



Reduction of *Vibrio Cholerae* Persistence with mutSMAP-18

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DESCRIPTION

Vibrio cholerae, the causative agent of cholera, poses a significant public health challenge, especially in regions with inadequate water and sanitation infrastructure. The emergence of antibiotic resistance in *V. cholerae* has further complicated its management, highlighting the need for alternative antimicrobial strategies. One such alternative is Mutational Signal Mutation Analysis Panel (mutSMAP-18), a synthetic peptide derived from Antimicrobial Peptides (AMPs) that has shown potential in combating bacterial infections.

Overview of *Vibrio cholerae* and its challenges

V. cholerae is a gram-negative bacterium responsible for cholera, a severe diarrheal disease characterized by acute dehydration and electrolyte imbalance. The bacterium's ability to form biofilms and its growing resistance to conventional antibiotics present challenges in treatment and prevention. Biofilms, complex communities of bacteria encased in an extracellular matrix, enhance bacterial survival in adverse environments and contribute to resistance against antimicrobial agents.

The study of Antimicrobial Peptides (AMPs) offers a potential direction in addressing these challenges. AMPs are naturally occurring molecules that exhibit broad-spectrum antibacterial activity and have demonstrated effectiveness against biofilm-forming pathogens. Synthetic derivatives, such as mutSMAP-18, have been designed to enhance these properties and provide targeted solutions against resistant strains.

Antibacterial activity of mutsmap-18 against *Vibrio cholerae*

MutSMAP-18 exhibits potent antibacterial activity against *V. cholerae*, targeting both planktonic cells and biofilm-associated bacteria. The peptide's mechanism of action involves disruption of the bacterial cell membrane, leading to rapid cell lysis. Studies have demonstrated that mutSMAP-18 interacts with negatively charged components of bacterial membranes, such as Teichoic

acids, Lipopolysaccharide (LPS), causing structural destabilization and loss of membrane integrity.

Minimal Inhibitory Concentration (MIC) assays have shown that mutSMAP-18 effectively inhibits the growth of *V. cholerae* at low concentrations, making it a potent agent against the pathogen. Comparative analyses with conventional antibiotics reveal that mutSMAP-18 retains its activity against antibiotic-resistant strains of *V. cholerae*, highlighting its potential utility in clinical settings.

Antibiofilm properties of mutsmap-18

Biofilms are a major factor in the persistence and virulence of *V. cholerae*, providing a protective barrier against environmental stresses and antimicrobial agents. MutSMAP-18 demonstrates significant antibiofilm activity, disrupting established biofilms and preventing their formation.

Inhibition of biofilm formation: mutSMAP-18 has been shown to interfere with the initial stages of biofilm formation by inhibiting bacterial adhesion to surfaces. This property is attributed to its interaction with bacterial surface components, which prevents the establishment of stable biofilm structures. Experimental data indicate a dose-dependent reduction in biofilm biomass when *V. cholerae* cultures are treated with mutSMAP-18.

Disruption of established biofilms: In addition to preventing biofilm formation, mutSMAP-18 is capable of penetrating and disrupting mature biofilms. The peptide destabilizes the Extracellular Polymeric Substances (EPS) matrix, the structural foundation of biofilms, exposing bacteria to external threats and antimicrobial agents. This dual activity inhibition and disruption makes mutSMAP-18 a versatile tool in managing biofilm-associated infections.

Mechanisms underlying the activity of mutsmap-18

The efficacy of mutSMAP-18 against *V. cholerae* can be attributed to its unique structural and functional properties. The peptide's

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amphipathic nature enables it to interact selectively with bacterial membranes, causing membrane disruption without significantly affecting mammalian cells. This selective toxicity reduces the risk of adverse effects, a common limitation of many conventional antibiotics.

Interaction with bacterial membranes: mutSMAP-18's mechanism involves electrostatic interactions with negatively charged bacterial membranes, followed by insertion into the lipid bilayer. This process results in pore formation, membrane depolarization and subsequent cell death. The rapid action of mutSMAP-18 minimizes the likelihood of bacterial adaptation and resistance development.

Interference with biofilm components: The peptide's ability to disrupt biofilms is linked to its interaction with EPS matrix components. By degrading the structural integrity of the matrix,

mutSMAP-18 enhances the susceptibility of biofilm-embedded bacteria to antimicrobial agents.

CONCLUSION

MutSMAP-18 offers a significant alternative to conventional antibiotics in combating *Vibrio cholerae*, particularly in addressing antibiotic resistance and biofilm-related challenges. Its potent antibacterial and antibiofilm activities, achieved through membrane disruption and interference with biofilm components, highlight its potential as an effective therapeutic agent. By targeting both planktonic and biofilm-associated bacteria, mutSMAP-18 could play a crucial role in managing cholera infections, especially in settings where antibiotic resistance is prevalent.