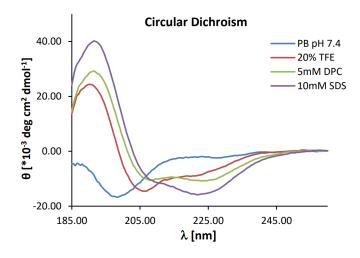
## α-Helical mixed peptide-peptoid antimicrobial peptides to control multidrug resistant Gram-negative bacteria

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Membrane disruptive antimicrobial peptides (AMPs) such as polymyxin B offer an opportunity to control multidrug resistant (MDR) Gram-negative bacteria, which are a leading cause of death in hospitals. Recently we discovered that inverting the chirality of lysine amino acids in an 11-residues  $\alpha$ -helical AMP with strong activity against these bacteria preserved its  $\alpha$ -helical folding and activity while abolishing its hemolytic properties and serum instability. Inspired by several reports of using peptoid building blocks to tune AMP activity, we investigate if our AMP activity might also be tolerant to peptoid substitutions. Our investigations revealed several peptide-peptoid hybrids with preserved  $\alpha$ -helical folding and antibacterial activity, but increased serum stability and reduced hemolysis compared to the parent all-L AMP sequence (Figure).

$$H_3N^+$$
 $H_3N^+$ 
 $H$ 



- [1] N. Mookherjee, M. A. Anderson, H. P. Haagsman and D. J. Davidson, *Nat Rev Drug Discov*, **2020**, *19*, 311–332.
- [2] C. J. Murray et al., The Lancet, **2022**, 399, 629–655.
- [3] S. Baeriswyl, H. Personne, I. D. Bonaventura, T. Köhler, C. van Delden, A. Stocker, S. Javor and J.-L. Reymond, *RSC Chemical Biology*, **2021**, *2*, 1608–1617.
- [4] T. Godballe, L. L. Nilsson, P. D. Petersen and H. Jenssen, *Chemical Biology & Drug Design*, **2011**, *77*, 107–116.