

## ELECTRONIC SUPPLEMENTARY INFORMATION

### Phenylalanine and Tryptophan-based surfactants as new antibacterial agents: characterization, self-aggregation properties, and DPPC/Surfactants vesicles formulation

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<sup>3</sup> Centre des Sciences et Technologies de la Formulation

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**Figure.S43** A) Scattered intensity patterns as a function of scattering vector modulus for DPPC and  $\text{C}_{14}\text{PC}_3\text{NH}_3\text{Cl}$  and their mixtures the curves correspond to the best fit of Gaussian bilayers or core-shell models. B) The corresponding electron density profiles of the bilayer models corresponding to the best fits of A).

**Figure.S44.** the core-shell model vesicle model

**Figure. S45.** Hemolysis (%) as a function of surfactants concentration.

**Figure S46:** Three-dimensional (3 D) closest interactions between active site residues of peptidoglycan glycosyltransferase (PDB ID:2OQO) With CnBenzalkonium (From C<sub>8</sub> to C<sub>14</sub> carbon atoms) derivatives.

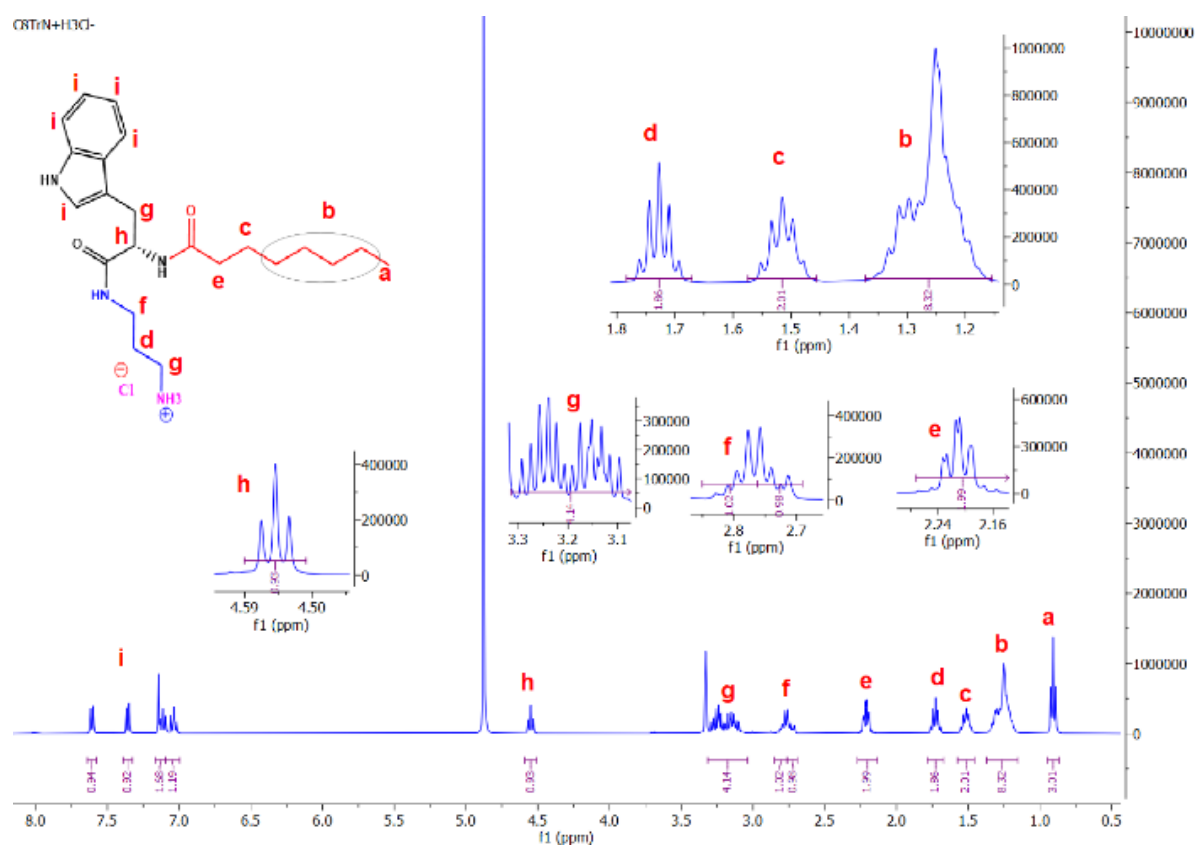
## Tables

**Table S1** Fitting parameters of Gaussian bilayers for DPPC/C<sub>12</sub>PC<sub>3</sub>NH<sub>3</sub>Cl mixtures.

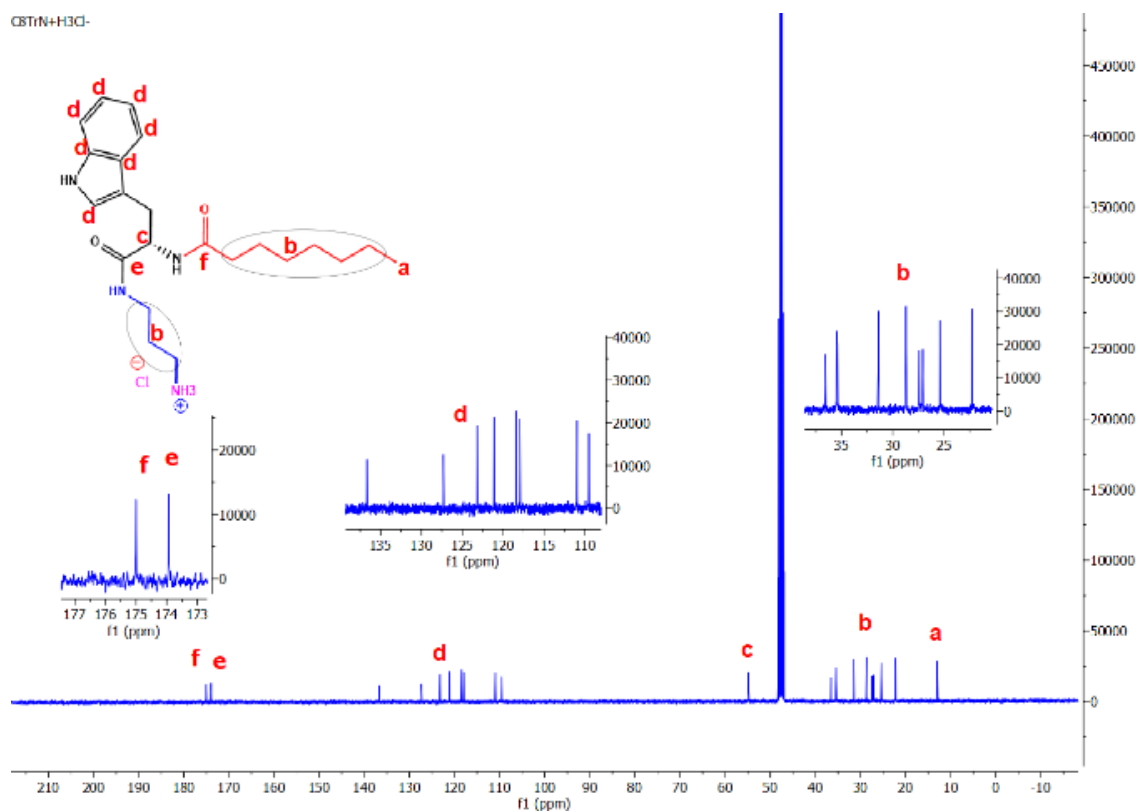
**Table S2** Fitting parameters of Gaussian bilayers for DPPC/C<sub>14</sub>PC<sub>3</sub>NH<sub>3</sub>Cl mixtures

**Table S3** Hemolytic activity of tryptophan and phenylalanine-based surfactants

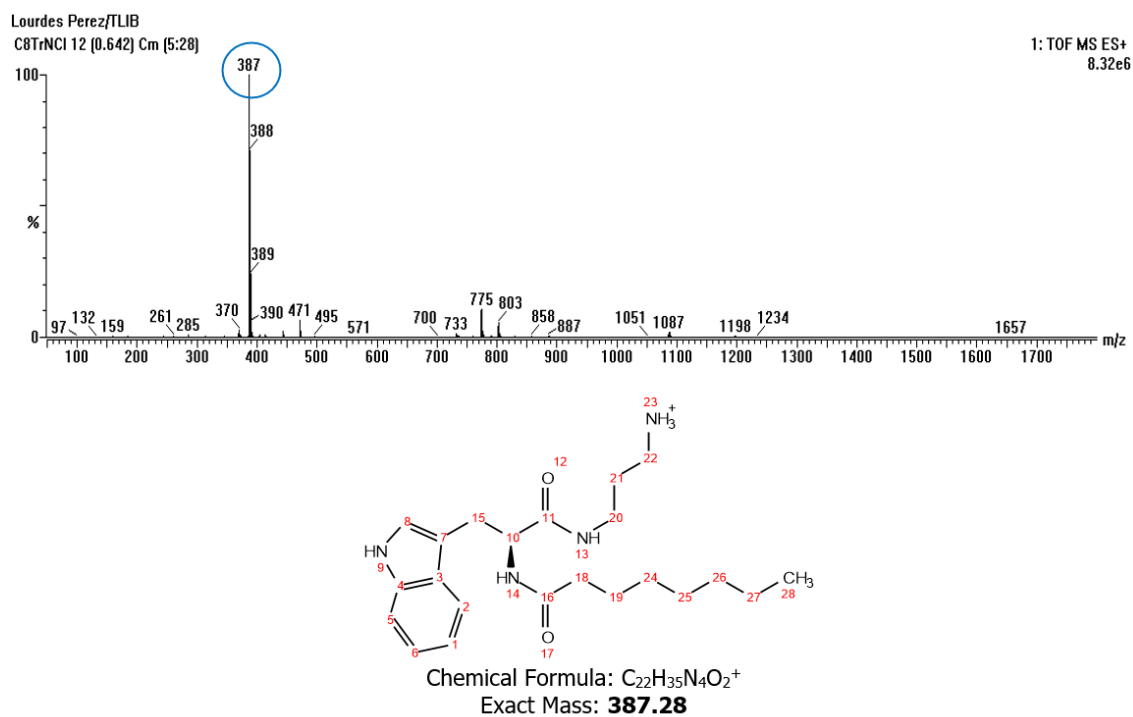
**Table S4** Results of the interaction details and docking score in (kcal/mol) of Benzalkonium derivative (From C<sub>8</sub> to C<sub>14</sub> carbon atoms) against the peptidoglycan glycosyltransferase (PDB ID:2OQO).



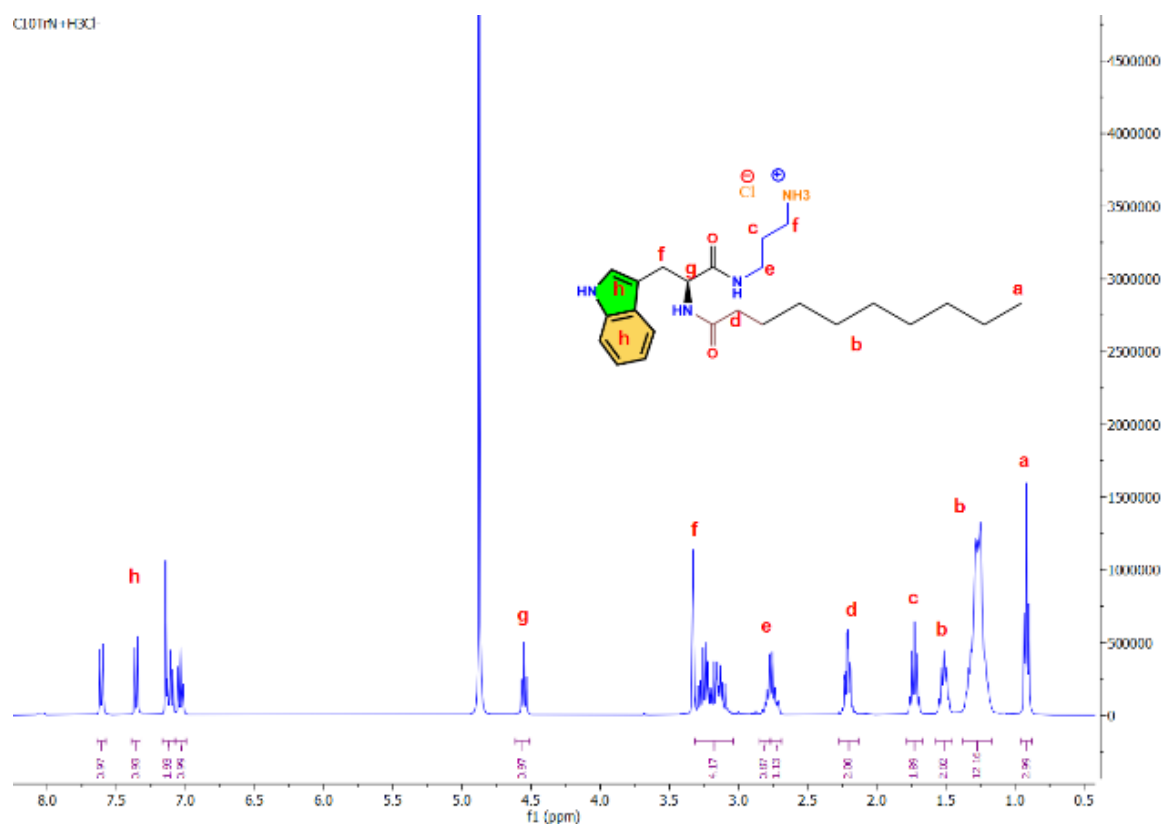
**Figure. S1:**  $^1\text{H}$  NMR spectrum of  $\text{C}_8\text{TC}_3\text{NH}_3\text{C}$



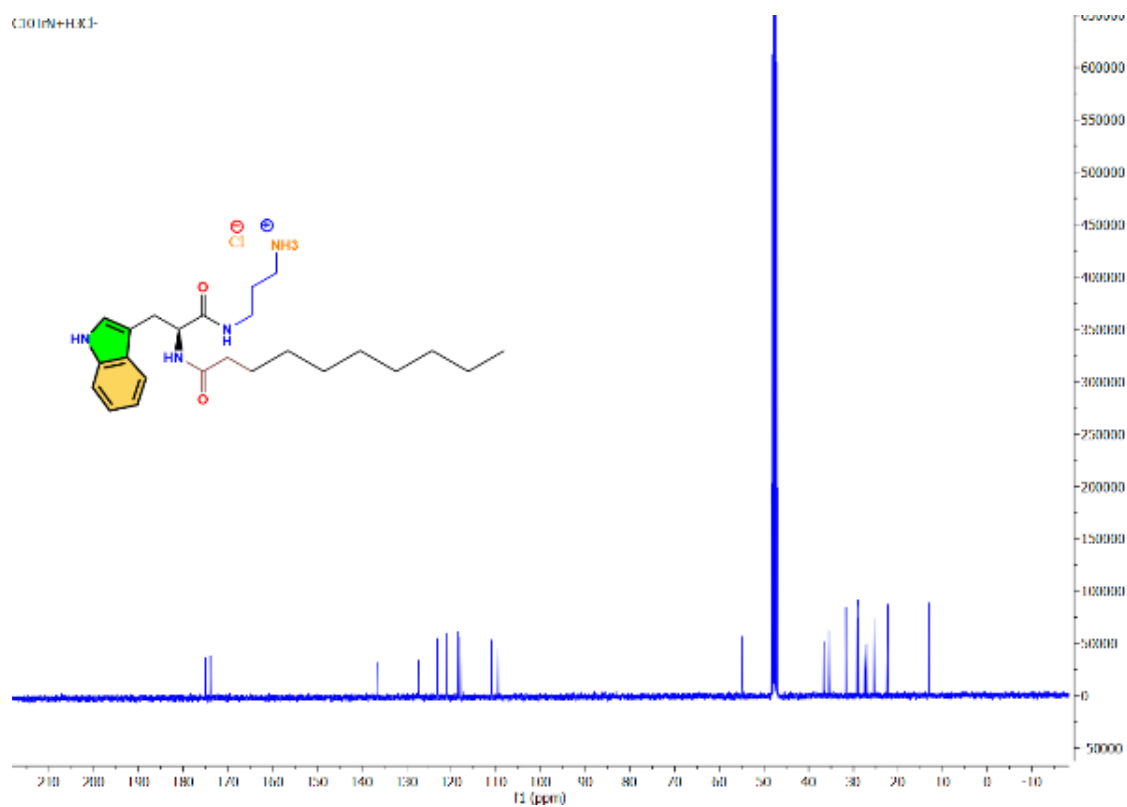
**Figure. S2:**  $^{13}\text{C}$  NMR spectrum of  $\text{C}_8\text{TC}_3\text{NH}_3\text{Cl}$



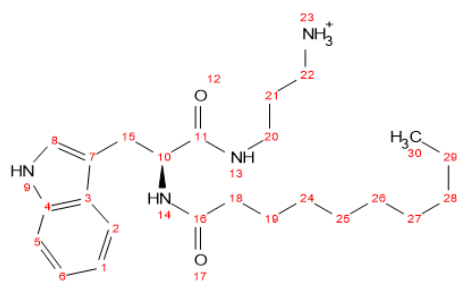
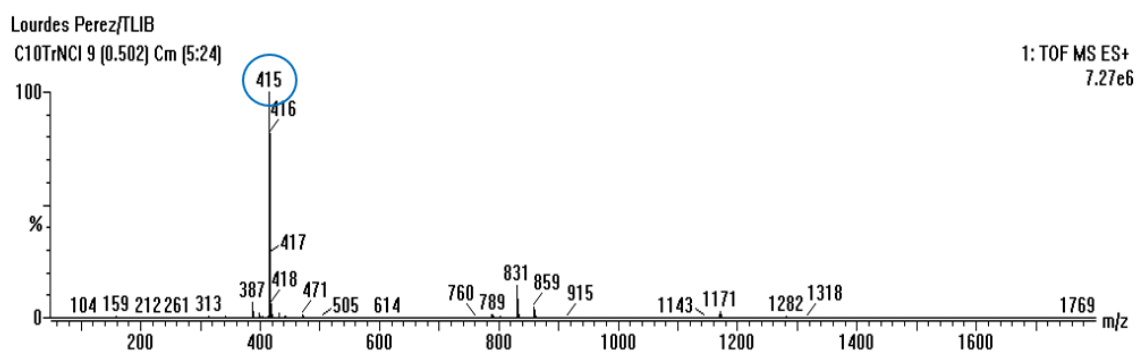
**Figure. S3:** ESI-MS spectrum of  $\text{C}_8\text{TC}_3\text{NH}_3\text{Cl}$



**Figure. S4:**  $^1H$  NMR spectrum of  $C_{10}TC_3NH_3C$



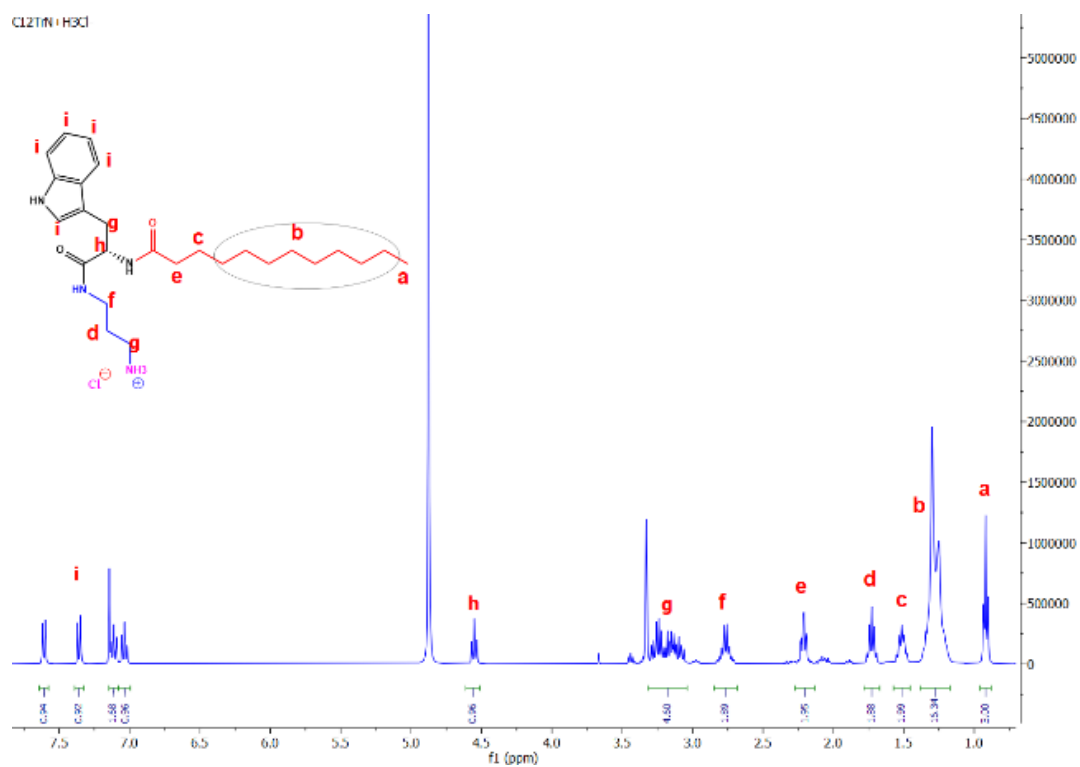
**Figure. S5**  $^{13}\text{C}$  NMR spectrum of  $\text{C}_{10}\text{TC}_3\text{NH}_3\text{C}$



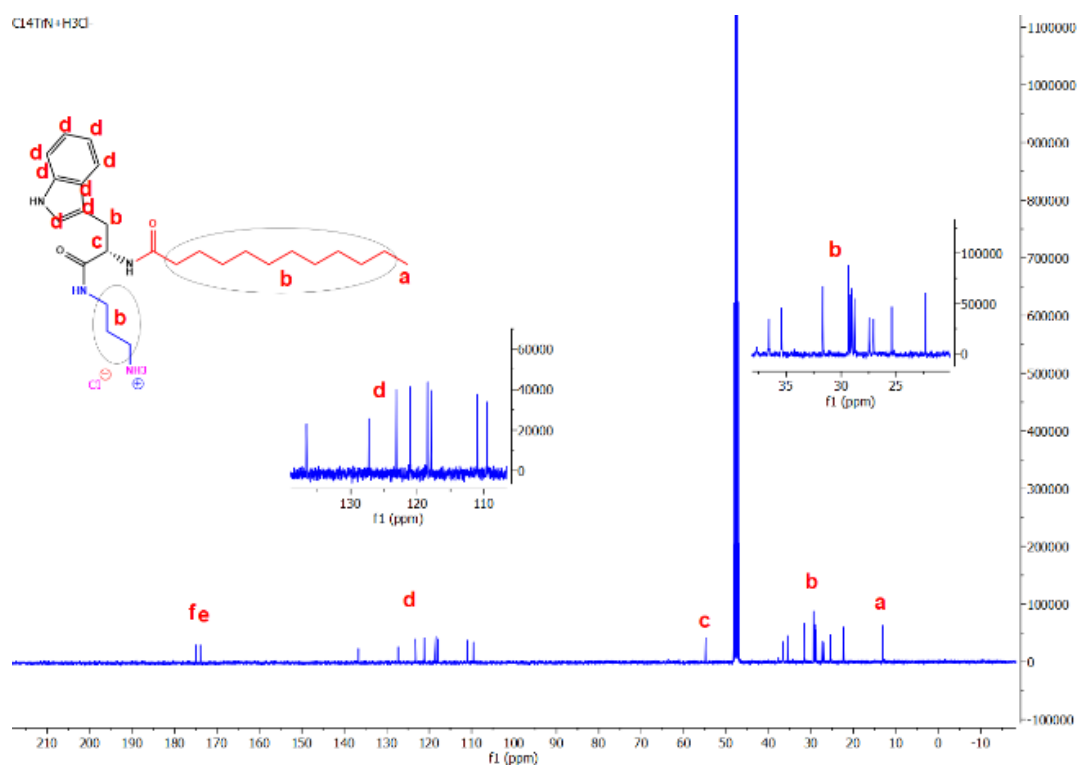
Chemical Formula:  $\text{C}_{24}\text{H}_{39}\text{N}_4\text{O}_2^+$

Exact Mass: **415.31**

**Figure. S6:** ESI-MS spectrum of  $\text{C}_{10}\text{TC}_3\text{NH}_3\text{Cl}$



**Figure. S7:**  $^1H$  NMR spectrum of  $C_{12}TC_3NH_3C$

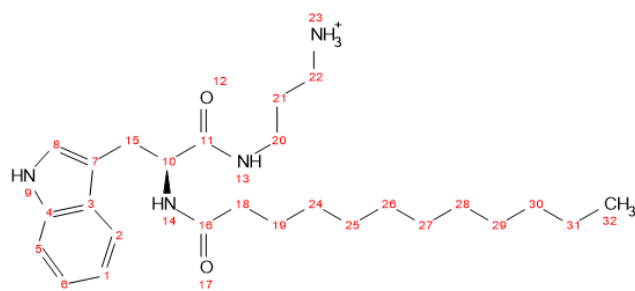
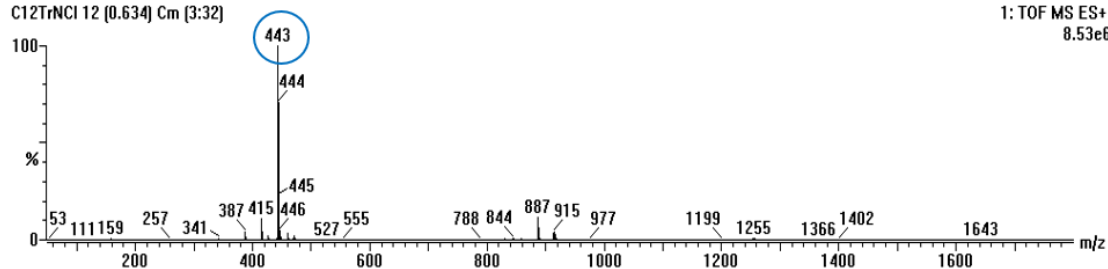


**Figure. S8:**  $^1H$  NMR spectrum of  $C_{12}TC_3NH$



Lourdes Perez/TLIB  
C12TrnCl 12 [0.634] Cm [3:32]

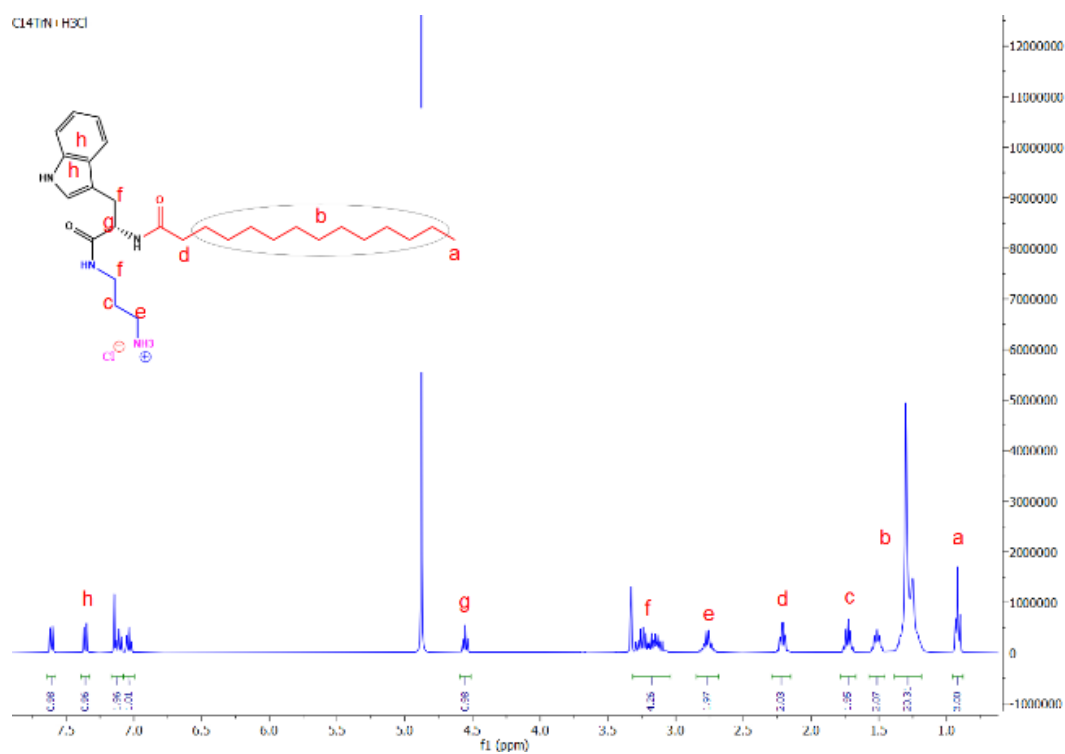
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8.53e6



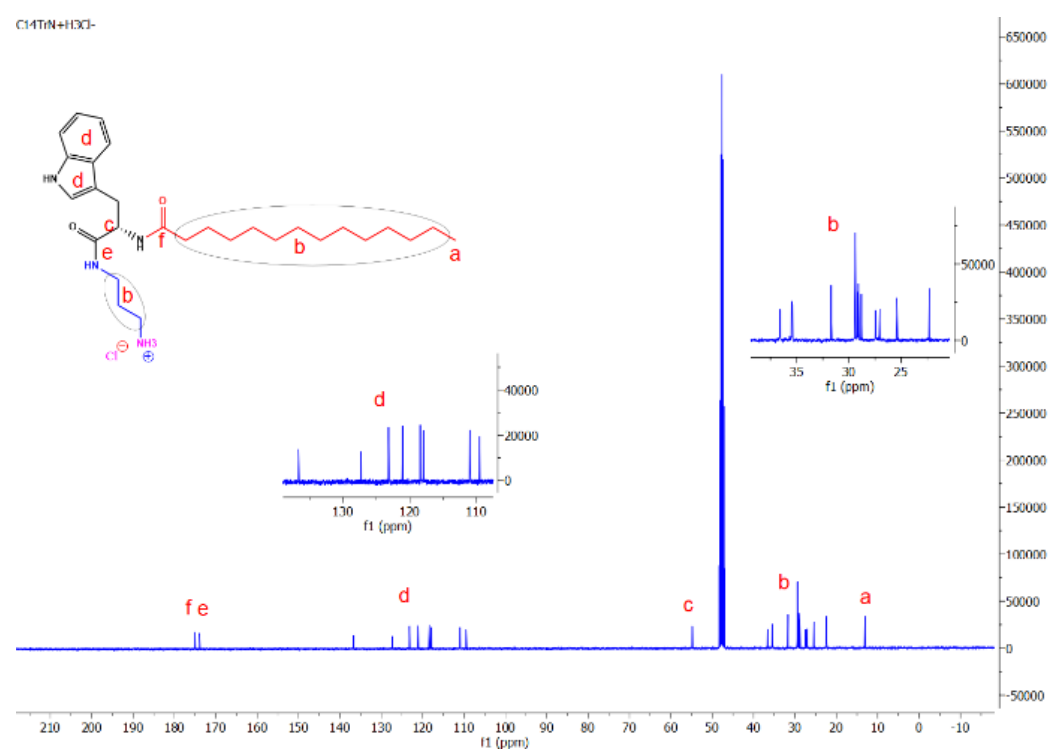
Chemical Formula: C<sub>26</sub>H<sub>43</sub>N<sub>4</sub>O<sub>2</sub><sup>+</sup>

Exact Mass: **443.34**

**Figure. S9:** ESI-MS spectrum of C<sub>12</sub>TC<sub>3</sub>NH<sub>3</sub>Cl



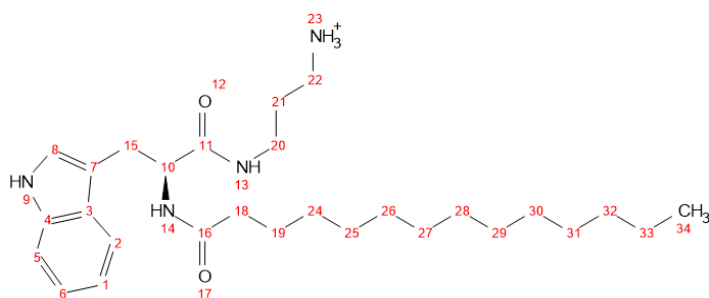
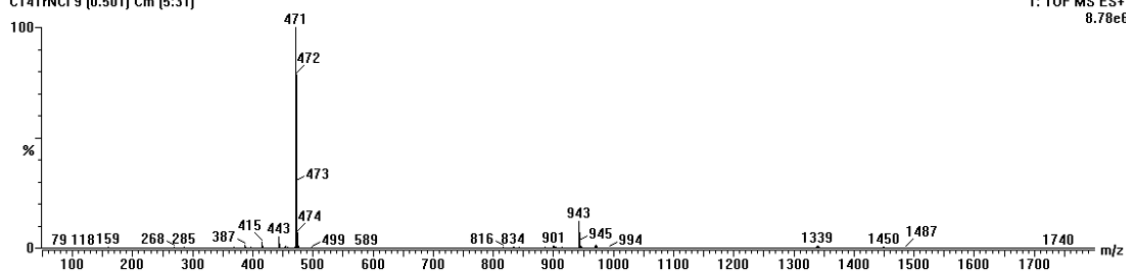
**Figure. S10:** <sup>1</sup>H NMR spectrum of C<sub>14</sub>TC<sub>3</sub>NH<sub>3</sub>C



**Figure. S11:** <sup>13</sup>C NMR spectrum of C<sub>12</sub>TC<sub>3</sub>NH

Lourdes Perez/TLIB  
C14TrNCI 9 [0.501] Cm [5:31]

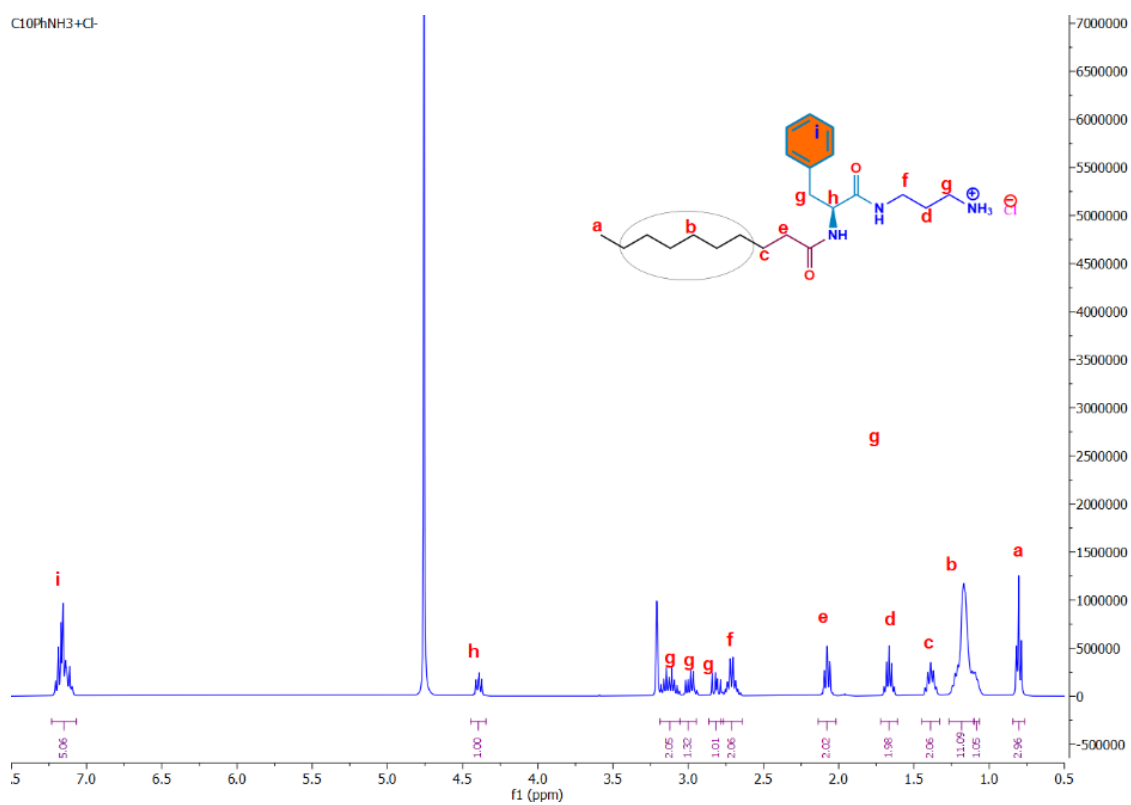
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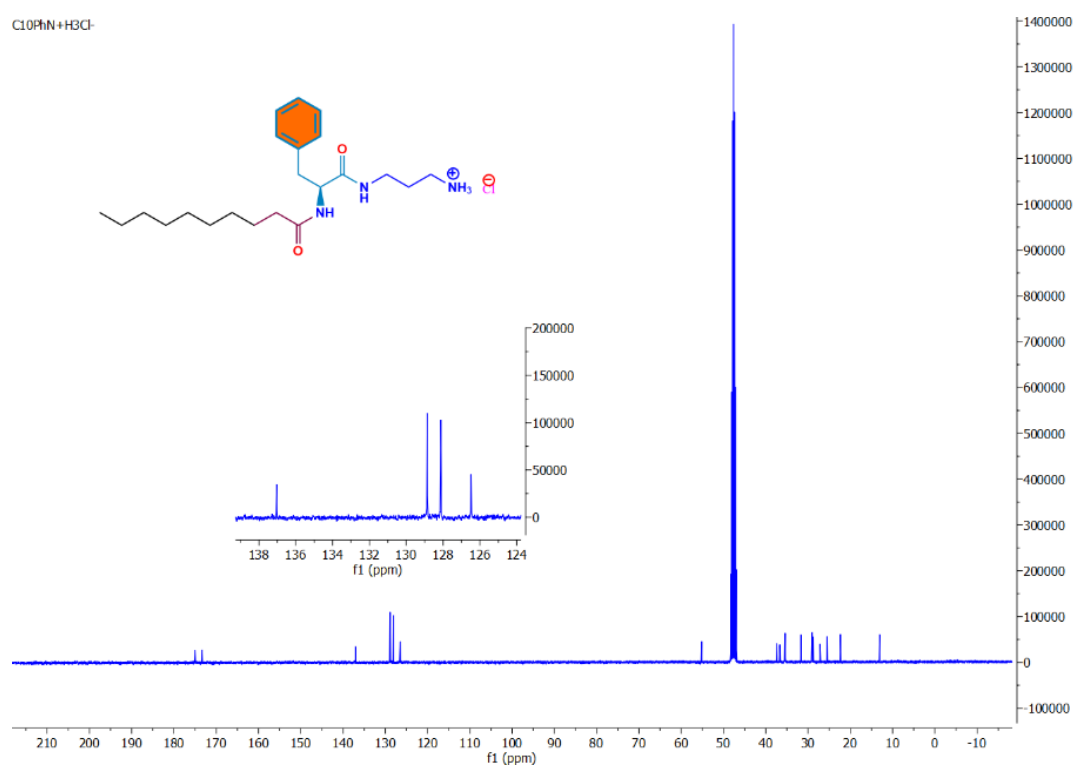
Chemical Formula:  $C_{28}H_{47}N_4O_2^+$

Exact Mass: **471.37**

**Figure. S12:** ESI-MS spectrum of  $C_{14}TC_3NH_3Cl$



**Figure. S13:**  $^1\text{H}$  NMR spectrum of  $\text{C}_{10}\text{PC}_3\text{NH}_3\text{C}$

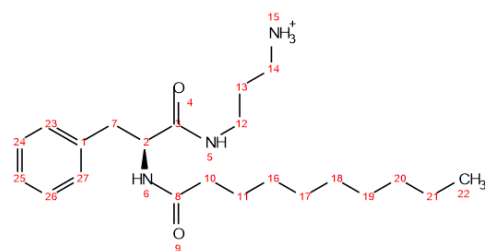
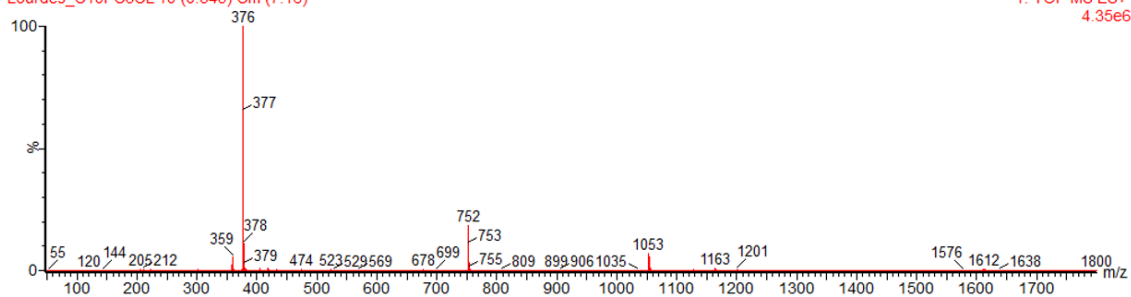


**Figure. S14:**  $^{13}\text{C}$  NMR spectrum of  $\text{C}_{10}\text{PC}_3\text{NH}_3\text{C}$

Lordes/TLIB

Lourdes\_C10PC3CL 10 (0.545) Cm (7:16)

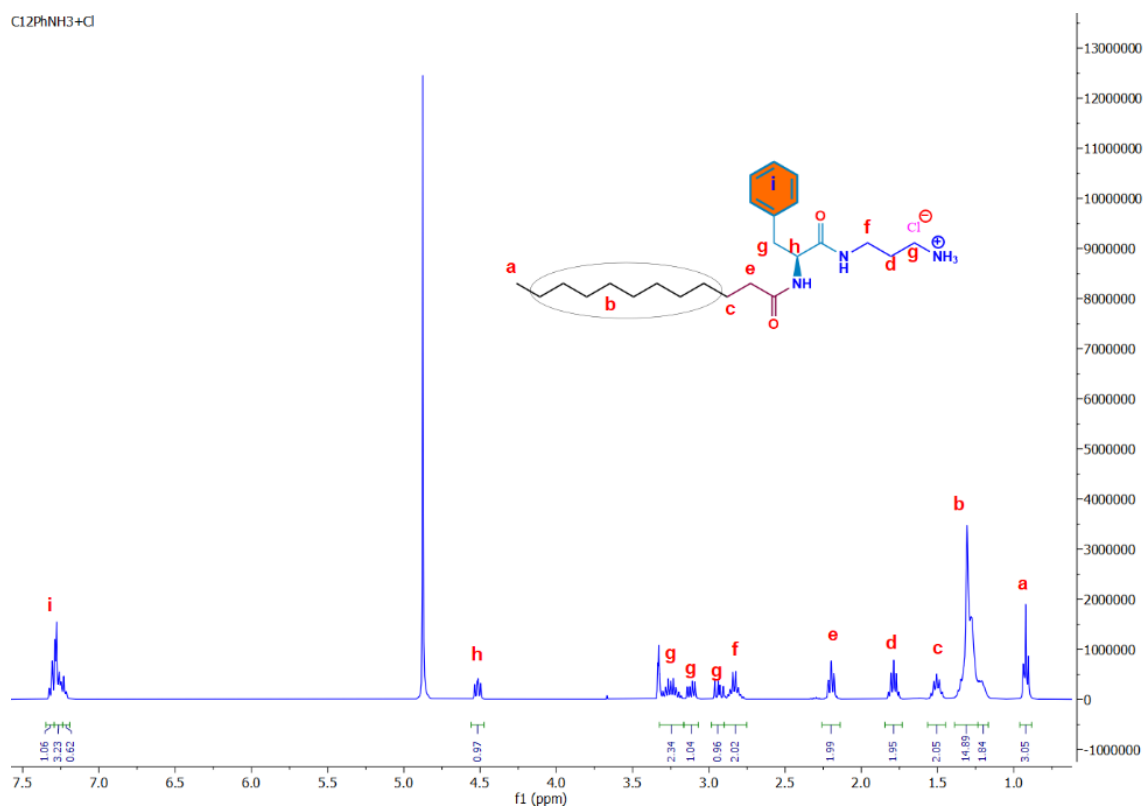
1: TOF MS ES+  
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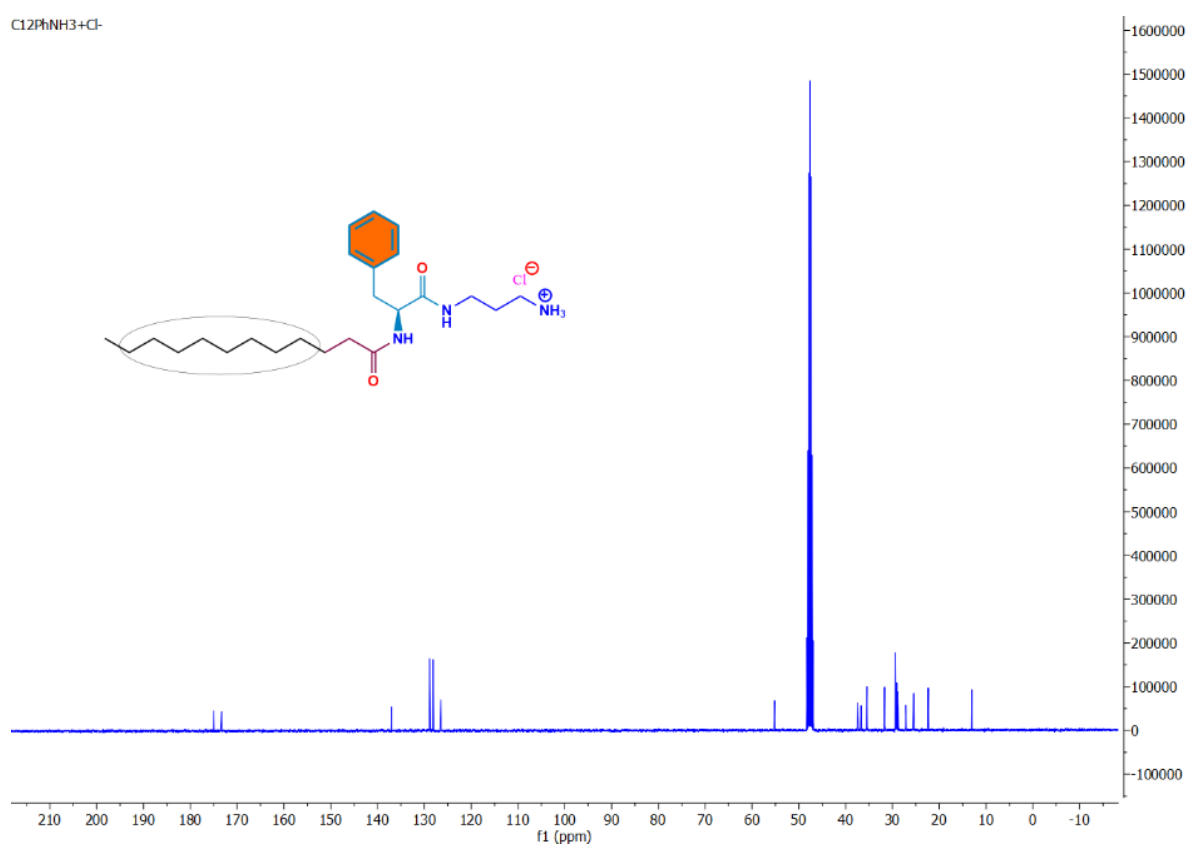
Chemical Formula:  $C_{22}H_{38}N_3O_2^+$

Exact Mass: **376.30**

**Figure. S15:** ESI-MS spectrum of  $C_{10}PC_3NH_3C$



**Figure. S16:**  $^1\text{H}$  NMR spectrum of  $\text{C}_{12}\text{PC}_3\text{NH}_3\text{C}$

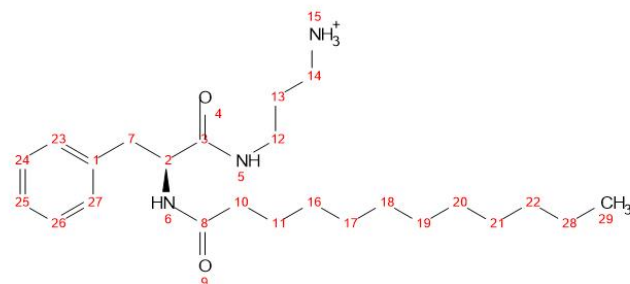
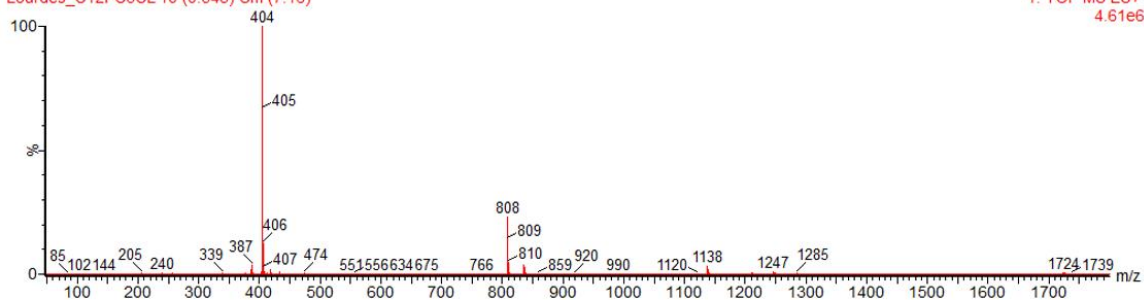


**Figure. S17:**  $^{13}\text{C}$  NMR spectrum of  $\text{C}_{12}\text{PC}_3\text{NH}_3\text{C}$

Lordes/TLIB

Lourdes\_C12PC3CL 10 (0.545) Cm (7:16)

1: TOF MS ES+  
4.61e6

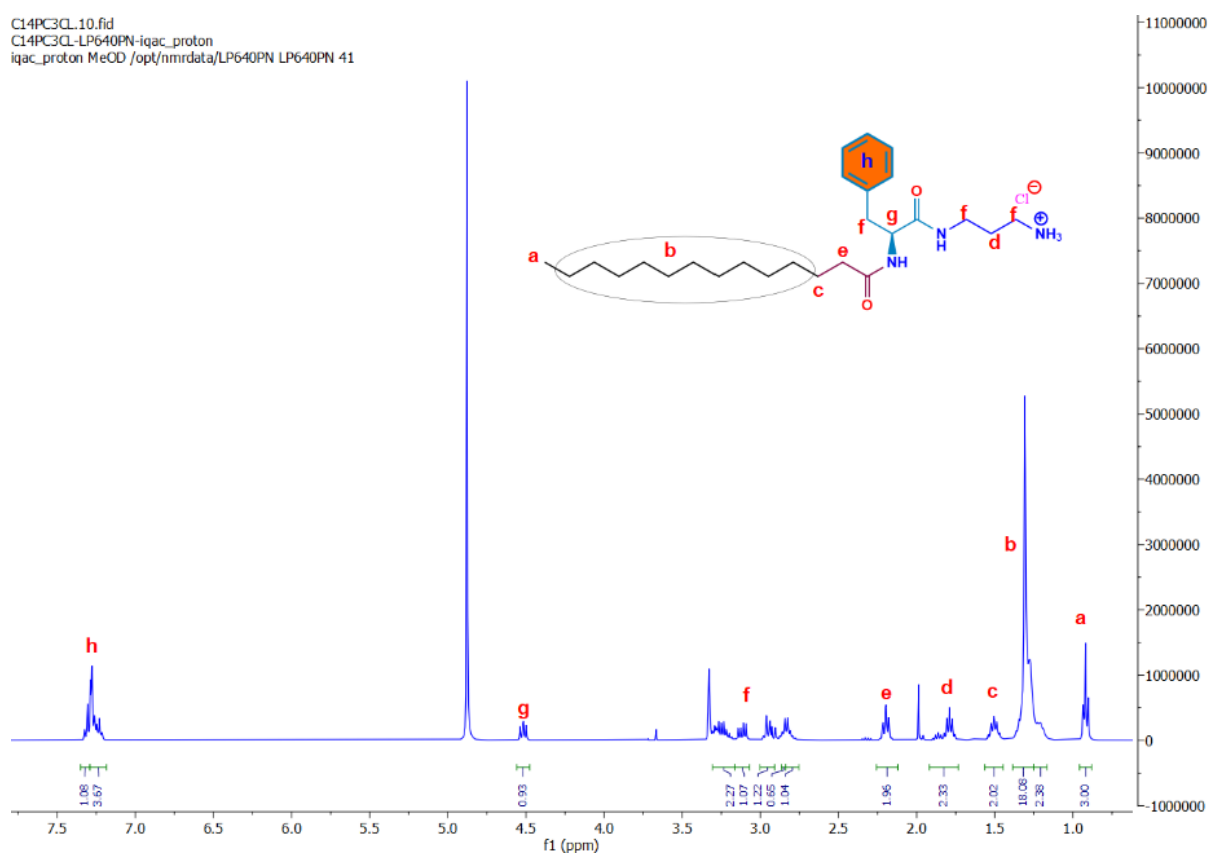


Chemical Formula: C<sub>24</sub>H<sub>42</sub>N<sub>3</sub>O<sub>2</sub><sup>+</sup>

Exact Mass: 404.33

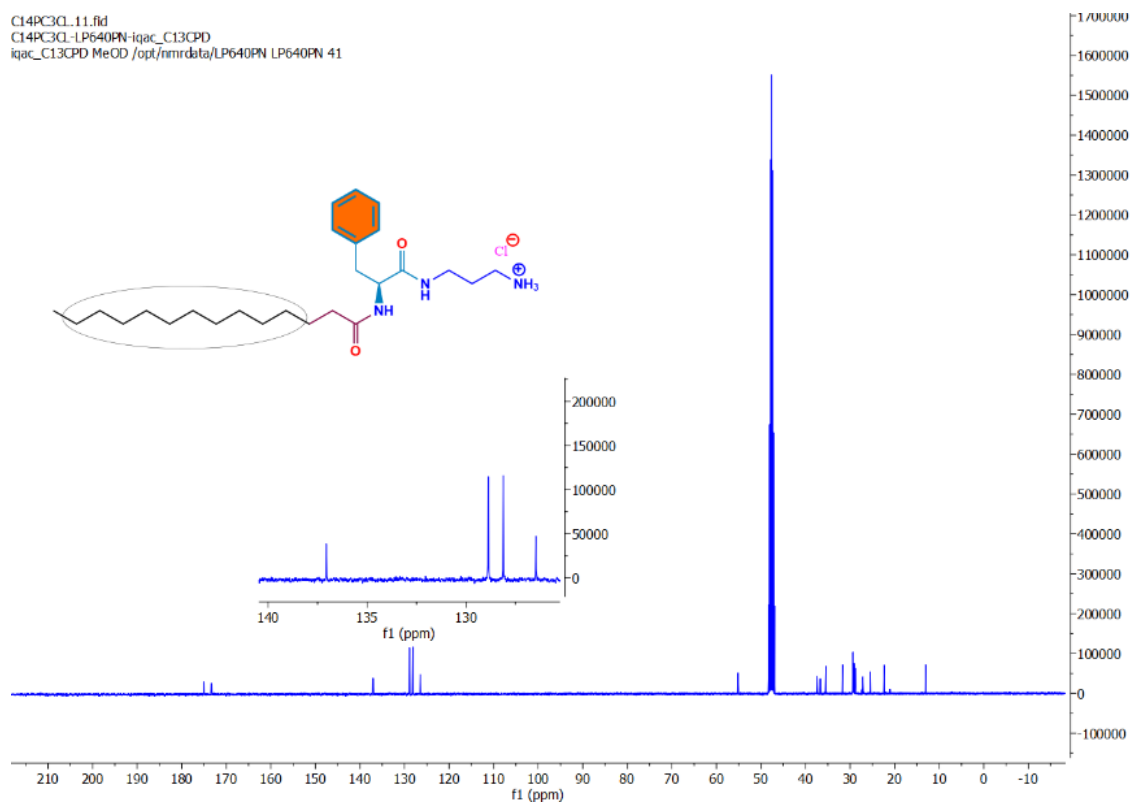
**Figure. S18:** ESI-MS spectrum of C<sub>12</sub>PC<sub>3</sub>NH<sub>3</sub>C

C14PC3CL10.fid  
C14PC3CL-LP640PN-igac\_proton  
igac\_proton MeOD /opt/nmrdata/LP640PN LP640PN 41



**Figure. S19:**  $^1\text{H}$  NMR spectrum of C<sub>14</sub>PC<sub>3</sub>NH<sub>3</sub>C

C14PC3CL11.fid  
C14PC3CL-LP640PN-igac\_C13CPD  
igac\_C13CPD MeOD /opt/nmrdata/LP640PN LP640PN 41



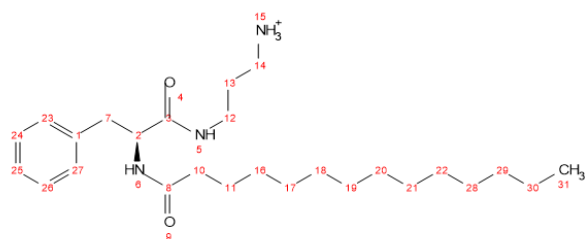
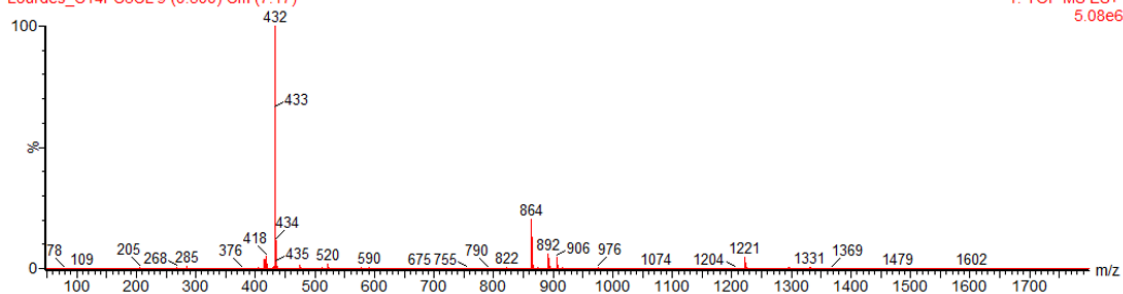
**Figure. S20:**  $^{13}\text{C}$  NMR spectrum of C<sub>14</sub>PC<sub>3</sub>NH<sub>3</sub>C



Lordes/TLIB

Lourdes\_C14PC3CL 9 (0.500) Cm (7:17)

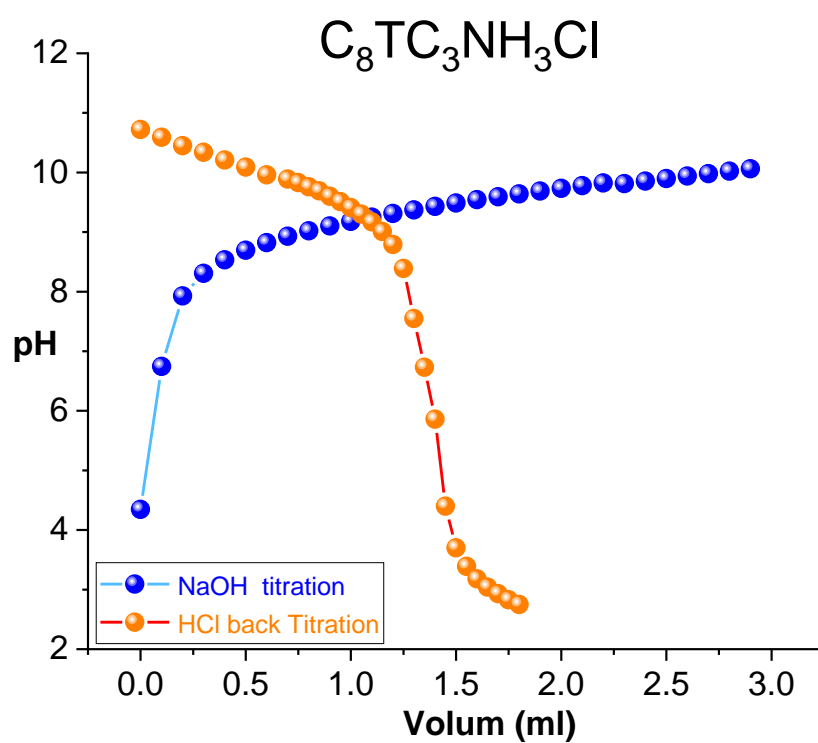
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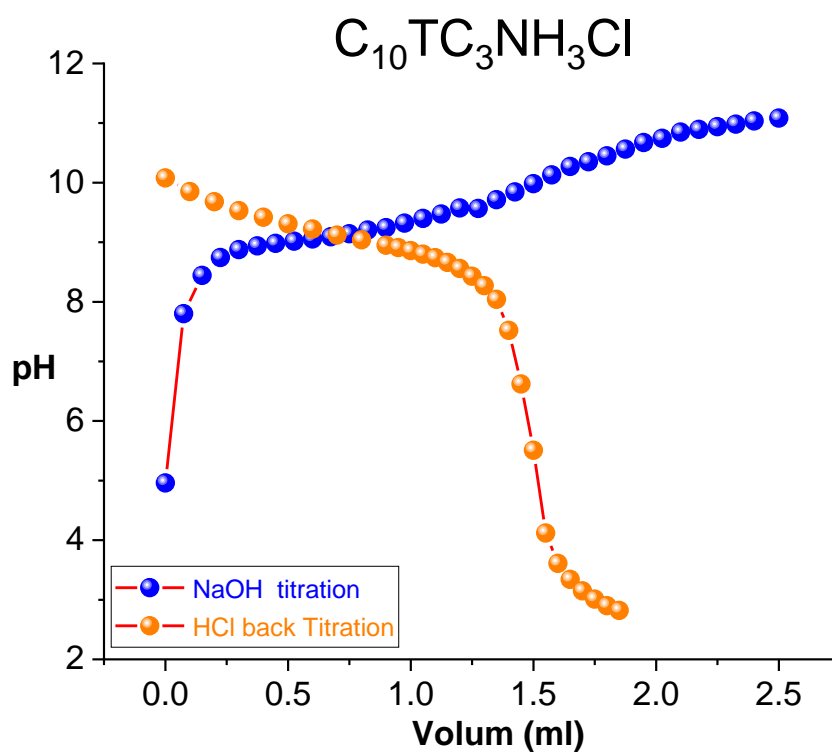
Chemical Formula: C<sub>26</sub>H<sub>46</sub>N<sub>3</sub>O<sub>2</sub><sup>+</sup>

Exact Mass: **432.36**

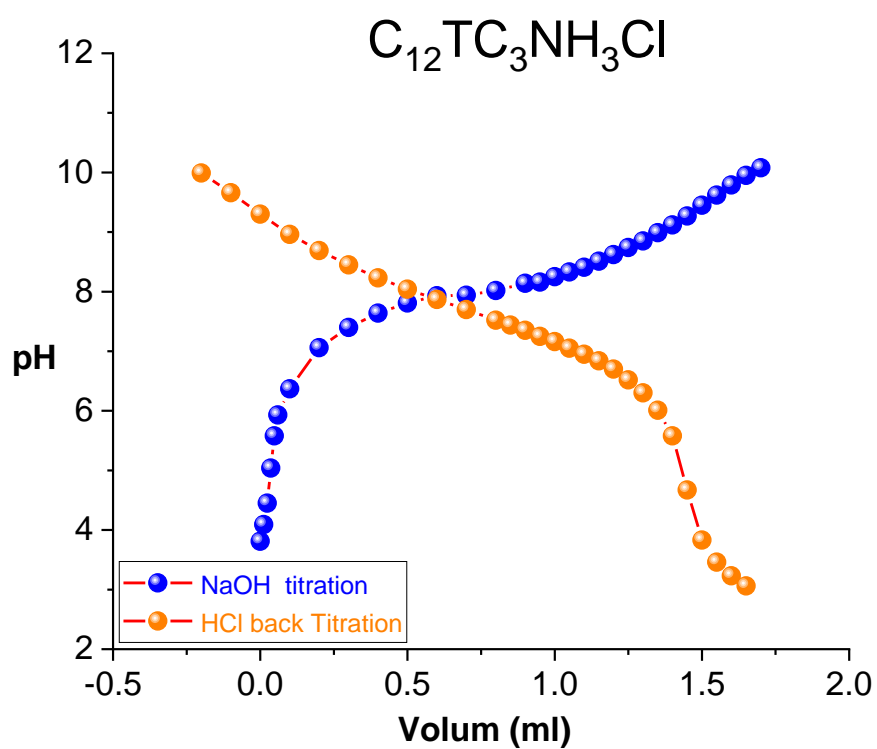
**Figure. S21:** ESI-MS spectrum of C<sub>14</sub>PC<sub>3</sub>NH<sub>3</sub>C



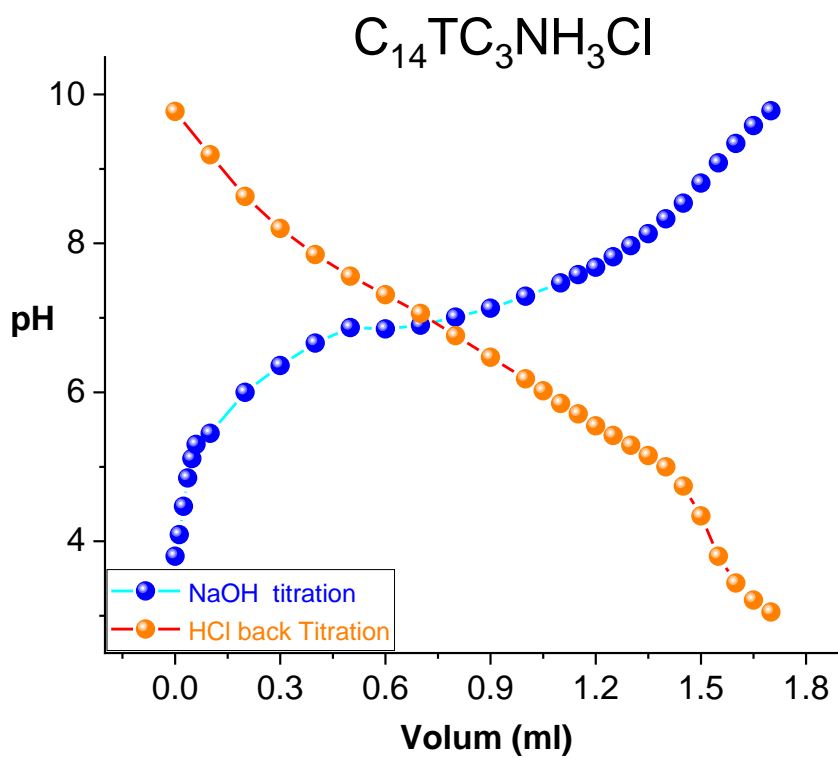
**Figure.S22** NaOH titration and HCl backtitration at 298.15 K for  $C_8TC_3NH_3Cl$



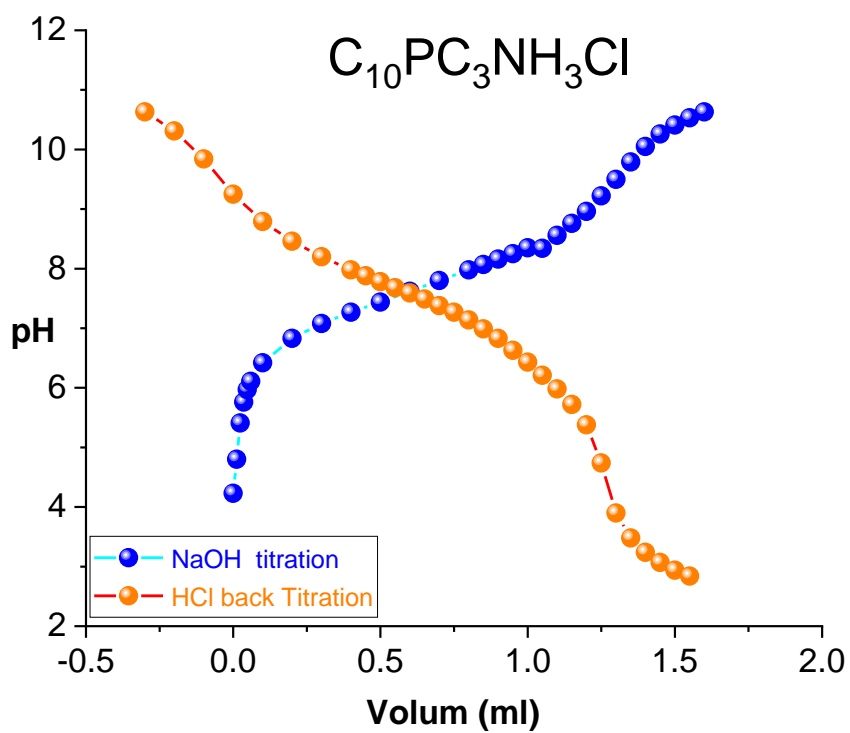
**Figure.S23** NaOH titration and HCl backtitration at 298.15 K for  $C_{10}TC_3NH_3Cl$



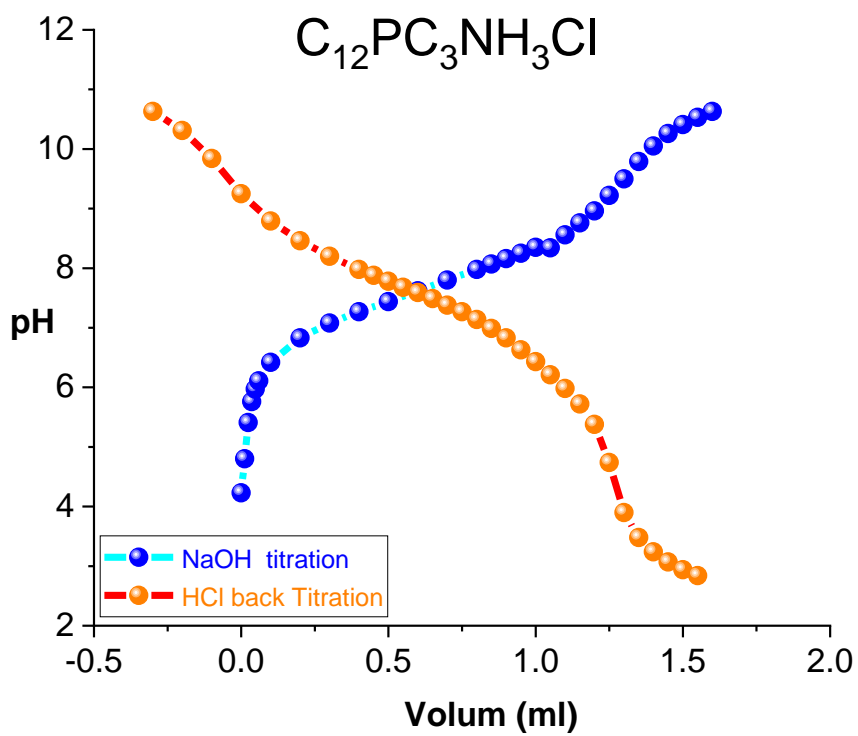
**Figure.S24** NaOH titration and HCl back titration at 298.15 K for  $C_{12}TC_3NH_3Cl$



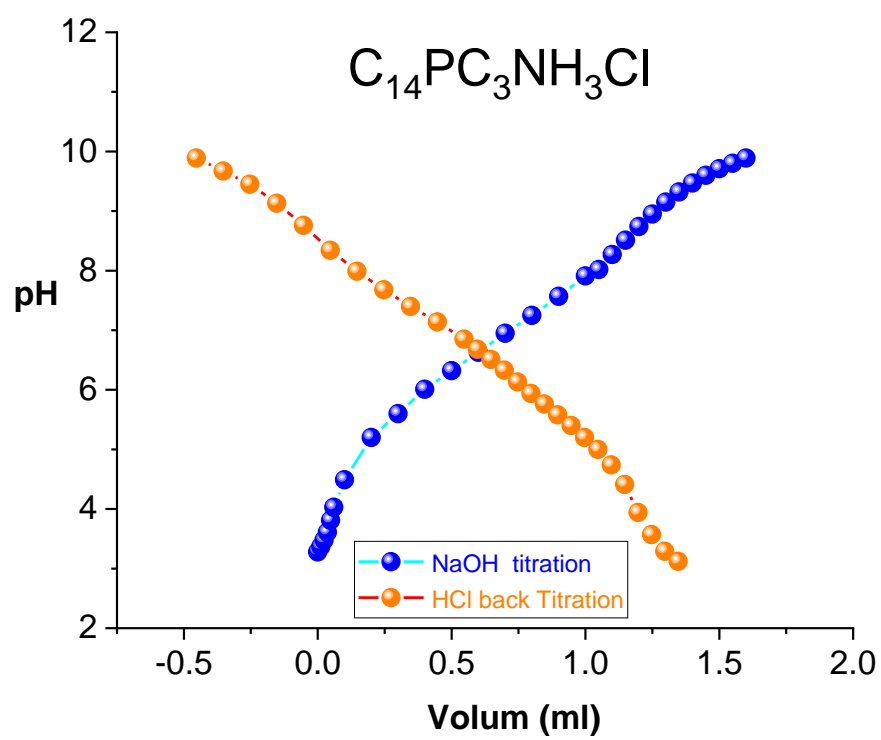
**Figure.S25** NaOH titration and HCl backtitration at 298.15 K for  $C_{14}TC_3NH_3Cl$



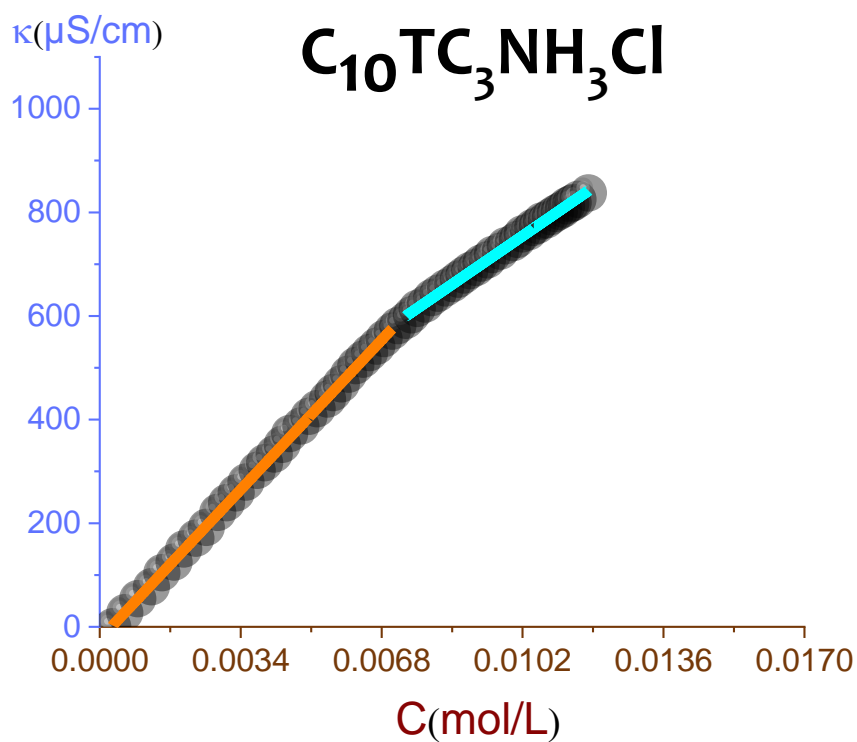
**Figure.S26** NaOH titration and HCl backtitration at 298.15 K for  $C_{10}PC_3NH_3Cl$



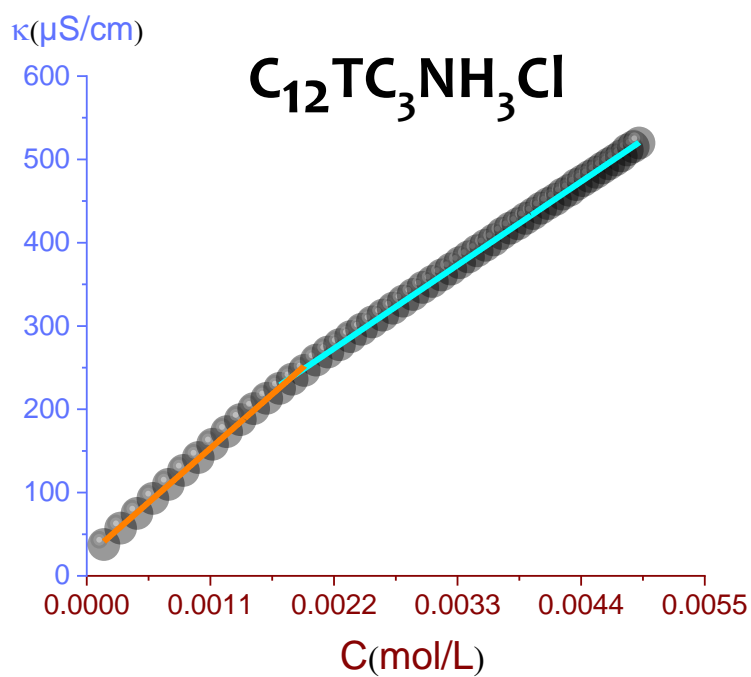
**Figure.S27** NaOH titration and HCl back titration at 298.15 K for  $C_{12}PC_3NH_3Cl$



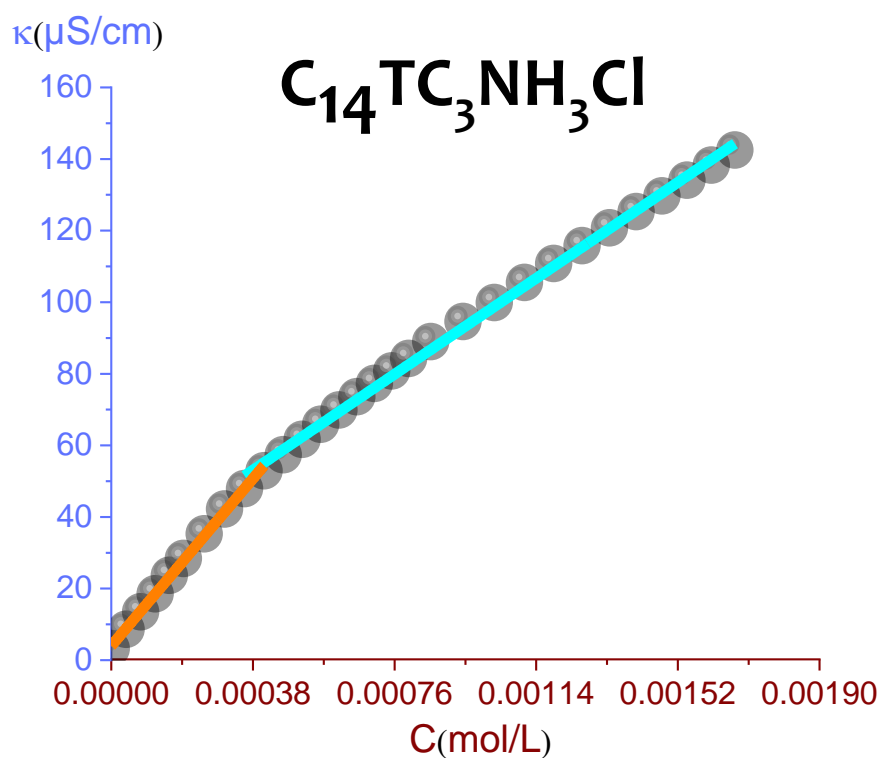
**Figure.S28** NaOH titration and HCl backtitration at 298.15 K for  $C_{14}PC_3NH_3Cl$



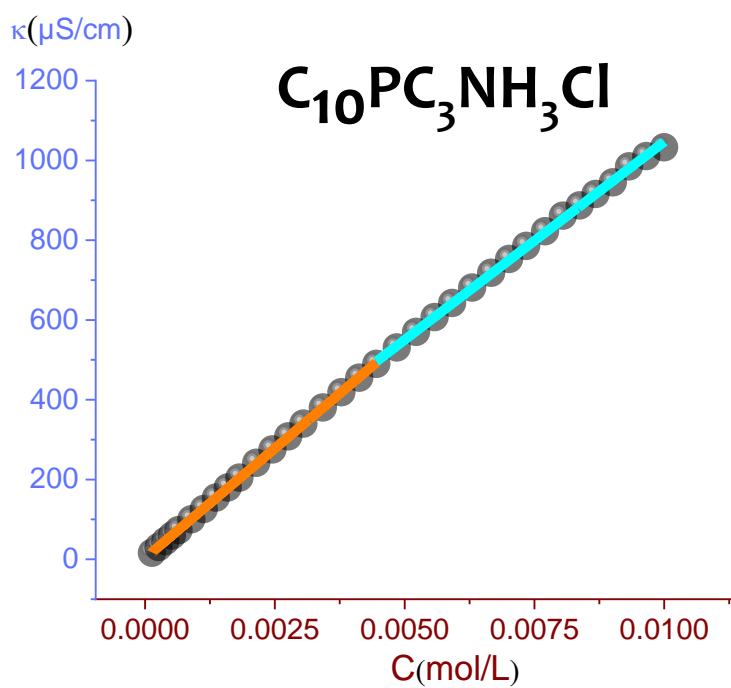
**Figure.S29** Specific conductivity ( $\kappa$ ) as a function of concentration of  $C_{10}TC_3NH_3Cl$



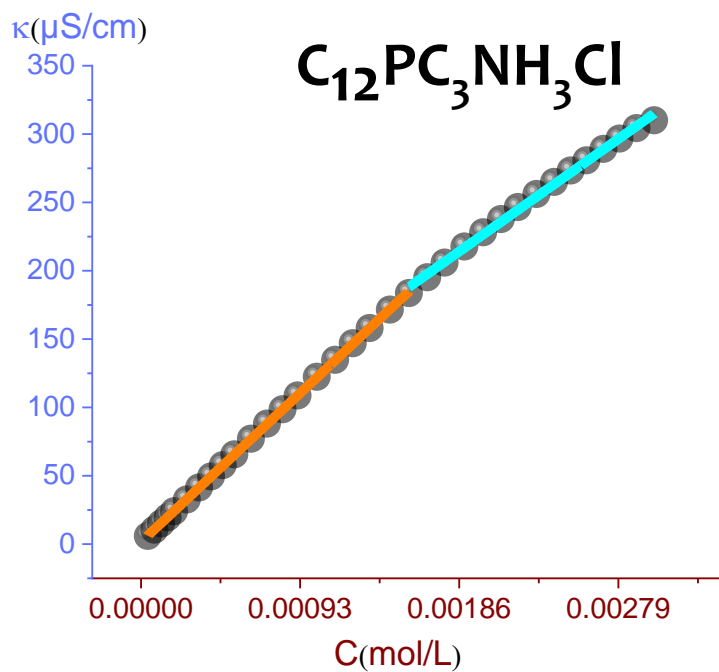
**Figure.S30** Specific conductivity ( $\kappa$ ) as a function of concentration of  $C_{12}TC_3NH_3Cl$



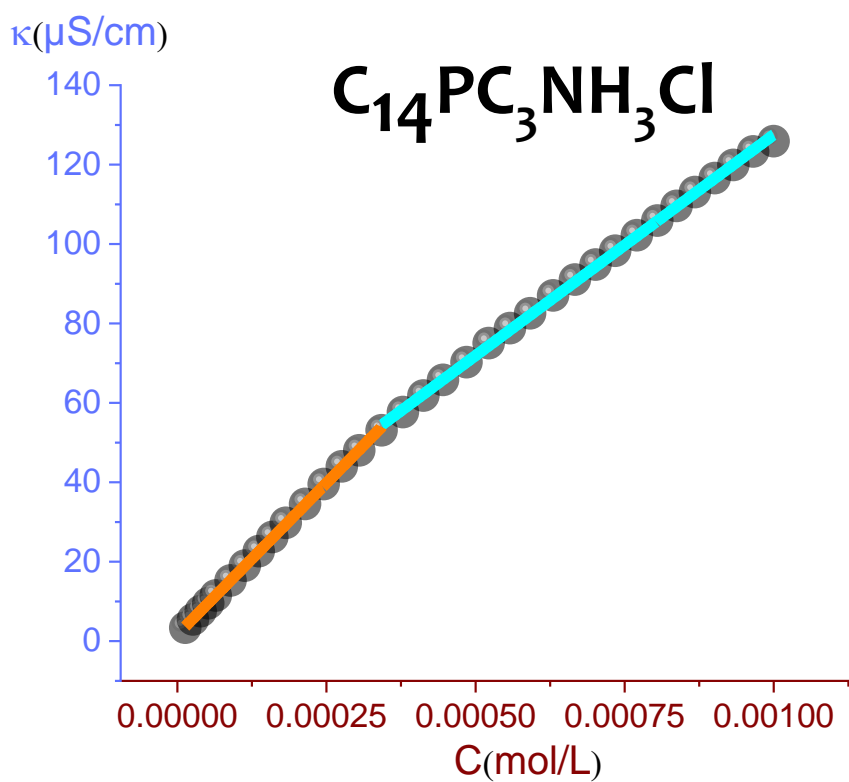
**Figure.S31** Specific conductivity ( $\kappa$ ) as a function of concentration of  $C_{14}TC_3NH_3Cl$



**Figure.S32** Specific conductivity ( $\kappa$ ) as a function of concentration of  $C_{10}PC_3NH_3Cl$

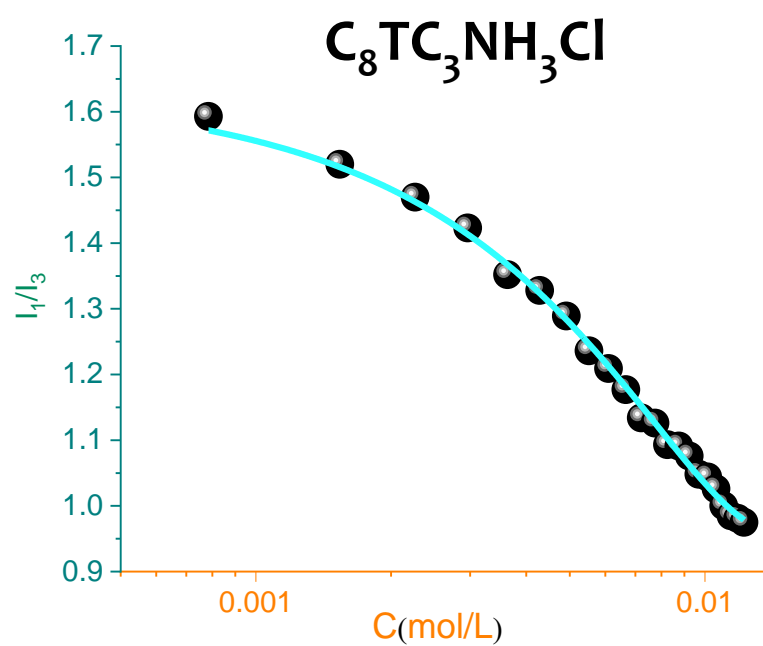


**Figure.S33** Specific conductivity ( $\kappa$ ) as a function of concentration of  $C_{12}PC_3NH_3Cl$

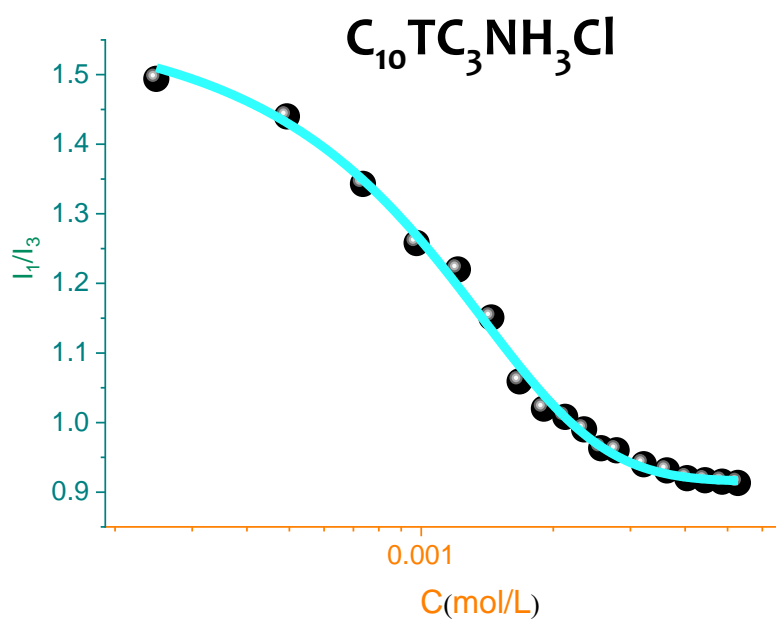


**Figure.S34** Specific conductivity ( $\kappa$ ) as a function of concentration of  $C_{14}PC_3NH_3Cl$

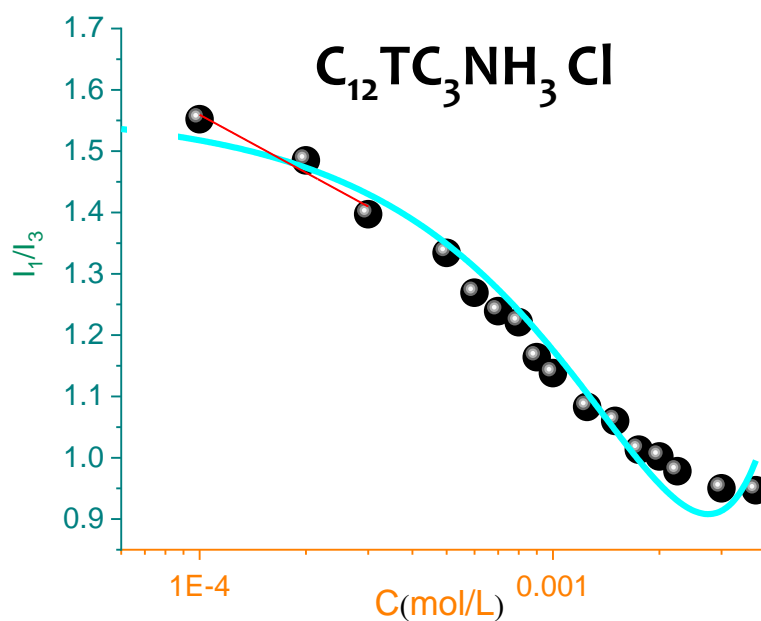




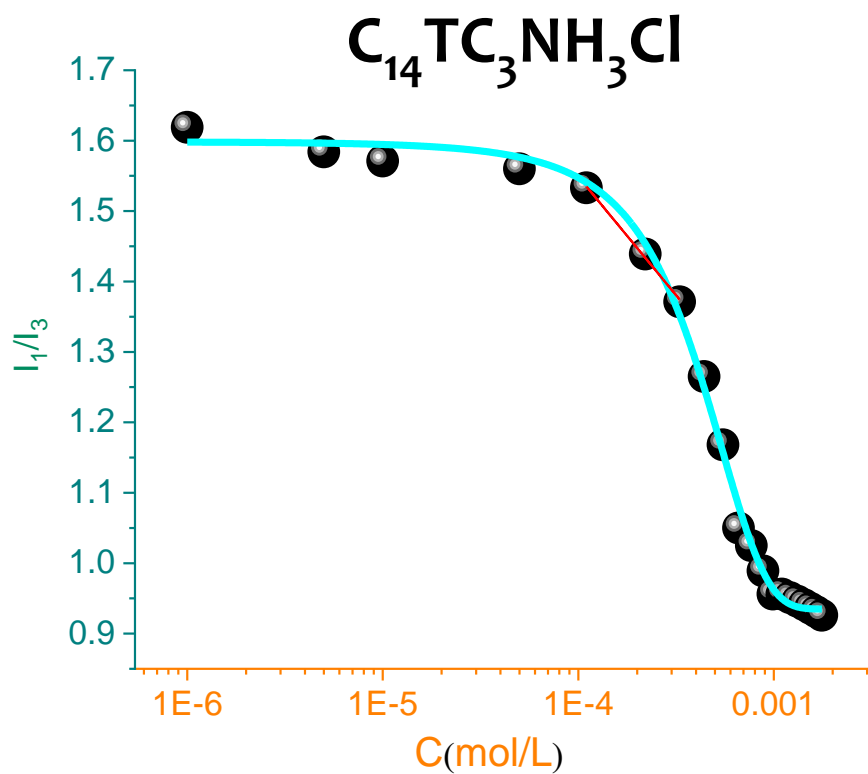
**Figure.S35** Variation of  $I_1/I_3$  ratio of pyrene fluorescence as a function of concentration  $C_8TC_3NH_3Cl$



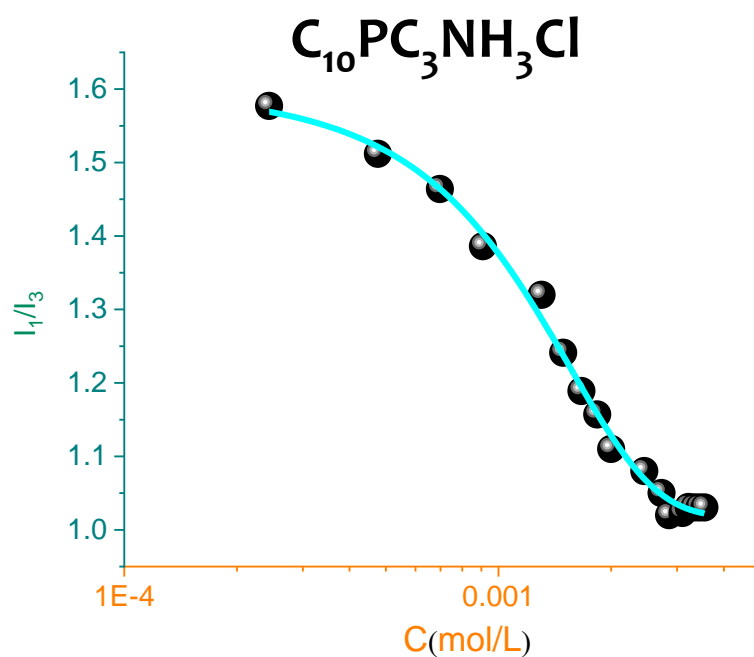
**Figure.S36** Variation of  $I_1/I_3$  ratio of pyrene fluorescence as a function of concentration  $C_{10}TC_3NH_3Cl$



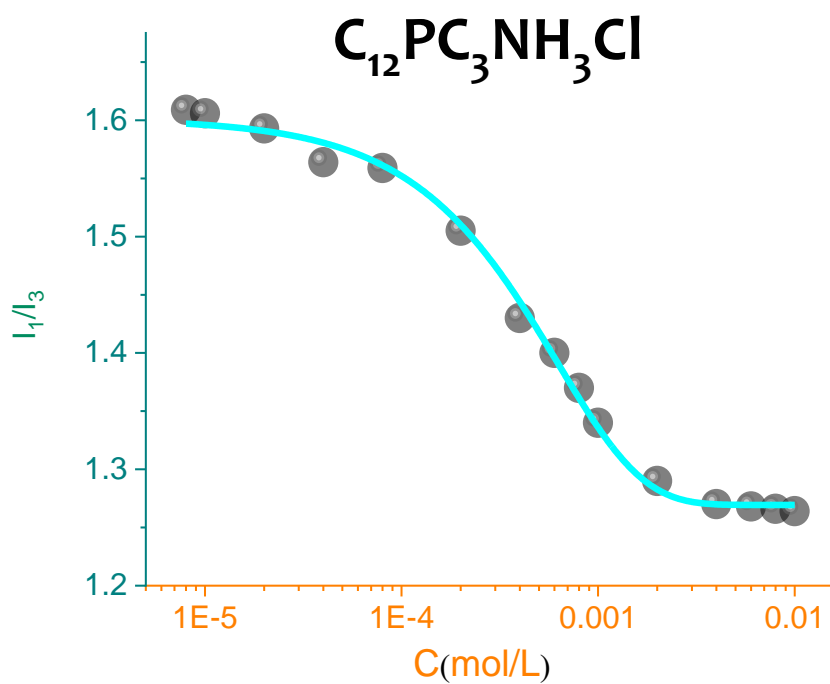
**Figure.S37** Variation of  $I_1/I_3$  ratio of pyrene fluorescence as a function of concentration  $C_{12}TC_3NH_3Cl$



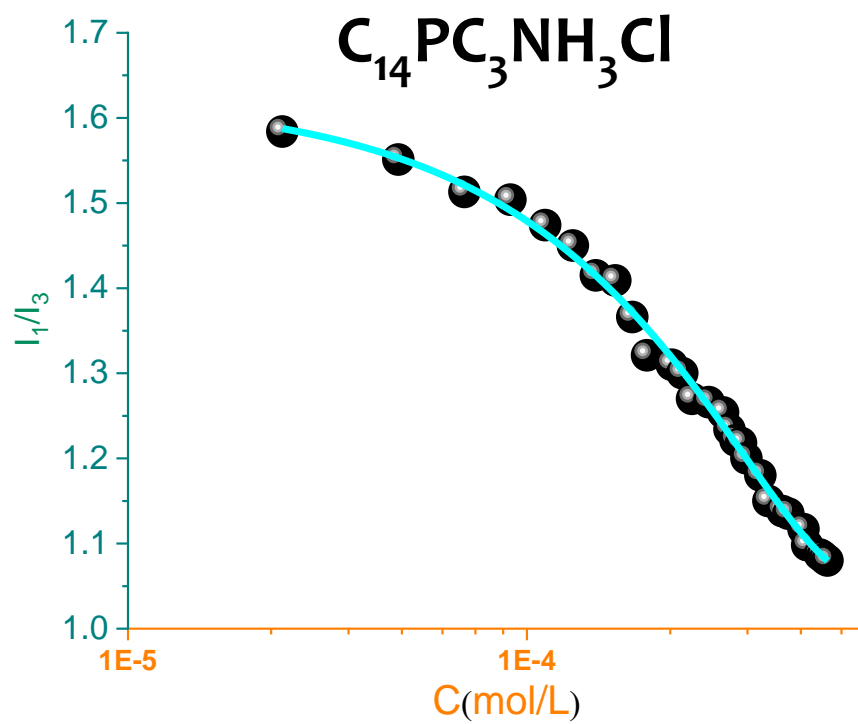
**Figure.S38** Variation of  $I_1/I_3$  ratio of pyrene fluorescence as a function of concentration  $C_{14}TC_3NH_3Cl$



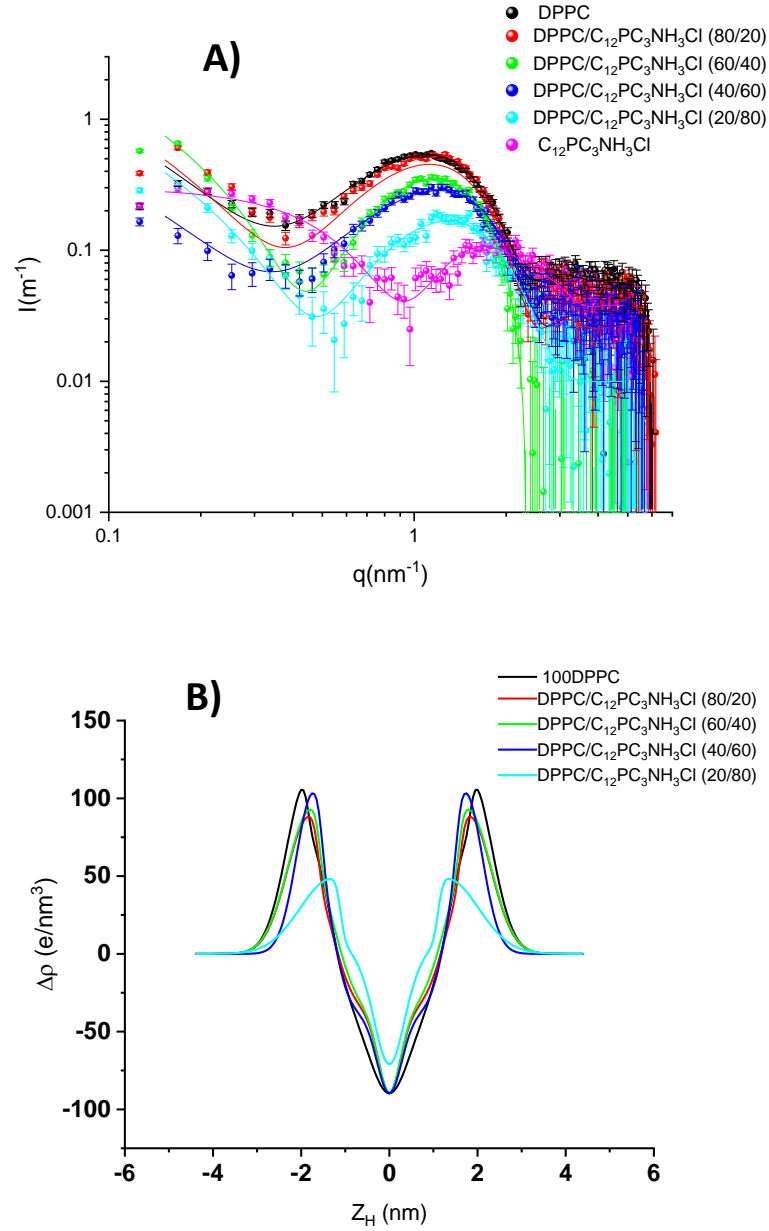
**Figure.S39** Variation of  $I_1/I_3$  ratio of pyrene fluorescence as a function of concentration  $C_{14}PC_3NH_3Cl$



**Figure.S40** Variation of  $I_1/I_3$  ratio of pyrene fluorescence as a function of concentration  $C_{12}PC_3NH_3Cl$



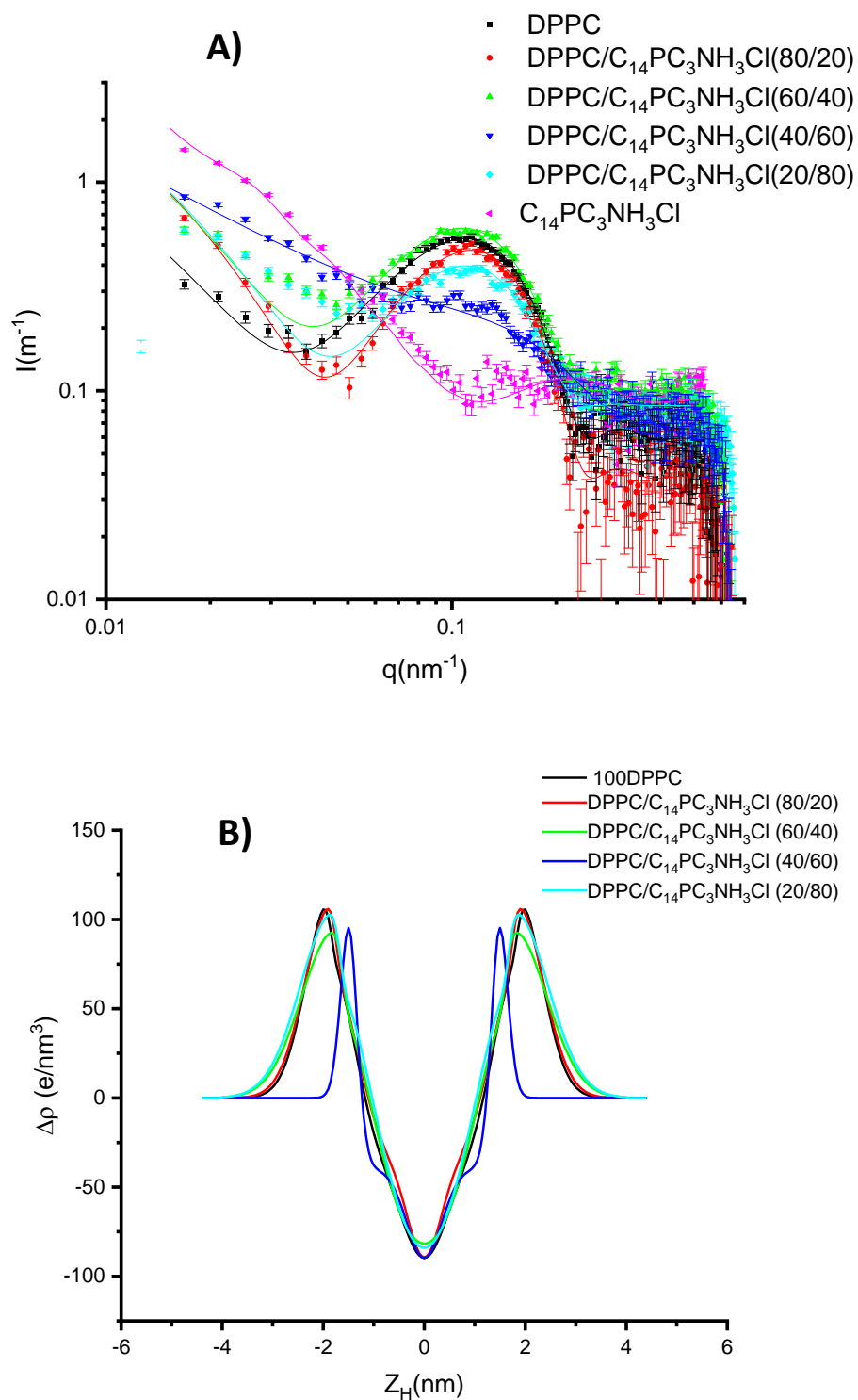
**Figure.S41** Variation of  $I_1/I_3$  ratio of pyrene fluorescence as a function of concentration  $C_{14}PC_3NH_3Cl$



**Figure.S42** A) Scattered intensity patterns as a function of scattering vector modulus for DPPC and  $C_{12}PC_3NH_3Cl$  and their mixtures the curves correspond to the best fit of Gaussian bilayers or core-shell models. B) The corresponding electron density profiles of the bilayer models corresponding to the best fits of A).

**Table S1** Fitting parameters of Gaussian bilayers for DPPC/C<sub>12</sub>PC<sub>3</sub>NH<sub>3</sub>Cl mixtures.

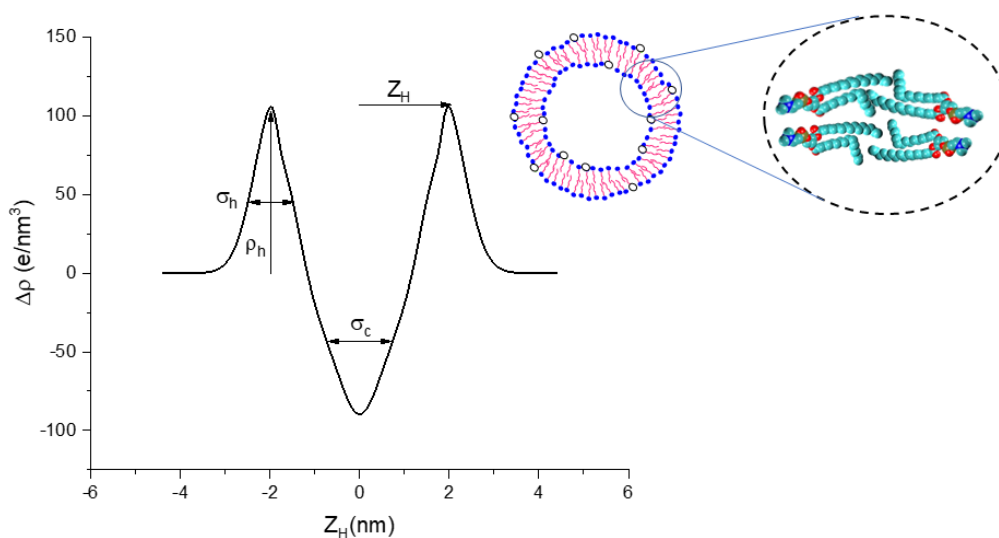
	DPPC/C <sub>12</sub> PC <sub>3</sub> NH <sub>3</sub> Cl				
	DPPC	80/20	60/40	40/60	20/80
$\chi^2_{red}$	1.19	0.75	1.24	1.02	1.05
$\sigma_h(\text{nm})$	0.47±0.05	0.50±0.05	0.52±0.05	0.38±0.05	0.71±0.05
$\rho_h(\text{e}/\text{nm}^3)$	107±10	87±10	92±10	103±10	47±5
$Z_h(\text{nm})$	1.87±0.05	1.81±0.05	1.77±0.05	1.72±0.05	1.30±0.05
$\sigma_c(\text{nm})$	0.43±0.10	0.26±0.10	0.26±0.10	0.28±0.10	0.31±0.10



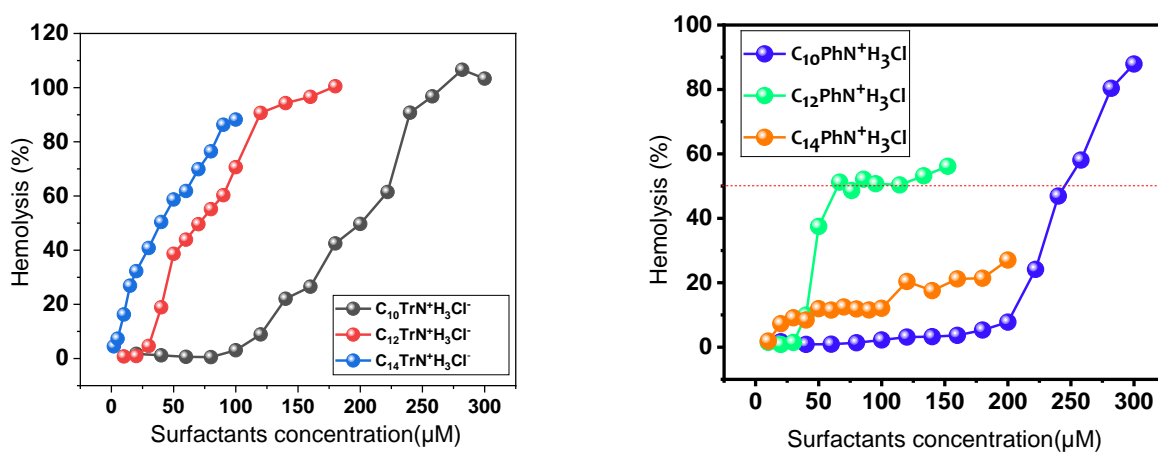
**Figure.S43** A) Scattered intensity patterns as a function of scattering vector modulus for DPPC and  $\text{C}_{14}\text{PC}_3\text{NH}_3\text{Cl}$  and their mixtures the curves correspond to the best fit of Gaussian bilayers or core-shell models. B) The corresponding electron density profiles of the bilayer models corresponding to the best fits of A).

**Table S2** Fitting parameters of Gaussian bilayers for DPPC/C<sub>14</sub>PC<sub>3</sub>NH<sub>3</sub>Cl mixtures.

	DPPC/C <sub>14</sub> PC <sub>3</sub> NH <sub>3</sub> Cl				
	DPPC	80/20	60/40	40/60	20/80
$\chi^2_{red}$	1.19	2.12	4.7	2.55	6.1
$\sigma_h(\text{nm})$	0.47±0.01	0.52±0.01	0.67±0.02	0.17±0.01	0.66±0.03
$\rho_h(\text{e}/\text{nm}^3)$	107±10	107±13	103±15	99±12	105±15
$Z_h(\text{nm})$	1.87±0.05	1.84±0.05	1.71±0.06	1.47±0.05	1.78±0.08
$\sigma_c(\text{nm})$	0.43±0.05	0.31±0.05	0.99±0.10	0.35±0.05	0.69±0.10



**Figure.S44.** Gaussian vesicle model



**Figure. S45.** Hemolysis (%) as a function of surfactants concentration.

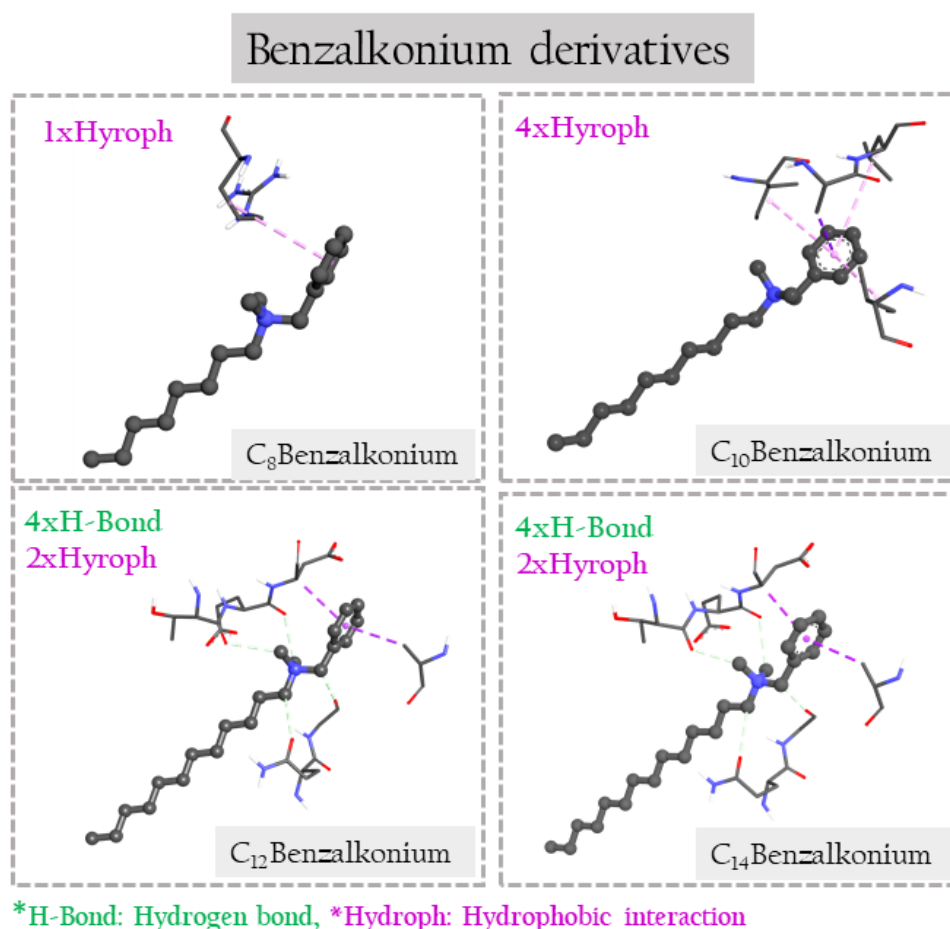


**Table S3** Hemolytic activity of tryptophan and phenylalanine-based surfactants.

	Hemolysis (HC <sub>50</sub> ) in $\mu$ M
C <sub>10</sub> TC <sub>3</sub> NH <sub>3</sub> Cl	200.51
C <sub>12</sub> TC <sub>3</sub> NH <sub>3</sub> Cl	69.88
C <sub>14</sub> TC <sub>3</sub> NH <sub>3</sub> Cl	40.27
C <sub>10</sub> PC <sub>3</sub> NH <sub>3</sub> Cl	244.88
C <sub>12</sub> PC <sub>3</sub> NH <sub>3</sub> Cl	64.55
C <sub>14</sub> PC <sub>3</sub> NH <sub>3</sub> Cl	Not Hemolytic

**Table S4** Results of the interaction details and docking score in (kcal/mol) of C<sub>n</sub>Benzalkonium derivative (From C<sub>8</sub> to C<sub>14</sub> carbon atoms) against the peptidoglycan glycosyltransferase (PDB ID:2OQO).

Inhibitors	Score Kcal/mol	Ligand	Receptor	Interactions		Distance Å
				Categories	Types	
C <sub>8</sub> Benzalkonium	-5.9	Ring(C6)	ARG225	Hydrophobic	Pi-Alkyl	5.0856
		Ring(C6)	ALA141	Hydrophobic	Pi-Sigma	3.52615
C <sub>10</sub> Benzalkonium	-5.8	Ring(C6)	ILE93	Hydrophobic	Pi-Alkyl	5.3345
		Ring(C6)	ILE138	Hydrophobic	Pi-Alkyl	5.14622
		Ring(C6)	LEU142	Hydrophobic	Pi-Alkyl	5.4331
		C(CH2)	GLY114	Hydrogen Bond	Carbon Hydrogen Bond	2.82665
		C(CH2)	GLN113	Hydrogen Bond	Carbon Hydrogen Bond	3.39365
C <sub>12</sub> Benzalkonium	-7.2	C(CH3)	THR82	Hydrogen Bond	Carbon Hydrogen Bond	3.34222
		C(CH3)	GLU83	Hydrogen Bond	Carbon Hydrogen Bond	2.99631
		Ring(C6)	ASP84	Hydrophobic	Pi-Sigma	3.91375
		Ring(C6)	ALA97	Hydrophobic	Pi-Sigma	3.50131
		C(CH2)	GLY114	Hydrogen Bond	Carbon Hydrogen Bond	2.77969
		C(CH2)	GLN113	Hydrogen Bond	Carbon Hydrogen Bond	3.37694
		C(CH3)	THR82	Hydrogen Bond	Carbon Hydrogen Bond	3.28951
C <sub>14</sub> Benzalkonium	-7.4	C(CH3)	GLU83	Hydrogen Bond	Carbon Hydrogen Bond	3.02791
		Ring(C6)	ASP84	Hydrophobic	Pi-Sigma	3.91929
		Ring(C6)	ALA97	Hydrophobic	Pi-Sigma	3.51456



**Figure S46:** Three-dimensional (3 D) closest interactions between active site residues of peptidoglycan glycosyltransferase (PDB ID:2OQO) With CnBenzalkonium (From C<sub>8</sub> to C<sub>14</sub> carbon atoms) derivatives