

Supporting Information for
Biosynthetic Tailoring of Microcin E492m: Post-Translational Modification
Affords an Antibacterial Siderophore-Peptide Conjugate

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Substrate Preparations.

Preparation of Ent and Glycosylated Ent. Enterobactin was synthesized as described in the literature.¹⁻³ IroB-catalyzed reactions were performed to obtain multi-milligram quantities of MGE and DGE for use in enzymatic assays.⁴ A 15 mL solution containing Ent (15.6 mg, 23 μ mol, dissolved in 500 μ L DMSO), UDP-Glc (124 mg, 1.64 mmol), 5 mM MgCl₂, 2.5 mM TCEP and 75 mM Tris-HCl buffered at pH 8 was incubated at room temperature for 3.5 h, quenched by addition of 7 mL of 2.5 N HCl in MeOH, vortexed and filtered through a 0.2 μ m membrane. Preparative HPLC using a gradient of 0 to 40% B (solvent A, 0.2% TFA/H₂O; solvent B, MeCN) over 40 min and a flow-rate of 10 mL/min afforded pure MGE (30.9 mg, 37.1 μ mol) and DGE (14.3 mg, 14.4 μ mol). Product purity was verified by analytical HPLC and identity by co-elution with known standards and MS. LRMS [M+H]⁺ *m/z* calc 832.2 (MGE), 994.3 (DGE); found, 832.1, 994.2.

Large-Scale Enzymatic Preparation of lin-Ent. Ent (20 mg, 30 μ mol) was dissolved in 400 μ L of DMSO and diluted with 15 mL of 75 mM Hepes pH 7.5 buffer. A portion (135 μ L of 71 μ M stock) of MceD was added and the solution was mixed and incubated at room temperature for 27 min, and quenched with 5 mL of 2.5 N HCl in MeOH. The mixture was vortexed and filtered through a 0.2 μ m membrane. Preparative HPLC using a solvent gradient of 15 to 60% B (solvent A, 0.1% TFA/H₂O; solvent B, MeCN) allowed for separation of the reaction components. The compounds were collected and lyophilized to dryness, which afforded white powders. The identity of lin-Ent was verified by MS and co-elution with lin-Ent prepared from a IroD-catalyzed reaction.⁵ LRMS: [M+H]⁺ *m/z* calc, 688.2; found, 688.3.

Preparation of Ferric Ent, MGE and DGE. The ferric complexes of Ent, MGE and DGE were prepared from FeCl₃ and siderophore in the presence of

base by modification of existing protocols.^{5,6} For $[\text{Fe}(\text{Ent})]^{3+}$: A 50.3-mL portion of Ent (10.5 mM in DMSO) was diluted with 49.7 μL of MeOH (5.28 mM final concentration) and aqueous FeCl_3 (5.28 mM, 110 μL) was added. The solution turned purple and was mixed with a pipette. A 7-mL aliquot of 1 M K_2CO_3 was added and the solution was mixed and turned wine-colored. The reaction was loaded onto a Sephadex LH-20 column packed with MeOH and eluted with MeOH. The $[\text{Fe}(\text{Ent})]^{3+}$ containing fractions were concentrated to dryness and the resulting dark-red residue was dissolved in 150 μL of DMSO. The product identity was verified and stock concentration determined by UV-vis. For $[\text{Fe}(\text{MGE})]^{3+}$: As described for $[\text{Fe}(\text{Ent})]^{3+}$ except that the 5.28 mM MGE solution was prepared in 1:1 MeOH/ H_2O . For $[\text{Fe}(\text{DGE})]^{3+}$: As described for $[\text{Fe}(\text{Ent})]^{3+}$ except that the DGE solution was prepared in MeOH (1.33 mM) and the FeCl_3 solution was also 1.33 mM.

References

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- (3) Deleuze, H.; Maillard, B. *J. Organomet. Chem.* **1995**, *490*, C14-C17.
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Table S1. Mass Data (m/z) for Hydrolysis Products from MceD-Catalyzed Reactions

Substrate	Product	Calc [M+H] ⁺	Found [M+H] ⁺	Calc [M-H] ⁻	Found [M-H] ⁻
Ent		670.2	670.2		
	lin-Ent	688.2	688.3		
	DHB-Ser Dimer	465.1	465.1		
	DHB-Ser Monomer			240.1	n.d. ^a
MGE		832.2	832.1		
	lin-MGE	850.2	850.1		
	Glc-Dimer	627.2	627.1		
	Glc-Monomer	404.1	n.d.		
	DHB-Ser			240.1	n.d.
DGE		994.3	994.1		
	lin-DGE	1012.3	1012.2		
	Glc-Dimer	627.2	627.3		
	Glc-Monomer	404.1	n.d.		
	DHB-Ser Monomer			240.1	n.d.

^a n.d. = not determined

Table S2. NOE Correlations for C₁₀-C6'-MGE (left) and C₁₀-C4'-MGE (right)

Group	Atom	Nuc	Shift	Group	Atom	Nuc	Shift
C6				C4			
Glc	H1	1H	3.987	Glc	H1	1H	4.077
Glc	H2	1H	3.462	Glc	H2	1H	3.474
Glc	H3	1H	3.575	Glc	H3	1H	3.590
Glc	H4	1H	3.715	Glc	H4	1H	4.960
Glc	H5	1H	3.510	Glc	H5	1H	3.545
Glc	H61	1H	4.454	Glc	Q6	1H	3.983
Glc	H62	1H	4.244	DHB1	H4	1H	7.033
DHB1	H4	1H	7.021	DHB1	H6	1H	7.184
DHB1	H6	1H	7.156	DHB1	H9	1H	4.999
DHB1	H9	1H	4.957	DHB1	HN	1H	9.027
DHB1	HN	1H	8.945	DHB2	H4	1H	6.933
DHB2	H4	1H	6.912	DHB2	H5	1H	6.647
DHB2	H5	1H	6.612	DHB2	H6	1H	7.000
DHB2	H6	1H	6.970	DHB2	H9	1H	4.890
DHB2	H9	1H	4.898	DHB2	HN	1H	8.805
DHB2	HN	1H	8.692	DHB3	H4	1H	6.933
DHB3	H4	1H	6.908	DHB3	H5	1H	6.645
DHB3	H5	1H	6.610	DHB3	H6	1H	7.002
DHB3	H6	1H	6.956	DHB3	H9	1H	4.915
DHB3	H9	1H	5.041	DHB3	HN	1H	8.778
DHB3	HN	1H	8.733	S1	HA	1H	4.117
S1	HA	1H	4.114	S1	HN	1H	8.410
S1	HN	1H	8.366	S1	QB	1H	3.953
S1	QB	1H	3.853	A2	HA	1H	4.390
A2	HA	1H	4.404	A2	HN	1H	8.713
A2	HN	1H	8.709	A2	QB	1H	1.371
A2	QB	1H	1.369	T3	HA	1H	4.324
T3	HA	1H	4.196	T3	HB	1H	4.188
T3	HB	1H	4.181	T3	QG2	1H	1.142
T3	QG2	1H	1.141	T3	HN	1H	8.154
T3	HN	1H	8.150	S4	HA	1H	4.632
S4	HA	1H	4.317	S4	HN	1H	8.238
S4	HN	1H	8.168	S4	QB	1H	3.968
S4	QB	1H	3.917	S5	HA	1H	4.452
S5	HN	1H	7.796	S5	HN	1H	8.280
S5	QB	1H	3.781	S5	QB	1H	3.804
S6	HN	1H	8.189	S6	HA	1H	4.463
S6	QB	1H	3.938	S6	HB2	1H	3.858
G7	HN	1H	8.324	S6	HB3	1H	3.790
G7	QA	1H	4.373	S6	HN	1H	8.375
S8	HA	1H	4.489	G7	HN	1H	7.132
S8	HN	1H	8.283	S8	HA	1H	4.420
S8	QB	1H	3.960	S8	HN	1H	8.313
G9	HN	1H	8.265	S8	QB	1H	3.825
G9	QA	1H	4.428	G9	HN	1H	8.203
S10	HA	1H	4.470	G9	QA	1H	4.405
S10	HN	1H	7.072	S10	HA	1H	4.001
S10	QB	1H	3.949	S10	HN	1H	7.067
				S10	QB	1H	3.496

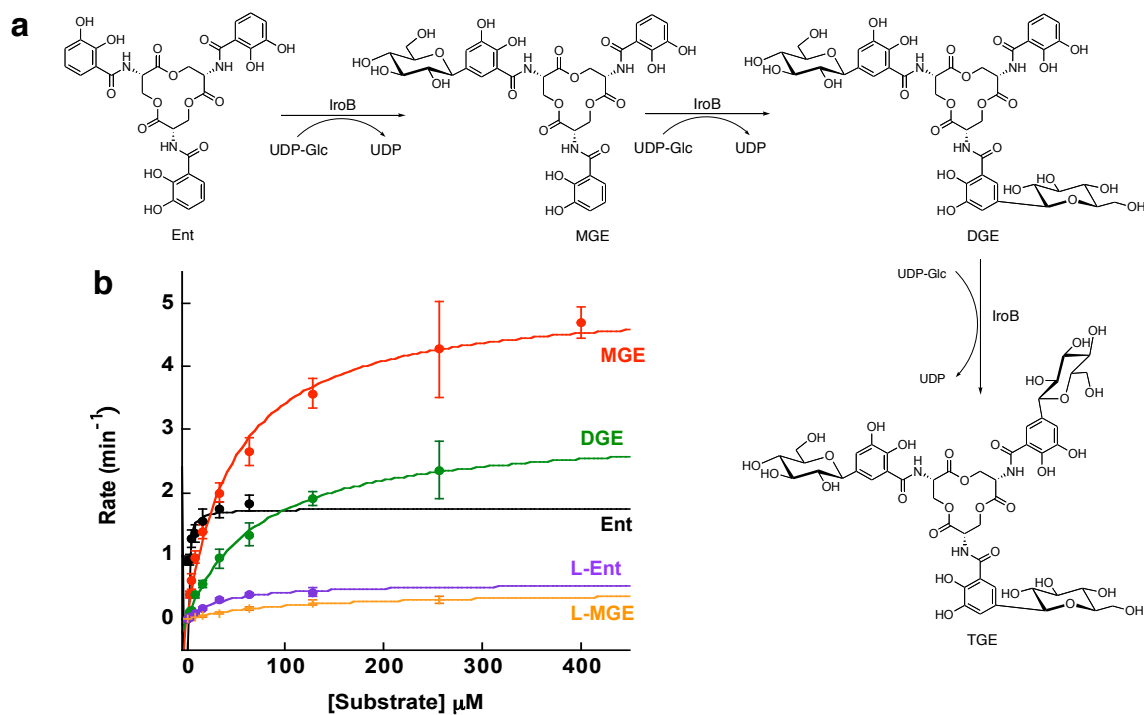


Figure S1. C-Glycosylation of Ent by IroB to give MGE, DGE and TGE. (b) Kinetic data for IroB-catalyzed C-glycosylation. Corresponding k_{cat} and K_m values are given in Table 1 of the main text.

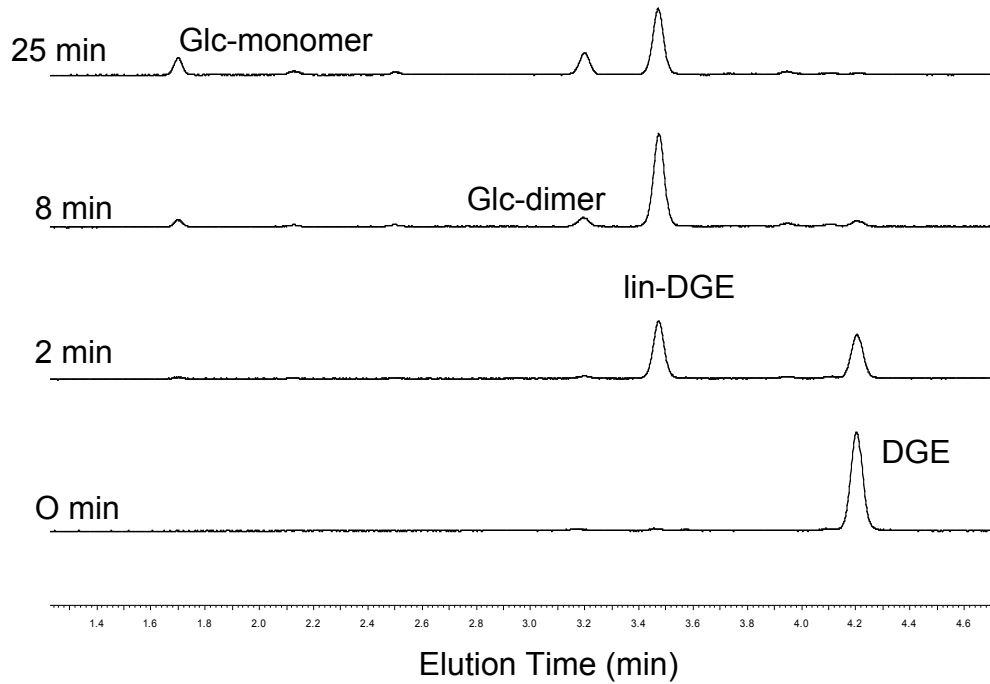


Figure S2. HPLC analysis of the MceD-catalyzed hydrolysis of DGE (75 mM HEPES, pH 7.5). The concentration of DGE was 32 μ M and 20 nM MceD was employed. MS values for the products are listed in Table S1.

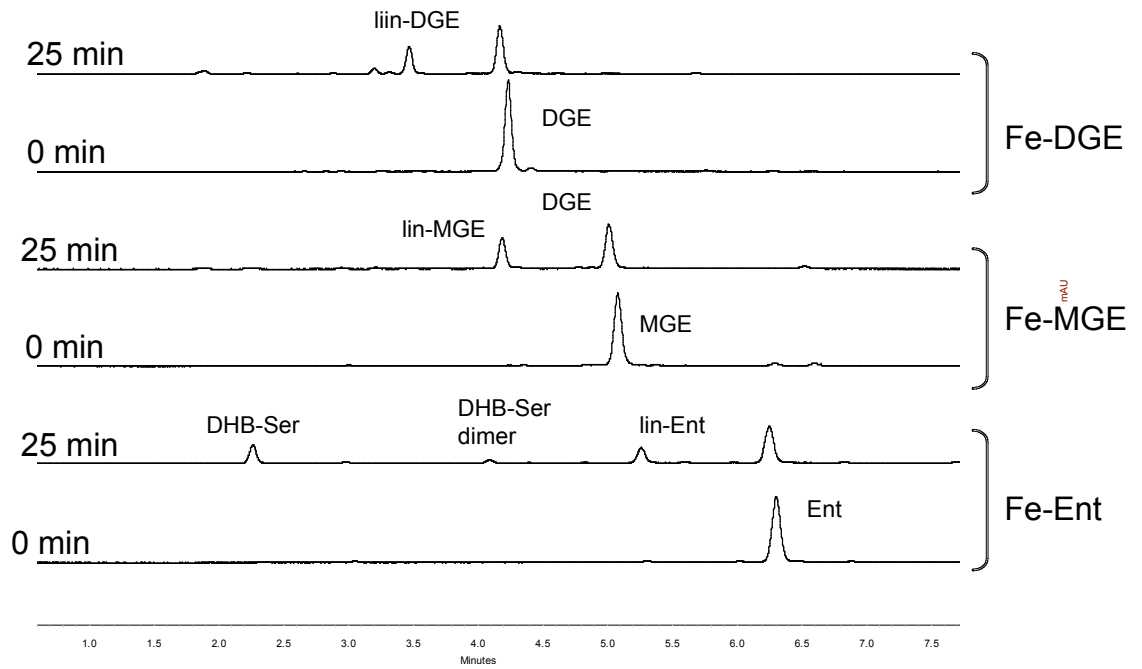


Figure S3. MceD-catalyzed hydrolysis of $[\text{Fe}(\text{Ent})]^{3-}$, $[\text{Fe}(\text{MGE})]^{3-}$ and $[\text{Fe}(\text{DGE})]^{3-}$ (75 mM HEPES, pH 7.5). The MceD concentration was 20 nM and the substrate concentrations were 32 μM .

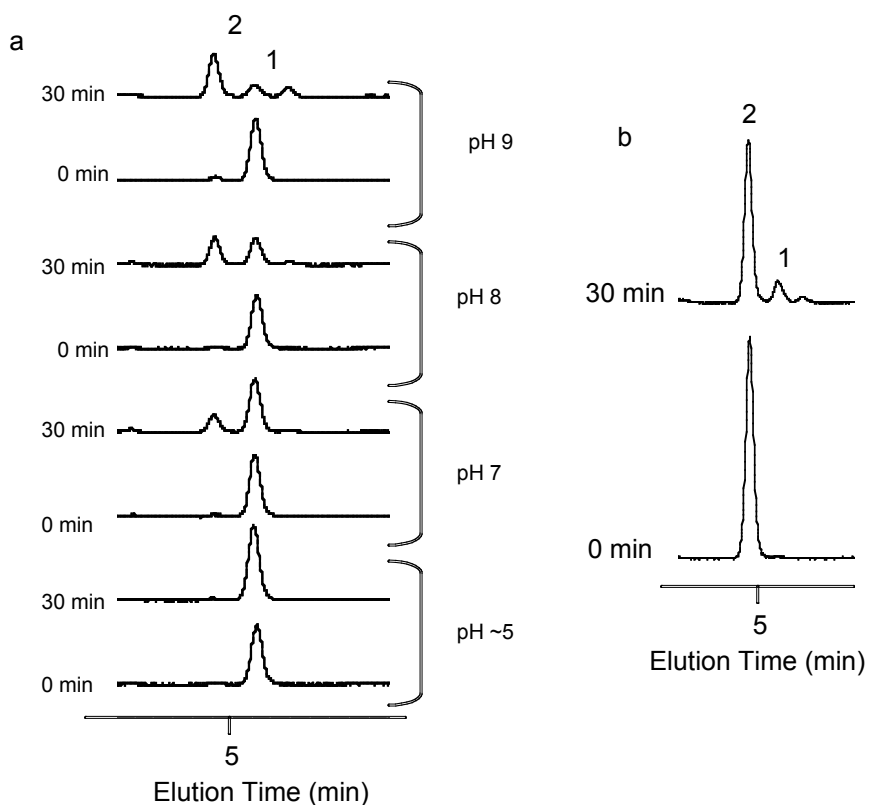


Figure S4. (a) Effect of pH on the migration of C_{10} from the $C4'$ to the $C6'$ position of the glucose moiety of MGE. Peak 1 is C_{10} - $C4'$ -MGE. Peak 2 is C_{10} - $C6'$ -MGE. Solutions of $50 \mu\text{M}$ C_{10} - $C6'$ -MGE were prepared and incubated at room temperature for 30 min in aqueous solution at pH ~5 (H_2O), pH 7 (75 mM Tris-HCl), pH 8 (75 mM Tris-HCl) and pH 9 (10 mM CHES). Migration to the $C6'$ position is accelerated by the presence of base. Some background hydrolysis of the ester bond between C_{10} and MGE is observed at pH 9. (b) Incubation of $50 \mu\text{M}$ C_{10} - $C6'$ -MGE at room temperature in pH 8 buffer (75 mM Tris-HCl). Some formation (~10%) of C_{10} - $C4'$ -MGE is observed during this timeframe.

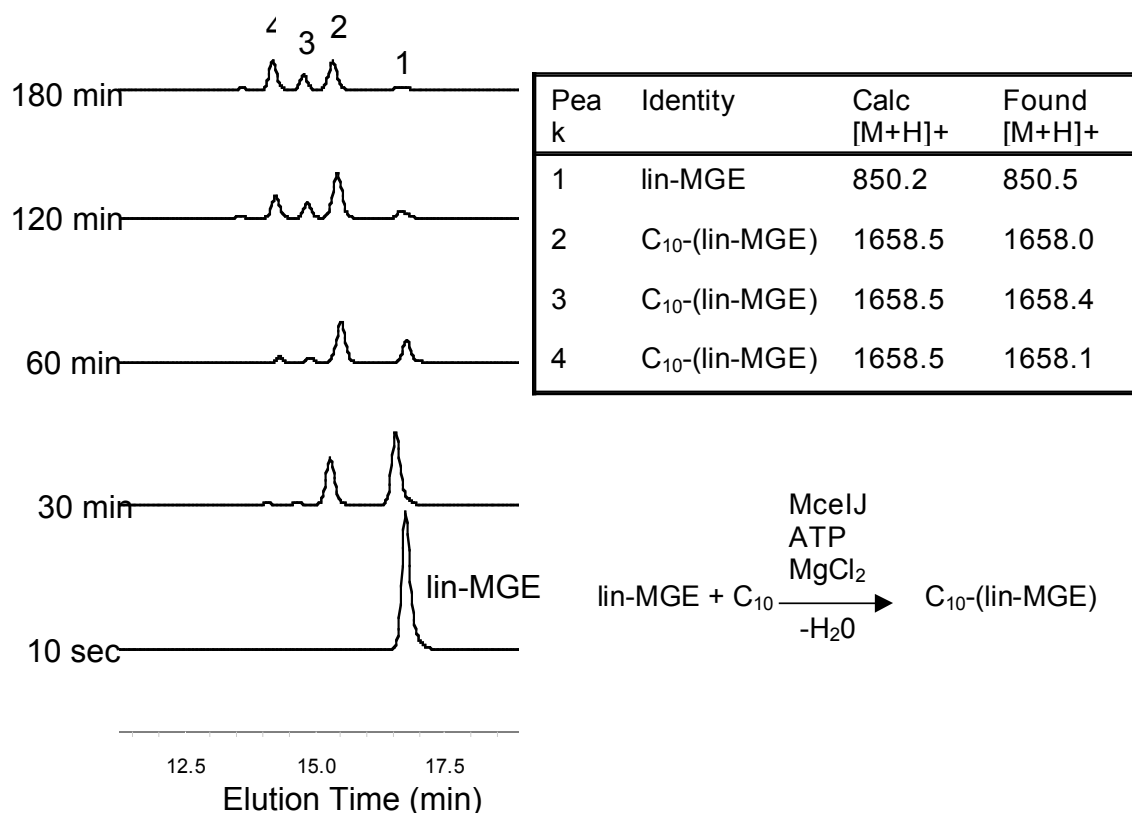


Figure S5. MceIJ-catalyzed attachment of lin-MGE to the C₁₀ peptide (2 μM MceIJ, 100 μM lin-MGE, 500 μM C₁₀, 5 mM ATP, 5 mM MgCl₂, 75 mM Tris-HCl pH 8).

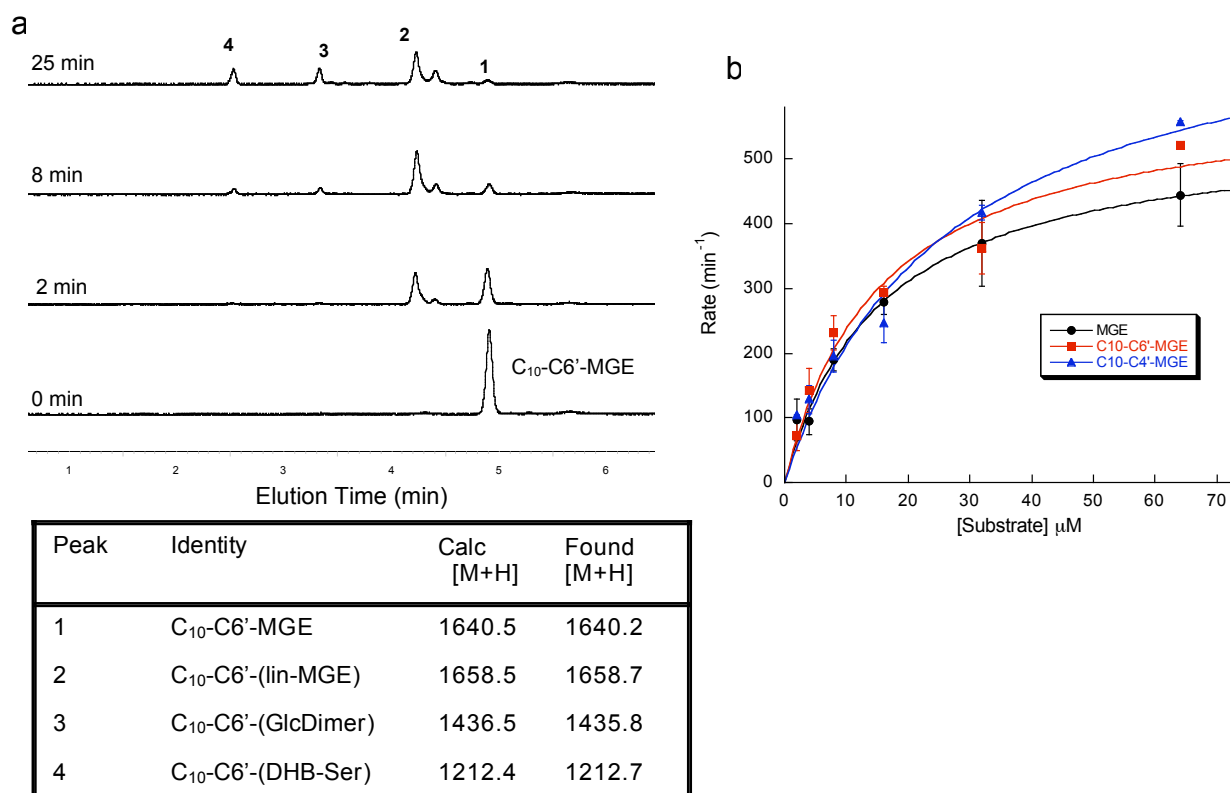
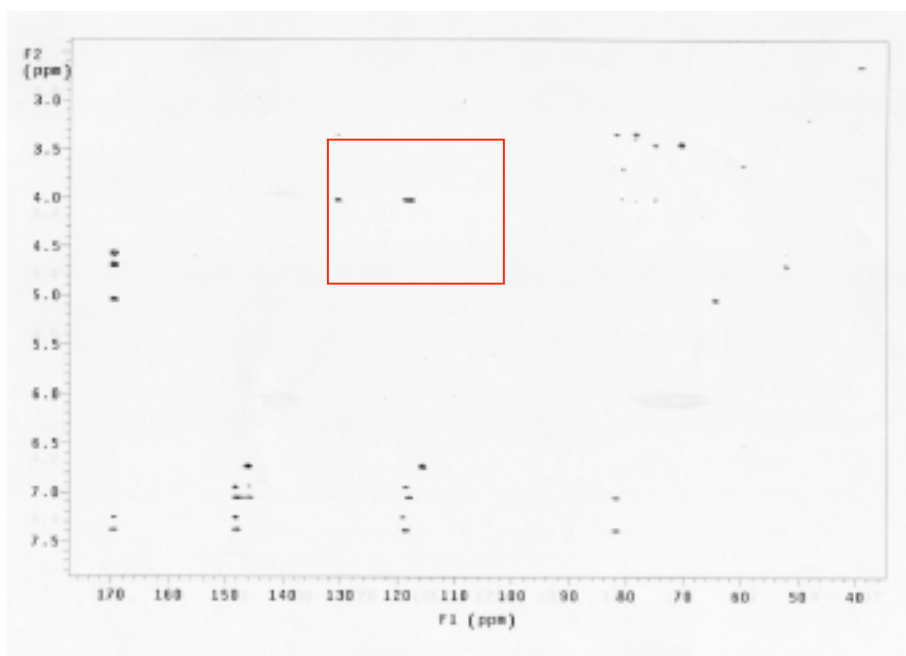
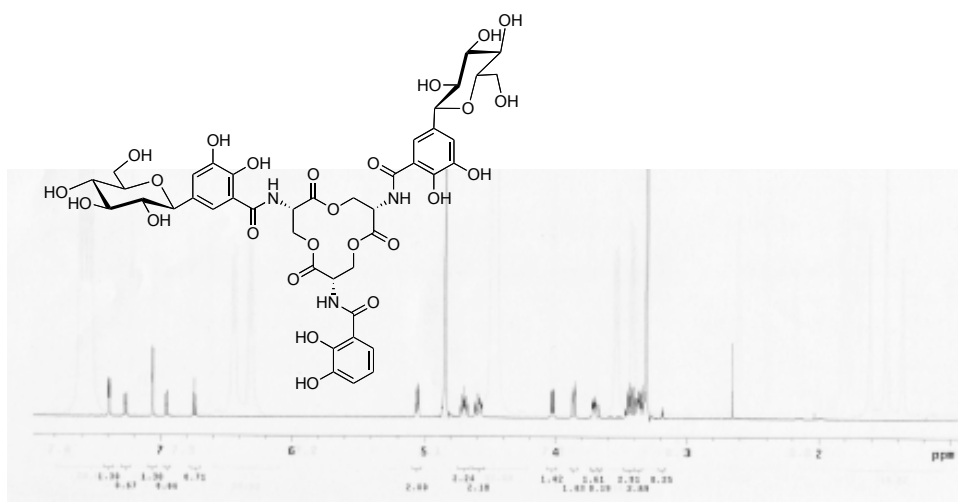


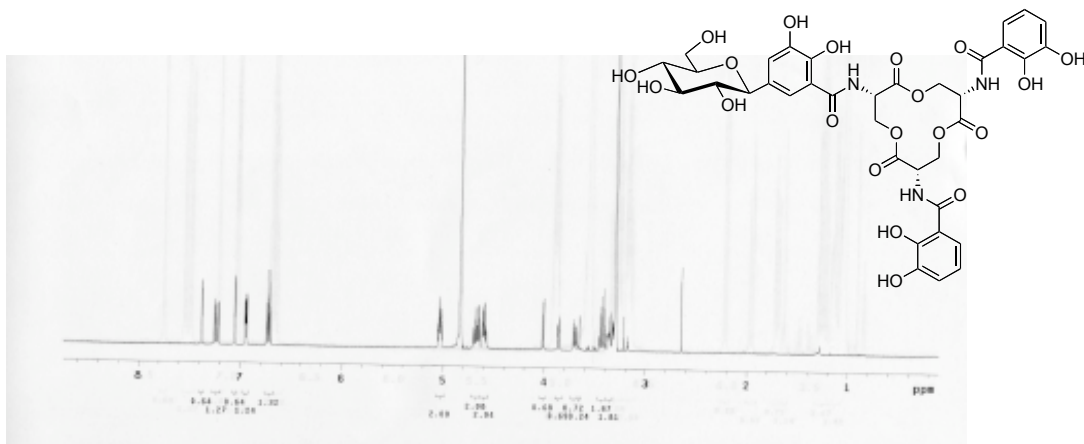
Figure S6. (a) MceD-catalyzed hydrolysis of the MGE moiety in C₁₀-C6'-MGE (20 nM MceD, 32 μM C₁₀-C6'-MGE, 75 mM Hepes pH 7.5). (b) Kinetic traces for the MceD-catalyzed hydrolysis of C₁₀-C6'-MGE, C₁₀-C4'-MGE and MGE. Corresponding kinetic parameters are given in Table 2 of the main text.

Representative NMR Spectra

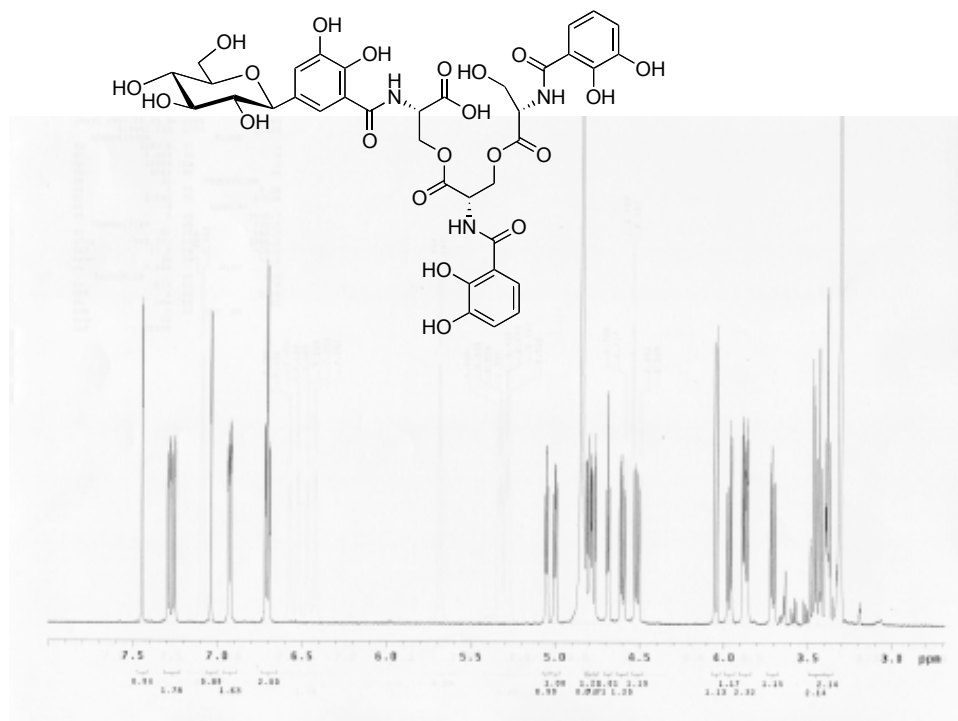
^1H NMR and HBMIC of DGE isolated from MceC-catalyzed glycosylation of Ent.



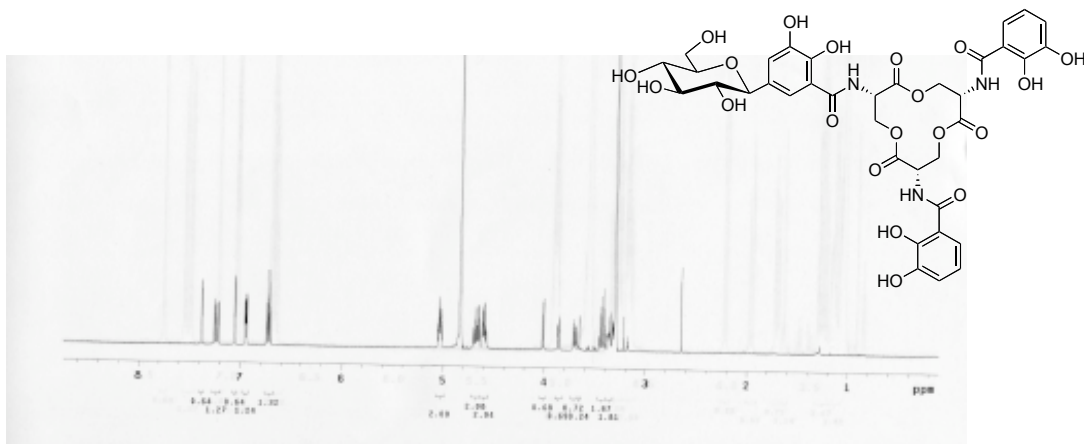
^1H NMR of MGE isolated from MceC-catalyzed glycosylation of Ent.



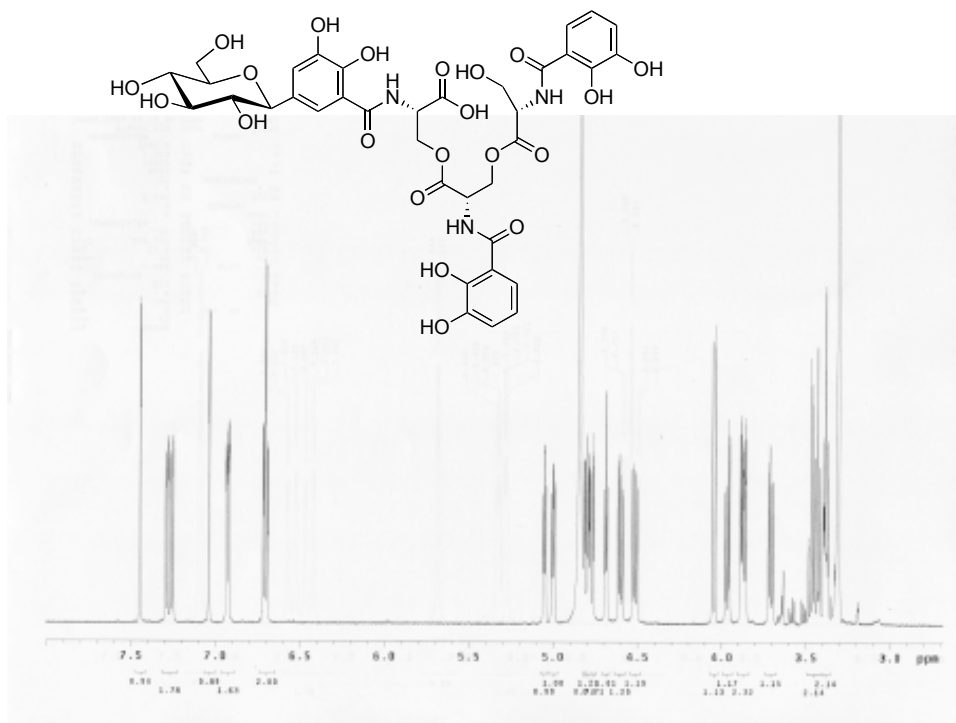
^1H NMR of linear MGE isolated from MceD-catalyzed hydrolysis of MGE.



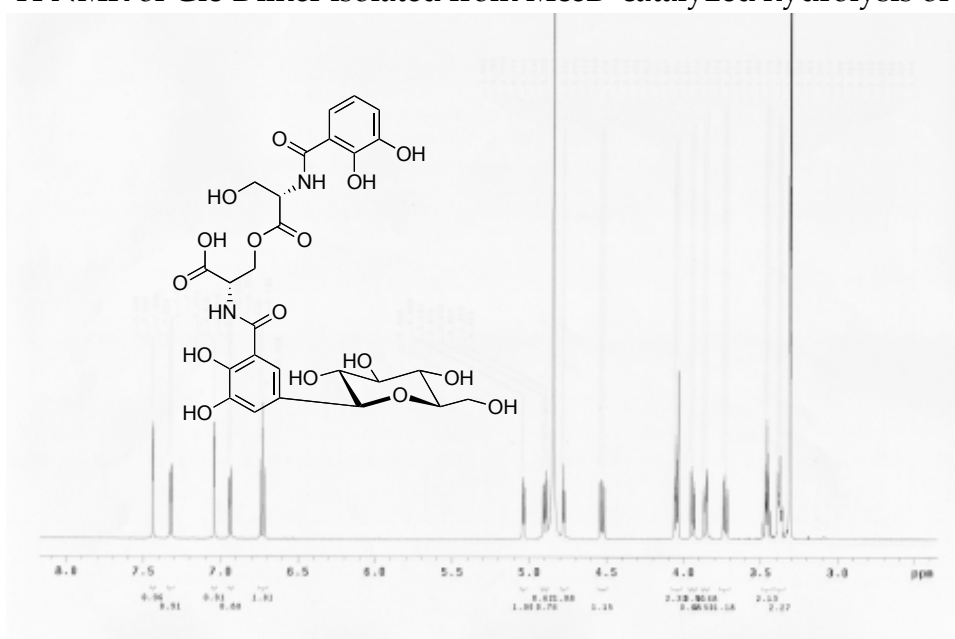
^1H NMR of MGE isolated from MceC-catalyzed glycosylation of Ent.



^1H NMR of linear MGE isolated from MceD-catalyzed hydrolysis of MGE.



^1H NMR of Glc-Dimer isolated from MceD-catalyzed hydrolysis of MGE.



^1H NMR of linear DGE from MceD-catalyzed hydrolysis of DGE.

