

Colistin: A Last Resort Antibiotic.

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ARTICLE INFORMATION

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ABSTRACT

Colistin is an antibiotic of last resort, works as a last line of defense against illness caused by the multiple drug resistant (MDR) gram negative bacteria. Polymyxins consist of aqua phobic acyl tail, linear tripeptide part, and heptapeptide ring. Chemical formula of colistin is C₅₂H₉₈N₁₆O₁₃. Colistin is given in two forms, colistin methane sulfonate (CMS) and colistin sulfate.

Findings: Colistin disturbs or destroys the outer membrane of bacteria, or stops the respiratory enzymes of the inner membrane of bacteria. Colistin also works as an antiendotoxin. Colistin has great bactericidal activity against gram negative rods. Colistin resistance is a serious concern of present time. High rate of the spread of MDR gram negative bacterial infections is leading to high use of colistin for treatment, due to which prevalence of colistin resistance is increasing worldwide. As bacterial outer membrane is the main target of colistin, resistance against it develop by modification of outer membrane LPS. There are several mechanisms of colistin resistance. Intrinsic mechanism involves expression of *arnBCADTEF* operon and *eptB* gene. Acquired non-transferable resistance involves *pmrAB*, *PhoPQ*, and *CrrAB* two component systems (TCSs), *mcrB* gene and *ramA*. Transferable resistance is more serious concern as it can be spread in population by horizontal gene transfer. *Mcr* genes are responsible for transferable or plasmid-mediated colistin resistance.

Conclusion: This review focuses on the chemistry of colistin, mechanism of action, spectrum of action, colistin resistance and different mechanisms behind it.

Keywords: Colistin, Last resort antibiotic, Multi drug resistance, Lipid A, Resistance, *mcr* gene



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Original Research Article

Introduction:

Colistin is an antibiotic of polymyxin group and is also known as “polymyxin E”. It exhibits significant antibacterial activity against Gram-negative bacteria⁽¹⁾. Polymyxin B and polymyxin E (Colistin) are two antibiotics among polymyxins which are being clinically used⁽²⁾. Colistin was isolated by *Bacillus polymyxa* in 1949 and use of colistin was started in 1950s as a therapeutic drug. It was reduced in 1980s due to high occurrence of nephrotoxicity. However, use of colistin is gradually increasing as an antibiotic of last resort because of the development and spread of multidrug resistance (MDR) in gram negative bacteria⁽³⁾. Resistance to multiple drugs is a growing threat. In fact, some bacteria have shown resistance against almost all antibiotics of general use. Multidrug resistant gram-negative bacteria are serious threat of present time, no commonly available antibacterial agent is effective against them⁽⁴⁾. Colistin is not commonly applied due to its side effects it is used when no other choice is left. Colistin is effective against MDR gram negative bacterial infections. It is used to treat these MDR infections as the last boundary of defense. If pan drug resistant bacteria become resistant to colistin, then such bacteria may cause untreatable infection. This is why

emergence of colistin resistance is a severe threat to the public health worldwide⁽⁵⁾.

Chemical Structure:

Polymyxins are composed of a cyclic heptapeptide ring, linear tripeptide part, and aquaphobic acyl tail⁽⁶⁾. Colistin is a cyclic polypeptide in structure. Its chemical formula is C₅₂H₉₈N₁₆O₁₃. Colistin has positive charge at several sites which means that it is polycationic. It has both hydrophobic and lipophilic parts⁽⁷⁾.

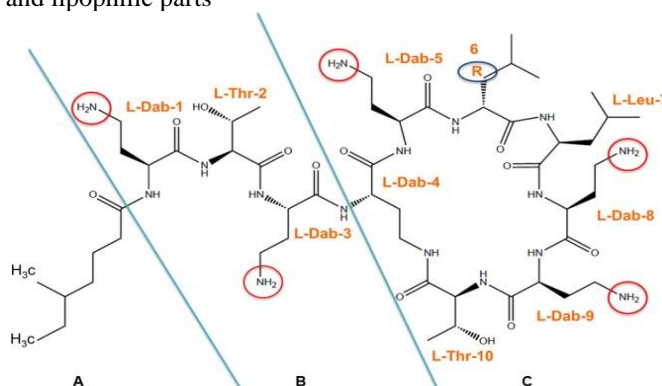


Figure 1 Chemical Structure of Colistin (Mohamed Rhouma, 2016)

Colistin is administered in either of two forms: 1) colistinmethanesulfonate (CMS) or 2) colistin sulfate. Interaction of sodium bisulfate and formaldehyde with colistin produces the CMS. CMS form is given parenterally. It is converted into active colistin in vivo, which is the actual antibacterial agent. Thus, CMS is regarded as an inactive prodrug⁽⁹⁾. Colistin sulfate is used topically. Both dosage forms of colistin can be inhaled in form of aerosols. Colistin sulfate is a stable and cationic form of colistin unlike CMS⁽¹⁰⁾.

Mechanism Of Antibacterial Activity

Colistin targets lipopolysaccharides (LPS) present in the external membrane of bacteria. Hydrophilic and lipophilic parts of colistin interact with the external membrane of gram-negative bacteria. These interactions interrupt the external membrane by displacing divalent cations, especially calcium and magnesium from the phosphate groups of membrane lipids. Lipopolysaccharides are released in result of these reactions. Disarrangement of the external membrane of bacteria directs to the alteration in the permeability, leakage of cell content and cell lysis⁽¹¹⁾.

One more method of that works for the colistin is inhibiting the respiratory enzymes of the bacterial inner membrane. Inhibition of these enzymes leads to the closure of many metabolic activities of bacteria⁽¹²⁾. Colistin also acts as an antiendotoxin by binding and neutralizing the lipopolysaccharide molecule of bacteria⁽¹³⁾.

Spectrum of action

Colistin is very effective drug against gram negative bacteria. It exhibits excellent bacteria-killing activity. *Acinetobacter* species, *P. aeruginosa*, *K. pneumoniae*, *E. coli*, and *Enterobacter* spp are some of the bacteria that are highly susceptible to colistin. While *Providentia* spp, *Serratia* spp, *Brucella* spp are resistant to colistin. Colistin is employed to cure infections caused by MDR-gram negative bacteria^(14, 15).

Resistance

Colistin is widely utilized as a remedy for diseases caused by MDR-gram negative bacteria, because of the increasing prevalence of multiple drug resistance⁽¹⁶⁾. Due to the enhancement in the utilization of colistin to cure MDR infections, colistin resistance is increasing worldwide. It is a matter of deep concern because colistin is the last line treatment option against these pathogens⁽¹⁷⁾.

Mechanisms Of Resistance

Outer membrane of the cell is the main place of work for colistin. Bacteria develop resistance against colistin mainly by alterations in the outer membrane. These changes may include loss of Lipopolysaccharides (LPS), loss of specific outer membrane proteins, and membrane lipid alterations⁽¹¹⁾. Various mechanisms have been reported by which bacteria develop resistance against colistin.

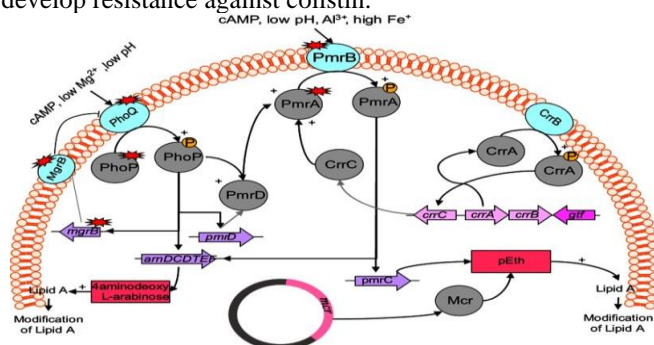


Figure 2 LPS modifications pathways in Enterobacteriaceae (al A. e., 2019)

Intrinsic Resistance:

Proteus mirabilis and *S. marcescens* are naturally resistant to colistin and this intrinsic resistance is associated with translation of *arnBCADTEF* operon and *eptB* gene. Expression of this operon and gene leads to alteration of the LPS by cationic replacement. The modification of LPS is done by the addition of phosphoethanolamine (pEtN) and 4-amino-4-deoxy-L-arabinose (L-Ara4N) to LPS⁽¹⁹⁾. A significant role is played by the *TherppA/rppB* two component systems (TCS) to activate *arnBCADTEF* operon (20). O-acetyltransferase gene involves the biosynthesis or transfer of amino arabinose⁽²¹⁾.

Acquired Resistance:

Pmrab And Phopq Two Component Systems

L-Ara4N is synthesized from uridinediphosphateglucuronic acid by the *pmrHFIJKLM* operon and *PmrE* gene⁽²²⁾. *PmrB* gene is activated by contact with aluminum, high level of iron, acidic pH, and macrophage phagosomes. *PmrB* activates *PmrA* by its kinase activity using phosphorylation. Then, *pmrABC* and *pmrHFIJKLM* operons and the *pmrE* gene are regulated by *pmrA*. The modification of LPS is done by fixing the L-Ara4N and PETN to lipid A, and this action is performed by these genes and operons⁽²³⁾. Changes in the genetic makeup of *pmrA/pmrB* upregulate the *pmrE* gene, and, *pmrHFIJKLM*, and *pmrABC* operons⁽²⁴⁾.

Low concentration of magnesium or calcium ions activates *phoQ*. *PhoQ* plays role in the activation of *phoP*. Transcription of the *pmrHFIJKLM* operon is activated by the activation of *PhoP* via phosphorylation. Lipid A is then modified. *PhoPQ* TCS can also modify lipid A by another way. The transcription of *pmrHFIJKLM* operon is activated by the indirect activation of *pmrA* by *PhoP* through evading *pmrD* connector protein. First synthesized PETN then added to lipid A. PETN can be added to LPS to modify them by the help of different PETN-coding genes⁽²⁵⁾.

mgrB gene

A little transmembrane protein having 47 amino acids encoded by *mgrB* gene gives negative feedback on *PhoPQ* TCS⁽²⁶⁾. This protein inhibits the phosphorylation via stopping the phosphorylation of *PhoQ*. Mutation or inactivation of *mgrB* gene upregulates the *PhoPQ* operon and activates *pmrHFIJKLM* operon. At the end, production of L-Ara4N modifies the lipid A, which leads to colistin resistance⁽²⁷⁾.

CrrAB two component system

The mutated *CrrB* protein leads to the regulation of *crrAB*-adjoining gene that translates a protein resembling glycosyltransferase, which modify the lipid A⁽²⁸⁾. *PmrC*, and *pmrE* genes, and *pmrHFIJKLM* operon are activated by the inactivation or change in the genetic makeup of *crrB* gene via upregulation of the *pmrAB* operon. Subsequently, colistin resistance is led by the synthesis and addition of L-Ara4N and PETN to lipid A. Change in the genetic makeup of *crrB* gene causes increased *CrrC* transcription. Amino acid replacements of the *CrrB* protein enhance self-phosphorylation of kinase, which results in resistance to colistin⁽²⁹⁾.

Regulator Rama

The *ramA* locus consists of *ramA*, *romA*, *ramR* genes. *RamR* acts as a repressor of the *ramA*, and *romA* genes. The important role is played by *ramA* regulator in LPS modification which results in the development of resistance to colistin. *lpxA*, *lpxC*, *lpxD*, *lpxB*, *lpxK*, *lpxL*, *lpxM*, and *lpxO* genes play role in lipid A biosynthesis. *RamA* activates *lpxC*, *lpxO*, and *lpxL2* genes, and creates changes in the lipid A⁽³⁰⁾.

Plasmid-mediated colistin resistance:

mcr genes

Plasmid-based colistin resistance is a serious concern, because colistin resistance genes are easily transferred to susceptible strains by horizontal gene transfer methods and make susceptible strains resistant. The mcr genes are the main culprits behind horizontal transmission of colistin resistance. Colistin resistance involving mcr gene was first observed in *E. coli* isolated from pigs in China. Expression of mcr genes directs to the inclusion of PETN to lipidA⁽³¹⁾.

Transferable colistin resistance due to mcr-1 gene has been reported in many regions. It is most frequently reported in *E. coli*. Mcr-2, mcr-3, mcr-4, mcr-5, mcr-6, mcr-7, and mcr-8 are other mcr genes, reported after mcr-1⁽³²⁾.

Role of Capsular Polysaccharides (CPS) in colistin resistance: CPS protects bacteria from cationic antibacterial peptides, including colistin. Resistance level is associated with number of capsule layers. More capsular layers lead to more colistin resistance. Cpx and Rcs are regulators of capsule formation. Ugd gene also contributes in CPS biosynthesis⁽³³⁾.

Conclusion:

Colistin is an antibiotic of last resort, used to cure MDR gram negative bacterial infections. It targets the LPS of bacterial outer membrane mainly. Bacteria have developed ways to protect itself from colistin. Prevalence of colistin resistance is increasing worldwide, which is a serious global concern. Bacteria can modify the lipid A region of bacterial outer membrane LPS. These modifications lead to resistance against colistin. Mcr gene is the main culprit of plasmid mediated colistin resistance. Other mechanisms involve pmrAB, phoPQ, and CrrAB two component systems. Plasmid mediated or transferable resistance is more serious concern than non-transferable, because it can spread in population by horizontal gene transfer.

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