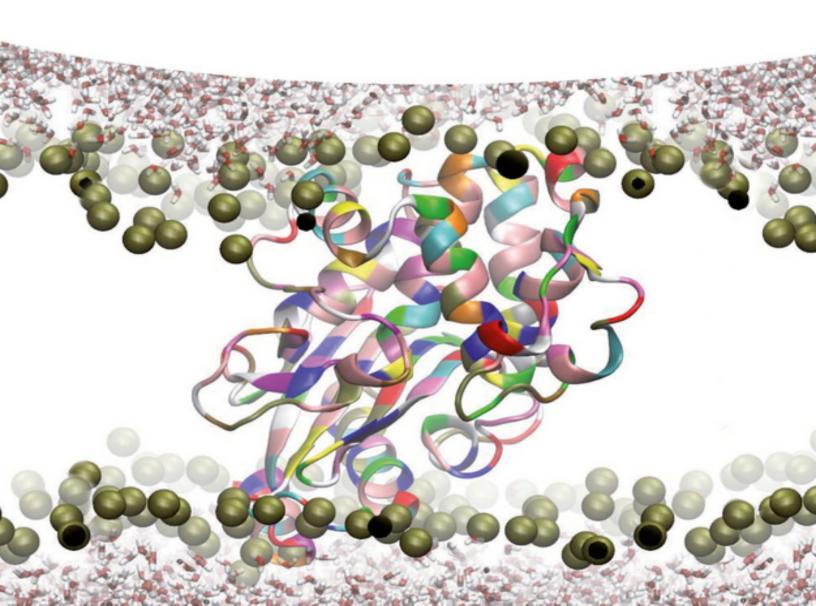
Edited by David A. Phoenix, Frederick Harris, and Sarah R. Dennison

## Novel Antimicrobial Agents and Strategies



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#### Cover design

The cover shows beta-lactamase, an enzyme produced by some bacteria, which provide bacterial resistance to beta-lactam antibiotics in the presence of a lipid bilayer. The image was created by Dr. Manuela Mura, University of Central Lancashire, UK.

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## **Preface**

The "Golden age of antibiotics" was between 1929 and the 1970s when over 20 antibiotic classes were marketed [1, 2]. Since the 1960s, the rise in the emergence of microbial pathogens with multiple drug resistance (MDR) has led to the realization that the "Golden age" had ended. The pharmaceutical industry has been constantly battling with MDR because of the overprescription and misuse of antibiotics [3-5]. In Chapter 1, Radecka and coworkers give an insight into bacterial resistance being a major threat to public health. They also discuss the implications arising from the threat posed by MDR pathogens in relation to factors such as medical practice and economics, along with an overview of recent practices and measures proposed to contain this threat, such as the introduction of stewardship programs. Concern regarding our future ability to combat infection has been further intensified by the decreasing supply of new agents [3, 6-8], and in the remainder of the book we review approaches being taken to identity and develop the antimicrobials of the future.

In response to the challenges outlined, in this book there has been increasing research into maximizing opportunities to develop and revitalize established classes of antibiotics. Coates and Hu consider this area in Chapter 2 where they look at opportunities to extend the life of old antibiotics such as  $\beta$ -lactams by the addition of agents that can overcome drug resistance factors, such as  $\beta$ -lactamase inhibitors. Identification of new, effective derivatives remains a challenge, and another approach in the search for new antibiotics has been to seek out new targets that would enable new classes of antibacterials to be developed. In Chapter 4, Capasso and Supuran review the cloning and

characterization of carbonic anhydrases (CAs). In this chapter, they make reference to the impact of inhibitors that target the  $\alpha$ -,  $\beta$ -, and  $\gamma$ -CAs from many pathogenic bacteria and suggest that this provides evidence that these proteins could provide novel antibacterial targets for the development of new antimicrobial compounds.

There remain concerns, though, that only a small number of drugs are currently under research and development as antibacterial agents [9]. It has been suggested that a further approach could be to revisit naturally occurring compounds with antibacterial potential. Due to the arrival of antibiotics, there has been a rapid loss of interest in the therapeutic potential of natural host antibiotics such as lysozyme [3, 4]. However, more recently, there has been an awakened interest in host defense molecules, such as antimicrobial peptides (AMPs) [10, 11]. Since the early 1990s, the potential of AMPs has been investigated using, for example, magainins isolated from the African clawed frog *Xenopus laevis*, to investigate the effect of the structural and physiochemical properties of these peptides on their antimicrobial action. These AMPs have the potency to target and kill a wide range of Gram-negative and Grampositive bacteria, fungi, viruses, and some tumor cells [12]. Based on this ability, AMPs are attractive propositions for development as therapeutically useful antimicrobial and anticancer agents [13]. The first clinical trials of these AMPs as potential novel antibiotics have been for topical treatments [14], and Dennison et al. review this area in Chapter 4. AMPs are not only produced by eukaryotes but are also generated by prokaryotes, and Lotfipour and coworkers review this class of peptides, generally known as bacteriocins, in Chapter 5. These prokaryotic peptides are produced by gene-encoded or ribosome-independent pathways [15]. Non-ribosomal prokaryotic AMPs generally include examples such as vancomycin and daptomycin,