#### Supplementary information

Title: "Influence of chemical modifications of sugar fatty acid esters on their antimicrobial properties"

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#### Figure S1. Nonanoic acid (1)



Product ion scan mode:  $m/z = 157 [M-H]^{-1}$ 



#### Figure S2. Methyl (3*R*)-3-hydroxynonanoate (3):







Predicted <sup>1</sup>HNMR of PHN methyl ester spectrum:



## Zoom in spectra of proton localized at C7 (experiment):



Zoom in spectra of proton localized at C7 (prediction):











Figure S4. Methyl (3R)-3-(2,2,2-trifluoroethoxy)nonanoate (10)



# <sup>1</sup>HNMR spectrum of PHN-O-CH2-CF3;



Prediction of <sup>1</sup>HNMR spectrum of PHN-O-CH2-CF3:



Zoom in protons localized at C7 and protons from -O- Ch2-CF3 (prediction):





## Zoom in protons localized at C7 and protons from -O-Ch2-CF3 (experiment):

Figure S5. Methyl (3R)-3-(2,2,2-trifluoroethoxy)heptanoate (11)

0.2 0.4 0.6 0.8 1 1.2 1.4 1.6 1.8 2 2.2 2.4 2.6 2.8 3 Counts vs. Acquisition Time (min)



(

20 40 60 80

100 120 140 160 180 Counts vs. Mass-to-Charge (m/z)

200 220 240 260

3.2 3.4

Scan mode:  $m/z = 227 [M-H]^{-1}$ 

0.

Predicted fragmentation: m/z = 163



Figure S6. [(35,65)-3,4,5,6-tetrahydroxyoxan-2-yl]methyl nonanoate (26)



Scan mode:  $m/z = 355 [M+Cl^{-}]^{-1}$ 

#### Predicted fragmentation: m/z= 229



<sup>1</sup>H NMR (400 MHz, Methanol-d4) δ 5.11 (dd, *J* = 10.3, 3.7 Hz, 1H), 4.38 (dd, *J* = 11.8, 2.2 Hz, 1H), 4.21 (dd, *J* = 11.8, 5.4 Hz, 1H), 3.98 (ddd, *J* = 10.1, 5.4, 2.2 Hz, 1H), 3.69 (t, *J* = 9.3 Hz, 1H), 3.41 – 3.24 (m, 2H), 2.35 (t, *J* = 7.4 Hz, 2H), 1.63 (p, *J* = 7.3, 6.8 Hz, 2H), 1.40 – 1.29 (m, 11H), 0.96 – 0.88 (m, 3H).

#### Prediction of <sup>1</sup>HNMR spectrum of C9-Gluc ester:



Figure S7. [(35,65)-3,4,5,6-tetrahydroxyoxan-2-yl]methyl (3R)-3-hydroxynanonoate (27)



Product ion scan mode: m/z = 335 [M-H]-

Scan mode:  $m/z = 371 [M+Cl^{-}]^{-1}$ 









m/z: 173.118318 intensity: 63.870538828638 

Scan mode: m/z = 343 [M+Cl<sup>-</sup>]

Figure S9. [(35,65)-3,4,5,6-tetrahydroxyoxan-2-yl]methyl (3R)- 3-(2,2,2-trifluoroethoxy)nonanoate (29)







# Figure S10. 5-phenylpentanoic acid (5)





Predicted fragmentation: m/z = 115



Figure S11. methyl (3R)-3-hydroxy-5-phenylpentanoate (7)

Scan mode: m/z =193 [M-H]<sup>-</sup> for (3R)-3-hydroxy-5-phenylpentanoic acid



Product ion scan mode: m/z =207 [M-H]-



## Predicted fragmentation: m/z = 177



Figure S12. [(35,65)-3,4,5,6-tetrahydroxyoxan-2-yl]methyl 5-phenylpentanoate (31)



Scan mode: m/z =339 [M-H]<sup>-</sup>





Figure S13. [(35,65)-3,4,5,6-tetrahydroxyoxan-2-yl]methyl (3R)-3-hydroxy-5-phenylpentanoate (33)



Product ion scan mode: m/z = 355 [M-H]<sup>-</sup>

