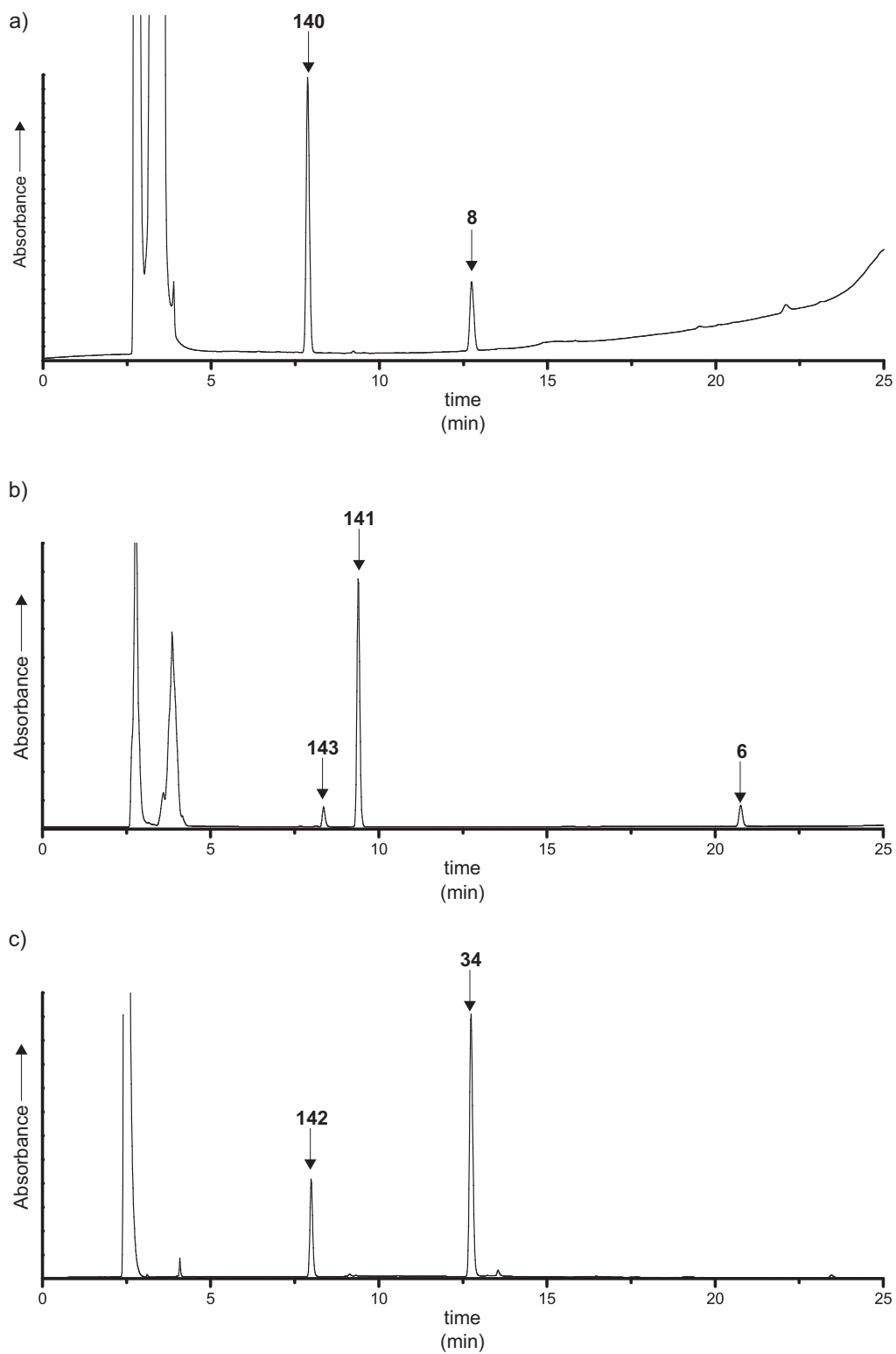
[a] % conversion is calculated as the sum of all observed products; [b] MONO – mass corresponds to monosaccharide product, DI – mass corresponds to disaccharide product; [c] refer to Supp. Info. 8.2 for HPLC methods; [d] n.f. – not found

8.2. HPLC methods.

Method	Solvent A ^[a]	Solvent B	Gradient	Detection (nm)
A	0.01% TFA	CH ₃ CN	10-75% B, 20 min; 75-100% B, 1min; 100% B, 5 min	254
B	0.01% TFA	CH ₃ CN	10-100% B, 20 min	280
C	50 mM Tris, pH 8.0	CH ₃ CN	5% B, 2.5 min; 5-60% B, 18.5 min; 60-100% B, 1 min; 100% B, 6 min	254
D	ddH ₂ O	CH ₃ CN	10-75% B, 20 min; 75-100% B, 1min; 100% B, 14 min	254
E	ddH ₂ O	CH ₃ CN	2% B, 2.5 min; 2-60% B, 17.5 min; 60-100% B, 1 min; 100% B, 4 min	254
F	0.01% TFA	CH ₃ CN	10-75% B, 20 min; 75-100% B, 1min; 100% B, 5 min	305
G	10 mM NH ₄ CO ₃ , pH 10.0	CH ₃ CN	10-75% B, 20 min; 75-100% B, 1min; 100% B, 5 min	254
H	ddH ₂ O	CH ₃ CN	10-75% B, 20 min; 75-100% B, 1min; 100% B, 5 min	254
J	ddH ₂ O	CH ₃ CN	5% B, 5 min; 5-75% B, 15 min; 75-100% B, 1 min; 100% B, 5 min	254
K	ddH ₂ O	CH ₃ CN	30-70% B, 5 min; 70-100% B, 30 min; 100% B, 3 min	243
L	0.01% TFA	CH ₃ CN	10-100% B, 15 min; 100% B, 20 min	254
M	0.01% TFA	CH ₃ CN	10-75% B, 20 min; 75-100% B, 1min; 100% B, 5 min	320

[a] 0.1% formic acid in ddH₂O was utilized for LC/ESI-MS methods when solvent A was ddH₂O or 0.1% trifluoroacetic acid in ddH₂O. 5 mM ammonium bicarbonate (pH 8.0) was utilized for LC/ESI-MS methods when solvent A was 50 mM Tris-HCl [pH 8.0] in ddH₂O or 10 mM ammonium bicarbonate (pH 10.0) in ddH₂O.

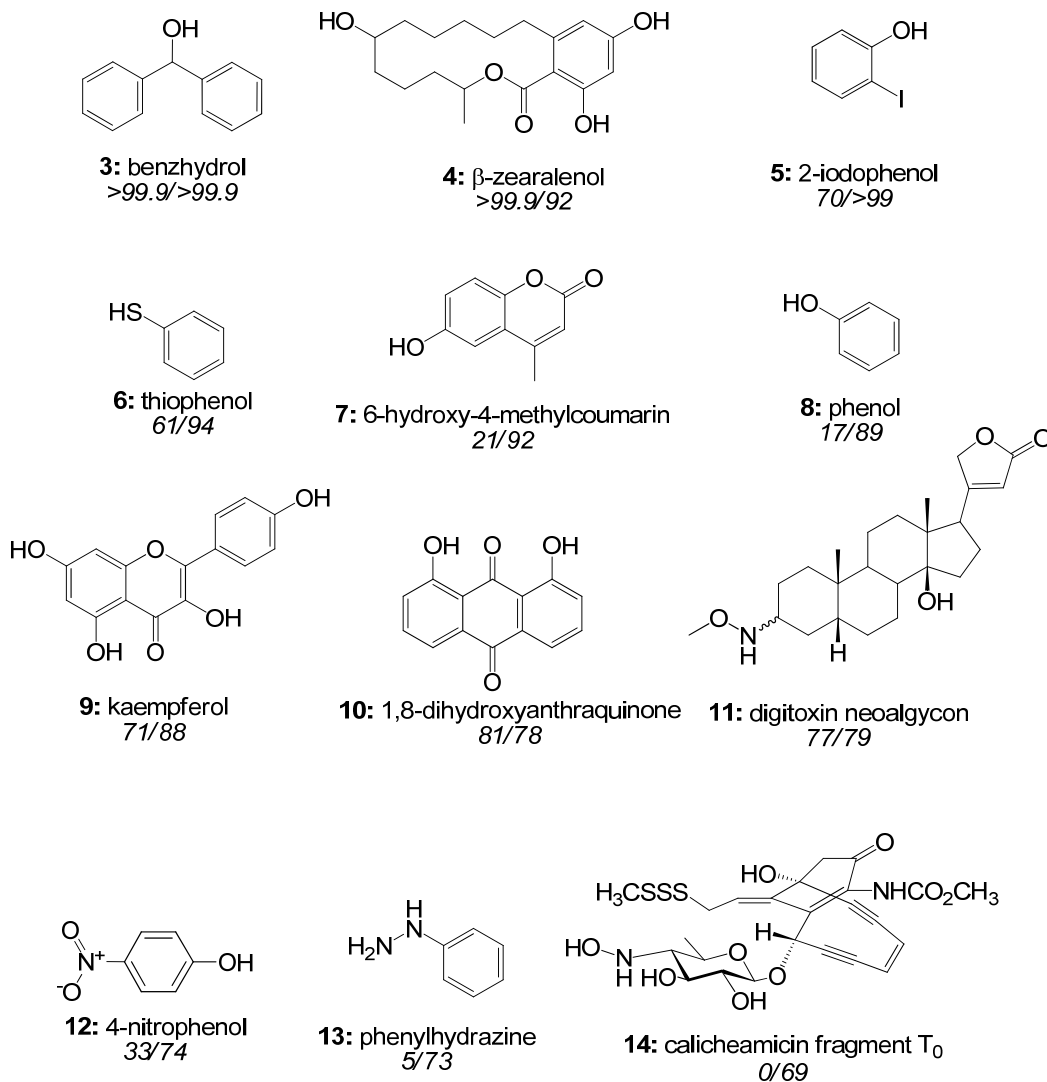
8.3 Representative HPLC traces.



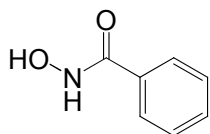
Supplementary Figure 3. Representative HPLC traces for OleD reactions with **a)** phenol (**8**), **b)** thiophenol (**6**), and **c)** aniline (**34**).

9. Structures of aglycon library members.

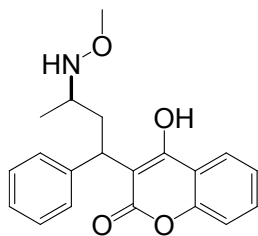
Supplementary Figure 4. Compounds that resulted in observable glucosylation (**3-73**) are listed in descending order of observed 'ASP' conversion. Compounds with no observable glucosylation (**74-139**) are listed in alphabetical order. *Italic numbers noted under compound number and name denote observed percent conversion for WT/ASP.*



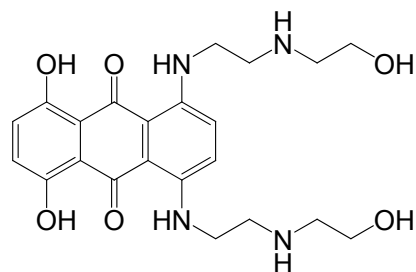
Structures of All Library members (con't)



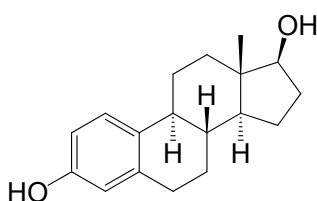
15: N-hydroxybenzamide
(benzohydroxamic acid)
25/66



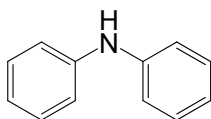
16: warfarin neoglycon
6/65



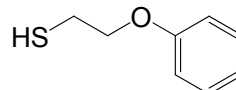
17: mitoxantrone
11/63



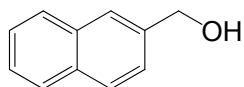
18: beta-estradiol
0/60



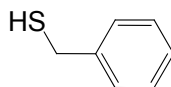
19: diphenylamine
51/49



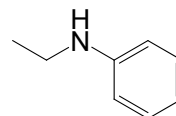
20: phenoxyethanethiol
32/49



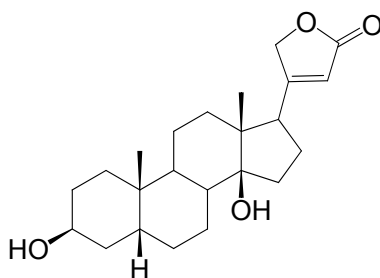
21: 2-naphthalenemethanol
80/46



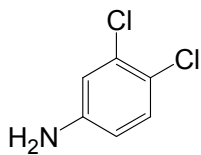
22: benzylmercaptan
37/45



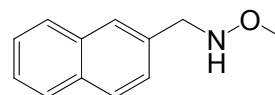
23: N-ethylaniline
10/41



24: digitoxigenin
9/40

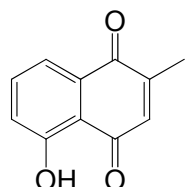


25: 3,4-dichloroaniline
9/39

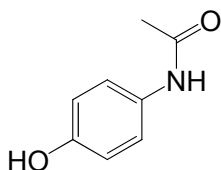


26: N-methoxy-2-
naphthalenemethanamine
64/38

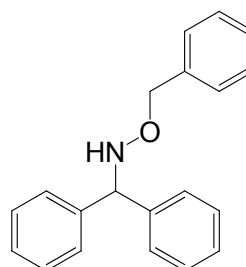
Structures of All Library members (con't)



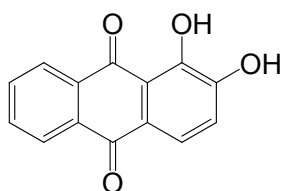
27: plumbagin
11/35



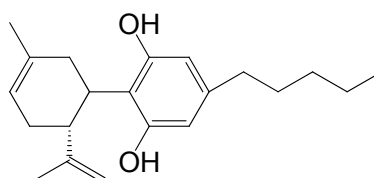
28: acetaminophen
3/35



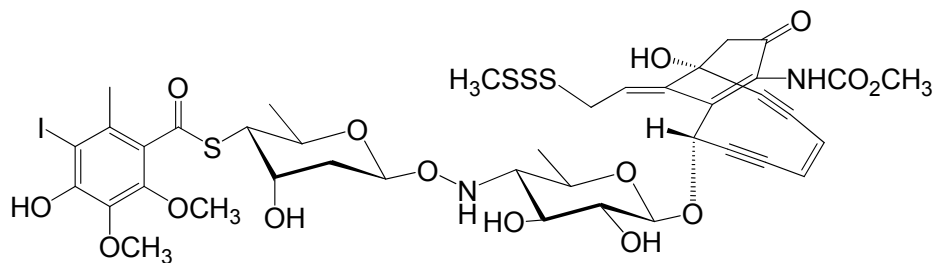
29: N-benzyloxybenzhydramine
87/32



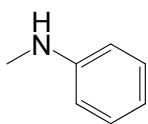
30: alizarin
27/31



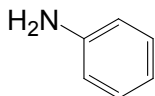
31: cannabidiol
49/29



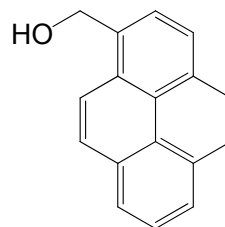
32: calicheamicin fragment pseudo-aglycon
43/27



33: N-methylaniline
4/27

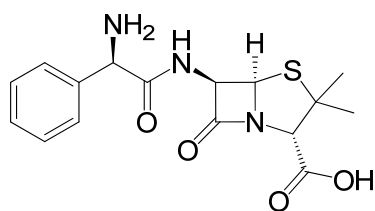


34: aniline
3/27

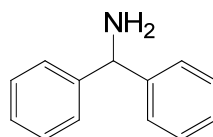


35: 1-pyrenemethanol
3/27

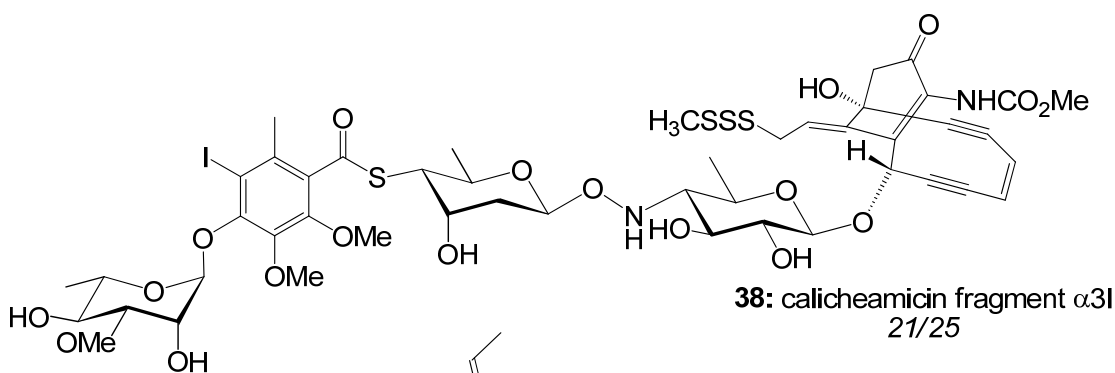
Structures of All Library members (con't)



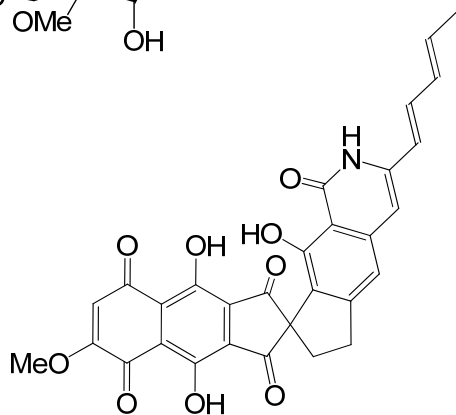
36: ampicillin
3/27



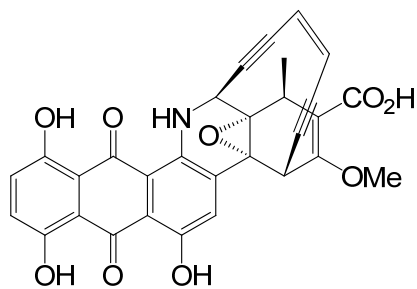
37: benzhydrylamine
(aminodiphenylmethane)
34/27



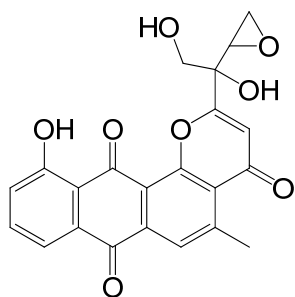
38: calicheamicin fragment α 3I
21/25



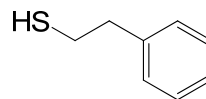
39: fredericamycin A
17/25



40: dynemicin A
24/25

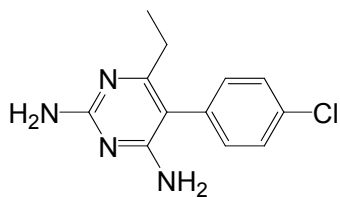


41: hydramycin
7/24

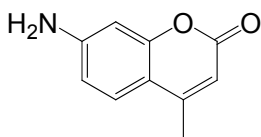


42: 2-phenylethanethiol
31/23

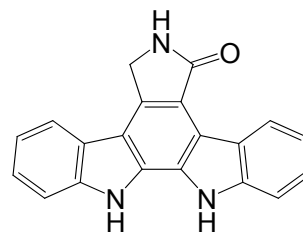
Structures of All Library members (con't)



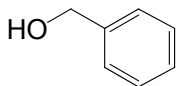
43: pyrimethamine
36/21



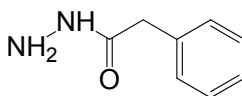
44: 7-amino-4-methylcoumarin
1/18



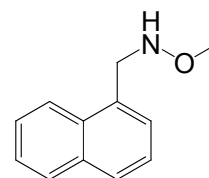
45: staurosporine aglycon
5/15



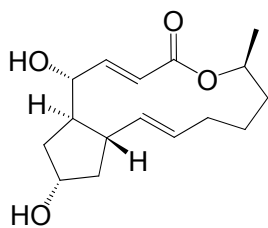
46: benzylalcohol
8/14



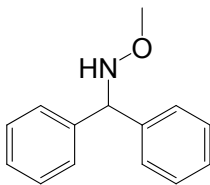
47: phenylacetic hydrazide
0/13



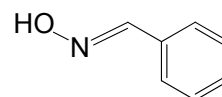
48: N-methoxy-1-naphthalenemethanamine
18/11



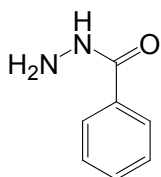
49: brefeldin A
1/10



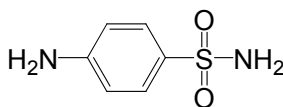
50: N-methoxybenzhydramine
9/2



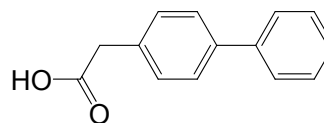
51: benzaldehyde oxime
3/9



52: benzoic hydrazide
3/7

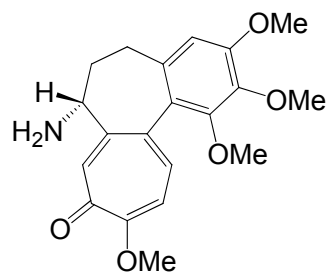


53: sulfanilamide
0/7

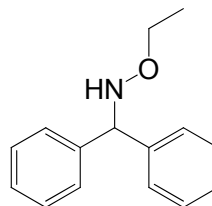


54: 4-biphenylacetic acid
1/6

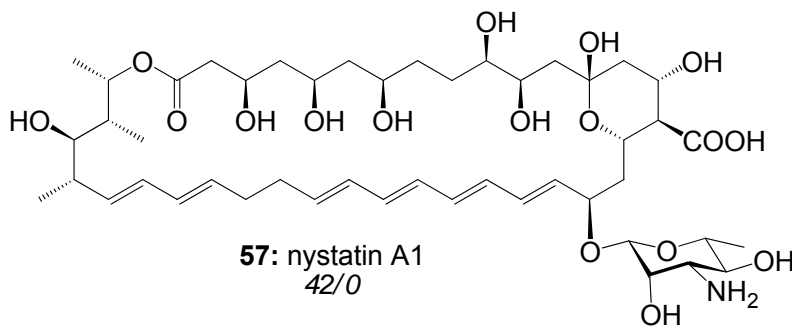
Structures of All Library members (con't)



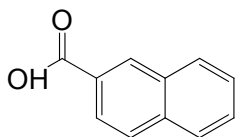
55: deacetyl-colchicine
21/3



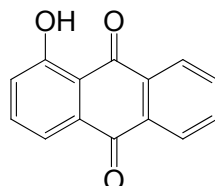
56: N-ethoxydiphenylmethanamine
41/10



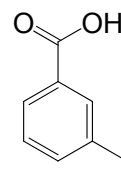
57: nystatin A1
42/0



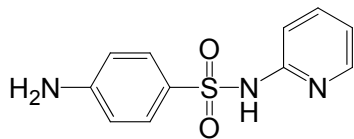
58: 2-naphthoic acid
0.2/5



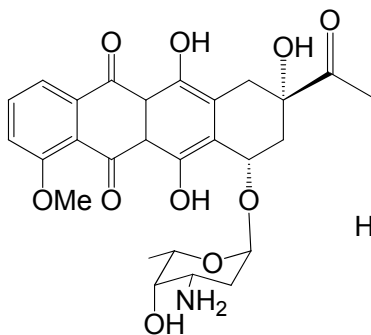
59: 1-hydroxyanthraquinone
3/4



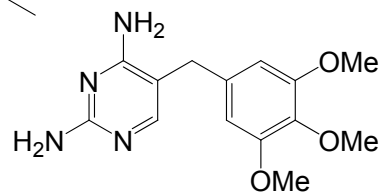
60: 3-iodobenzoic acid
1/3



61: sulfapyridine
1/3

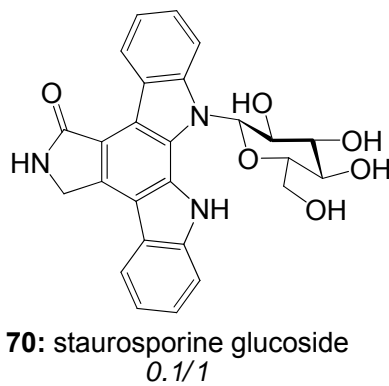
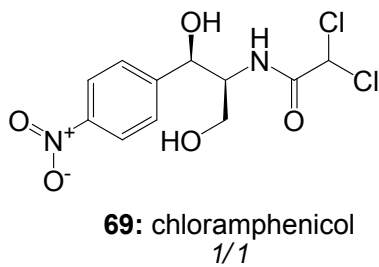
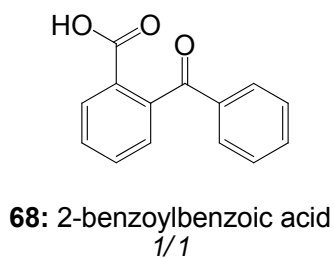
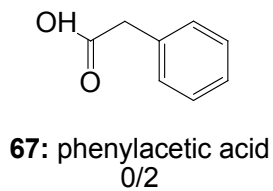
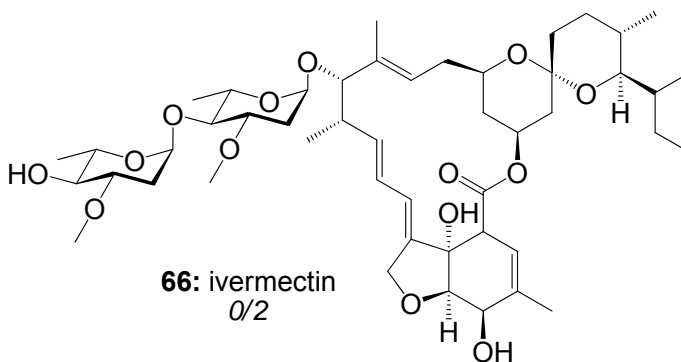
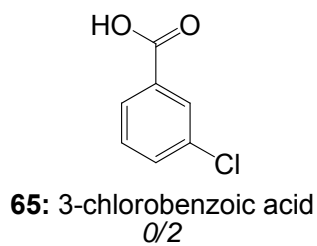
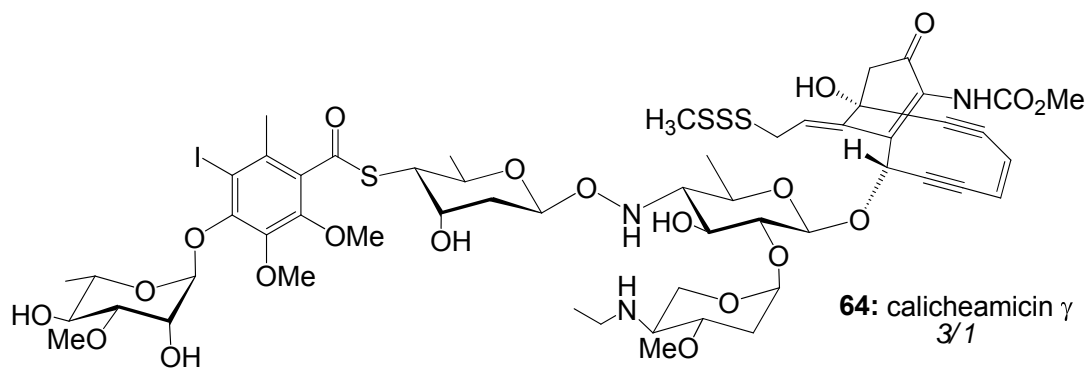


62: daunorubicin
0/3

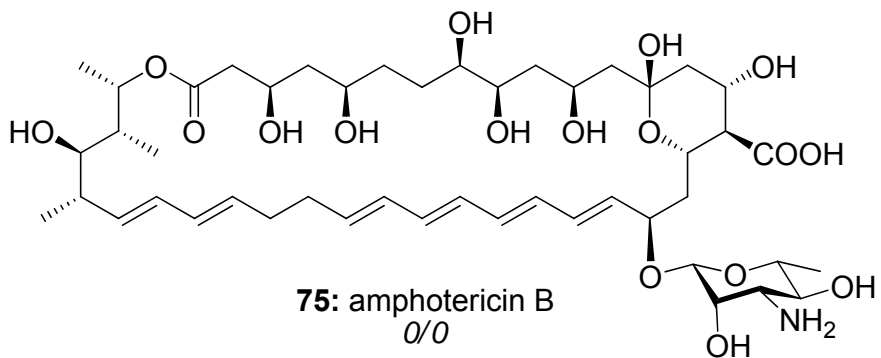
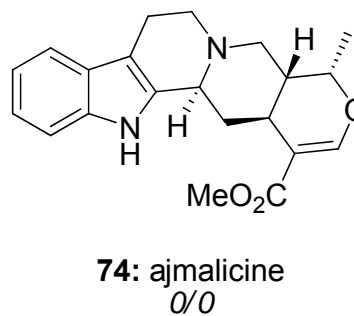
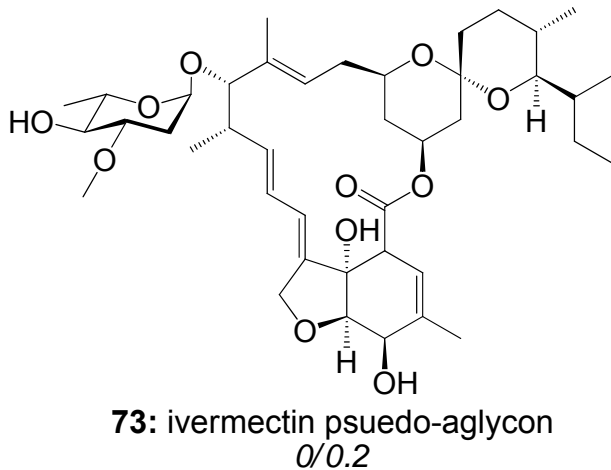
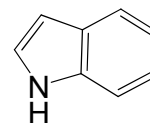
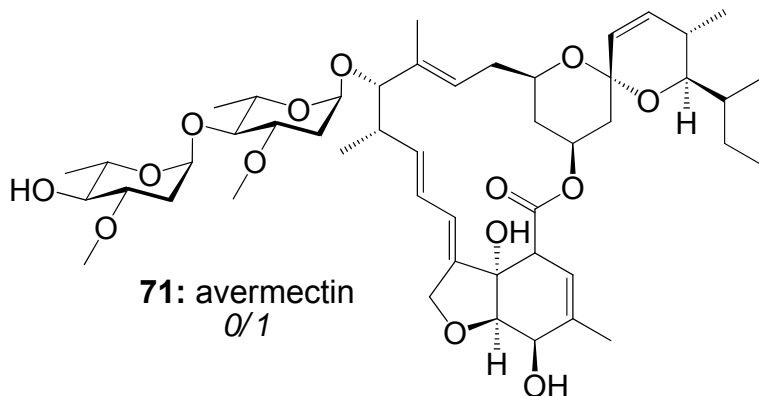


63: trimethoprim
2/2

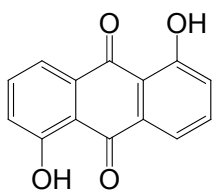
Structures of All Library members (con't)



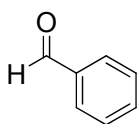
Structures of All Library members (con't)



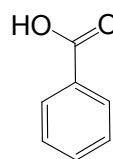
Structures of All Library members (con't)



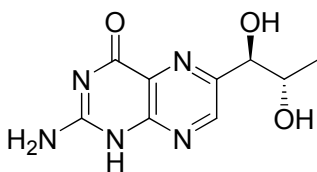
76: anthrurufin
0/0



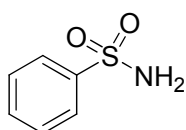
77: benzaldehyde
0/0



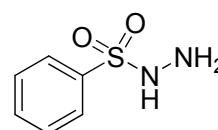
78: benzoic acid
0/0



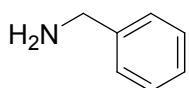
79: 6-biopterin
0/0



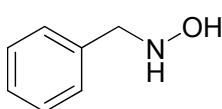
80: benzenesulfonamide
0/0



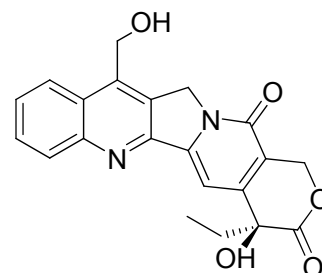
81: benzenesulfonyl hydrazide
0/0



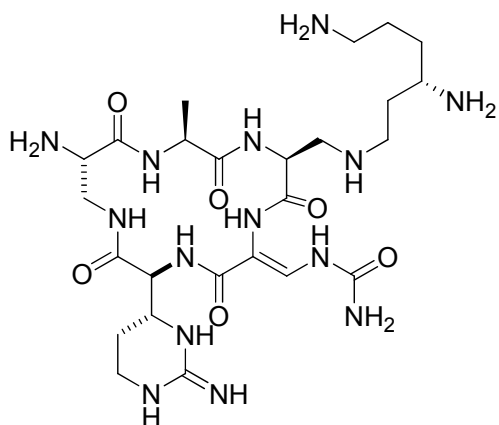
82: benzylamine
0/0



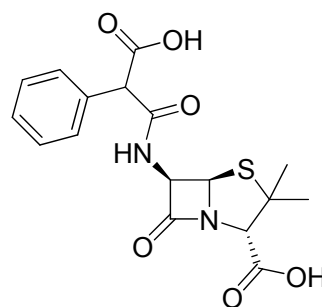
83: O-benzylhydroxylamine
0/0



84: camptothecin
0/0

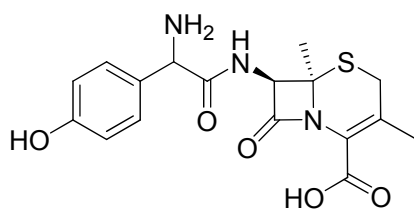


85: capreomycin 1B
0/0

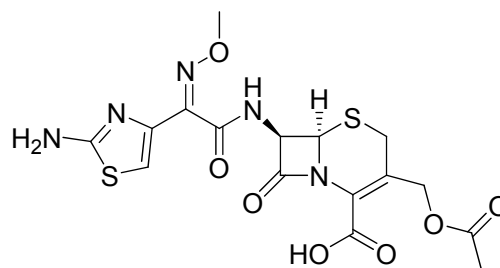


86: carbenicillin
0/0

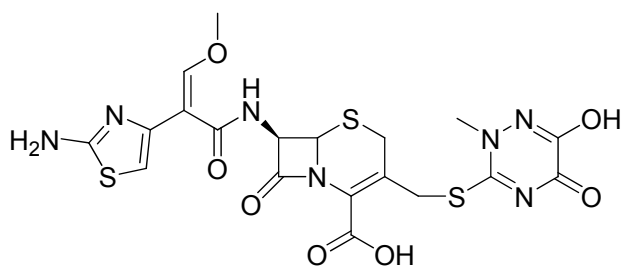
Structures of All Library members (con't)



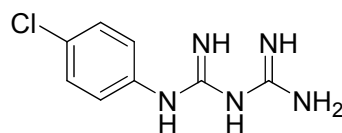
87: cefadroxil
0/0



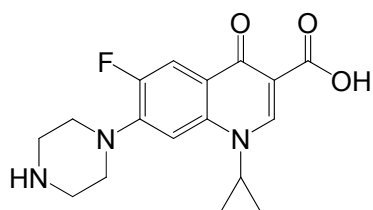
88: cefotaxime
0/0



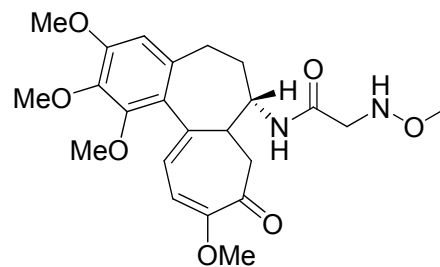
89: ceftriaxone
0/0



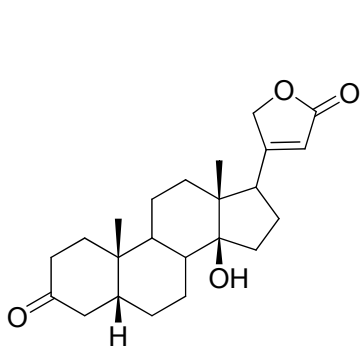
90: 1-(4-chlorophenyl)biguanide
0/0



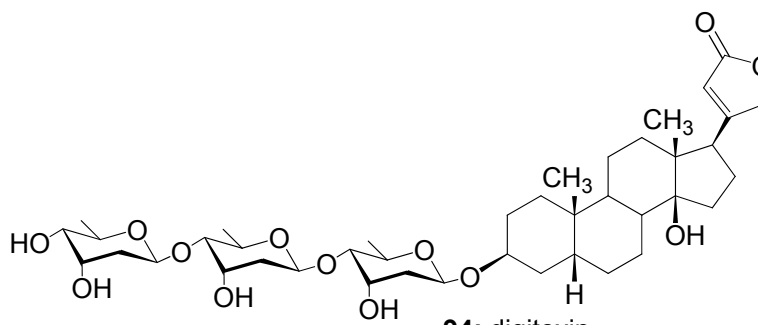
91: ciprofloxacin
0/0



92: colchicine neoaglycon
0/0

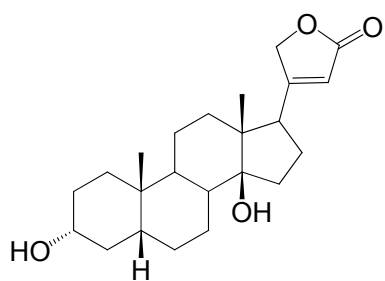


93: digitoxone
0/0

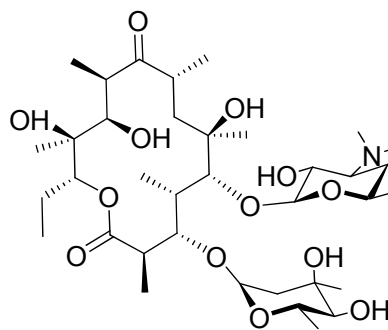


**94: digitoxin
(digoxin)**
0/0

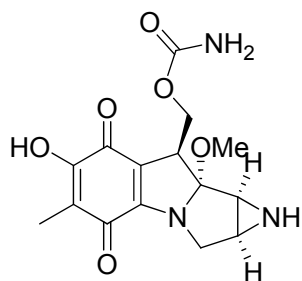
Structures of All Library members (con't)



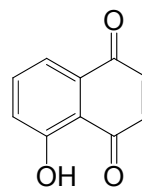
95: epi-digitoxigenin
0/0



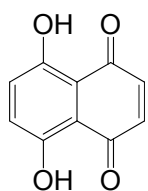
96: erythromycin
0/0



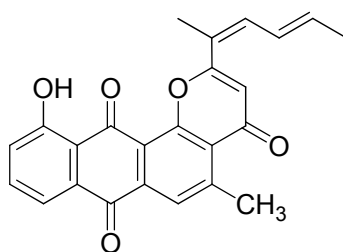
97: hydroxymitomycin C
0/0



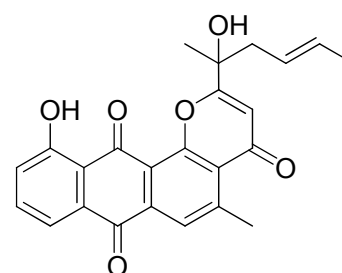
98: 5-hydroxy-1,4-naphthaquinone
0/0



99: 5,8-dihydroxy-1,4-naphthaquinone
0/0

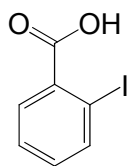


100: α -indomycinone
0/0

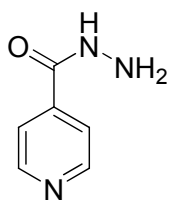


101: β -indomycinone
0/0

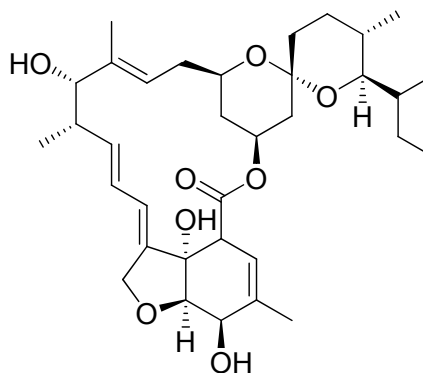
Structures of All Library members (con't)



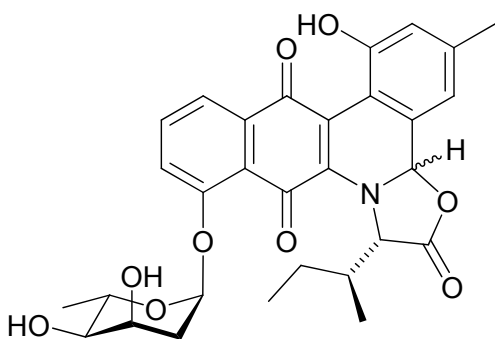
102: 2-iodobenzoic acid
0/0



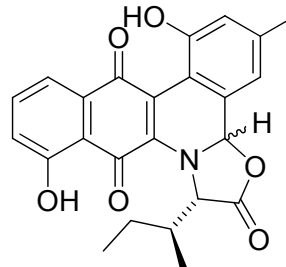
103: isoniazid
0/0



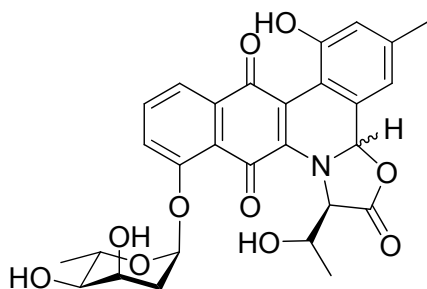
104: ivermectin aglycon
0/0



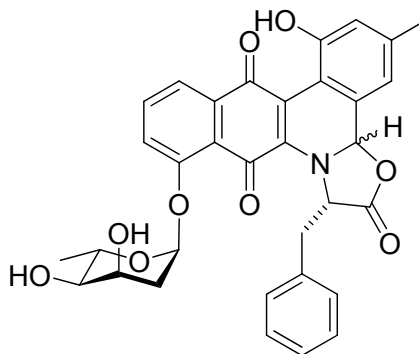
105: jadomycin B
0/0



106: jadomycin B aglycone
0/0

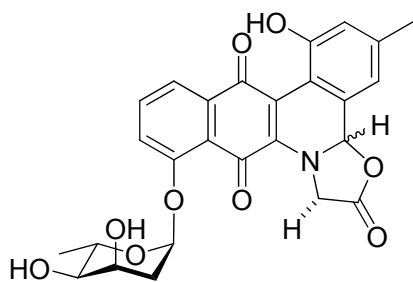


107: jadomycin DT

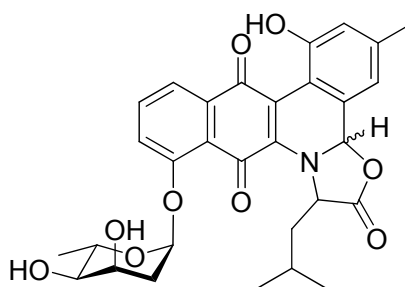


108: jadomycin F

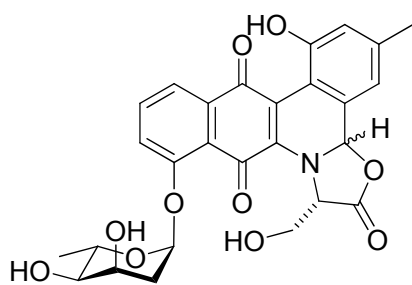
Structures of All Library members (con't)



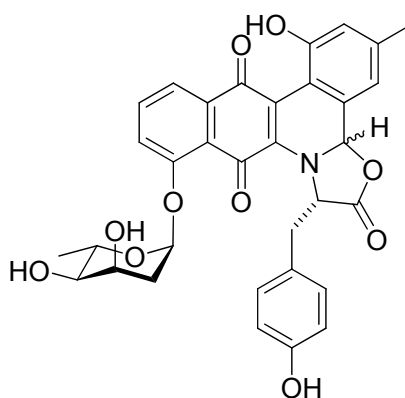
109: jadomycin G



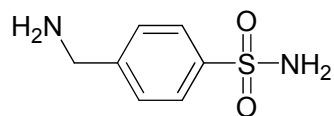
110: jadomycin L



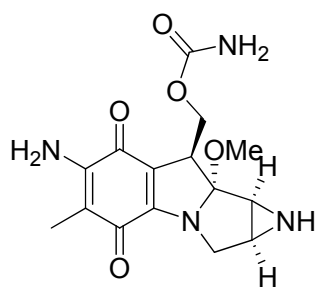
111: jadomycin S



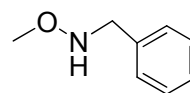
112: jadomycin Y



113: mafenide
0/0

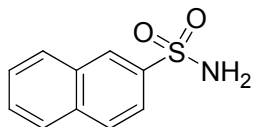


114: mitomycin C
0/0

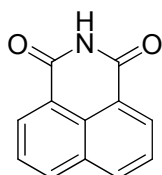


115: N-methoxybenzylamine
0/0

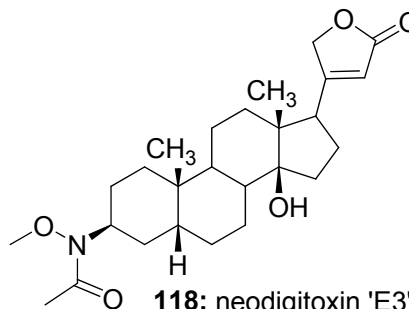
Structures of All Library members (con't)



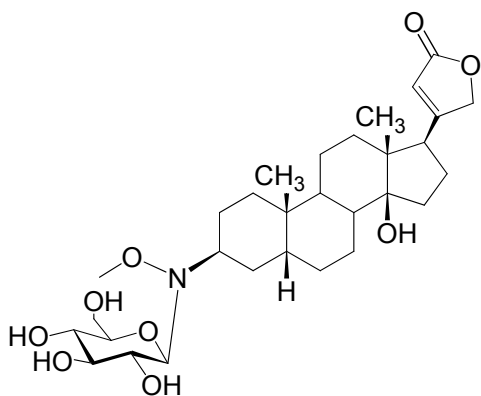
116: naphthalene-2-sulfonamide
0/0



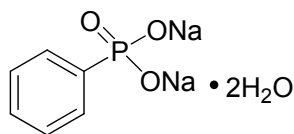
117: 1,8-naphthalimide
0/0



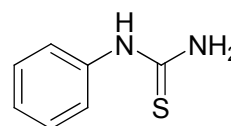
118: neodigitoxin 'E3'
0/0



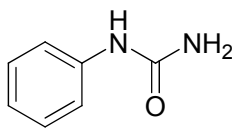
119: neodigitoxin glucoside
0/0



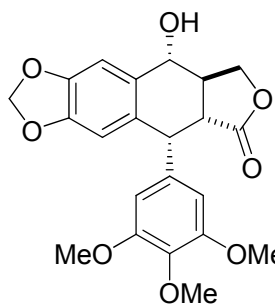
120: sodium phenyl phosphate
dibasic dihydrate
0/0



121: N-phenylthiourea
0/0

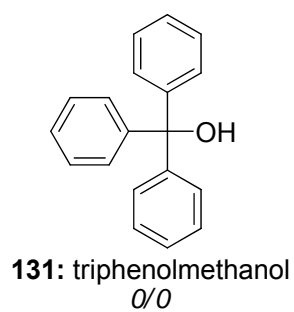
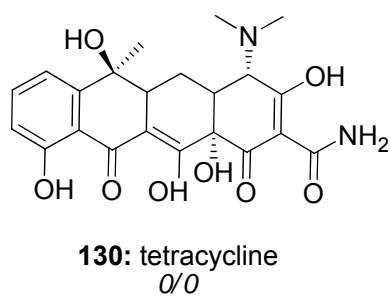
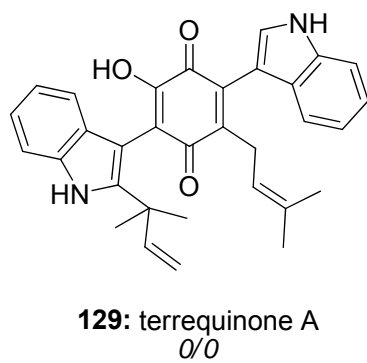
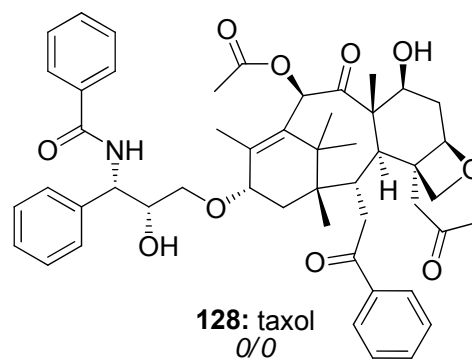
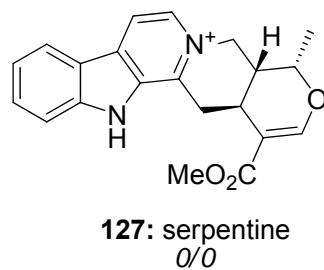
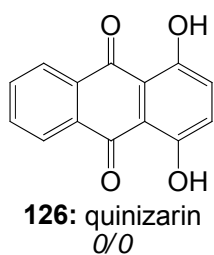
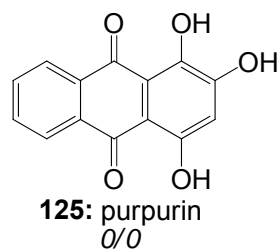
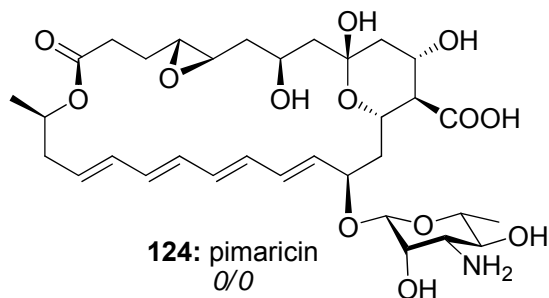


122: N-phenylurea
0/0

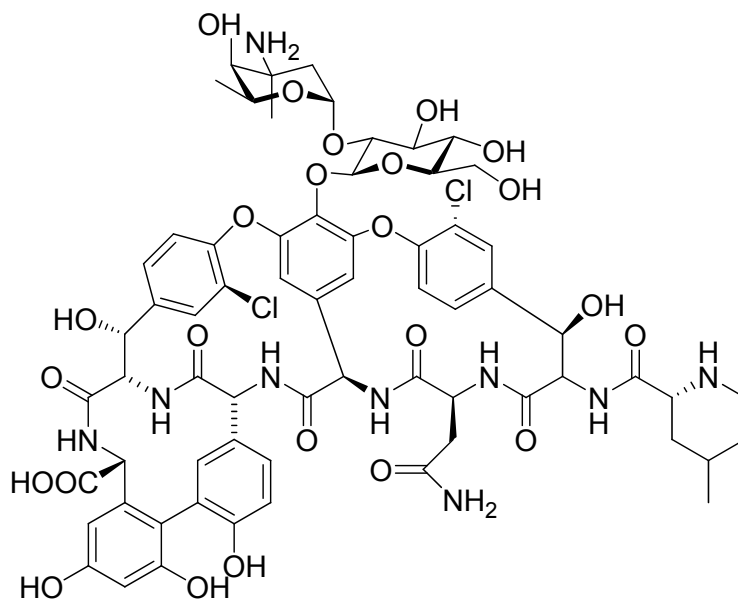


123: podophyllotoxin
0/0

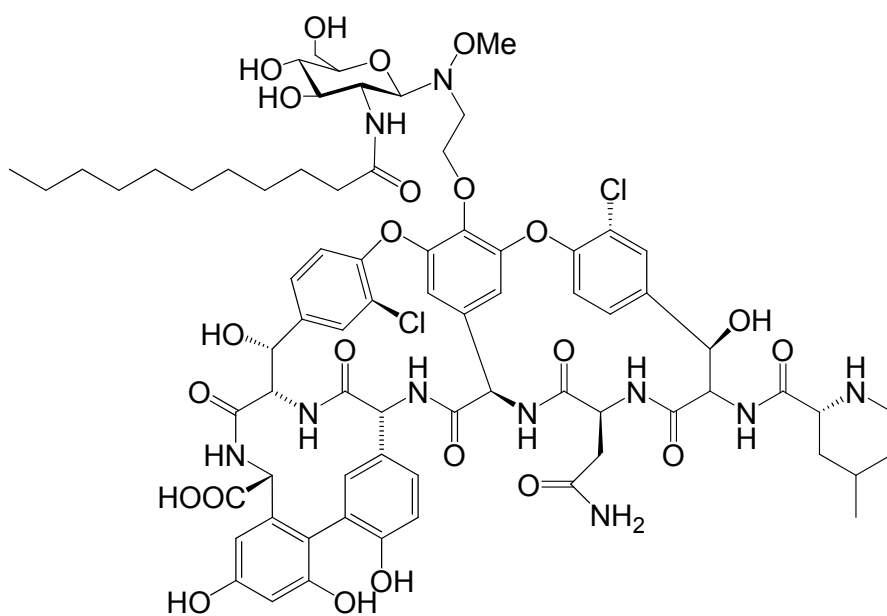
Structures of All Library members (con't)



Structures of All Library members (con't)

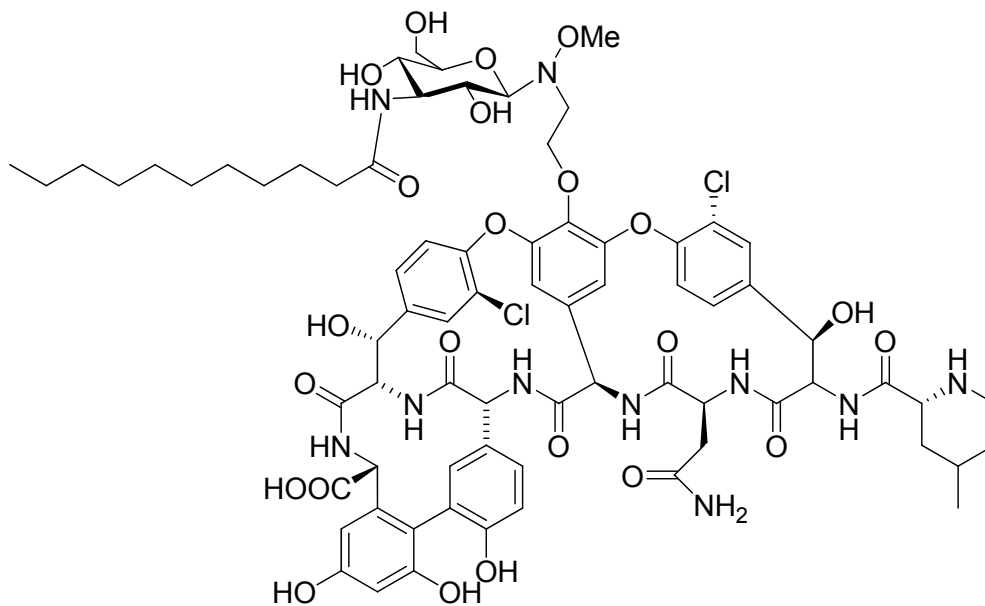


132: vancomycin
0/0

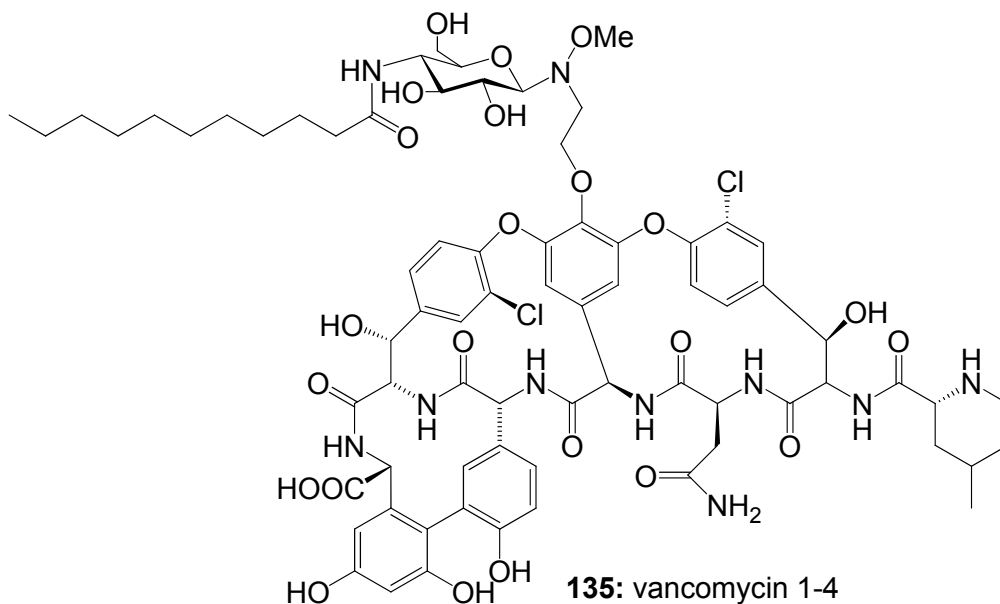


133: vancomycin 1-1
0/0

Structures of All Library members (con't)

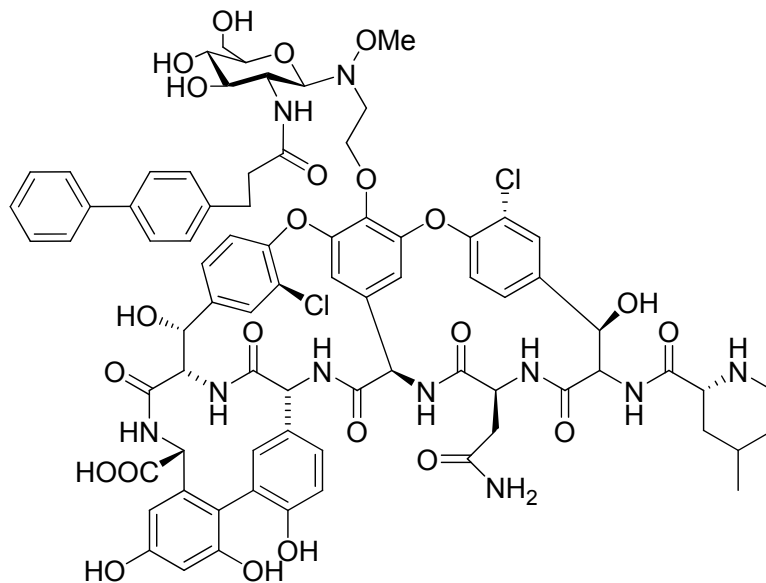


134: vancomycin 1-2
0/0

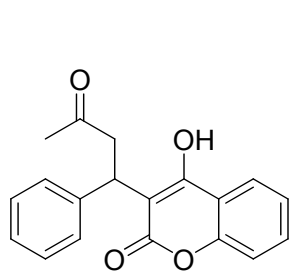


135: vancomycin 1-4
0/0

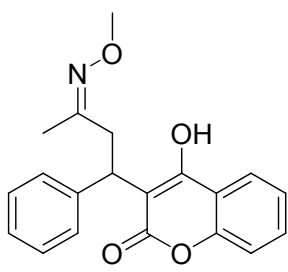
Structures of All Library members (con't)



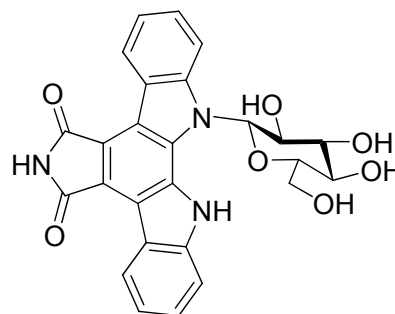
134: vancomycin 1-5
0/0



135: warfarin
0/0



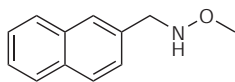
136: warfarin oxime
0/0



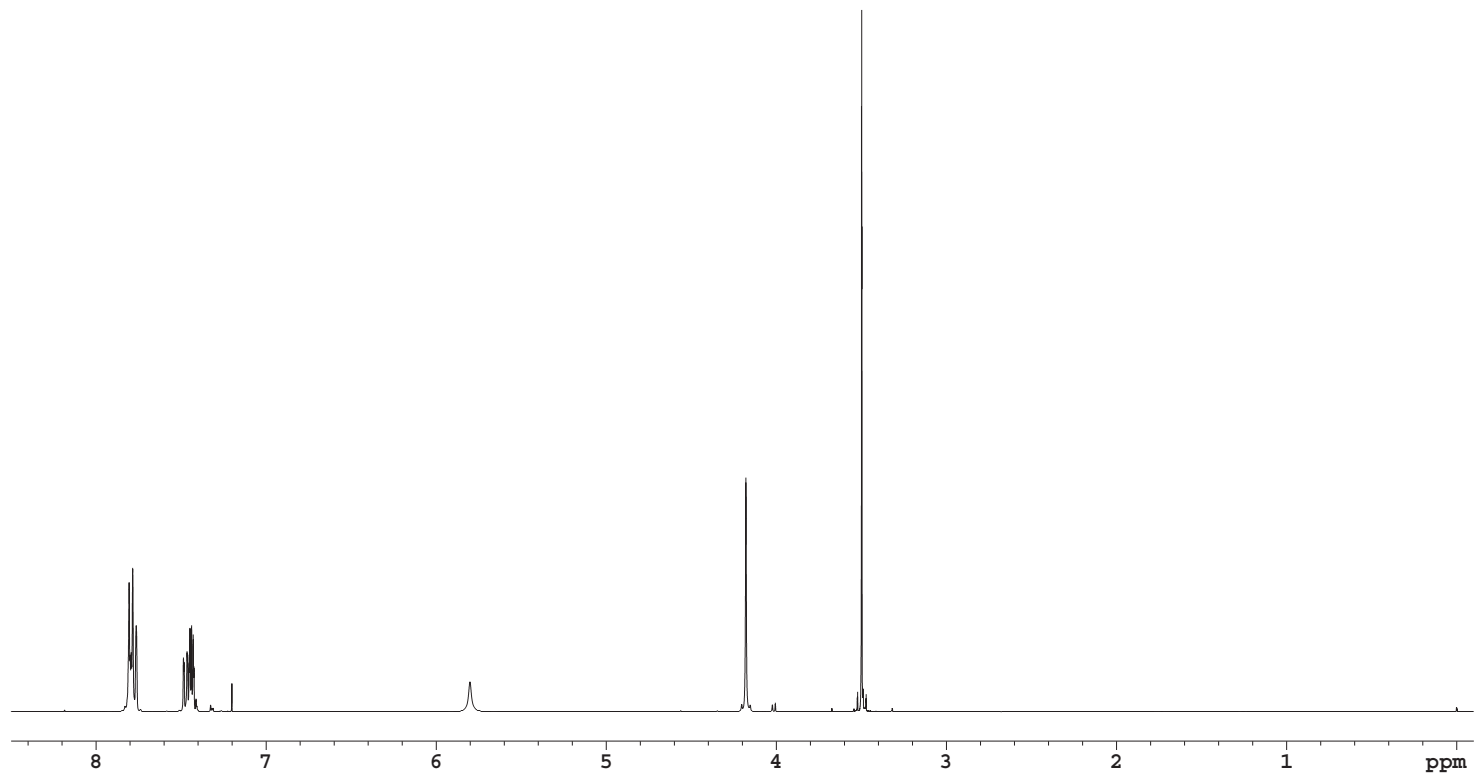
137: '131'
0/0

10. ^1H and ^{13}C NMR Spectra.

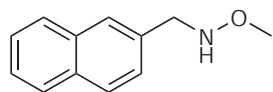
^1H , 400 MHz, CDCl_3



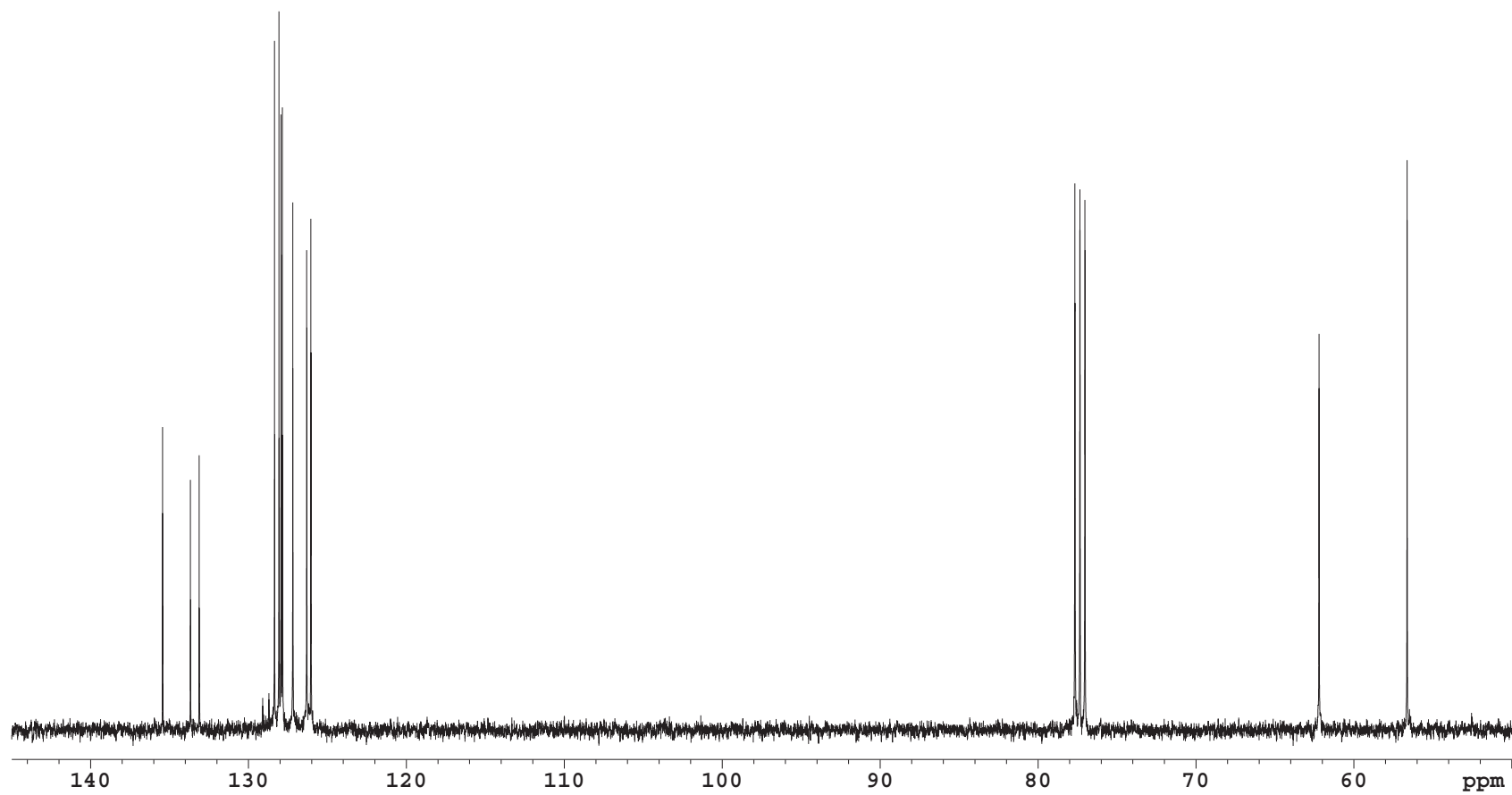
26



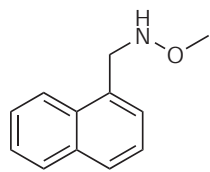
^{13}C , 100 MHz, CDCl_3



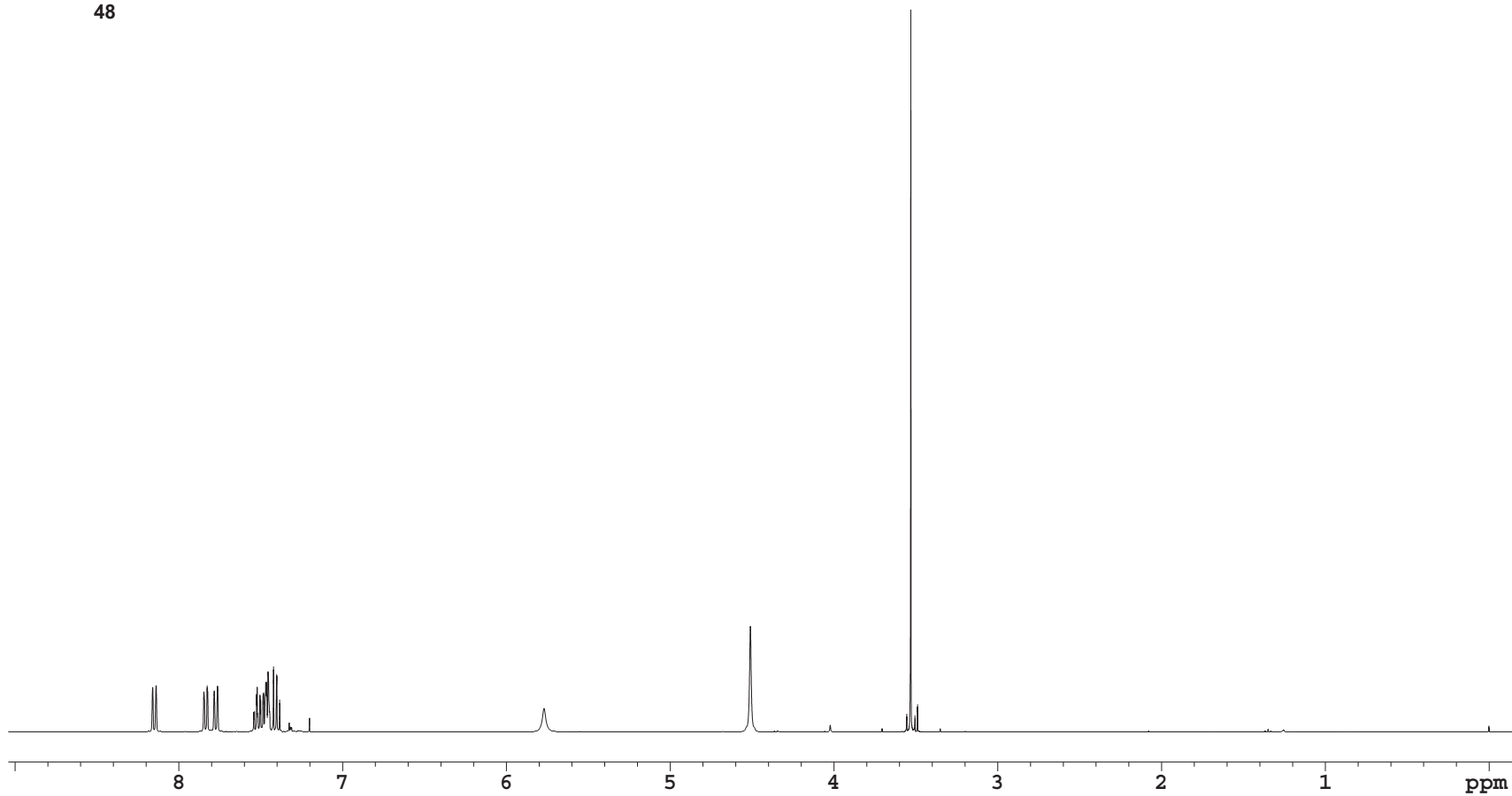
26



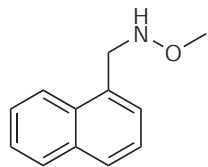
^1H , 400 MHz, CDCl_3



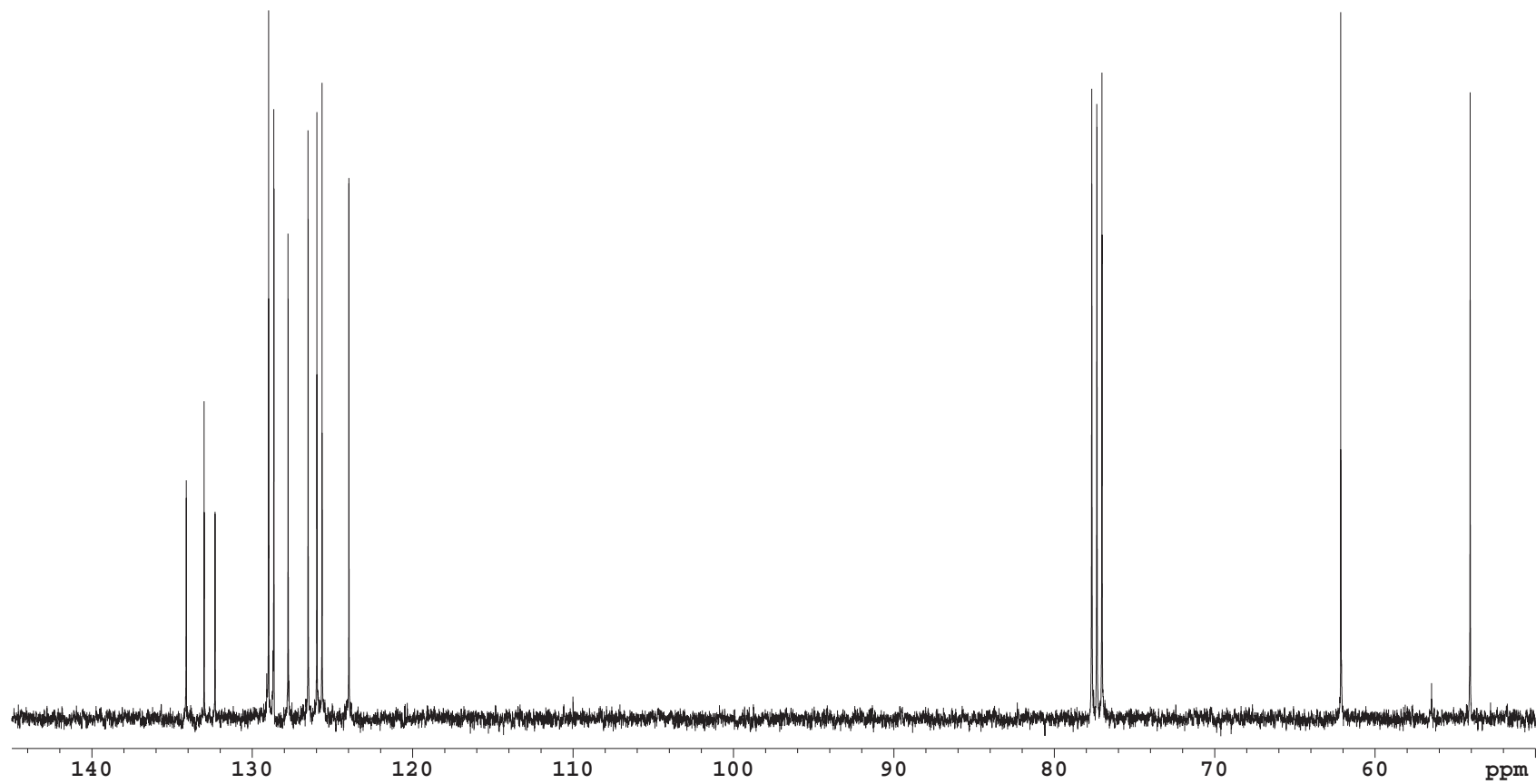
48



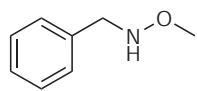
^{13}C , 100 MHz, CDCl_3



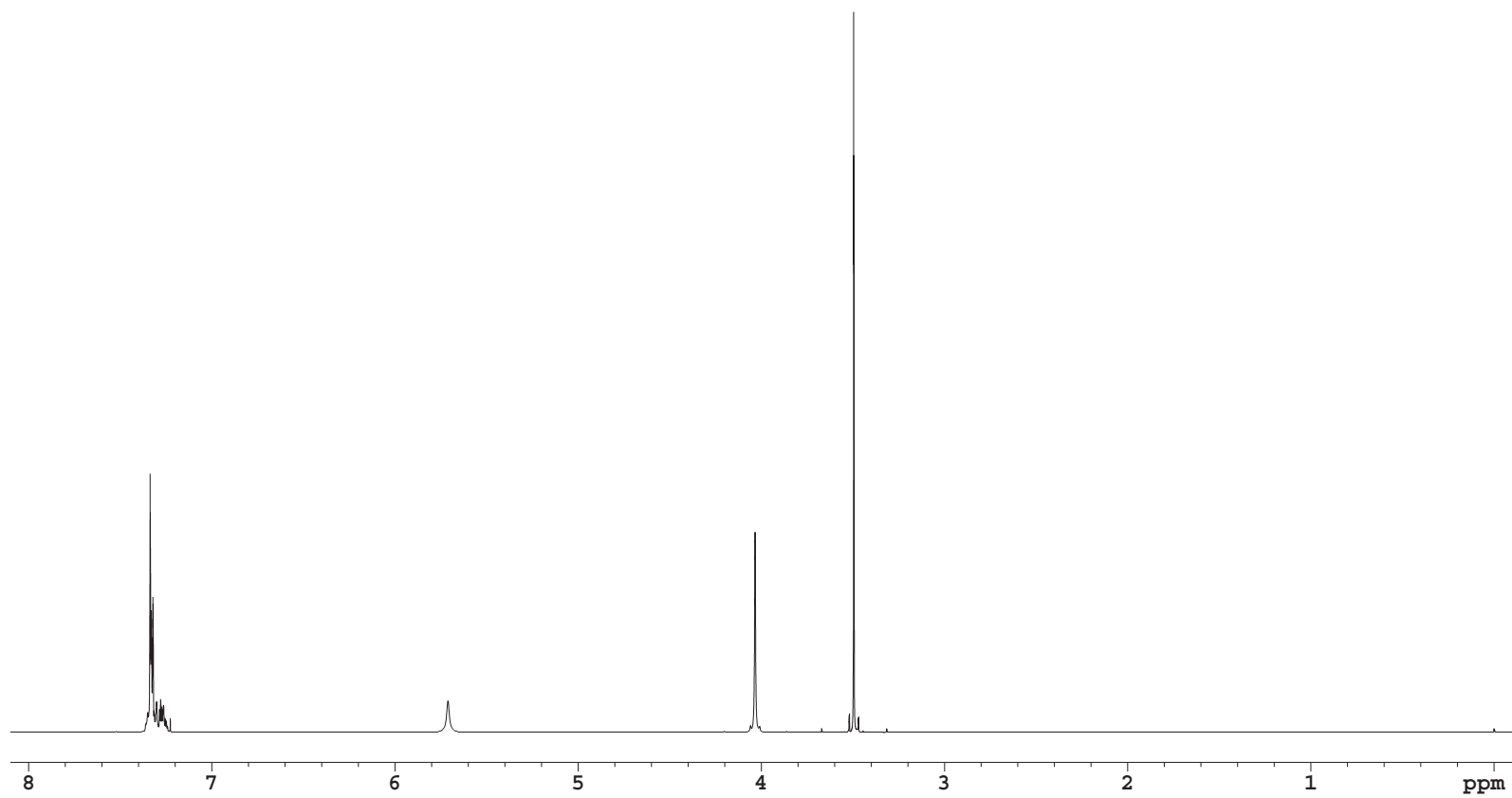
48



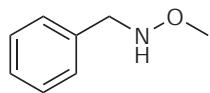
^1H , 400 MHz, CDCl_3



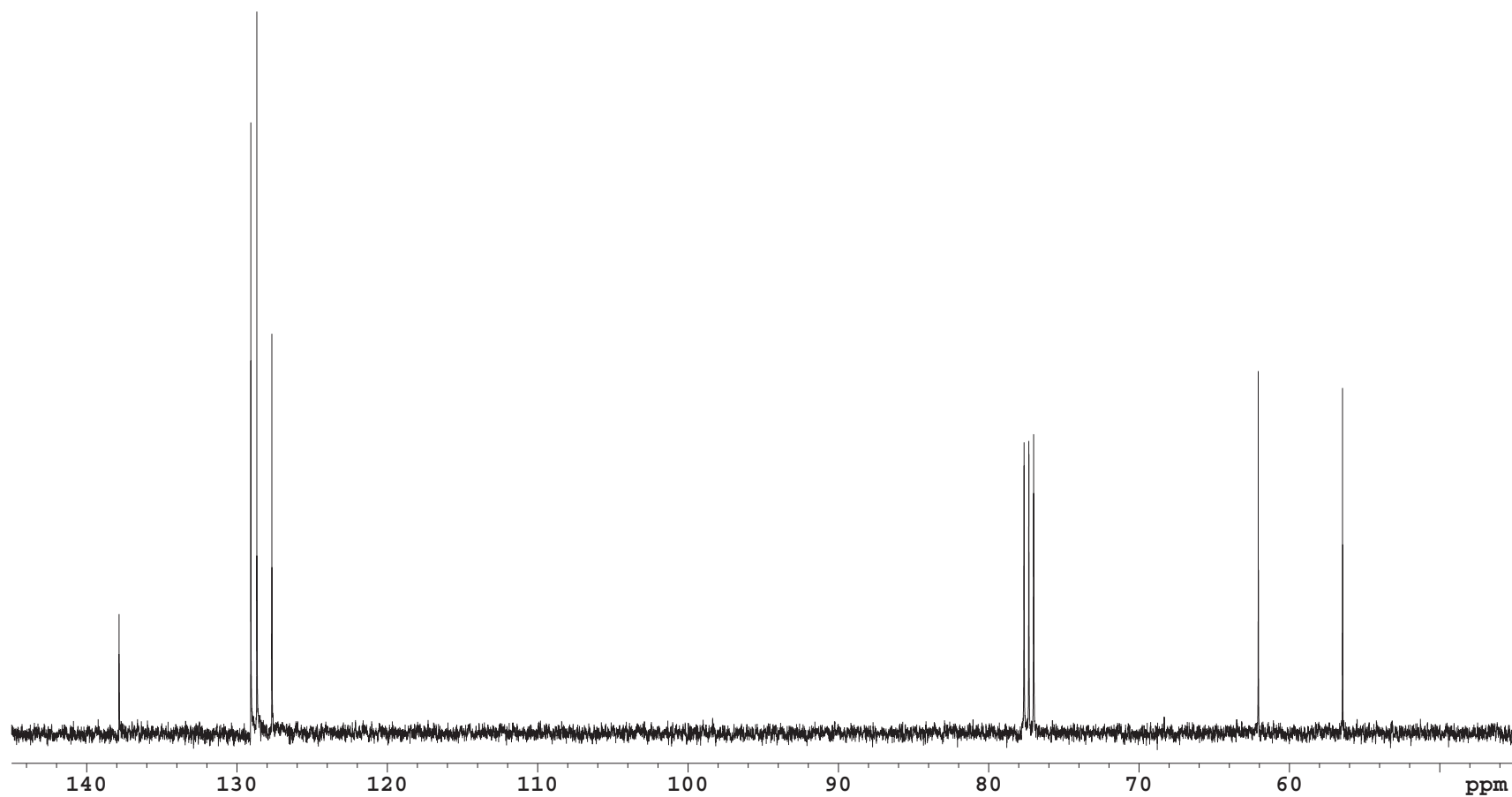
115



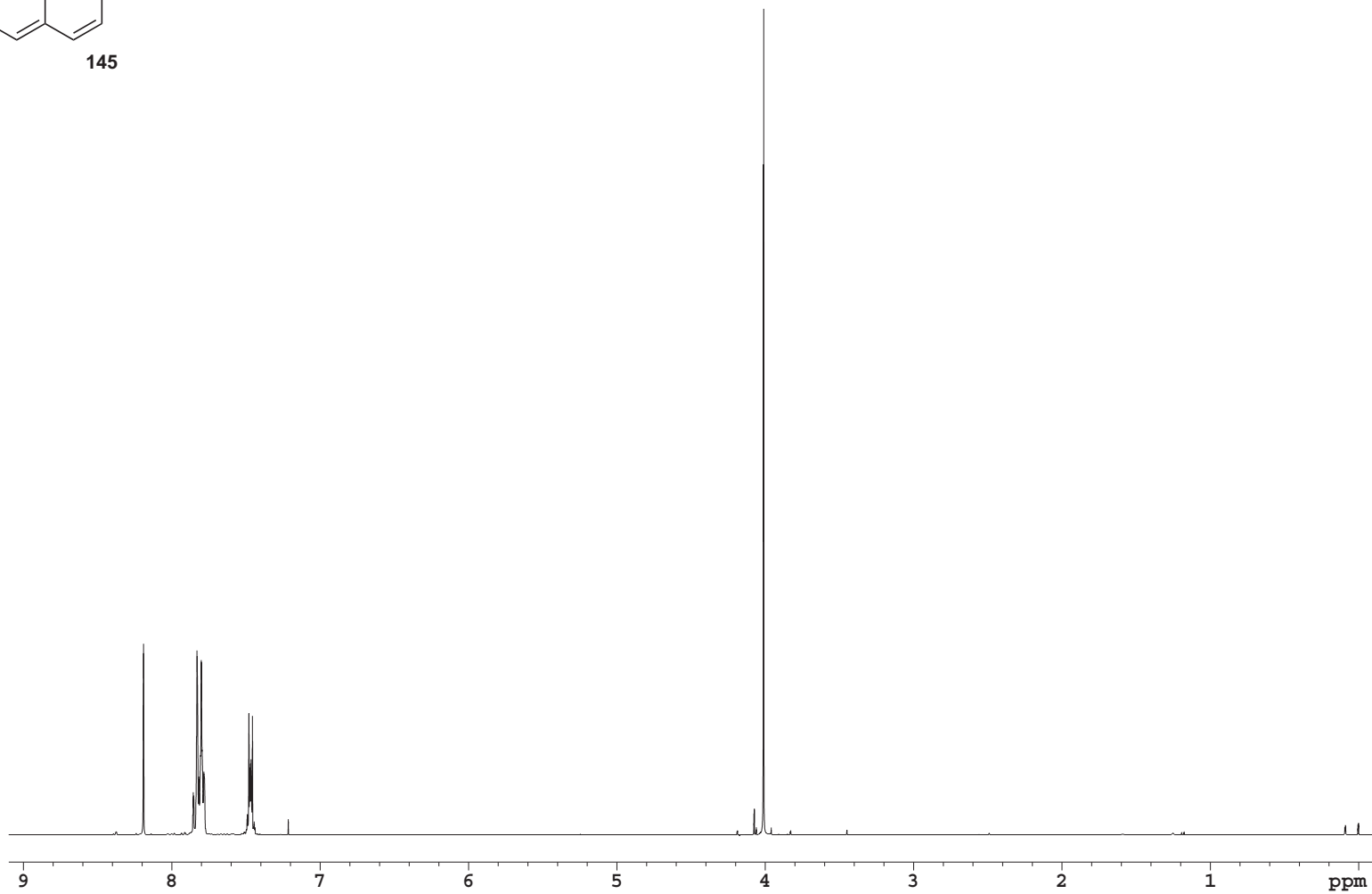
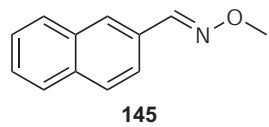
^{13}C , 100 MHz, CDCl_3



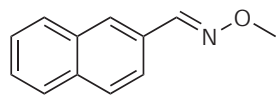
115



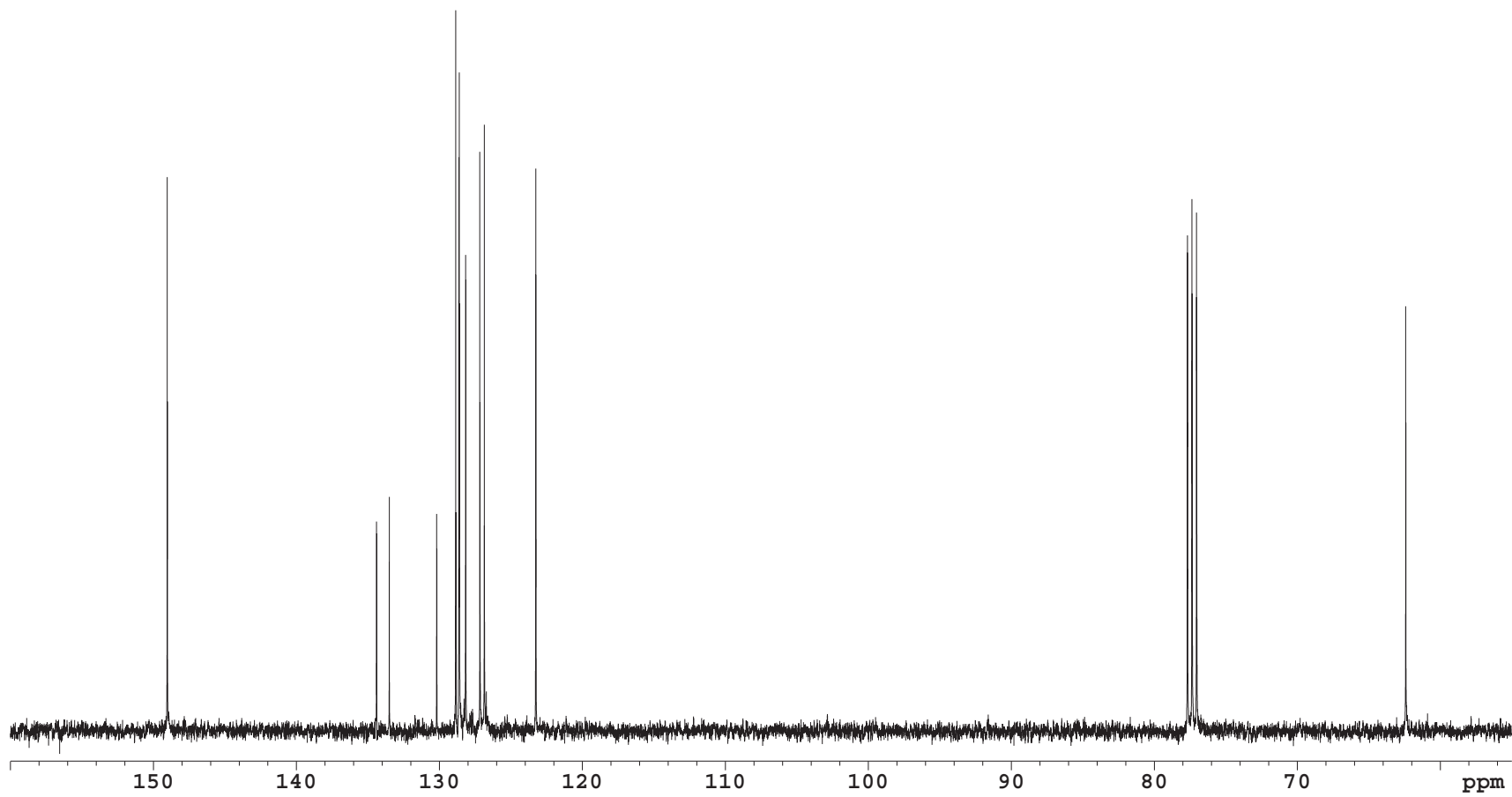
^1H , 400 MHz, CDCl_3



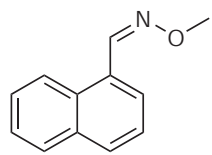
^{13}C , 100 MHz, CDCl_3



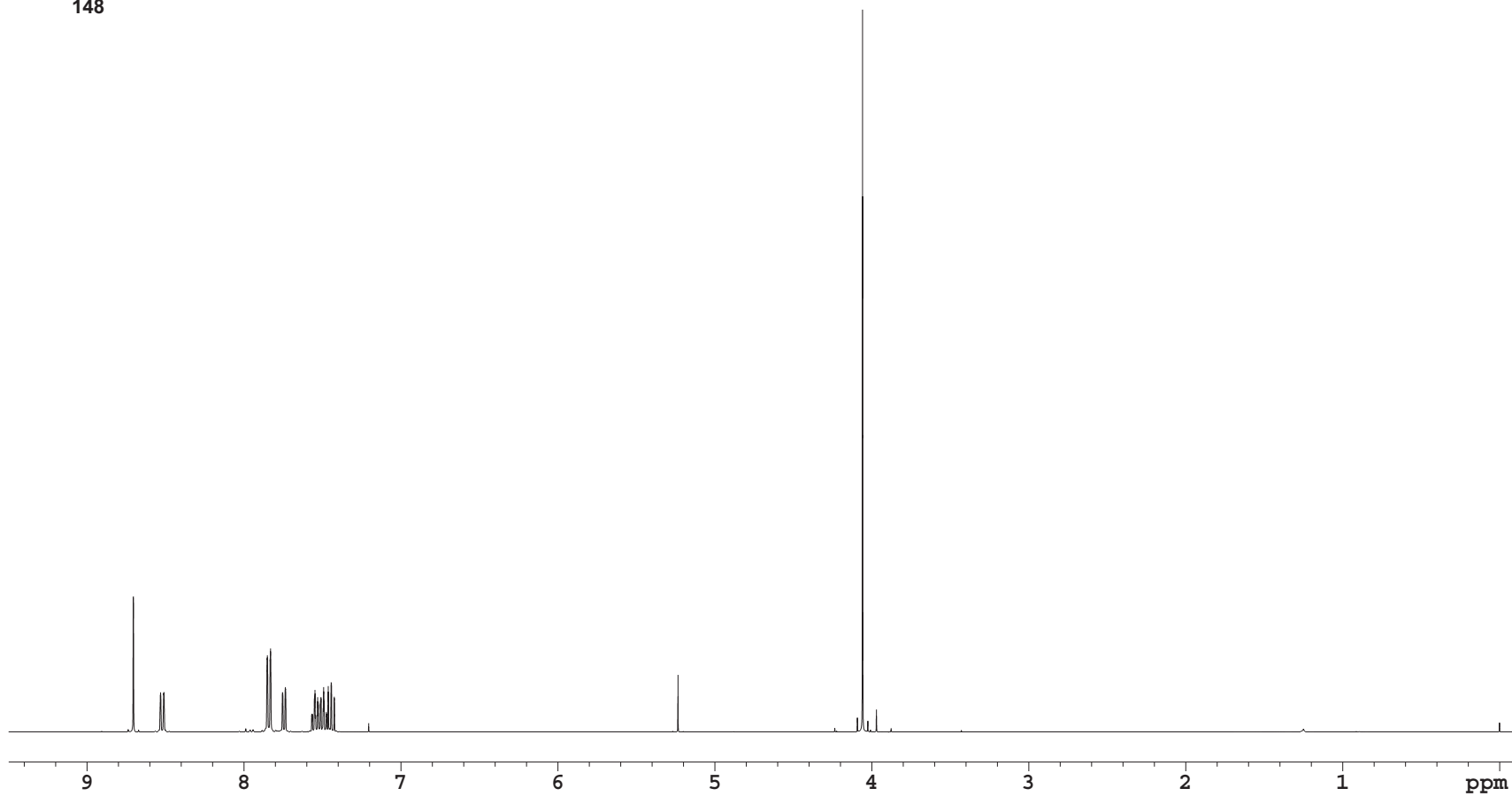
145



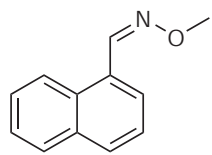
^1H , 400 MHz, CDCl_3



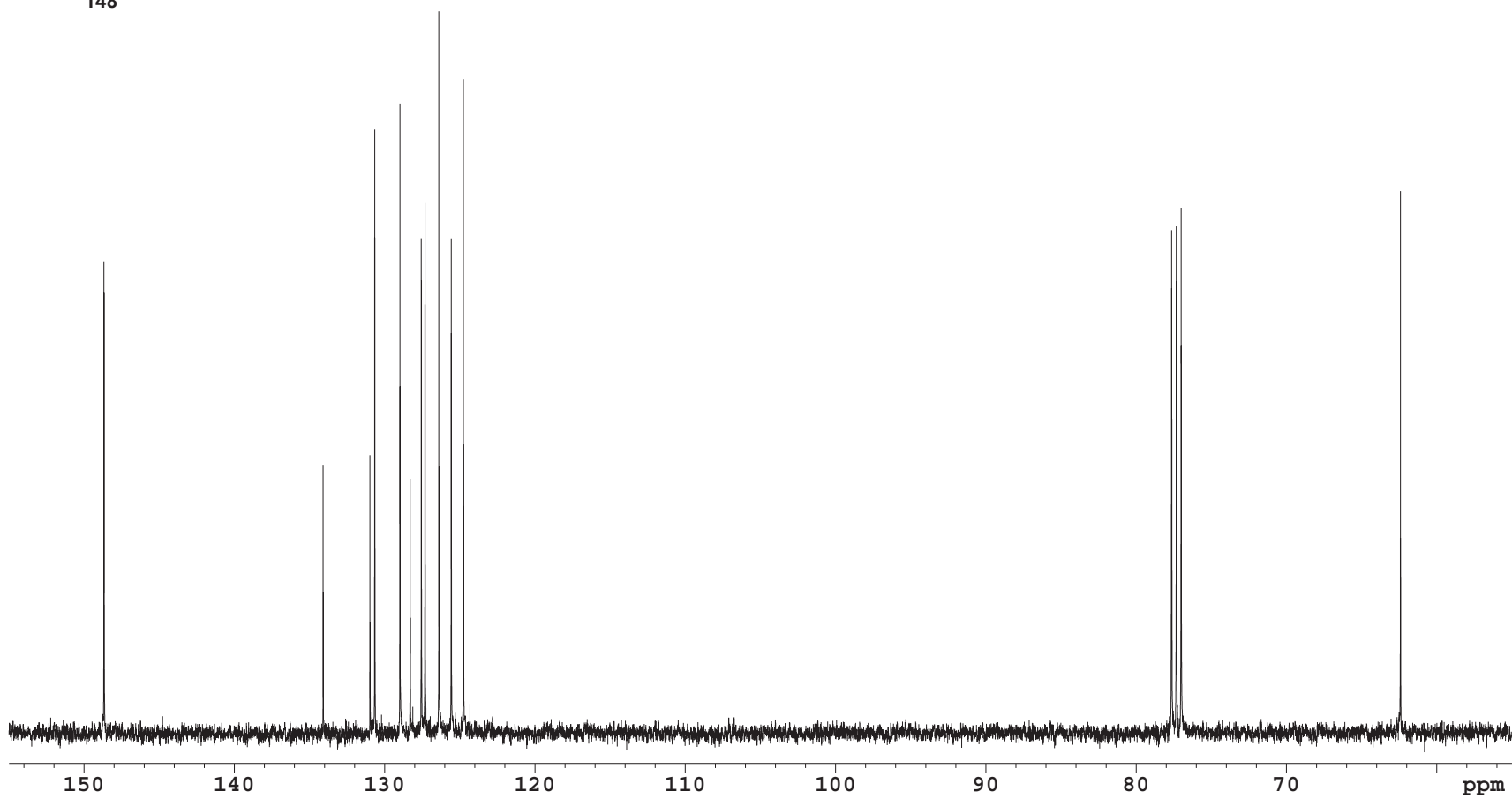
148



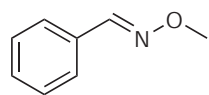
^{13}C , 100 MHz, CDCl_3



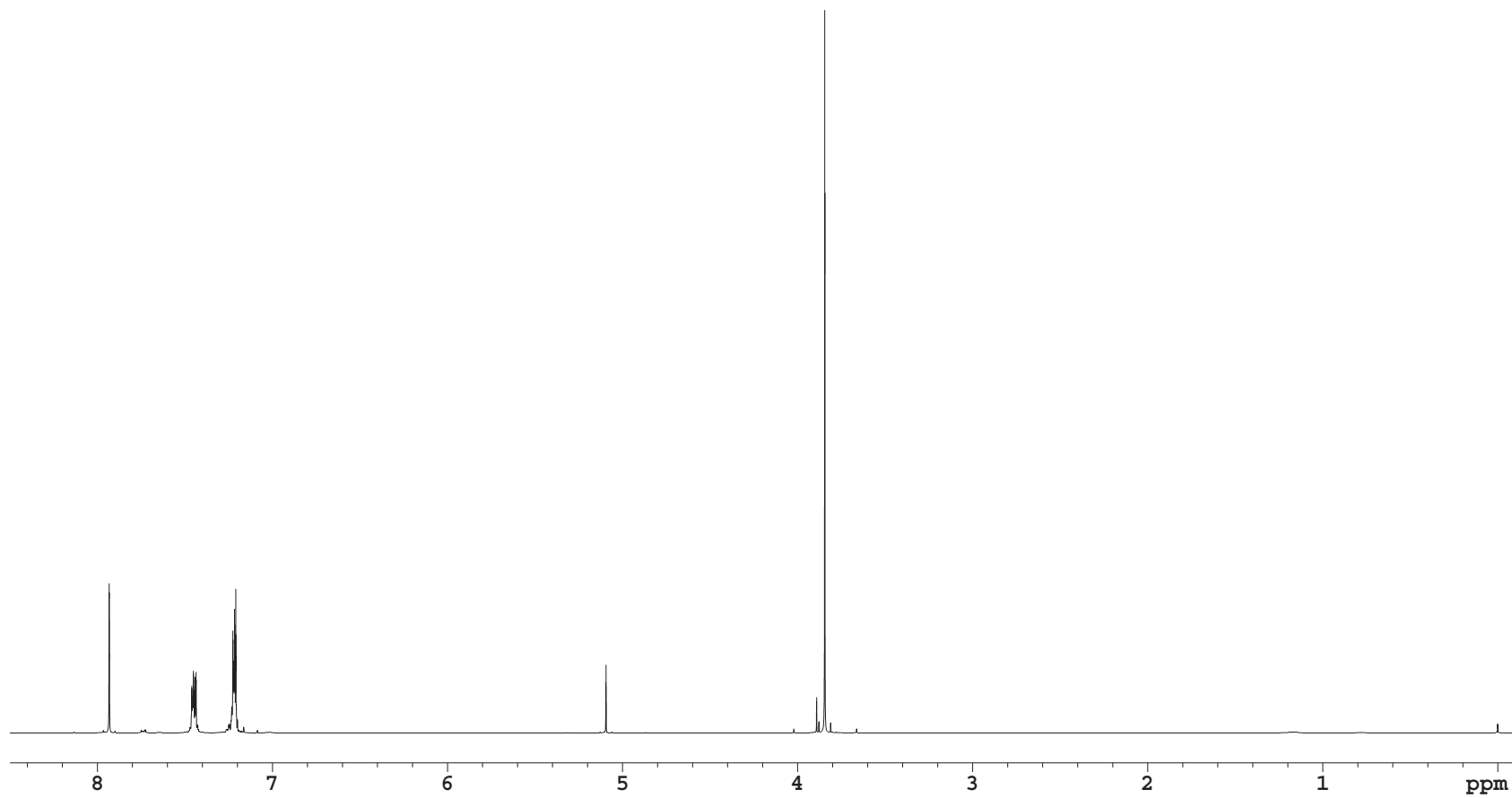
148



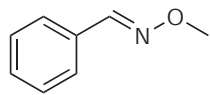
^1H , 400 MHz, CDCl_3



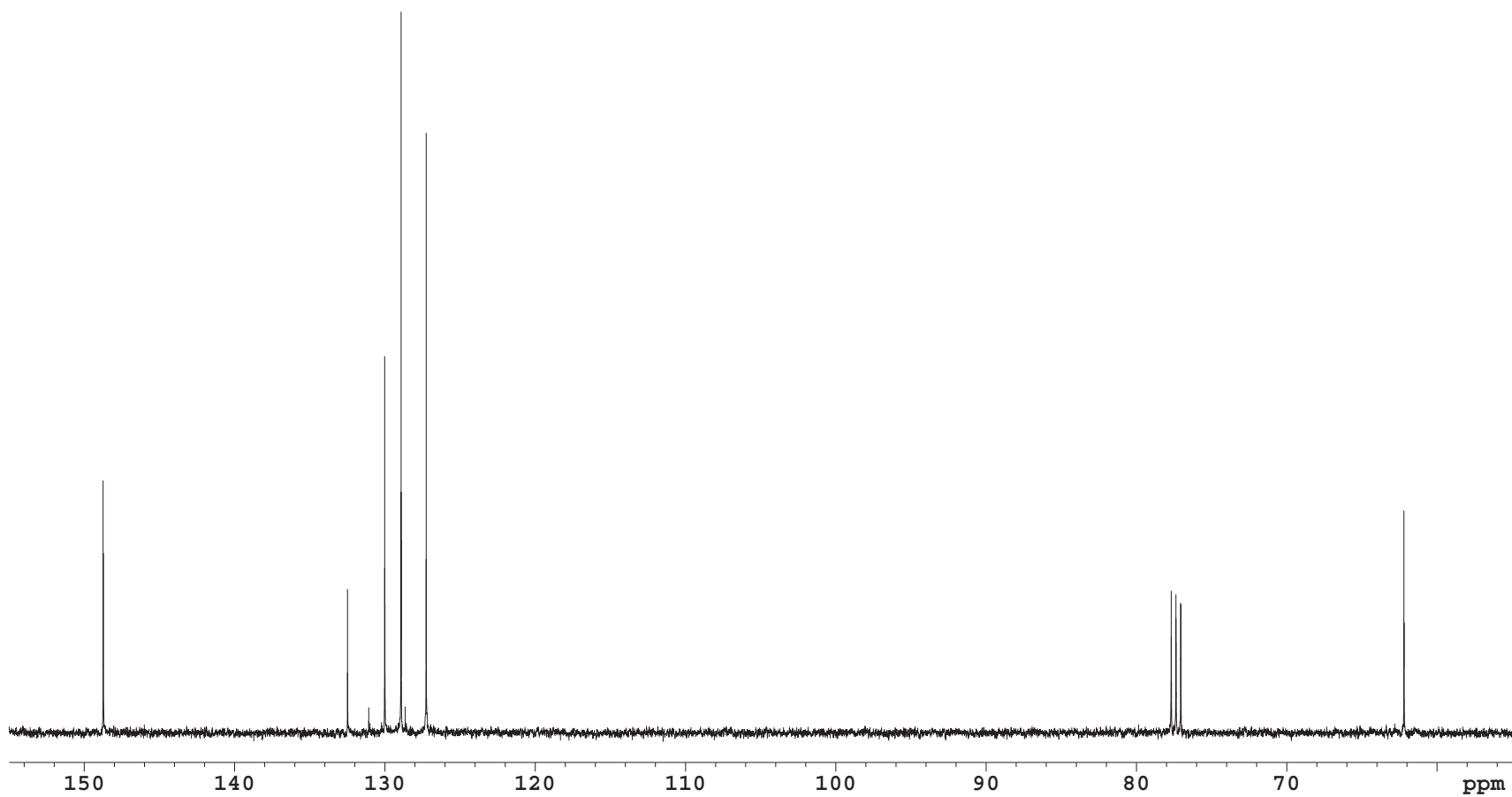
150



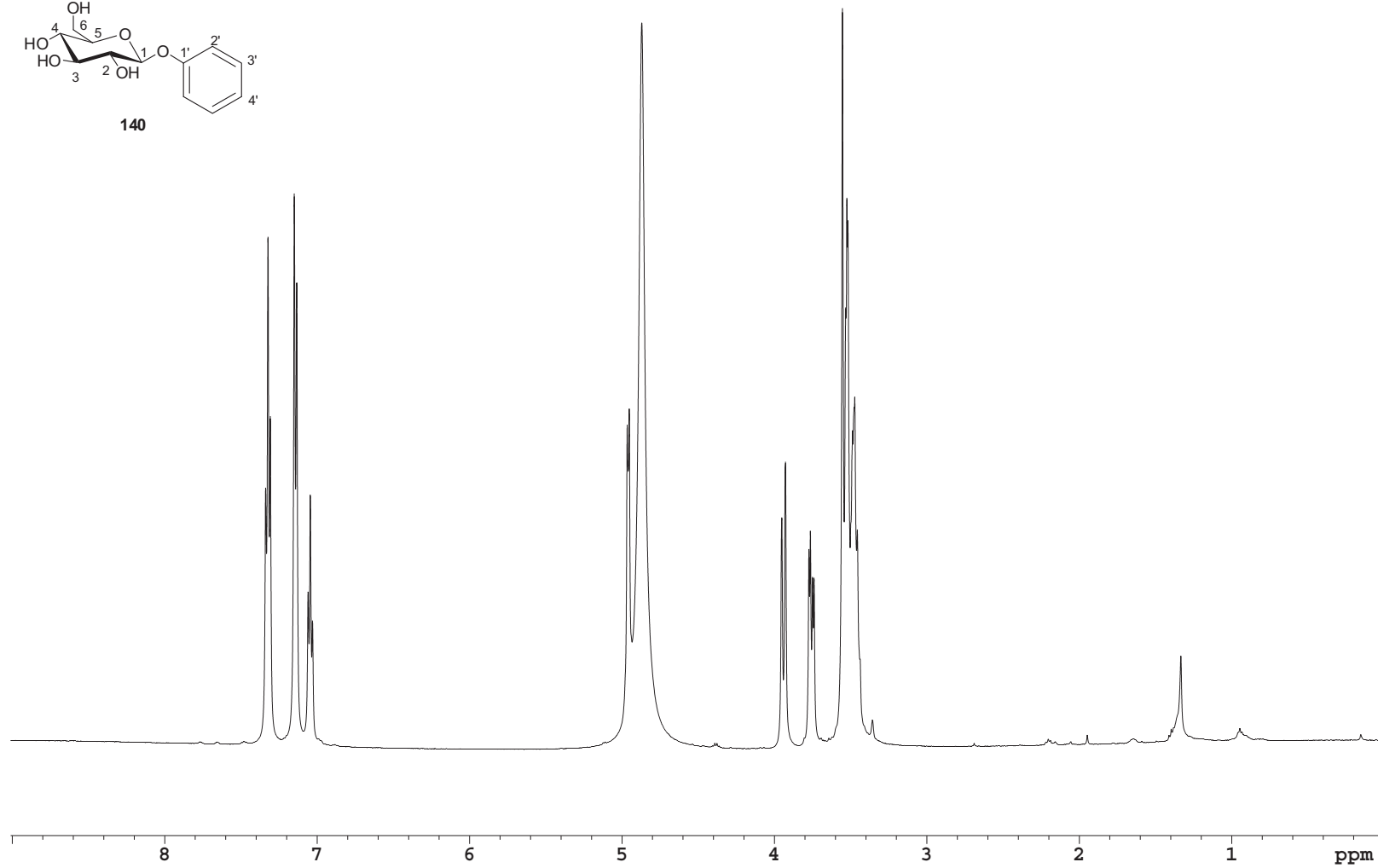
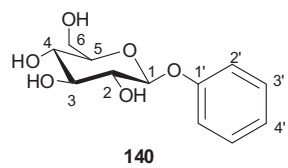
^{13}C , 100 MHz, CDCl_3



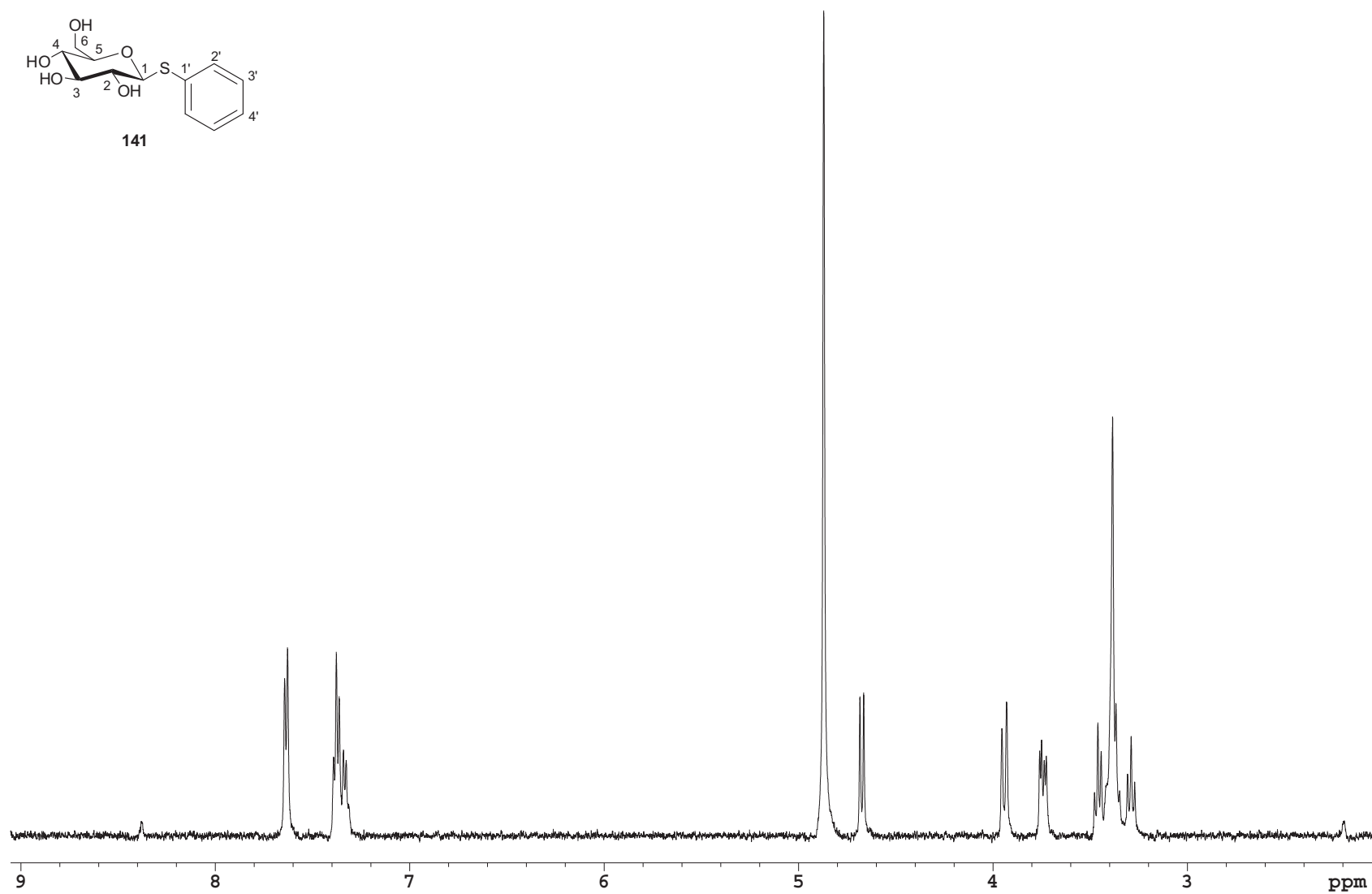
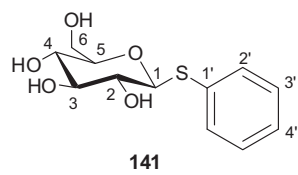
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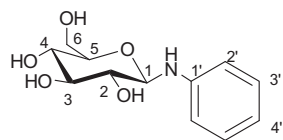
^1H , 500 MHz, CD_3OD



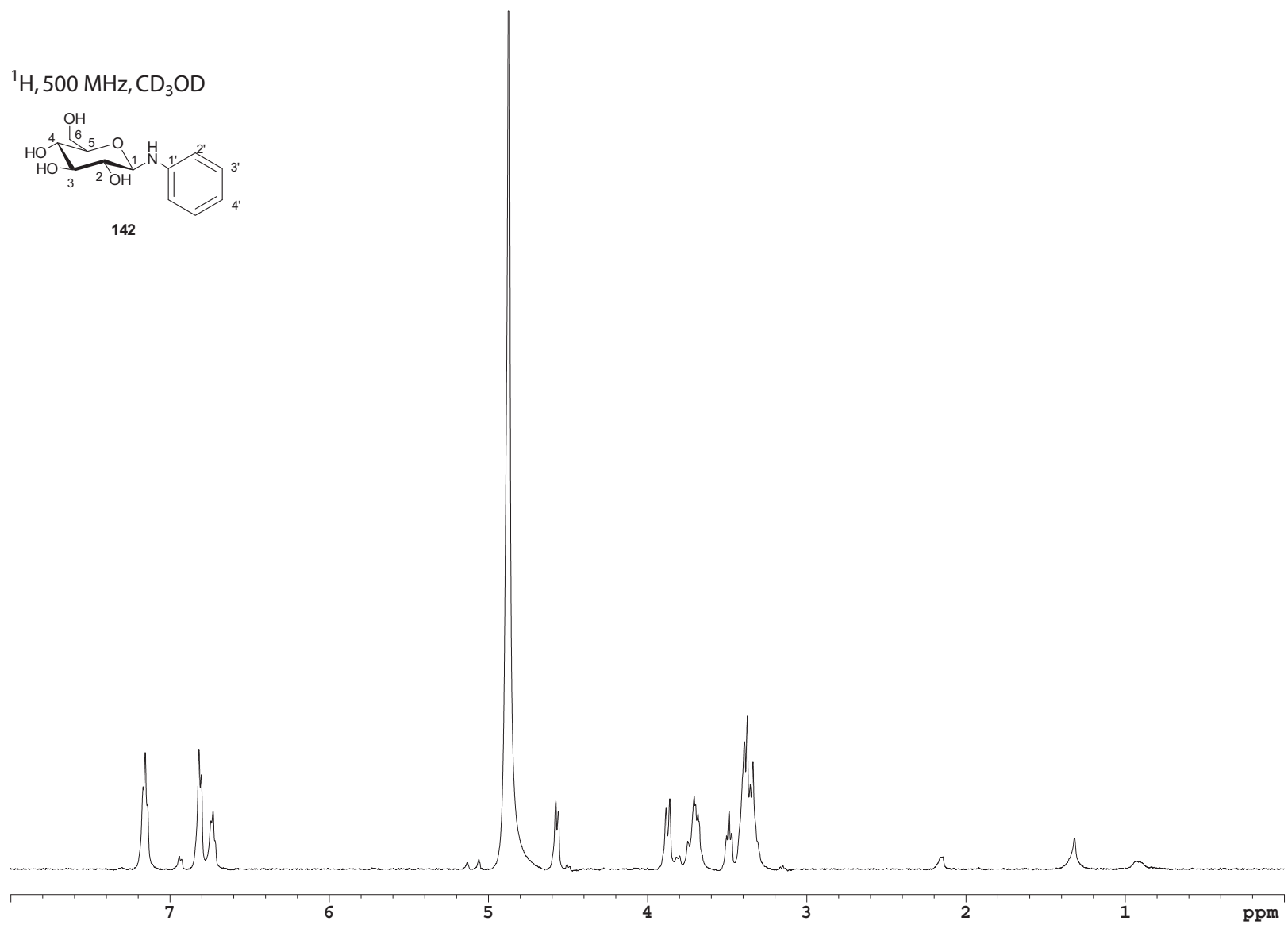
^1H , 500 MHz, CD_3OD



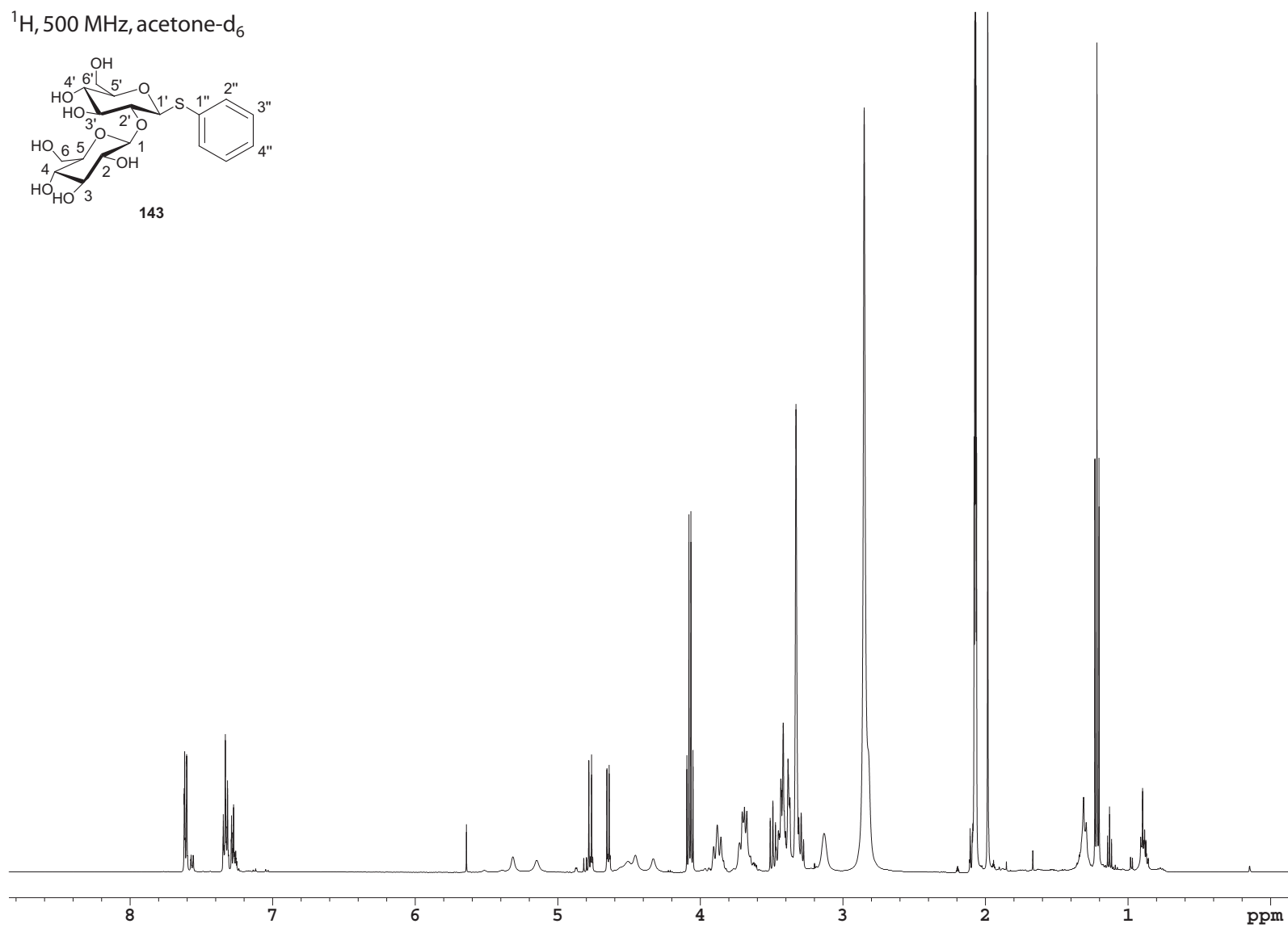
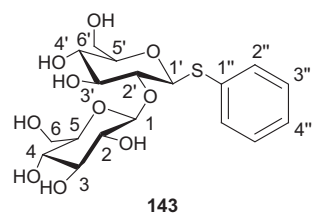
^1H , 500 MHz, CD_3OD



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^1H , 500 MHz, acetone- d_6



11. Supplementary references.

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