

## Supporting information

### Weakly acidic pH-responsive liposomal content release induced by histidine-modified agents

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- Schematic representation of Cn-His solid phase synthesis (example for C8-His)
- Identification of a series of histidine-modified pH-sensitive Cn-His ( $n = 8, 12, 18$ ) agents

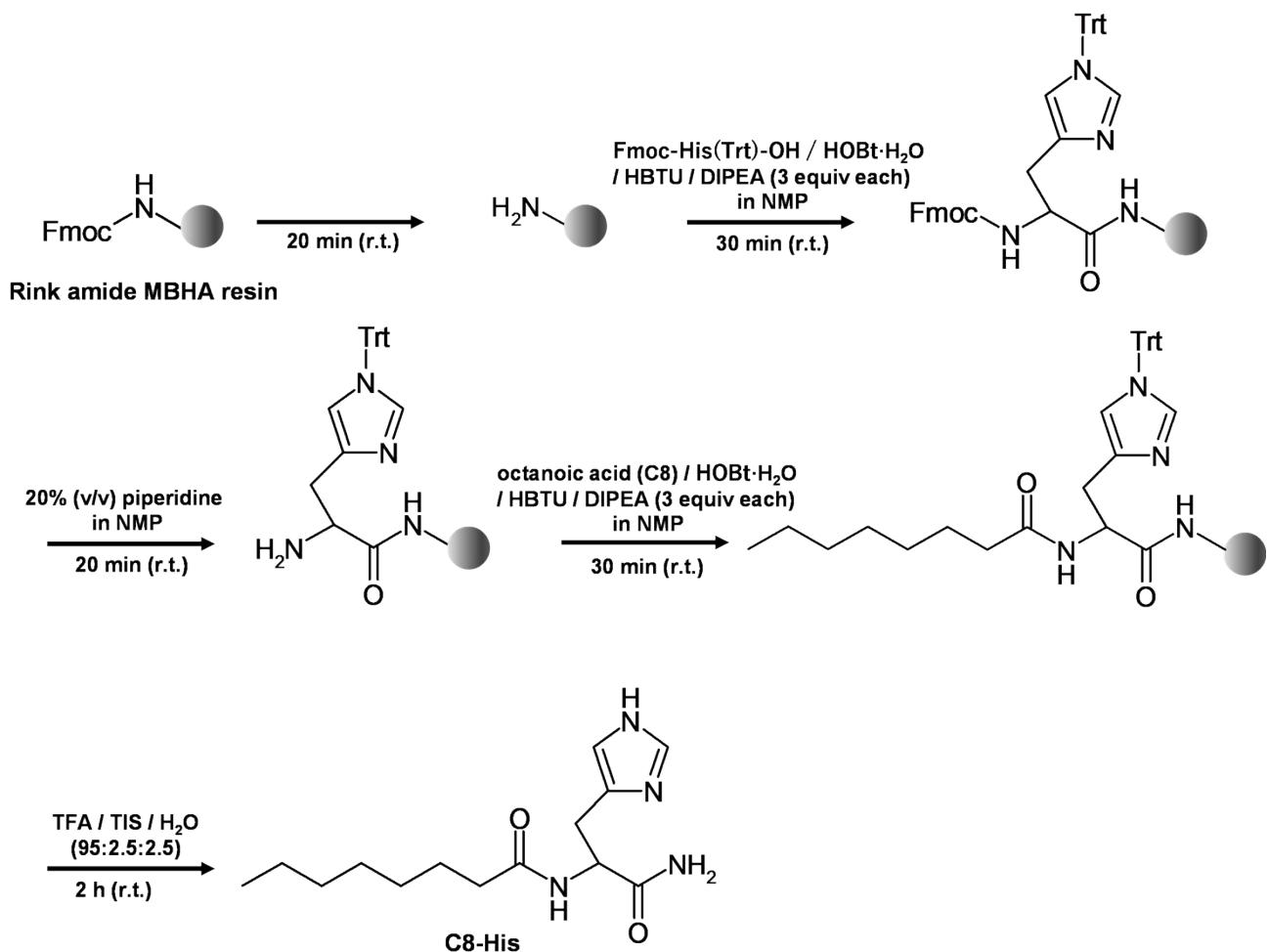
#### HPLC chromatogram profile

#### <sup>1</sup>HNMR

#### High-Resolution ESI-TOF MS

- Characterization of Cn-His incorporated liposomes
  - Hydrodynamic diameter (Dh) and  $\zeta$ -potential values (Table S1)
- pH-switch mode for content release from C8-His-incorporated liposome
  - Release profiles of CF and  $\zeta$ -potential (Fig. S1)

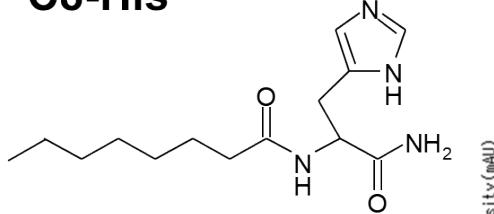
## Schematic representation of Cn-His solid phase synthesis (example for C8-His)



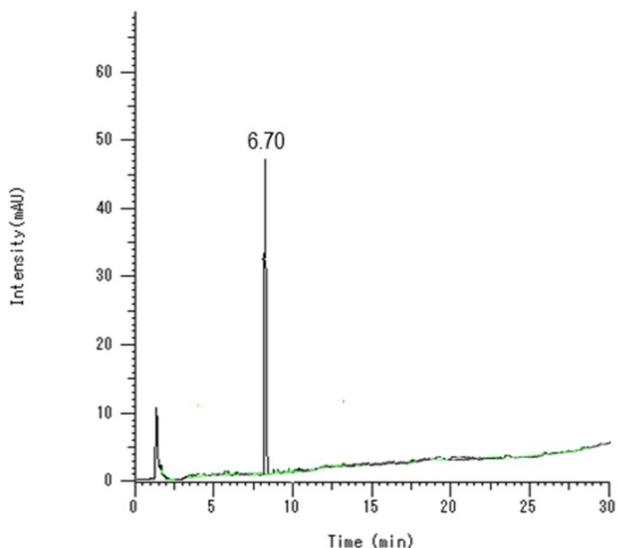
## Identification of C8-His

HPLC chromatogram profile of purified C8-His

### C8-His

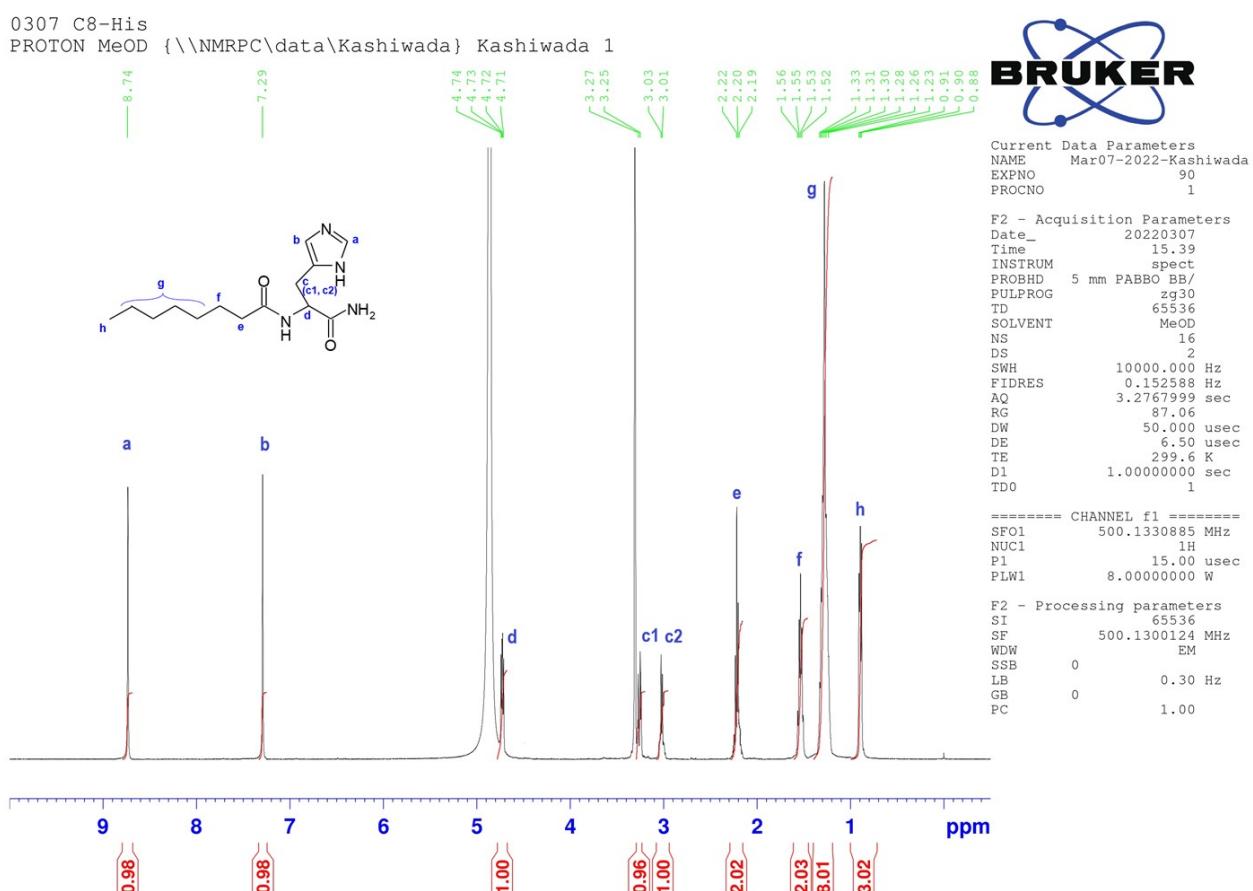


**Column:** Inertsil ODS-3  
(5  $\mu$ m, 250  $\times$  4.6 mm i.d.)  
**Eluent:**  
A: 0.1% TFA / H<sub>2</sub>O  
B: 0.1% TFA / acetonitrile  
Linear gradient of 5–95% (v/v)  
eluent B over 30 min  
**Flow rate:** 1.0 mL/min  
**Temp.:** 25°C  
**Detection:** UV at 230 nm



### <sup>1</sup>H NMR (500 MHz, MeOD, $\delta$ in ppm)

8.74 (1 H, s), 7.29 (1 H, s), 4.73 (1 H, m), 3.26 (1 H, d), 3.02 (1 H, d), 2.20 (2 H, t), 1.54 (2 H, q), 1.33 – 1.23 (8 H, m), 0.90 (3 H, t).



## High-Resolution ESI-TOF MS

### Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

888 formula(e) evaluated with 8 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-500 H: 0-1000 N: 0-200 O: 0-200 Na: 0-1

25-Jan-2024

C8-His 2ppm nega

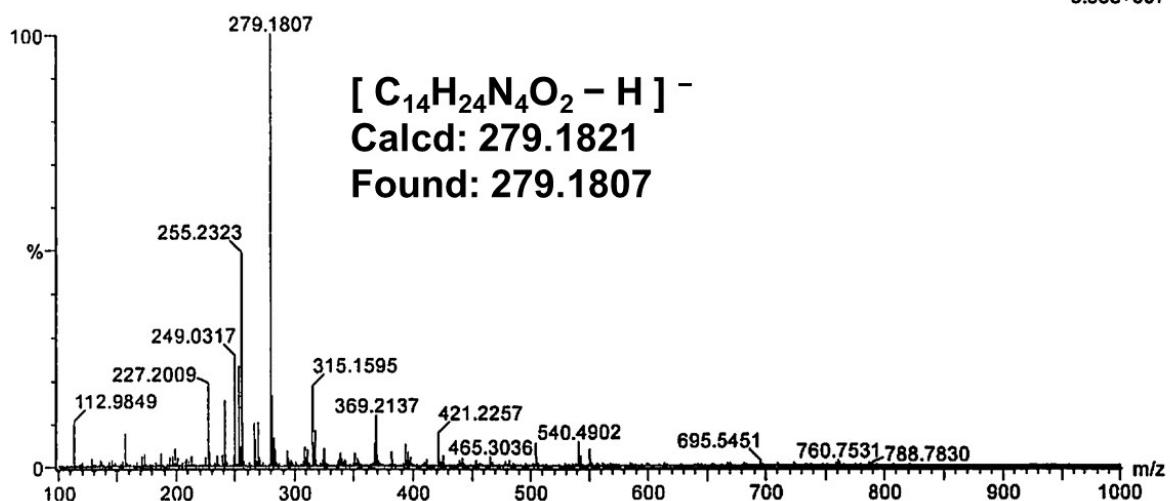
XEVO-G2SQTOF#YDA064

2024\_0125\_01 3 (0.129) AM2 (Ar,30000.0,0.00,0.00); Cr (2:15)

1: TOF MS ES-

18:03:10

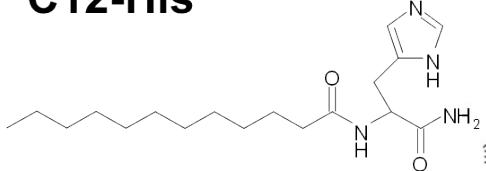
6.88e+007



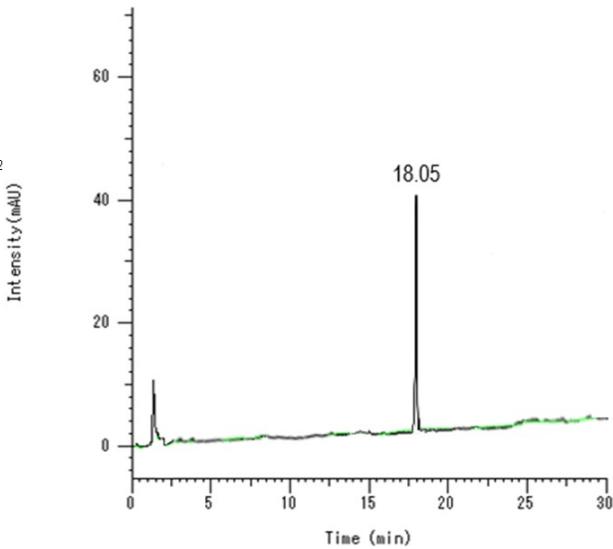
## Identification of C12-His

## HPLC chromatogram profile of purified C12-His

## C12-His

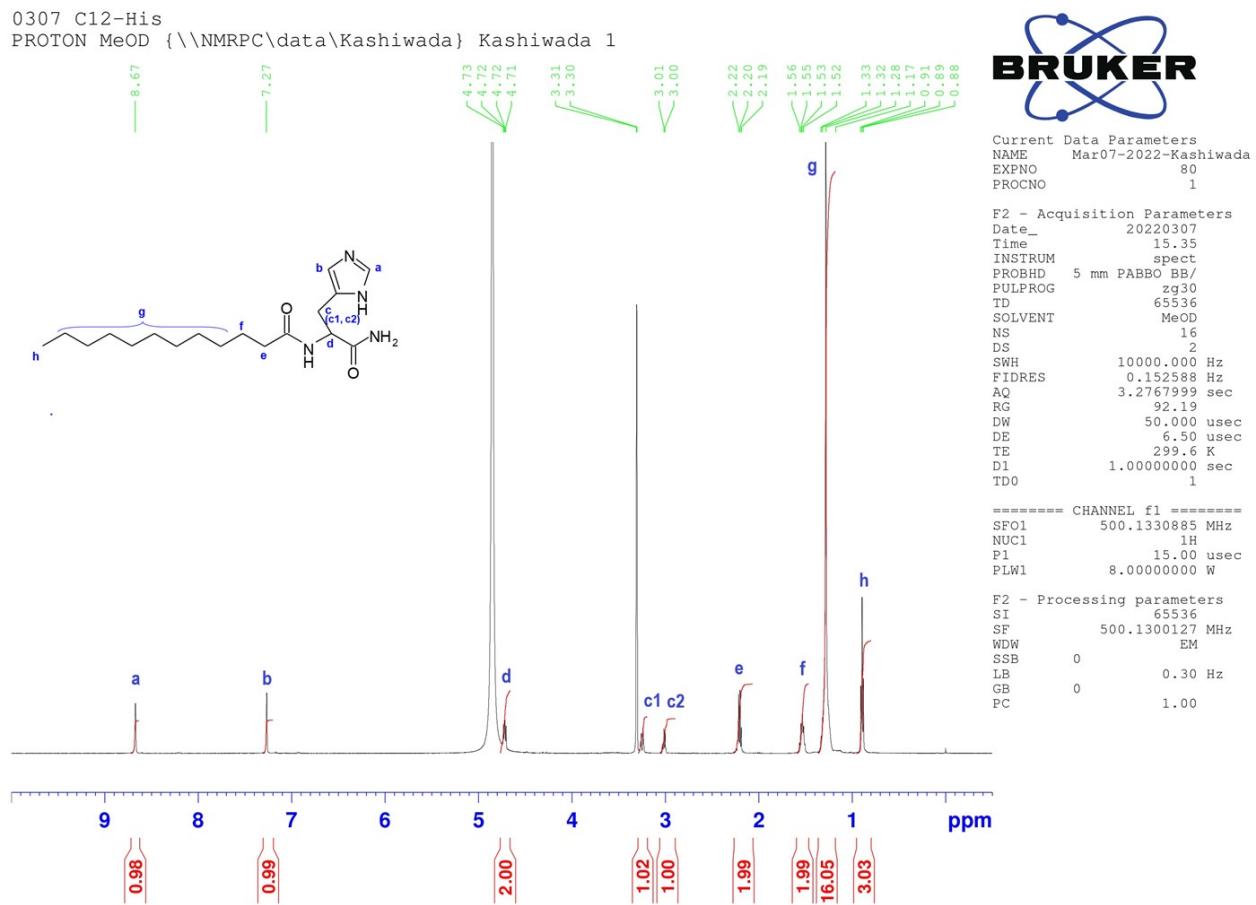


Column	Inertsil ODS-3 (5 µm, 250 × 4.6 mm i.d.)
Eluent	A: 0.1% TFA / H <sub>2</sub> O B: 0.1% TFA / acetonitrile Linear gradient of 5–95% (v/v) eluent B over 30 min
Flow rate	1.0 mL/min
Temp.	25 °C
Detection	UV at 230 nm



**<sup>1</sup>H NMR (500 MHz, MeOD, δ in ppm)**

8.67 (1 H, s), 7.27 (1 H, s), 4.72 (1 H, m), 3.31 (1 H, d), 3.01 (1 H, d), 2.21 (2 H, t), 1.54 (2 H, q), 1.33 – 1.17 (16 H, m), 0.89 (3 H, t).



## High-Resolution ESI-TOF MS

### Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

1423 formula(e) evaluated with 11 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-500 H: 0-1000 N: 0-200 O: 0-200 Na: 0-1

20-Jan-2024

15:18:03

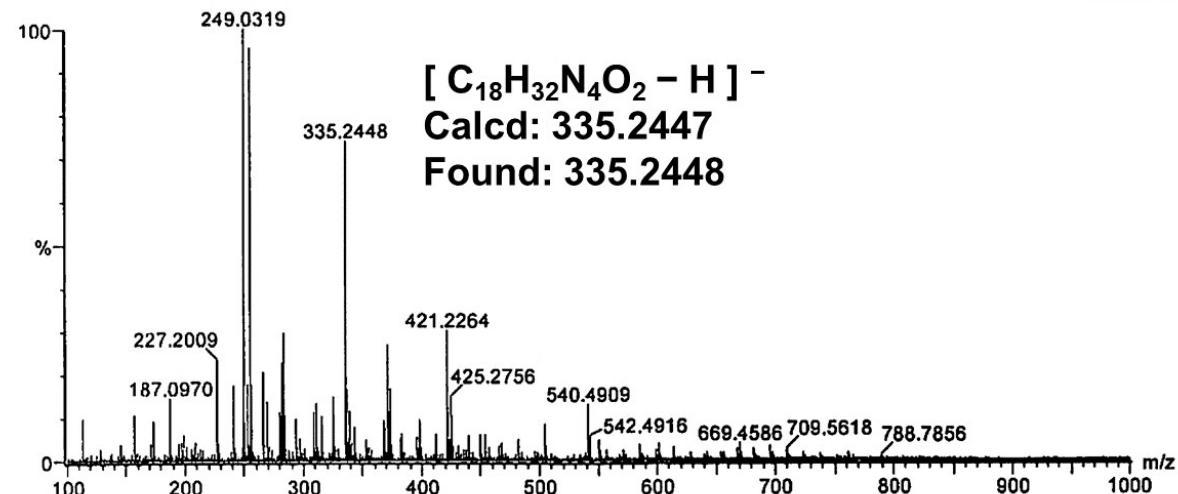
C12-His 2ppm nega

XEVO-G2SQTOF#YDA064

2024\_0120\_05 4 (0.174) AM2 (Ar,30000.0,0.00,0.00); Cm (2:14)

1: TOF MS ES-

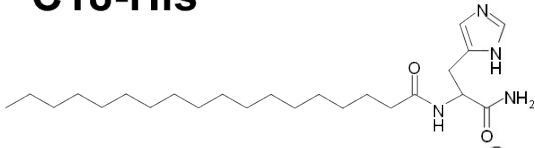
4.87e+007



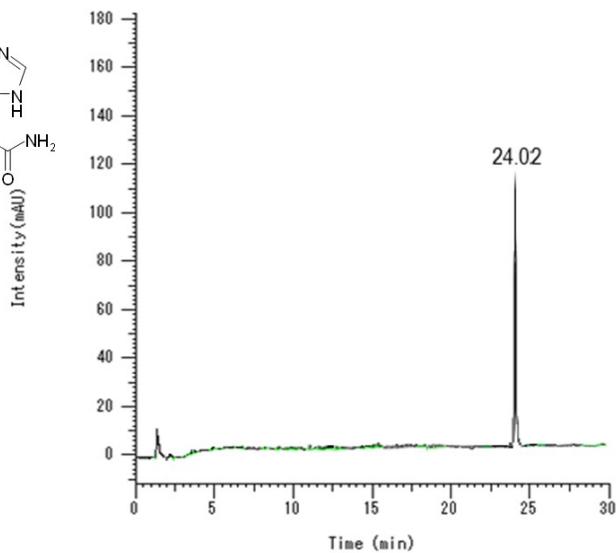
## Identification of C18-His

HPLC chromatogram profile of purified C18-His

### C18-His

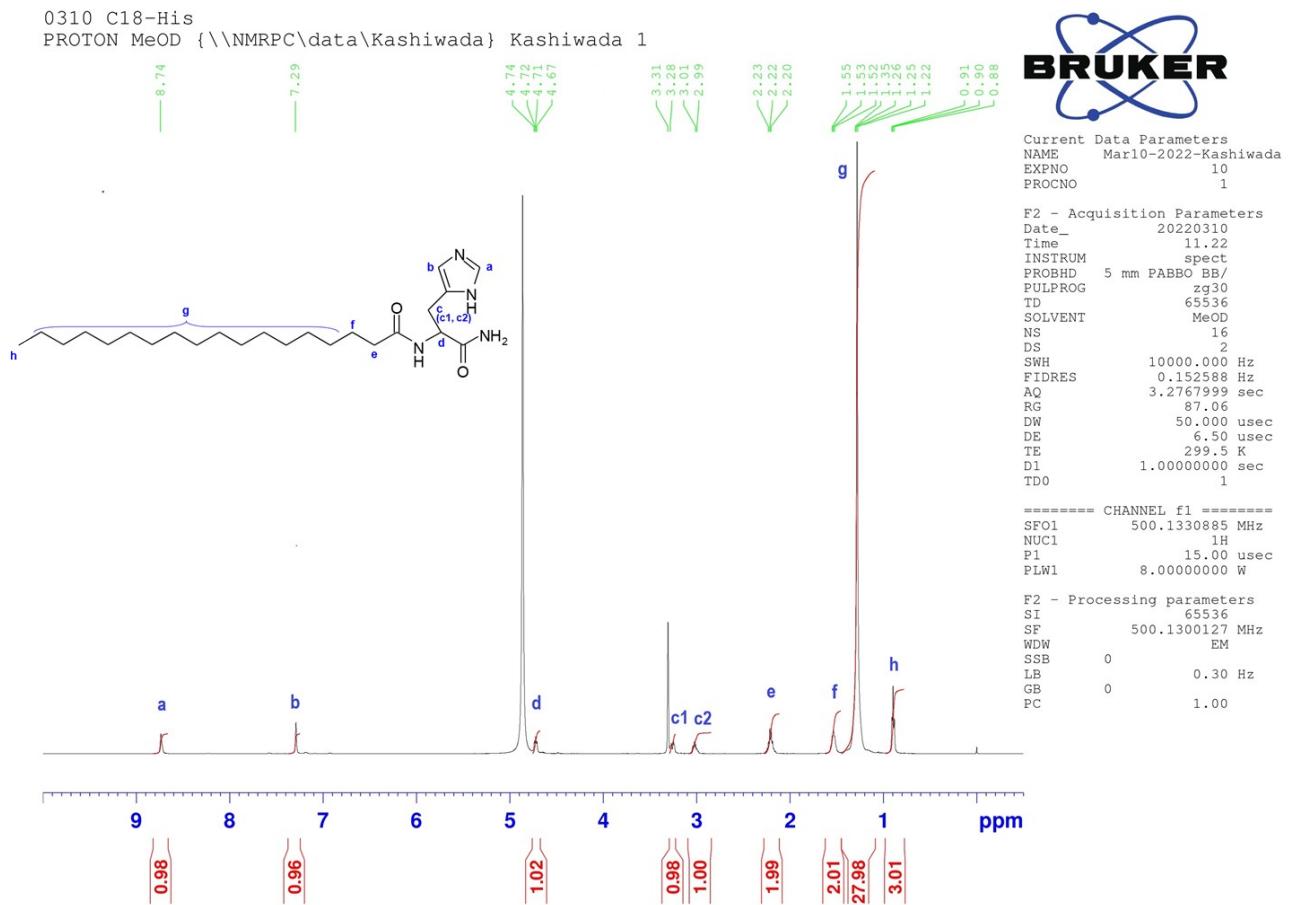


**Column** Inertsil ODS-3  
(5  $\mu$ m, 250  $\times$  4.6 mm i.d.)  
**Eluent** A: 0.1% TFA / H<sub>2</sub>O  
B: 0.1% TFA / acetonitrile  
Linear gradient of 5–95% (v/v)  
eluent B over 30 min  
**Flow rate** 1.0 mL/min  
**Temp.** 25°C  
**Detection** UV at 230 nm



### <sup>1</sup>H NMR (500 MHz, MeOD, $\delta$ in ppm)

8.74 (1 H, s), 7.29 (1 H, s), 4.71 (1 H, m), 3.30 (1 H, d), 3.00 (1 H, d), 2.22 (2 H, t), 1.53 (2 H, q), 1.35 – 1.22 (28 H, m), 0.90 (3 H, t).



## High-Resolution ESI-TOF MS

### Single Mass Analysis

Tolerance = 5.0 mDa / DBE: min = -1.5, max = 50.0

Element prediction: Off

Number of isotope peaks used for i-FIT = 3

Monoisotopic Mass, Even Electron Ions

3003 formula(e) evaluated with 16 results within limits (up to 50 closest results for each mass)

Elements Used:

C: 0-500 H: 0-1000 N: 0-200 O: 0-200 Na: 0-1

30-Jan-2024 pos?

C18-His 2ppm nega HFIP/AN/MeOH

10:04:50

XEVO-G2SQTOF#YDA064

2024\_0130\_02 7 (0.286) AM2 (Ar,30000.0,0.00,0.00); Cm (2:13)

1: TOF MS ES+

2.59e+008

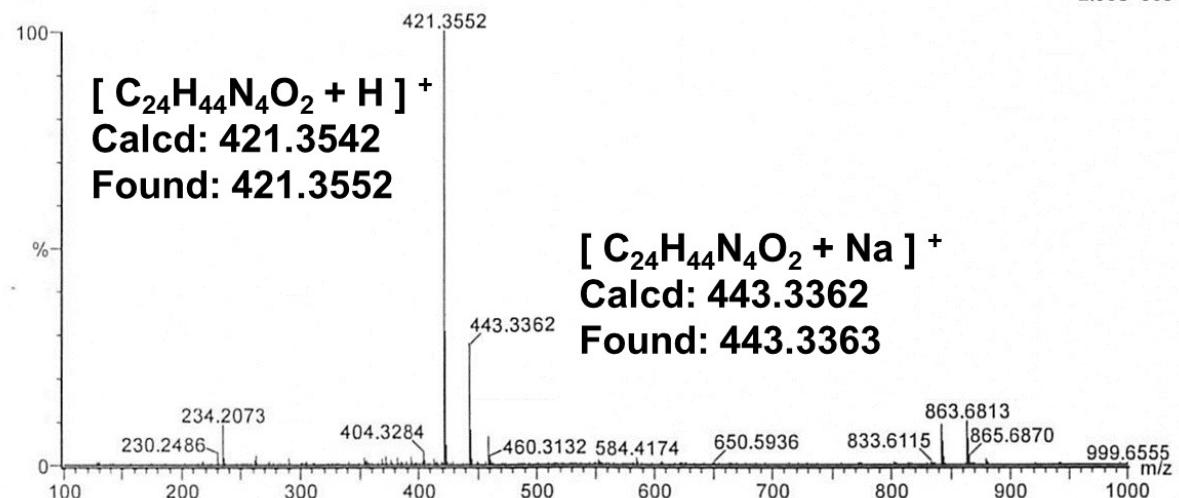


Table S1 Hydrodynamic diameter (Dh) from DLS measurements and  $\zeta$ -potential values of bare DOPC liposomes and 1.0% of Cn-His incorporated liposomes

Liposome	pH	Dh / nm	PDI	$\zeta$ -potential / mV
DOPC (bare)	5.5	119.8 ± 9.6	0.101 ± 0.027	-0.48 ± 0.40
	6.0	121.4 ± 1.2	0.144 ± 0.018	-2.12 ± 0.81
	6.5	123.2 ± 0.9	0.102 ± 0.004	-3.32 ± 0.66
	7.0	121.7 ± 2.8	0.150 ± 0.031	-4.45 ± 0.77
	7.5	124.2 ± 1.0	0.164 ± 0.025	-5.60 ± 0.76
	8.0	121.4 ± 1.8	0.163 ± 0.025	-8.55 ± 0.46
C8-His / DOPC	5.5	122.6 ± 5.9	0.106 ± 0.043	+3.06 ± 0.72
	6.0	110.0 ± 2.7	0.155 ± 0.034	+0.42 ± 0.28
	6.5	113.8 ± 1.1	0.125 ± 0.022	-1.19 ± 0.39
	7.0	116.5 ± 1.0	0.062 ± 0.018	-2.47 ± 0.56
	7.5	116.6 ± 4.6	0.126 ± 0.022	-4.57 ± 0.57
	8.0	122.4 ± 8.7	0.138 ± 0.051	-8.28 ± 1.04
C12-His / DOPC	5.5	119.6 ± 1.5	0.088 ± 0.027	+4.83 ± 0.73
	6.0	116.3 ± 7.1	0.120 ± 0.034	+1.76 ± 0.85
	6.5	115.0 ± 1.2	0.152 ± 0.033	+0.16 ± 0.88
	7.0	113.9 ± 1.5	0.170 ± 0.023	-1.51 ± 0.48
	7.5	115.7 ± 1.0	0.150 ± 0.061	-2.79 ± 0.62
	8.0	115.3 ± 0.5	0.115 ± 0.006	-7.59 ± 0.84
C18-His / DOPC	5.5	110.4 ± 1.3	0.126 ± 0.040	+8.24 ± 0.84
	6.0	105.7 ± 1.7	0.098 ± 0.015	+5.43 ± 0.61
	6.5	108.8 ± 0.6	0.109 ± 0.041	+0.48 ± 0.62
	7.0	113.8 ± 2.1	0.122 ± 0.026	-0.75 ± 0.23
	7.5	113.4 ± 1.3	0.145 ± 0.037	-2.30 ± 0.33
	8.0	113.8 ± 3.2	0.133 ± 0.051	-6.24 ± 0.51

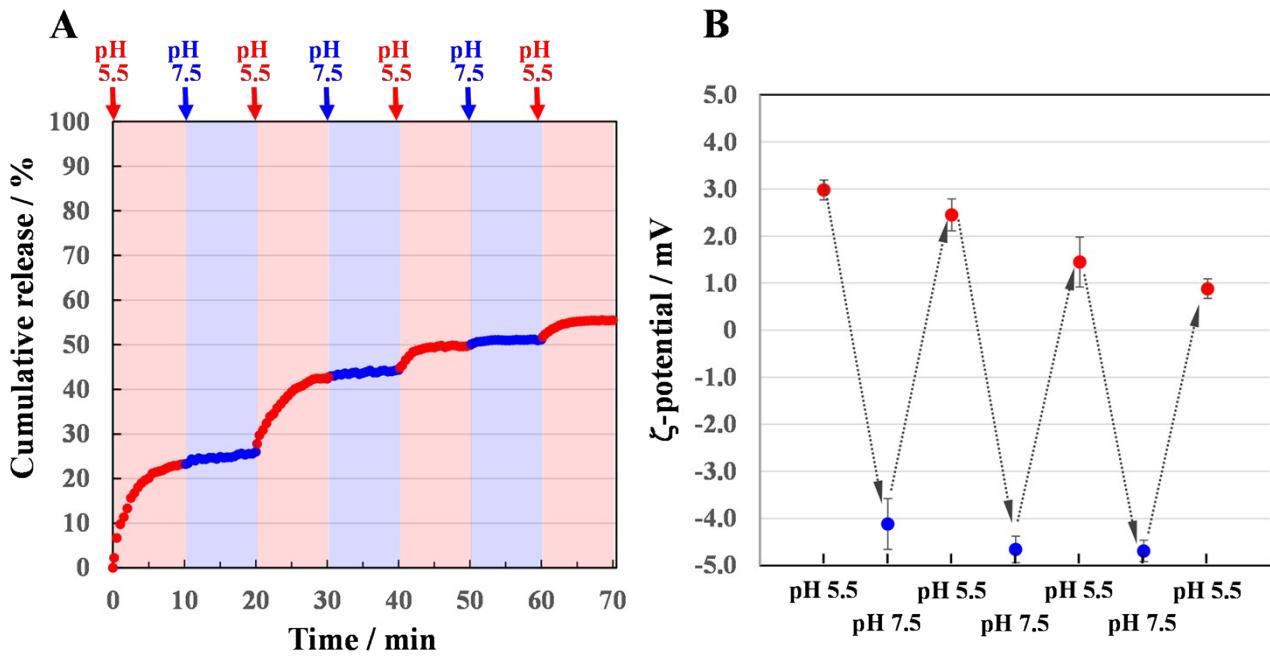


Fig. S1 A: Release profiles of CF from 1.0% of C8-His incorporated liposomes under intermittent pH between 5.5 and 7.5 (Red area: pH 5.5; Blue area: pH 7.5). B: The  $\zeta$ -potential values of 1.0% of C8-His incorporated liposomes under intermittent pH between 5.5 and 7.5. Error bars indicate standard error over three measurements.