

Electronic Supporting Information for

Genomics-driven discovery of chiral triscatechol siderophores with enantiomeric Fe(III) coordination

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Figure S1. Structures of select triscatecholate siderophores

Scheme S1. Synthetic scheme for cyclic (DHB^{L/D}Lys^{L/D}Ser)₃ diastereomers

Figure S2. ¹H NMR of cyclic (DHB^LLys^LSer)₃

Figure S3. ¹³C NMR of cyclic (DHB^LLys^LSer)₃

Figure S4. ¹H-¹³C HMBC NMR of cyclic (DHB^LLys^LSer)₃

Table S1. NMR data for cyclic (DHB^LLys^LSer)₃

Table S2. NMR data for cyclic (DHB^DLys^LSer)₃

Figure S5. HPLC chromatograms of FDAA-derivatized (DHB^LLys^LSer)₃ and (DHB^LLys^DSer)₃ hydrolysate

Figure S6. HPLC chromatograms of FDAA-derivatized (DHB^DLys^LSer)₃ and (DHB^DLys^DSer)₃ hydrolysate

Table S3. Sequencing and assembly statistics for *Dickeya chrysanthemi* EC16

Table S4. Annotation of cyclic trichrysobactin gene cluster in *Dickeya chrysanthemi* EC16

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Figure S7. HPLC of the MeOH XAD-4 extract from the supernatant of a *Y. frederiksenii* ATCC 33641 culture and ESI-MS overlays of peaks 1 and 2

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Figure S11. ¹H NMR of frederiksenibactin (expansion of 1.2 ppm – 5.2 ppm region)

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Table S7. NMR data for frederiksenibactin

Table S8. Optimized Cartesian coordinates (Å) of Δ-Fe(III)-[(DHB^LLys^LSer)₃]

Table S9. Optimized Cartesian coordinates (Å) of Δ-Fe(III)-[(DHB^DLys^DSer)₃]

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Table S11. Optimized Cartesian coordinates (Å) of Δ-Fe(III)-[(DHB^DLys^LSer)₃]

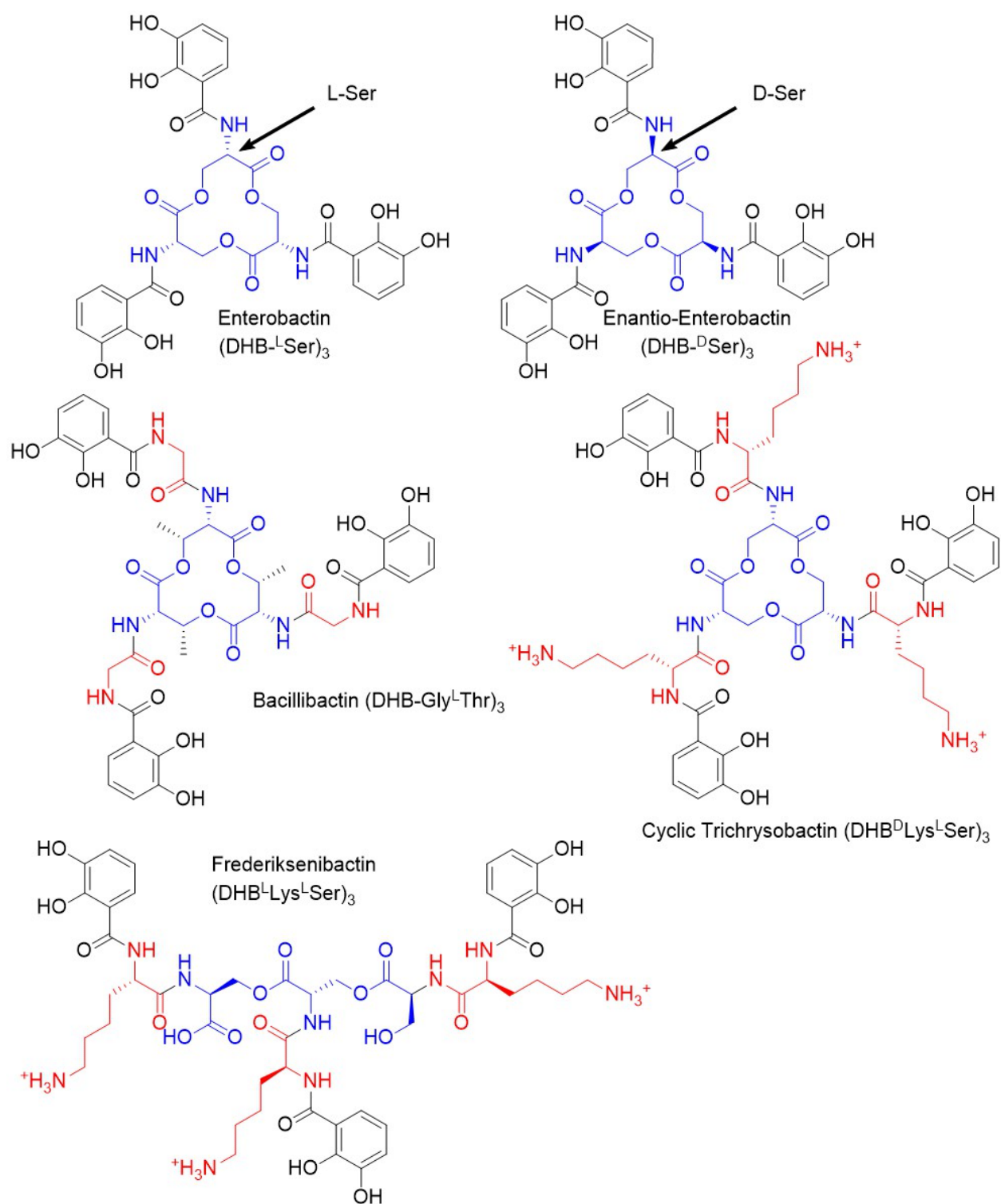
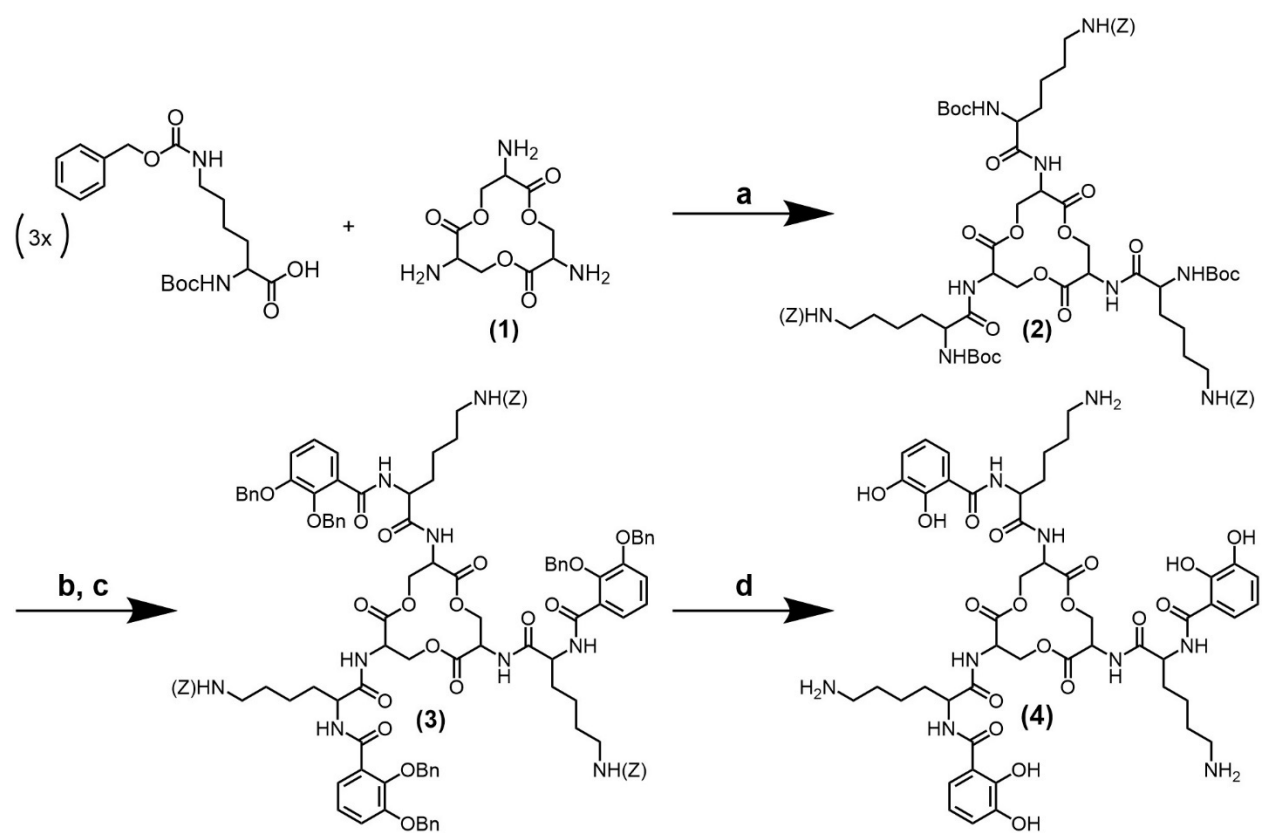
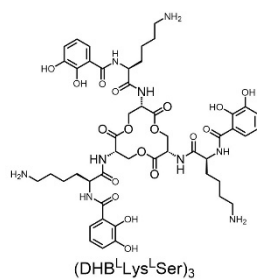


Figure S1. Structures of select triscatechol siderophores



Scheme S1. Synthetic scheme for cyclic (DHB^{L/D}Lys^{L/D}Ser)₃ diastereomers.



LSerLLysCAM_1H_070720

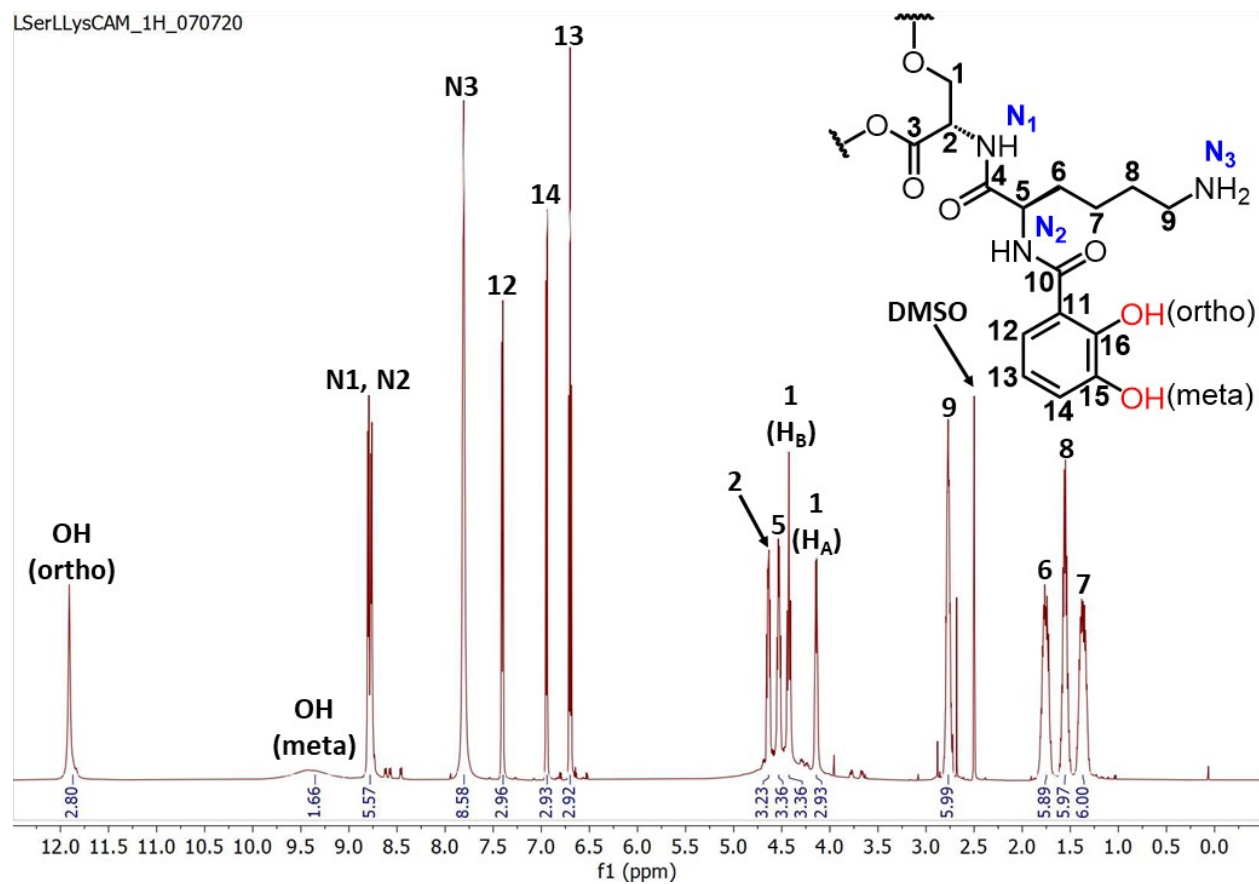
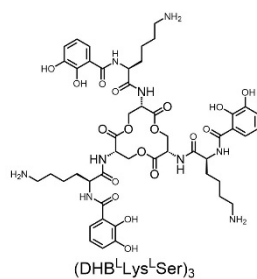


Figure S2: ¹H NMR of cyclic (DHB^L-Lys^L-Ser)₃ in (CD₃)₂SO.



LSerLLysCAM_13C_070720
Std carbon

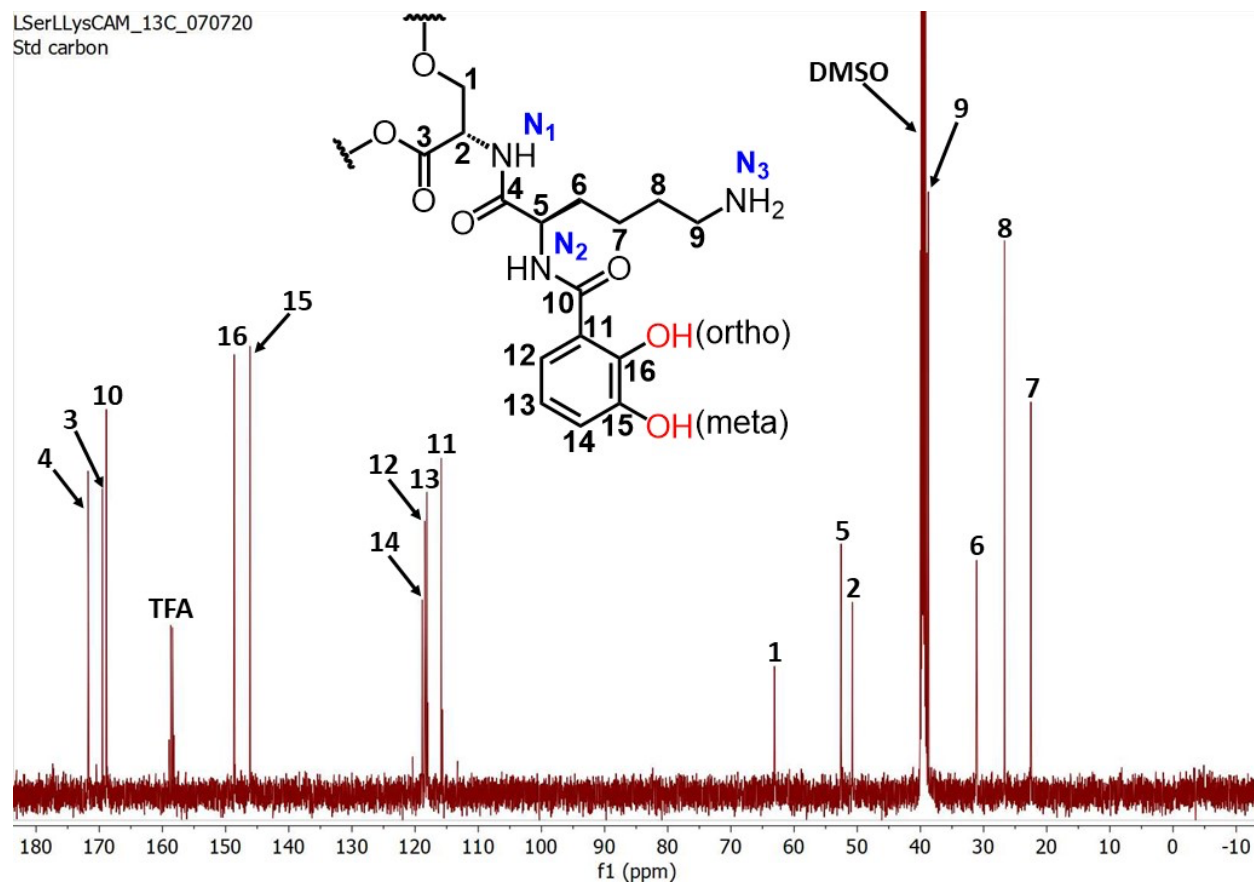


Figure S3. ¹³C NMR of cyclic (DHB^L-Lys^L-Ser)₃ in (CD₃)₂SO.

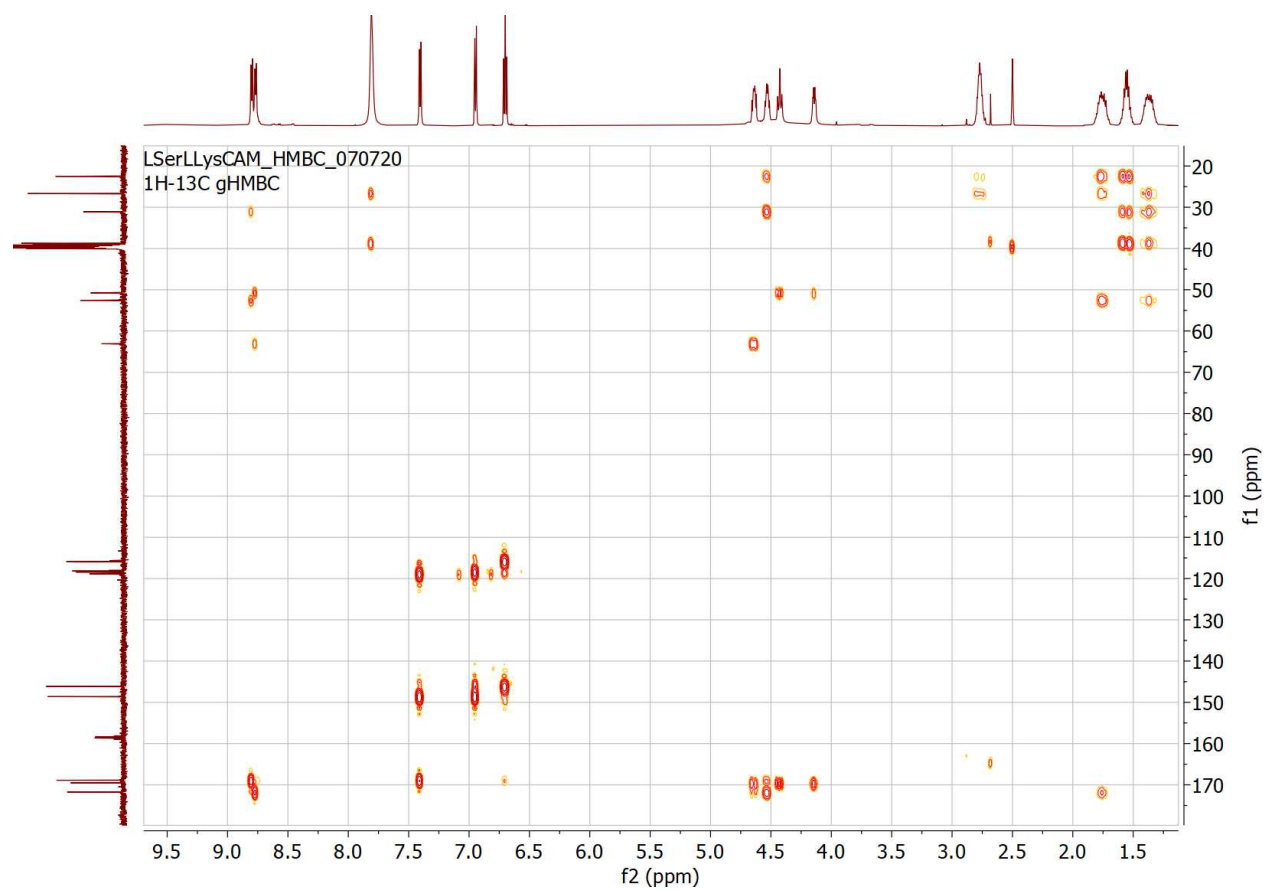


Figure S4. ^1H - ^{13}C HMBC NMR of cyclic $(\text{DHB}^{\text{L}}\text{Lys}^{\text{L}}\text{Ser})_3$ in $(\text{CD}_3)_2\text{SO}$.

Table S1. NMR data for cyclic (DHB^LLys^LSer)₃ in (CD₃)₂SO. HMBC correlations are from proton(s) stated to the indicated carbon.

(DHB ^L Lys ^L Ser) ₃			
Position	δ _C , type	δ _H (J in Hz)	HMBC
1 (H _A)	63.1, CH ₂	4.43, t (10.6)	2, 3
1 (H _B)	63.1, CH ₂	4.14, dd (10.8, 6.2)	2, 3
2	50.8, CH	4.64, ddd (10.1, 7.2, 4.7)	1, 3
3	169.5, C		
4	171.6, C		
5	52.6, CH	4.53, td (8.5, 4.8)	4, 6, 7, 10
6	31.1, CH ₂	1.75, m	4, 5, 7, 8
7	22.5, CH ₂	1.37, m	5, 6, 8, 9
8	26.7, CH ₂	1.55, m	6, 7, 9
9	38.7, CH ₂	2.77, m	8
10	168.9, C		
11	115.9, C		
12	118.4, CH	7.40, d (8.2)	10, 11, 13, 16
13	118.2, CH	6.70, t (7.9)	10, 11, 12, 15
14	118.9, CH	6.95, d (7.8)	12, 15, 16
15	146.1, C		
16	148.6, C		
N1		8.77, d (7.2)	1, 2, 4
N2		8.80, d (7.6)	5, 6, 10
N3		7.81, s	8, 9
OH (meta)		9.40, s	-
OH (ortho)		11.91, s	-

Table S2. NMR data for cyclic (DHB^DLys^LSer)₃ in (CD₃)₂SO. HMBC correlations are from proton(s) stated to the indicated carbon. Indicated positions correspond to the analogous positions in (DHB^LLys^LSer)₃ (Table S1).

(DHB ^D Lys ^L Ser) ₃			
Position	δ _C , type	δ _H (J in Hz)	HMBC
1 (H _A)	63.1, CH ₂	4.44, t (10.5)	2, 3
1 (H _B)	63.1, CH ₂	4.12, dd (10.8, 4.7)	2, 3
2	50.7, CH	4.65, ddd (10.2, 7.5, 4.7)	1, 3
3	169.5, C		
4	171.7, C		
5	52.5, CH	4.55, td (8.4, 5.2)	4, 6, 7, 10
6	31.3, CH ₂	1.75, m	4, 5, 7, 8
7	22.5, CH ₂	1.34, m	5, 6, 8, 9
8	26.7, CH ₂	1.54, m	6, 7, 9
9	38.7, CH ₂	2.77, s	-
10	168.7, C		
11	115.9, C		
12	118.5, CH	7.41, d (8.2)	10, 11, 14, 16
13	118.2, CH	6.70, t (7.9)	10, 11, 14, 15
14	118.8, CH	6.94, d (7.9)	12, 16
15	146.1, C		
16	148.5, C		
N1		8.80, d (overlapping with N2)	1, 2, 4
N2		8.80, d (overlapping with N1)	5, 6, 10
N3		7.78, s	-
OH (meta)		9.44, s	-
OH (ortho)		11.86, s	11, 15

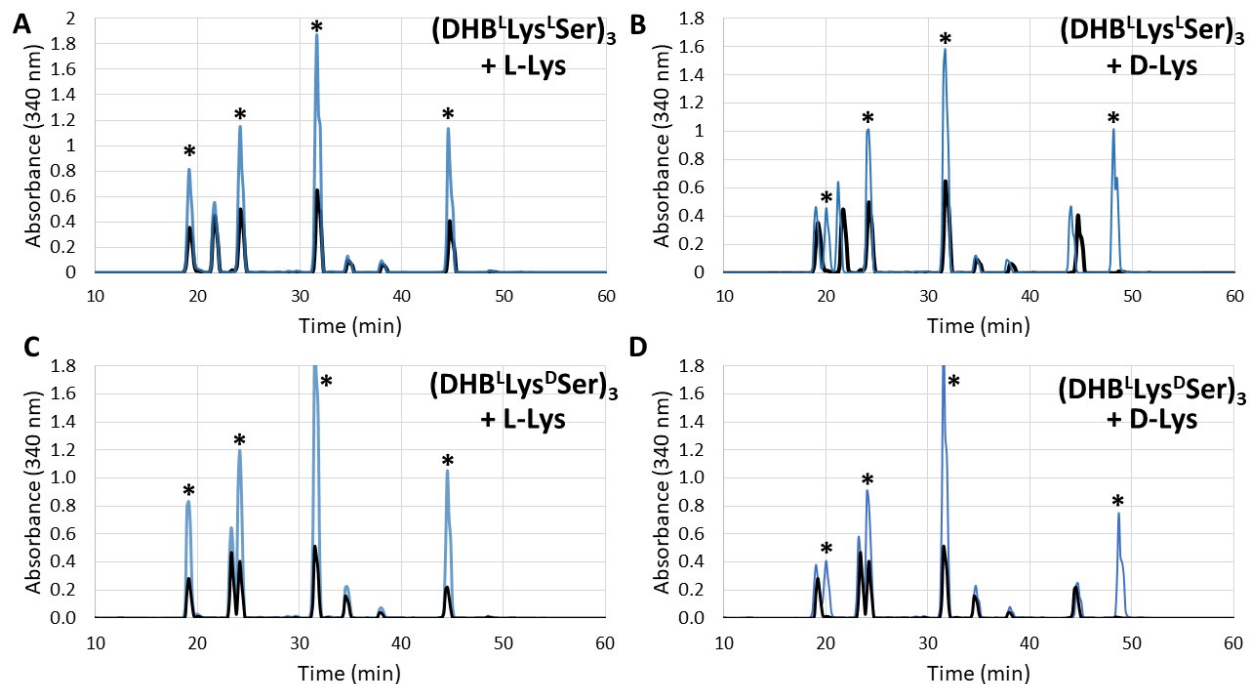


Figure S5. HPLC chromatograms of $(\text{DHB}^{\text{L}}\text{Lys}^{\text{L}}\text{Ser})_3$ and $(\text{DHB}^{\text{L}}\text{Lys}^{\text{D}}\text{Ser})_3$ hydrolysate derivatized with FDAA (light blue trace) overlayed with FDAA-derivatized amino acid standards (black trace). Derivatized hydrolysis products were separated by HPLC on a YMC 4.6 x 250 mm C18-A column using a gradient from 10% CH_3CN in TEAP buffer (50 mM, pH 3.00) to 40% CH_3CN in TEAP over 60 min. Asterisks indicate the peaks that increase upon addition of the designated amino acid standard.

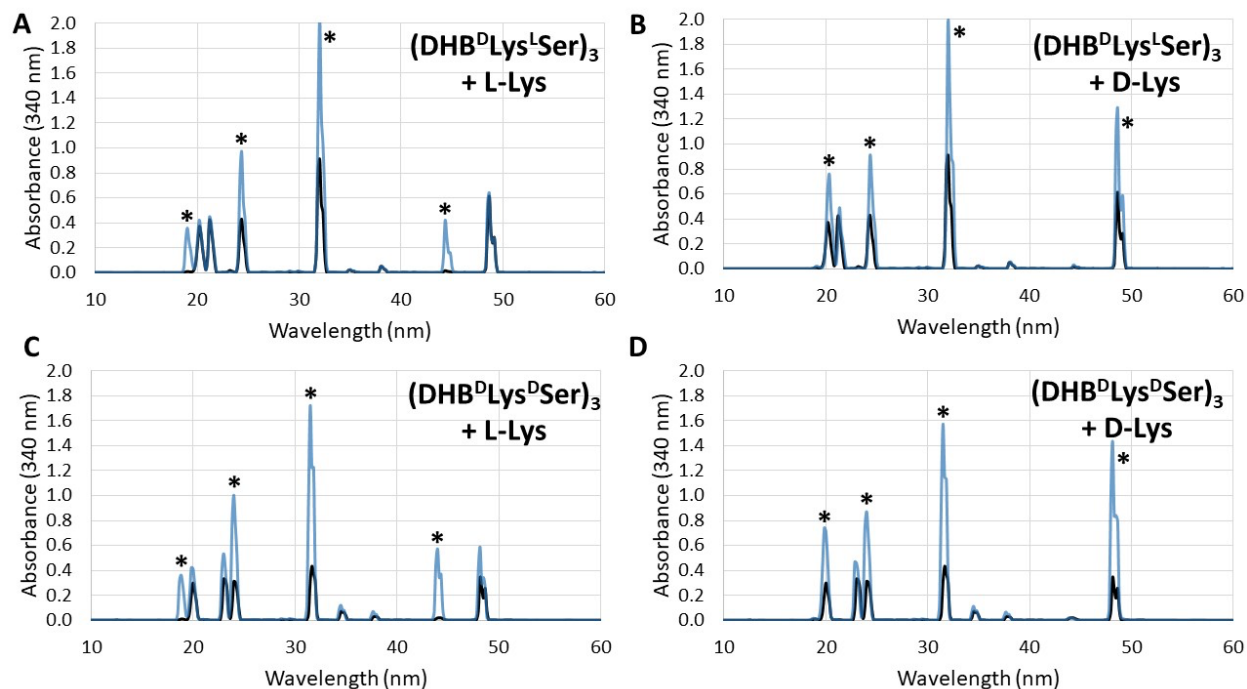


Figure S6. HPLC chromatograms of $(\text{DHB}^{\text{D}}\text{Lys}^{\text{L}}\text{Ser})_3$ and $(\text{DHB}^{\text{D}}\text{Lys}^{\text{D}}\text{Ser})_3$ hydrolysate derivatized with FDAA (light blue trace) overlayed with FDAA-derivatized amino acid standards (black trace). Derivatized hydrolysis products were separated by HPLC on a YMC 4.6 x 250 mm C18-A column using a gradient from 10% CH_3CN in TEAP buffer (50 mM, pH 3.00) to 40% CH_3CN in TEAP over 60 min. Asterisks indicate the peaks that increase upon addition of the designated amino acid standard.

Table S3. Sequencing and assembly statistics for *Dickeya chrysanthemi* EC16 (=ATCC 11662)

Statistic	Data for <i>Dickeya chrysanthemi</i> EC16
Total filtered read length (bp)	865,361,754
Assembly size (bp)	4,685,942
Coverage (×)	185
No. of scaffolds	70
No. of contigs	76
Scaffold N_{50} (bp)	190,138
Contig N_{50} (bp)	186,974
G+C content (%)	54.36
No. of identified genes (total)	4,205
No. of identified CDSs (total)	4,136
No. of rRNAs (5S, 16S, 23S)	1, 2, 1
No. of predicted tRNAs	58
No. of predicted ncRNAs	7
Genome accession no.	JAFCAF000000000
SRA accession no.	SRR13439815

Table S4. Annotation of cyclic trichrysobactin gene cluster in *Dickeya chrysanthemi* EC16, including predicted protein functions based on sequence analysis using Pfam and BLAST. The sequence similarity of each gene to its corresponding homolog from the chrysobactin locus (*Dickeya dadantii* 3937) is indicated.

Name		Locus tag	Homologs in <i>D. dadantii</i> 3937			Accession
			%ID	%SIM	Locus tag	
<i>cbsA</i>	2,3-dihydro-2,3-DHB dehydrogenase	JJO56_07665	90	92	DDA3937_RS14720	WP_013318793
<i>cbsB</i>	Isochorismatase / Aryl carrier protein	JJO56_07670	95	96	DDA3937_RS14715	WP_013318792
<i>cbsE</i>	2,3-DHB-AMP ligase	JJO56_07675	92	95	DDA3937_RS14710	WP_013318791
<i>cbsC</i>	Isochorismate synthase	JJO56_07680	92	94	DDA3937_RS14705	WP_013318790
<i>fctA</i>	TonB-dependent receptor	JJO56_07685	81	88	DDA3937_RS14700	WP_013318789
<i>cbsH</i>	Esterase	JJO56_07690	88	92	DDA3937_RS14695	WP_033111969
<i>cbsI</i>	MbtH-like protein	JJO56_07695	86	90	DDA3937_RS14690	WP_013318786
<i>cbsF</i>	NRPS	JJO56_07700	91	94	DDA3937_RS14685	WP_033112377
<i>fctC</i>	ABC ATP-binding component	JJO56_07705	96	96	DDA3937_RS14675	WP_033111968
<i>fctG</i>	ABC permease	JJO56_07710	90	94	DDA3937_RS14670	WP_033112376
<i>fctD</i>	ABC permease	JJO56_07715	92	95	DDA3937_RS14665	WP_013318781
<i>cbsS</i>	MFS transporter	JJO56_07720	96	97	DDA3937_RS14660	WP_013318779
<i>fctB</i>	ABC periplasmic binding protein	JJO56_07725	84	90	DDA3937_RS14655	WP_033111967

Table S5. Annotation of frederiksenibactin gene cluster of *Yersinia frederiksenii* ATCC 33641. The predicted protein functions based on sequence analysis using Pfam and BLAST are included. The sequence similarity of each gene to its corresponding homolog in the cyclic trichrysobactin locus (*D. chrysanthemi* EC16, Table S2) is indicated.

Name	Putative function	Locus tag	Accession	Homologs in <i>D. chrysanthemi</i> EC16		
				Name	%ID	%SIM
<i>ffuA</i>	TonB-dependent receptor	DJ58_RS13925	WP_004709694	<i>fctA</i>	54	71
<i>freH</i>	Esterase	DJ58_RS13920	WP_080544642	<i>cbsH</i>	49	62
<i>freI</i>	MbtH-like protein	DJ58_RS13915	WP_032911217	<i>cbsI</i>	63	76
<i>freF</i>	NRPS	DJ58_RS13910	WP_050504533	<i>cbsF</i>	58	71
<i>ffuC</i>	ABC ATP-binding component	DJ58_RS13905	WP_080544678	<i>fctC</i>	71	84
<i>ffuG</i>	ABC permease	DJ58_RS13900	WP_032911802	<i>fctG</i>	63	77
<i>ffuD</i>	ABC permease	DJ58_RS13895	WP_004711942	<i>fctD</i>	59	77
<i>freS</i>	MFS transporter	DJ58_RS13890	WP_004711941	<i>cbsS</i>	64	77
<i>ffuB</i>	ABC periplasmic binding protein	DJ58_RS13885	WP_004711939	<i>fctB</i>	54	72
<i>freC</i>	Isochorismate synthase	DJ58_RS13880	WP_004711937	<i>cbsC</i>	47	62
<i>freE</i>	2,3-DHB-AMP ligase	DJ58_RS13875	WP_032911801	<i>cbsE</i>	64	74
<i>freB</i>	Isochorismatase / Aryl carrier protein	DJ58_RS13870	WP_004711933	<i>cbsB</i>	58	74
<i>freA</i>	2,3-dihydro-2,3-DHB dehydrogenase	DJ58_RS13865	WP_032911814	<i>cbsA</i>	65	77

Table S6. Complete genomes with a putative (DHB^LLys^LSer)₃ siderophore.

Strain	RefSeq Accession
<i>Yersinia enterocolitica</i> LC20	GCF_000597945.1
<i>Y. kristensenii</i> Y231	GCF_000834865.1
<i>Y. frederiksenii</i> Y225	GCF_000834215.1
<i>Y. frederiksenii</i> FDAARGOS_417	GCF_002591095.1
<i>Y. frederiksenii</i> FDAARGOS_418	GCF_002591195.1
<i>Y. massiliensis</i> GTA	GCF_003048255.1
<i>Y. hibernica</i> CFS1934	GCF_004124235.1

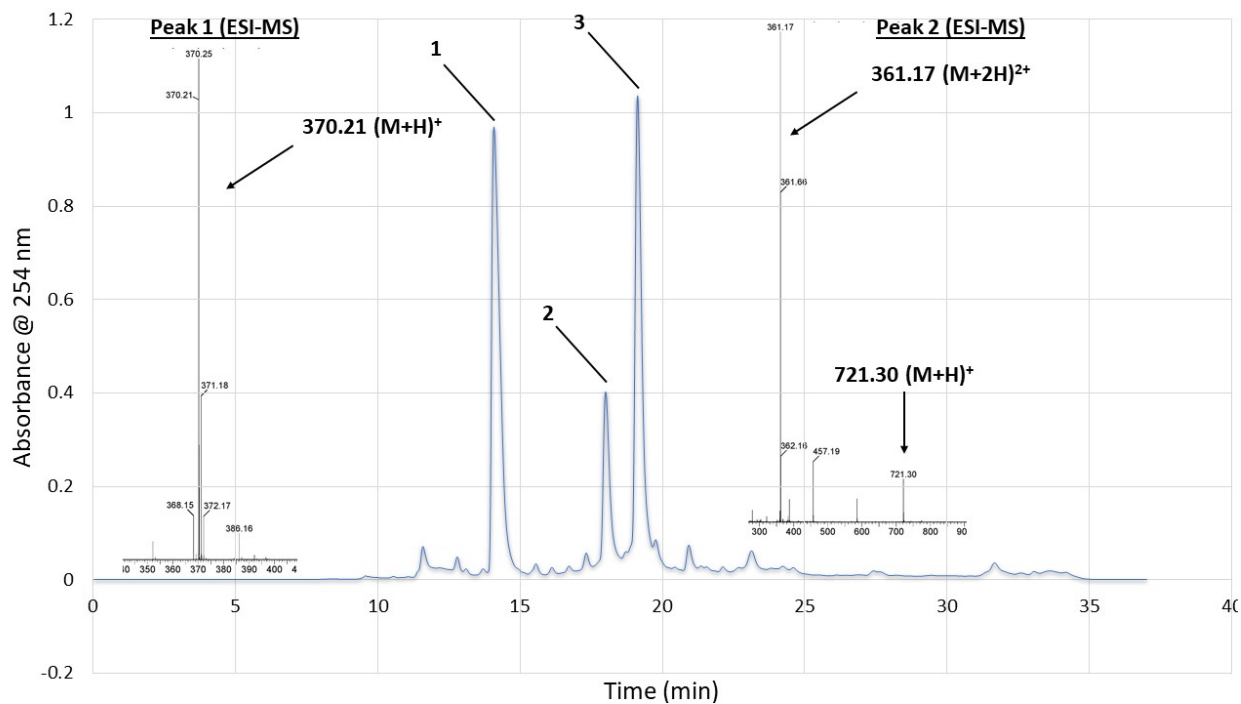


Figure S7. HPLC of the MeOH XAD-4 extract from the supernatant of a *Y. frederiksenii* ATCC 33641 culture. ESI-MS of isolated peaks **1** and **2** corresponding to the related monocatechol and dicatechol fragments of frederiksenibactin, respectively. MS data for peak **3** corresponding to frederiksenibactin shown in Figure S8.

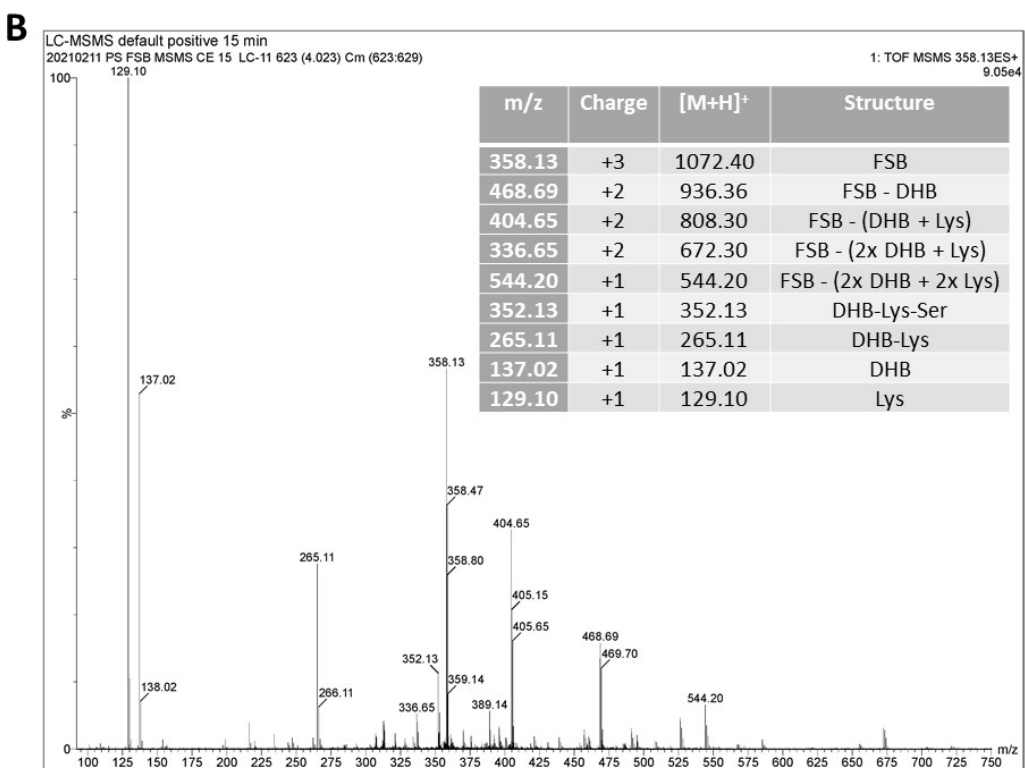
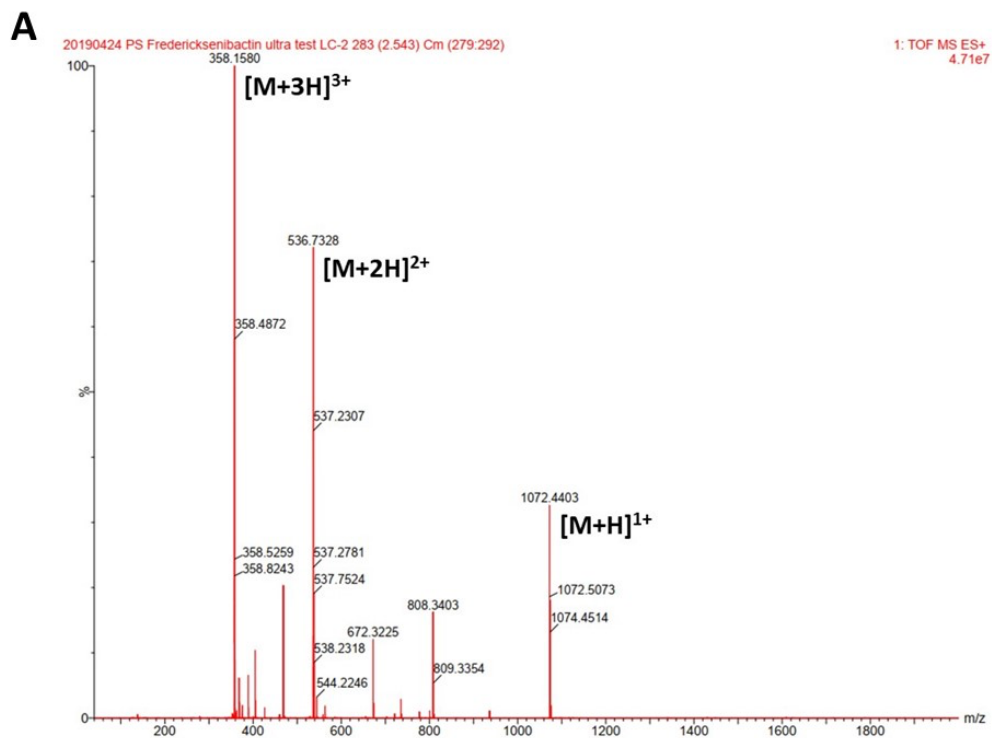


Figure S8. (a) HR-ESI-MS of frederiksenibactin. Observed m/z 1072.4403 $[M+H]^+$. Calculated exact mass for frederiksenibactin $[M+H]^+$ is 1072.4475 ($C_{48}H_{66}N_9O_{19}$). (b) ESI-MSMS of frederiksenibactin.

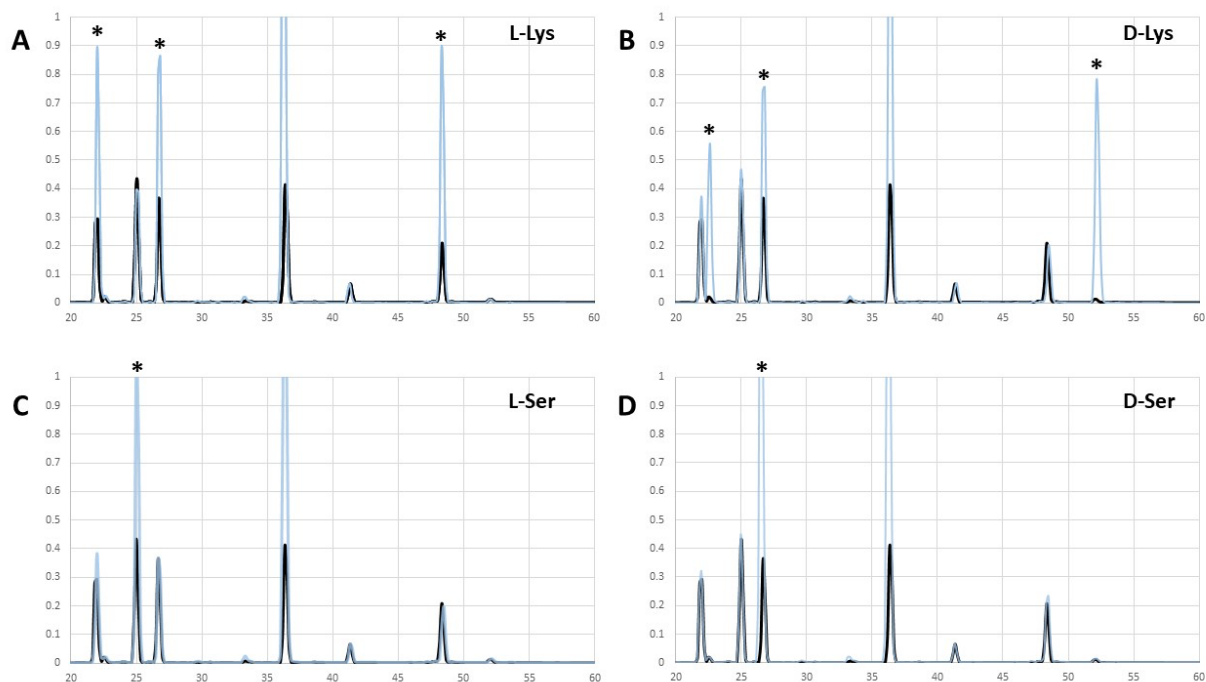


Figure S9. HPLC chromatograms of frederiksenibactin hydrolysate derivatized with FDAA (light blue trace) overlaid with FDAA-derivatized amino acid standards (black traces). Derivatized hydrolysis products were separated by HPLC on a YMC 4.6 x 250 mm C18-AQ column using a gradient from 10% CH₃CN in ddH₂O (0.05% trifluoroacetic acid) to 40% CH₃CN in ddH₂O (0.05% trifluoroacetic acid) over 60 min. Asterisks indicate the peaks that increase upon addition of the designated amino acid standard.

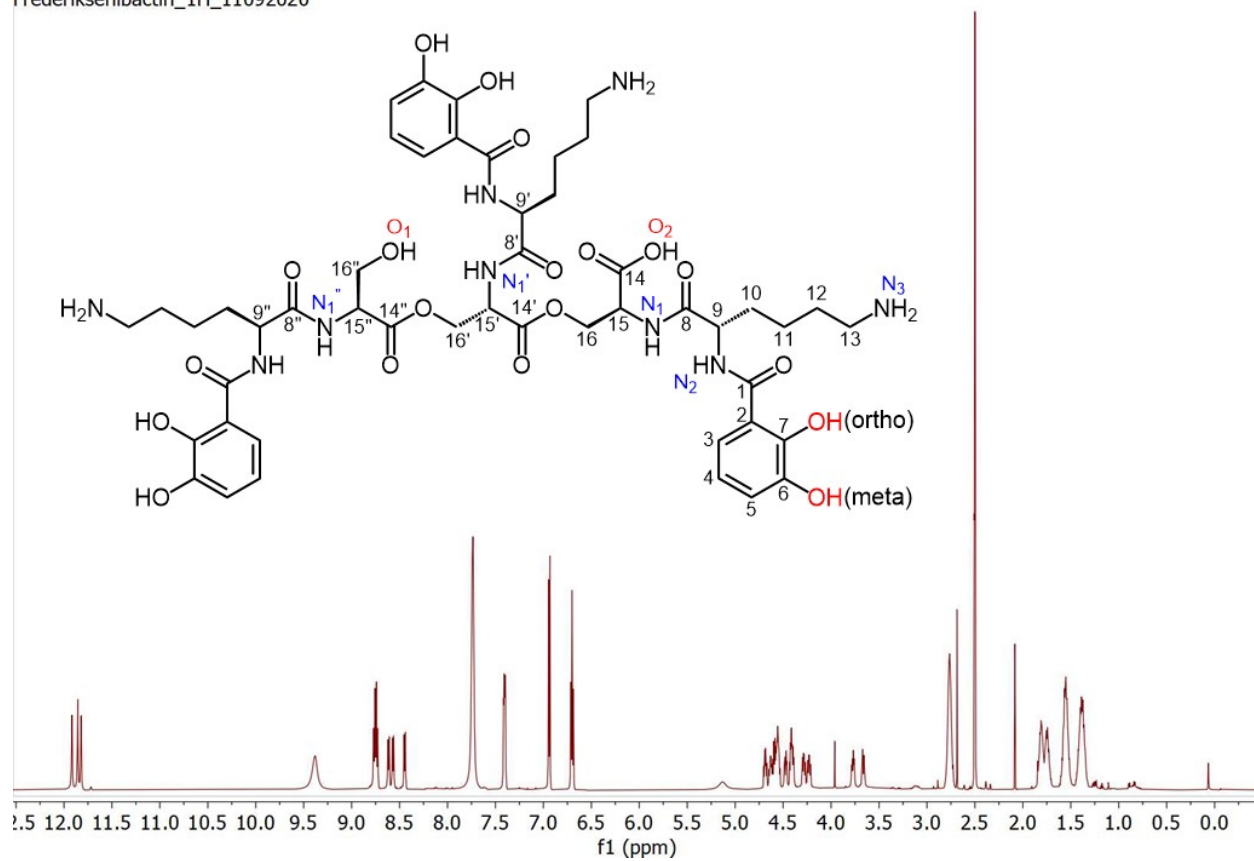


Figure S10. ^1H NMR of frederiksenibactin, FSB, in $(\text{CD}_3)_2\text{SO}$.

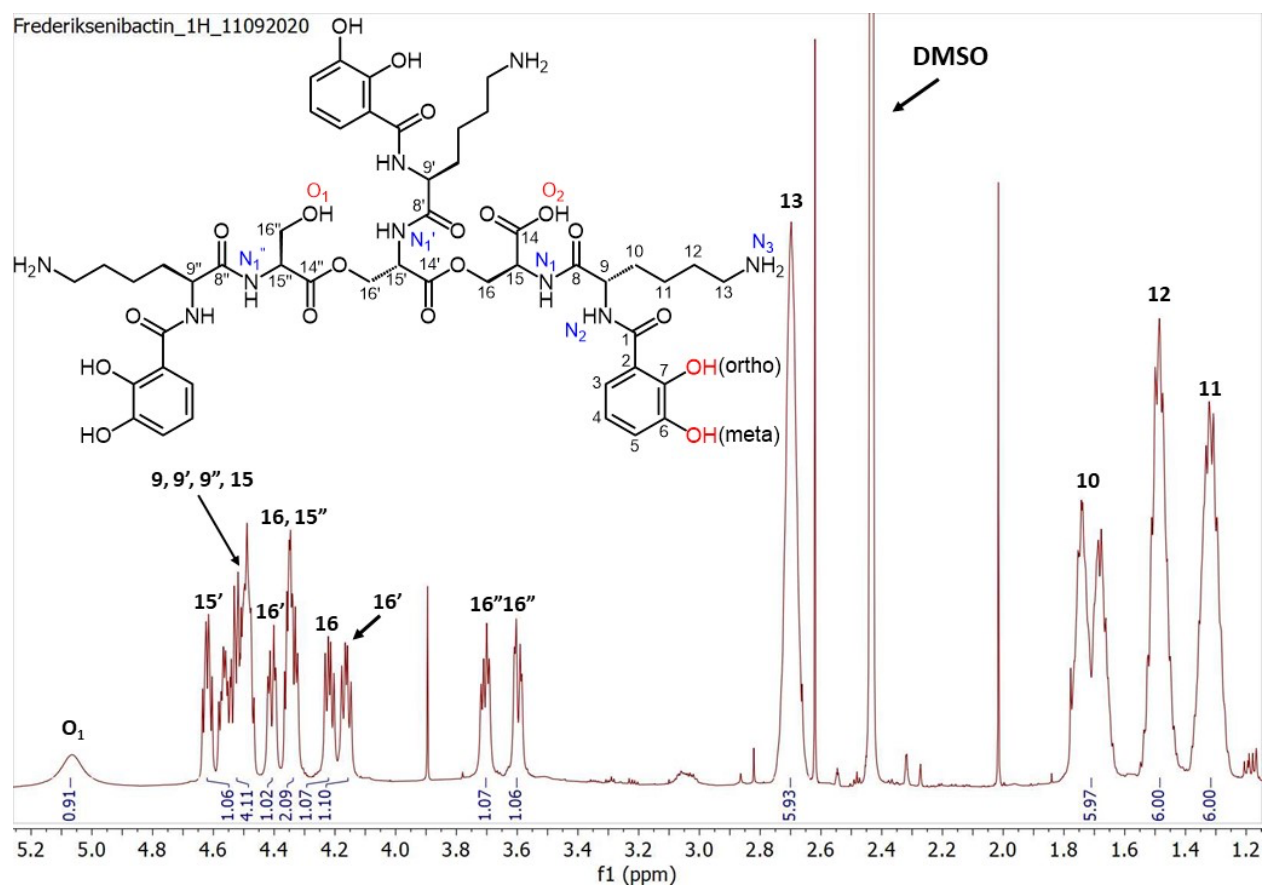


Figure S11. 1H NMR of frederiksenibactin (expansion of 1.2 ppm – 5.2 ppm region) in $(CD_3)_2SO$.

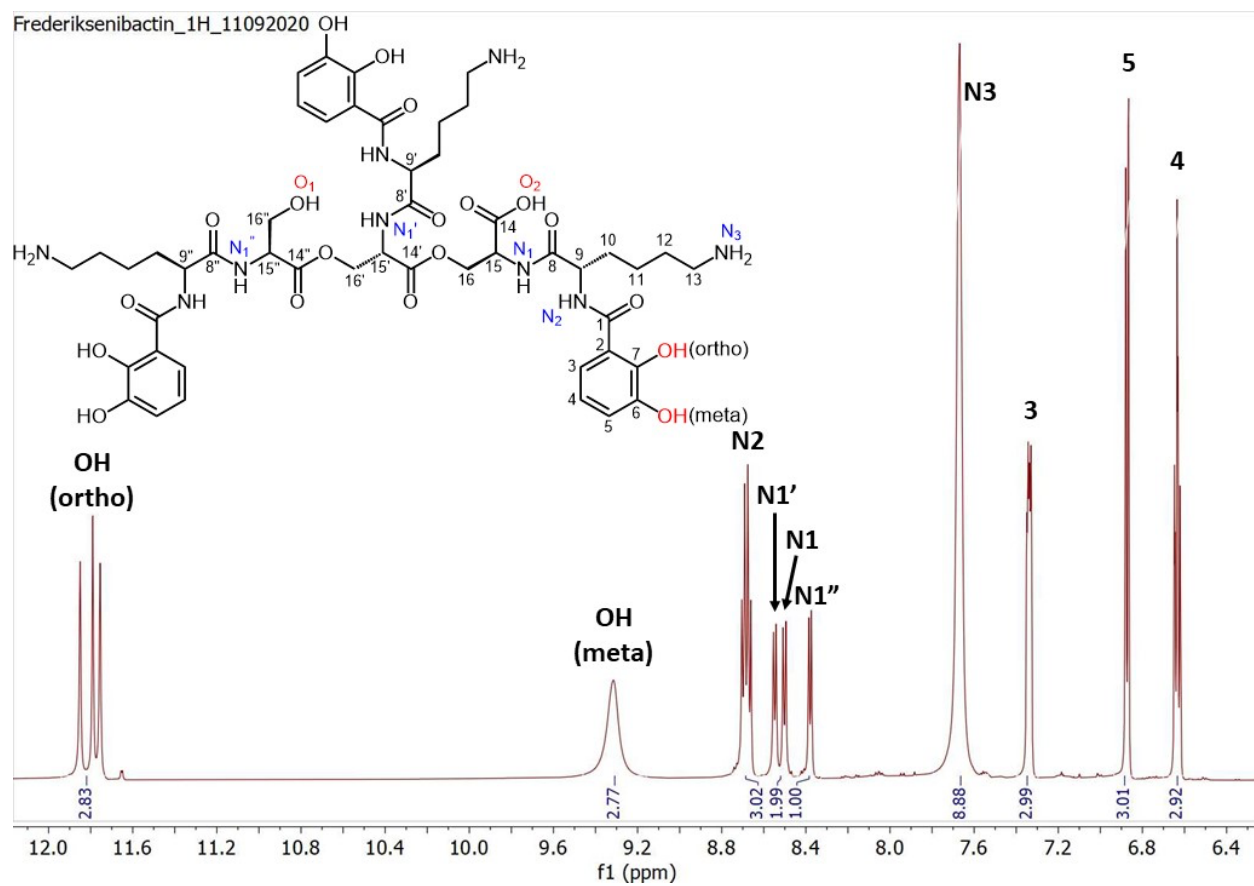
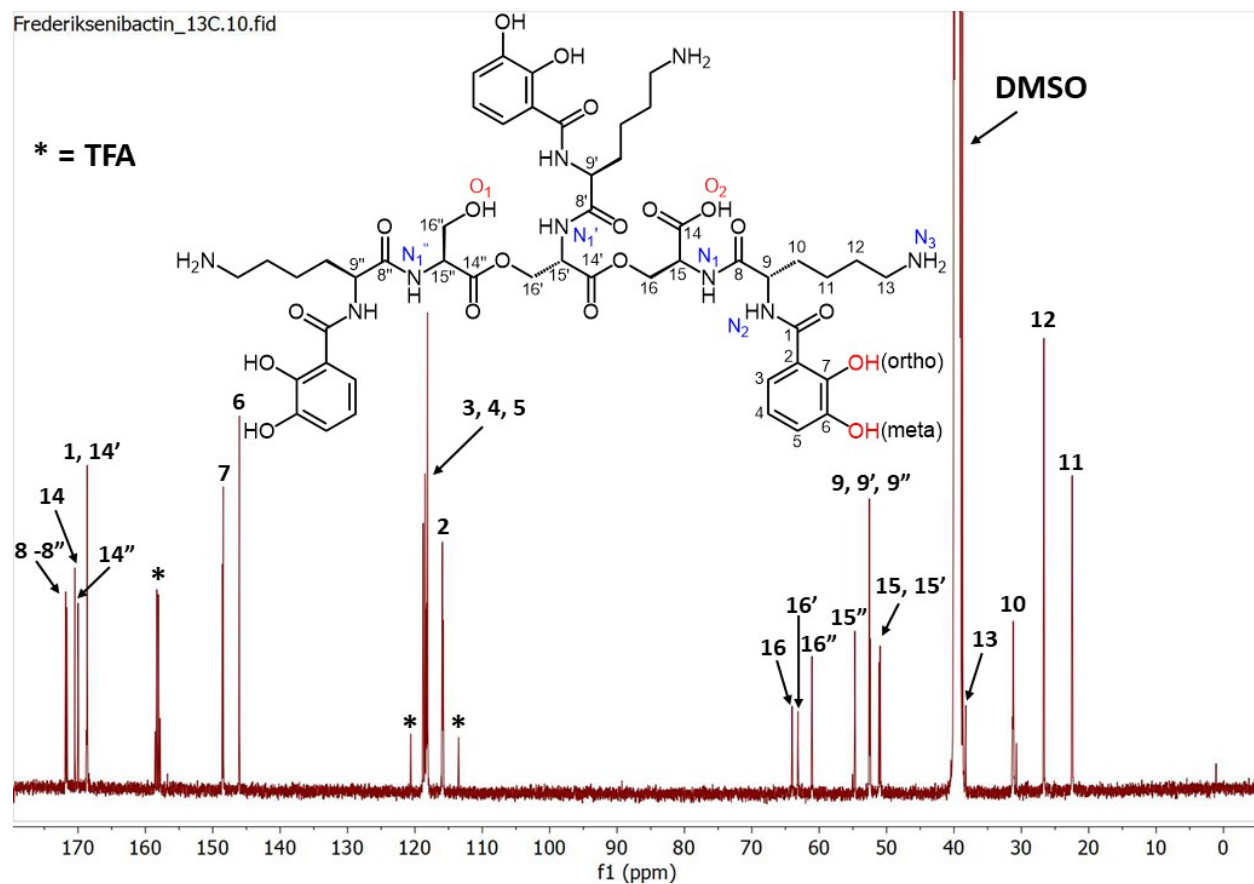


Figure S12. ¹H NMR of frederiksenibactin (expansion of 6.4 ppm – 12.0 ppm region) in (CD₃)₂SO.



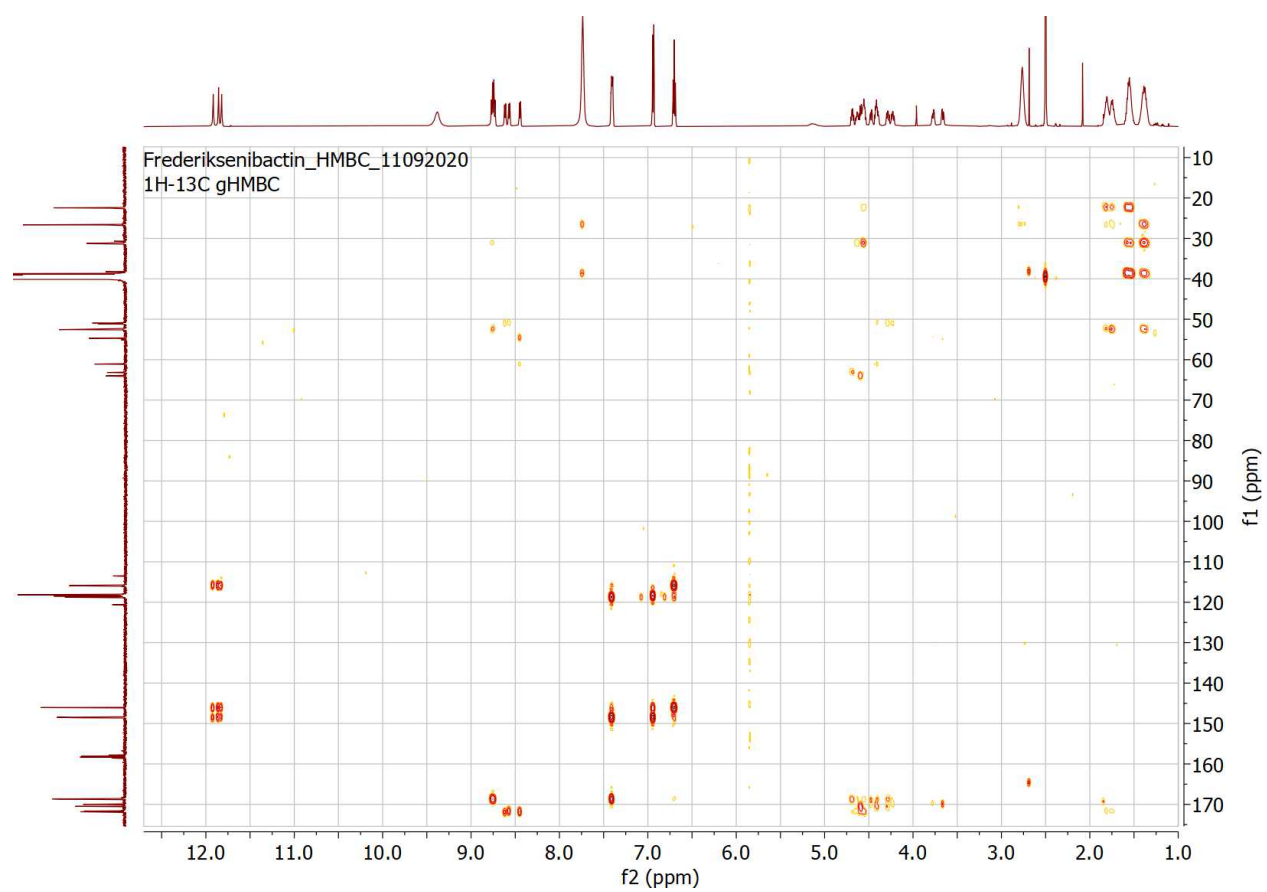


Figure S14. ^1H - ^{13}C HMBC of frederiksenibactin (FSB) in $(\text{CD}_3)_2\text{SO}$.

Table S7. NMR data for frederiksenibactin in (CD₃)₂SO. HMBC correlations are from proton(s) stated to the indicated carbon.

Frederiksenibactin (1)			
Position	δ_c , type	δ_H (J in Hz)	HMBC
1	168.7, C		
2	115.9, C		
3	118.5, CH	7.40, m	1, 2, 5, 6, 7
4	118.1, CH	6.70, td (7.9, 2.2)	3, 5, 6, 7
5	118.8, CH	6.94, d (7.8)	3, 6, 7
6	146.1, C		
7	148.5, C		
8	171.7, C		
8'	171.9, C		
8''	171.8, C		
9	52.5, CH	4.65, m	8, 10, 11
9' / 9''	52.6, CH	4.50 - 4.60, m	8', 8'', 10, 11
10 (H _A)	31.2, CH ₂	1.75, m	9, 11, 12
10 (H _B)	31.2, CH ₂	1.80, m	9, 11, 12
11	22.4, CH ₂	1.39, m	9, 10, 12, 13
12	26.7, CH ₂	1.55, m	10, 11, 13
13	38.7, CH ₂	2.77, m	
14	170.5, C		
15	51.0, CH	4.59, m	14, 16
16 (H _A)	64.0, CH ₂	4.28, dd(11.1, 6.4)	14, 15, 17
16 (H _B)	64.0, CH ₂	4.40, m	15, 17
14'	168.7, C		
15'	51.1, CH	4.69, td (7.2, 4.6)	
16' (H _A)	63.2, CH ₂	4.23, dd (11.4, 7.0)	17, 18, 20
16' (H _B)	63.2, CH ₂	4.47, dd (11.4, 4.3)	17, 20
14''	170.0, C		
15''	54.7, CH	4.41, m	20, 22
16'' (H _A)	61.1, CH	3.66, dd (11.4, 4.0)	20
16'' (H _B)	61.1, CH	3.77, dd (11.3, 5.3)	20
N1		8.57, d (7.7)	8, 15
N1'		8.61, d (7.9)	8', 18
N1''		8.45, d (8.4)	8'', 21
N2		8.75, m	1, 9-9''
N3		7.73, br s	12, 13
O1		13.13, br s	
O2		5.13, br s	
OH (meta)		9.38, br s	
OH (ortho)		11.82 - 11.92, s	2, 6, 7

Table S8. Optimized Cartesian coordinates (Å) of Δ -Fe(III)-[(DHB^LLys^LSer)₃].

O	-1.732100	0.793600	5.333300	H	-5.268300	-3.093900	0.464700
O	-0.385900	-2.339000	5.168900	H	-2.253300	-3.347100	0.859000
O	1.583300	0.399500	5.420400	H	-3.399400	-4.166100	1.932300
C	-1.791800	0.807200	4.132000	C	1.909800	4.587700	-0.109800
C	-0.223700	-2.275600	3.978800	C	0.414500	4.282400	0.017800
C	1.629100	0.607700	4.236700	H	2.203800	4.645600	-1.161600
C	-2.437800	-1.527400	3.625600	H	2.125000	5.561300	0.341400
C	2.166200	-1.645600	3.771000	C	-0.166400	4.906500	1.285200
C	-0.057800	2.403900	3.976700	H	-0.115000	4.653400	-0.868900
H	-2.461700	-1.735000	4.695100	H	0.239800	3.198500	0.033800
H	-3.153200	-2.164000	3.110400	C	-1.611200	4.527400	1.532500
H	2.245000	-1.641600	4.857900	H	0.424200	4.613000	2.152000
H	3.122800	-1.919900	3.332800	H	-0.078100	5.998600	1.231200
H	-0.220200	2.388000	5.054000	H	-1.786600	3.474000	1.302600
H	-0.241600	3.403200	3.589400	H	-1.921300	4.711500	2.561700
O	1.861200	-0.319400	3.301800	C	-0.523000	-4.224900	-1.100400
O	-0.999800	1.527100	3.331400	O	-0.829600	-5.325400	-0.589800
O	-1.136800	-1.849700	3.099800	C	4.011500	1.719800	-0.557800
C	1.380700	1.996600	3.682600	O	5.059200	1.962300	0.082500
H	1.987700	2.661800	4.305600	C	-3.419700	2.680300	-0.746100
C	1.103500	-2.654100	3.352100	O	-4.218500	3.424400	-0.133100
H	1.396900	-3.583400	3.850800	C	-2.481100	3.203300	-1.750100
C	-2.790800	-0.063300	3.395800	C	-1.494400	2.389200	-2.336500
H	-3.736100	0.077900	3.930100	C	-2.653400	4.527900	-2.221400
N	1.072900	-2.938500	1.940900	C	-0.722700	2.885800	-3.446000
N	-3.043800	0.285000	2.021900	C	-1.842700	5.029300	-3.218100
N	1.795000	2.215500	2.321200	H	-3.475800	5.119100	-1.832300
H	1.654700	1.501800	1.620300	C	-0.875400	4.212200	-3.831300
H	0.527700	-2.365400	1.312800	H	-1.982400	6.047600	-3.566900
H	-2.282600	0.534900	1.406800	H	-0.268200	4.595000	-4.647000
C	2.476900	3.340100	2.001800	C	3.989700	0.782500	-1.690800
O	2.746500	4.208900	2.823200	C	2.788200	0.405900	-2.319400
C	2.838200	3.547800	0.539200	C	5.221100	0.339300	-2.232900
H	3.851700	3.955000	0.558300	C	2.827700	-0.368000	-3.533400
N	2.854200	2.344300	-0.253400	C	5.245900	-0.484600	-3.338600
H	2.048700	2.072000	-0.828600	H	6.146400	0.710300	-1.804400
C	1.685000	-4.048700	1.466600	C	4.051500	-0.842200	-3.989900
O	2.299100	-4.828400	2.185500	H	6.196500	-0.821400	-3.740100
C	1.666600	-4.264800	-0.038500	H	4.075700	-1.454900	-4.886800
H	1.517300	-5.339600	-0.163700	C	-1.337300	-3.599100	-2.153200
N	0.604300	-3.580500	-0.730800	C	-1.068200	-2.301000	-2.625500
H	0.764000	-2.680000	-1.196700	C	-2.339900	-4.371300	-2.789600
C	-4.319600	0.377200	1.577300	C	-1.762200	-1.803100	-3.784900
O	-5.289900	0.154700	2.291800	C	-3.070800	-3.848600	-3.835900
C	-4.525700	0.686100	0.102200	H	-2.478500	-5.403100	-2.484100
H	-5.383100	1.362600	0.077300	C	-2.786300	-2.564500	-4.334800
N	-3.415500	1.347400	-0.534600	H	-3.839700	-4.449700	-4.311000
H	-2.718800	0.823300	-1.076400	H	-3.332100	-2.171800	-5.188200
C	3.022100	-3.886500	-0.658900	O	-1.250000	1.158800	-1.969600
C	3.502200	-2.468000	-0.338300	O	0.076900	2.021400	-4.030200
H	2.913400	-4.022700	-1.738600	O	1.601500	0.755800	-1.897900
H	3.765800	-4.611600	-0.313900	O	1.676200	-0.560600	-4.137100
C	4.350400	-2.450000	0.932100	O	-0.172100	-1.505000	-2.104900
H	4.074300	-2.071300	-1.186500	O	-1.354100	-0.644300	-4.251900
H	2.648900	-1.788400	-0.213700	H	-3.664500	-4.468800	-0.964800
C	4.744600	-1.054400	1.368400	H	-2.470400	-5.297500	-0.196300
H	3.813800	-2.932500	1.748600	H	-4.106900	-5.516600	0.240800
H	5.252400	-3.055900	0.780500	H	5.792200	-0.891700	-0.423500
H	3.915900	-0.353700	1.243000	H	5.812300	0.501400	0.450600
H	5.073800	-1.018200	2.407500	H	6.775300	-0.833400	0.908300
C	-4.891300	-0.588600	-0.676500	N	5.871300	-0.546000	0.538500
C	-3.900200	-1.744700	-0.518000	N	-3.404200	-4.836800	-0.043400
H	-4.975500	-0.292300	-1.725800	H	-3.379000	4.760300	0.426300
H	-5.884800	-0.911000	-0.349500	H	-2.762300	6.216100	1.076500
C	-4.288000	-2.640400	0.656700	H	-2.073200	5.499700	-0.249400
H	-3.861000	-2.327300	-1.447100	N	-2.517300	5.321200	0.657100
H	-2.881200	-1.364800	-0.365900	Fe	0.100800	0.212000	-3.175300
C	-3.273800	-3.728100	0.941700				
H	-4.419700	-2.041200	1.557500				

Table S9. Optimized Cartesian coordinates (Å) of Δ -Fe(III)-[(DHB^DLys^DSer)₃].

N	-2.494900	-2.457300	-0.621000	H	1.816100	5.008500	-2.825500
N	-2.547800	0.674600	2.491100	C	4.666000	4.015000	-1.229000
H	-2.203000	-0.011500	1.835100	H	4.004300	5.891900	-2.065500
H	-2.162000	-1.536900	-0.870600	H	5.616600	4.411000	-0.881800
C	-1.866300	1.866900	2.477400	C	2.195300	0.305700	3.684700
O	-2.336200	2.879600	2.953400	C	3.132300	0.254600	2.629500
C	-0.563200	1.840400	1.671900	C	2.572900	-0.164600	4.960700
H	-0.850400	1.401300	0.707300	C	4.412700	-0.359300	2.825300
N	0.465800	0.959400	2.176900	C	3.821900	-0.718800	5.156700
H	1.265700	0.846400	1.527600	H	1.862300	-0.088200	5.777000
C	-1.821000	-3.035900	0.425800	C	4.738900	-0.827900	4.092600
O	-2.284600	-3.964000	1.055400	H	4.101900	-1.086500	6.139100
C	-0.525900	-2.317300	0.809200	H	5.710600	-1.288200	4.251000
H	-0.822600	-1.267200	0.948500	C	2.209100	-3.226500	-1.572700
N	0.478200	-2.270700	-0.230800	C	3.145500	-2.288400	-1.086700
H	1.268900	-1.643000	-0.007700	C	2.579600	-4.080800	-2.634000
C	-1.827700	1.155200	-2.908700	C	4.418000	-2.134300	-1.728500
O	-2.323400	1.085200	-4.014200	C	3.820200	-3.955400	-3.226200
C	-0.528200	0.451000	-2.516000	H	1.870800	-4.827700	-2.975800
H	-0.803400	-0.185900	-1.665300	C	4.736100	-2.979000	-2.785600
N	0.498300	1.321900	-1.992700	H	4.094900	-4.608500	-4.048900
H	1.287000	0.829900	-1.535000	H	5.701100	-2.872300	-3.274200
C	0.045500	-2.783300	2.144600	O	2.922700	0.885300	-1.275700
C	-0.809200	-2.291500	3.299600	O	5.144300	1.813200	-0.372200
H	1.048900	-2.350800	2.232000	O	2.916100	0.730600	1.430900
H	0.139700	-3.873200	2.155600	O	5.181800	-0.440100	1.755200
C	-0.145000	-2.458300	4.654700	O	2.935200	-1.507700	-0.060500
H	-0.982500	-1.220300	3.135000	O	5.188800	-1.172300	-1.255700
H	-1.780900	-2.796800	3.283400	H	-2.197700	4.240000	-1.965900
C	-1.028300	-1.995800	5.797900	H	-2.093000	5.471700	-3.053000
H	0.097400	-3.509900	4.838000	H	-0.785500	4.441000	-2.758600
H	0.798300	-1.903700	4.665300	H	-2.226400	-3.859500	-2.728100
H	-1.905200	-2.635800	5.907800	H	-2.079200	-5.430200	-3.193600
H	-0.479600	-1.979500	6.739700	H	-0.784800	-4.597300	-2.494900
C	-0.020400	-0.500700	-3.593600	H	-2.205600	-0.579500	4.802900
C	-0.899600	-1.741600	-3.653500	H	-2.009700	-0.251300	6.401900
H	0.997000	-0.796200	-3.314100	H	-0.785500	0.070800	5.284300
H	0.022400	0.008100	-4.561300	N	-1.536900	-0.608900	5.573400
C	-0.292400	-2.870400	-4.469300	N	-1.584800	4.975500	-2.322400
H	-1.021700	-2.094000	-2.622200	N	-1.596400	-4.533500	-3.164600
H	-1.892900	-1.479900	-4.035800	N	-2.472300	1.761900	-1.858500
C	-1.187900	-4.093300	-4.531700	H	-2.116900	1.508800	-0.947300
H	-0.111900	-2.545400	-5.499000	C	-4.650700	0.574200	-2.121400
H	0.678500	-3.143100	-4.041100	C	-4.412400	2.547300	-0.631000
H	-2.106500	-3.890300	-5.085300	O	-5.788400	0.515500	-2.499900
H	-0.675800	-4.937800	-4.993500	O	-3.899700	-0.473400	-1.778600
C	-0.031300	3.237800	1.366000	H	-5.502600	2.565900	-0.608200
C	-0.880600	3.916100	0.301600	H	-4.027300	3.564100	-0.531800
H	0.990100	3.120500	0.986400	O	-3.895400	1.774700	0.461600
H	0.008200	3.835500	2.281700	C	-4.449200	-1.793500	-1.887500
C	-0.268600	5.204100	-0.223000	C	-4.663200	1.598400	1.538800
H	-0.962200	3.215100	-0.535400	H	-5.539500	-1.760000	-1.889400
H	-1.891000	4.095700	0.688100	H	-4.094600	-2.222800	-2.827800
C	-1.158700	5.904500	-1.232800	C	-3.933700	-2.591200	-0.699300
H	-0.093600	5.908400	0.596700	O	-5.774200	2.034300	1.671700
H	0.704400	4.982500	-0.673900	C	-3.991900	0.708900	2.586600
H	-2.072300	6.276800	-0.765600	C	-4.679100	-2.139600	0.560500
H	-0.640400	6.736200	-1.711500	H	-4.211400	-3.637300	-0.846400
C	0.868400	-3.330200	-0.979200	H	-4.300800	1.110300	3.554200
O	0.110600	-4.303800	-1.170400	O	-5.790400	-2.504700	0.828700
C	0.843000	0.844500	3.469800	O	-3.973800	-1.257600	1.268000
O	0.066900	1.134600	4.403700	C	-4.552700	-0.697600	2.456900
C	0.860100	2.518100	-2.507200	H	-5.640900	-0.675900	2.383100
O	0.096000	3.165300	-3.255700	H	-4.258400	-1.315700	3.306400
C	2.180000	3.016200	-2.090300	C	-3.910800	1.897100	-1.910200
C	3.115700	2.162300	-1.470800	H	-4.199600	2.534400	-2.748700
C	2.527300	4.365100	-2.318400	Fe	4.257500	0.054900	0.038700
C	4.369000	2.676000	-1.002100				
C	3.747600	4.851200	-1.893800				

Table S10. Optimized Cartesian coordinates (Å) of Δ -Fe(III)-[(DHB^LLys^DSer)₃].

N	-1.226600	2.549900	2.237700	H	5.278400	-2.955300	-2.047700
N	-1.607900	-2.331700	2.254200	C	2.864100	-2.983600	-4.456500
H	-1.443100	-1.560700	1.621800	H	4.620900	-4.185600	-4.093000
H	-0.649900	2.009200	1.607900	H	2.598000	-3.503000	-5.373300
C	-1.234500	-3.571400	1.846100	C	-3.201600	-2.270300	-2.096100
O	-1.167100	-4.522200	2.610500	C	-2.286300	-1.464000	-2.810000
C	-0.807100	-3.694100	0.396800	C	-4.521900	-2.401700	-2.579600
H	-0.937500	-4.747800	0.129200	C	-2.685900	-0.823400	-4.031800
N	-1.522100	-2.835700	-0.510500	C	-4.914200	-1.751400	-3.730900
H	-0.981700	-2.216200	-1.139500	H	-5.210300	-3.046200	-2.043400
C	-2.487400	2.848200	1.832400	C	-4.000300	-0.965500	-2.460400
O	-3.340400	3.273000	2.596800	H	-5.932200	-1.857600	-4.092900
C	-2.811100	2.535000	0.384700	H	-4.310900	-0.472500	-5.377800
H	-3.656300	3.177200	0.116400	C	-0.363700	3.894000	-2.095100
N	-1.709600	2.719900	-0.523000	C	-0.113300	2.699400	-2.807200
H	-1.438300	1.940600	-1.147500	C	0.175800	5.106600	-2.577800
C	3.713600	0.727600	1.843200	C	0.646600	2.729400	-4.025900
O	4.499300	1.262600	2.610700	C	0.937800	5.125800	-3.727200
C	3.608700	1.162200	0.394300	H	-0.047000	6.024300	-2.043900
H	4.587500	1.575000	0.129200	C	1.172900	3.942300	-4.454400
N	3.224900	0.115100	-0.515700	H	1.348900	6.063300	-4.089000
H	2.414400	0.266300	-1.141000	H	1.758100	3.968100	-5.369800
C	-3.191800	1.058500	0.260400	O	1.608900	-0.289200	-2.448600
C	-4.353700	0.554200	1.097800	O	1.002400	-1.471100	-4.653200
H	-2.296800	0.467600	0.496200	O	-1.041700	-1.260100	-2.454600
H	-3.409200	0.872700	-0.798200	O	-1.749300	-0.139300	-4.660300
C	-4.545400	-0.908300	0.730100	O	-0.555500	1.517600	-2.454000
H	-4.138600	0.675800	2.164300	O	0.781500	1.576900	-4.653200
H	-5.262300	1.126600	0.880900	H	6.299200	-3.702900	1.340900
C	-5.628400	-1.645100	1.485100	H	5.623600	-4.230300	-0.060700
H	-4.758800	-0.967300	-0.343600	H	-0.015100	7.250300	1.436600
H	-3.603000	-1.442300	0.887400	H	0.784300	7.038900	0.019000
H	-5.329400	-1.884200	2.506900	H	-6.329300	-3.625800	1.391000
H	-6.568600	-1.092600	1.515800	H	-6.500100	-2.792900	-0.014700
C	2.518200	2.227800	0.265800	N	-5.882400	-2.941500	0.782700
C	2.663000	3.490000	1.096900	N	5.473300	-3.628800	0.748700
H	1.559800	1.749300	0.506300	N	0.369000	6.546000	0.808600
H	2.464300	2.504200	-0.794000	N	2.830000	-0.220900	2.247100
C	1.493500	4.387700	0.726000	H	2.077600	-0.456200	1.614900
H	2.661700	3.248000	2.164700	C	1.869000	0.702400	4.341100
H	3.613700	3.988500	0.877800	C	2.014600	-1.762000	3.984900
C	1.395900	5.692000	1.483800	O	1.837800	0.829500	5.535500
H	1.549700	4.603500	-0.347300	O	1.241400	1.493200	3.462200
H	0.559200	3.839000	0.881400	H	2.010300	-1.912100	5.064500
H	1.067300	5.545200	2.513900	H	2.575400	-2.558200	3.494300
H	2.337000	6.244100	1.492400	O	0.675800	-1.825900	3.471500
C	0.660000	-3.279300	0.268200	C	0.519600	2.625900	3.969200
C	1.683200	-4.031000	1.100200	C	-0.328000	-1.950200	4.348100
H	0.720600	-2.208900	0.505400	H	0.655700	2.705800	5.047700
H	0.926700	-3.372700	-0.791400	H	0.926300	3.505400	3.469300
C	3.045100	-3.480600	0.708600	C	-0.954000	2.495600	3.646600
H	1.483300	-3.895000	2.167800	O	-0.208000	-1.959800	5.543700
H	1.631100	-5.105800	0.894700	C	-1.686700	-2.068000	3.663300
C	4.227900	-4.040000	1.466300	C	-1.536200	1.269000	4.342600
H	3.192400	-3.658800	-0.363400	H	-1.448900	3.349900	4.117700
H	3.043700	-2.394500	0.841900	H	-2.169900	-2.926100	4.139400
H	4.293700	-3.641800	2.480000	O	-1.608800	1.180300	5.538800
H	4.216500	-5.129900	1.517500	O	-1.927700	0.330500	3.472600
C	-1.228400	3.914900	-0.904600	C	-2.538300	-0.859800	3.992800
O	-1.519600	4.957800	-0.274900	H	-2.660700	-0.778300	5.072800
C	-2.794200	-3.024600	-0.899900	H	-3.510200	-0.951100	3.507000
O	-3.553900	-3.797800	-0.272500	C	2.640700	-0.422500	3.656500
C	4.025700	-0.893400	-0.899500	H	3.626000	-0.419100	4.131700
O	5.073200	-1.162400	-0.267600	Fe	0.007600	-0.011100	-3.699400
C	3.581400	-1.628800	-2.094000	H	5.386900	-2.637800	0.379700
C	2.419100	-1.254900	-2.805600	H	-0.420900	5.955500	0.416900
C	4.369100	-2.695700	-2.579100	H	-4.980200	-3.345900	0.397800
C	2.070400	-1.926500	-4.026600				
C	4.008800	-3.365900	-3.729400				

Table S11. Optimized Cartesian coordinates (Å) of Δ -Fe(III)-[(DHB^DLys^LSer)₃].

O	2.708200	-0.681000	5.512500	H	-0.021500	3.464500	1.525400
O	-0.777000	2.348600	5.647900	H	-0.332200	5.891900	2.140400
O	-1.859900	-1.974000	5.567700	H	-0.872600	6.193200	0.473900
C	2.231900	-0.365600	4.454600	C	2.005500	-3.755900	-0.251000
C	-0.803700	2.001500	4.497100	C	3.291000	-3.608000	0.561700
C	-1.444100	-1.767400	4.458800	H	2.186500	-3.402600	-1.272200
C	1.564700	2.028000	4.314200	H	1.704700	-4.805800	-0.304700
C	-2.588800	0.306600	4.293600	C	3.737700	-2.155000	0.679700
C	0.949800	-2.352400	4.330800	H	3.156400	-4.034800	1.560500
H	1.506100	1.990500	5.402900	C	4.078800	-4.187200	0.066700
H	1.904800	3.013500	3.993000	C	5.120300	-2.031100	1.328000
H	-2.551400	0.263600	5.382800	H	3.726100	-1.685400	-0.310200
H	-3.606700	0.126200	3.946800	H	3.046900	-1.592500	1.317200
H	0.933400	-2.298300	5.420500	H	5.205500	-2.593300	2.214100
H	1.663000	-3.111700	4.006300	H	5.893300	-2.370700	0.590700
O	-1.731800	-0.696000	3.719900	C	-2.470300	2.788300	-1.069900
O	1.343700	-1.087200	3.772500	O	-3.179400	3.544800	-0.366900
O	0.269100	1.799300	3.731700	C	-1.145700	-3.539300	-1.044900
C	-0.439600	-2.681900	3.783900	O	-1.452100	-4.521300	-0.328800
H	-0.665200	-3.698800	4.110700	C	3.635900	0.765900	-1.072300
C	-2.105100	1.667300	3.790900	O	4.644000	1.007500	-0.370000
H	-2.842600	2.396700	4.132600	C	3.705800	0.021100	-2.338700
C	2.526500	0.962800	3.782600	C	2.609300	-0.019600	-3.226800
H	3.524000	1.262800	4.108400	C	4.897200	-0.656700	-2.679600
N	-2.014800	1.712500	2.341100	C	2.700600	-0.755400	-4.456700
N	2.498800	0.881500	2.330900	C	4.981300	-1.362000	-3.862600
N	-0.482200	-2.620700	2.332700	H	5.736000	-0.609100	-1.993100
H	-0.571800	-1.691000	1.948600	C	3.888000	-1.414300	-4.751200
H	-1.164600	1.330800	1.952700	H	5.897000	-1.887600	-4.116000
H	1.734000	0.341700	1.951900	H	3.961600	-1.977300	-5.678100
C	0.341000	-3.473400	1.639200	C	-1.839400	-3.222600	-2.302600
O	0.726600	-4.520700	2.120900	C	-1.330900	-2.256900	-3.198700
C	0.847000	-2.929900	0.301900	C	-3.028200	-3.911200	-2.630500
H	1.230600	-1.931800	0.532100	C	-2.019900	-1.974300	-4.426700
N	-0.178300	-2.681000	-0.682300	C	-3.690000	-3.633200	-3.808800
H	0.057900	-1.997900	-1.420800	H	-3.403200	-4.658900	-1.939400
C	-3.158200	1.458000	1.624200	C	-3.189900	-2.669200	-4.707800
O	-4.262700	1.662100	2.089100	H	-4.607500	-4.160600	-4.052200
C	-2.928300	0.763500	0.283000	H	-3.719200	-2.456200	-5.633100
H	-2.265900	-0.076400	0.514100	C	-1.863000	3.214900	-2.338900
N	-2.180600	1.533500	-0.681700	C	-1.291800	2.277900	-3.227700
H	-1.710700	0.982800	-1.419400	C	-1.869000	4.583600	-2.687200
C	2.832300	2.012200	1.624500	C	-0.713700	2.714800	-4.467600
O	3.553100	2.869500	2.095800	C	-1.310400	4.999600	-3.878100
C	2.108400	2.165000	0.288000	H	-2.319300	5.292100	-1.999800
H	1.052800	2.002200	0.526600	C	-0.734400	4.071000	-4.769300
N	2.403300	1.141800	-0.686200	H	-1.309700	6.054100	-4.137000
H	1.691900	1.009400	-1.423900	H	-0.293200	4.409600	-5.703200
C	-4.215600	0.190900	-0.301600	O	1.467700	0.582700	-3.014300
C	-4.775300	-0.983300	0.500500	O	1.618400	-0.754400	-5.213600
H	-3.969200	-0.156100	-1.311400	O	-0.236700	-1.569500	-2.996700
H	-4.958600	0.988400	-0.389600	O	-1.476500	-1.048100	-5.195700
C	-3.722700	-2.059100	0.755600	O	-1.241800	0.989600	-3.006700
H	-5.181700	-0.631800	1.453600	O	-0.187300	1.769500	-5.225100
H	-5.606500	-1.417900	-0.066100	H	4.762300	-0.262500	2.311800
C	-4.344100	-3.360600	1.210000	H	6.353700	-0.461800	1.918900
H	-3.148500	-2.231400	-0.162200	H	5.218700	0.015900	0.749200
H	-3.018300	-1.726500	1.528000	H	-2.102900	4.286900	2.355000
H	-4.977800	-3.218200	2.087400	H	-2.734100	5.745200	1.908300
H	-4.935800	-3.818100	0.415200	H	-2.597300	4.482100	0.783100
C	2.246000	3.572100	-0.285100	H	-2.710200	-3.964900	2.342800
C	1.507200	4.632400	0.530000	H	-3.659100	-5.226100	1.864000
H	1.819400	3.539900	-1.293800	N	-3.273600	-4.328500	1.574600
H	3.306600	3.822800	-0.374500	N	5.399500	-0.600900	1.590400
C	0.048400	4.256800	0.770700	N	-2.141400	4.974900	1.603300
H	2.009600	4.791500	1.488600	Fe	-0.009400	-0.006800	-4.307200
H	1.550500	5.578800	-0.021000	H	-2.601100	-4.477400	0.767500
C	-0.769900	5.434100	1.251300				
H	-0.385000	3.866200	-0.157300				

