

Molecular uptake of antibiotics through the silent chitoporin from *Escherichia coli* (*Ec*ChiP)

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Abstract: EcChiP is a monomeric protein channel found in the outer membrane (OM) of Escherichia coli (E. coli) and used for the transport of small chitooligosaccharides across the OM. In this study, we identified antibiotic transport through the *Ec*ChiP-reconstituted in lipid membrane. The antibiotics that can inhibit bacterial growth were further chosen for evaluating their specific interactions with *Ec*ChiP channel. The results show that 2 mM of gentamycin, minocycline, and tigecycline could occlude the ion flow through the EcChiP channel, indicating that these antibiotics could enter and interact with the channel lumen. Crystallizations of *Ec*ChiP in complex with gentamycin, minocycline, and tigecycline were grown in the optimized condition G8 (33% PEG400, 0.1 M sodium chloride, 0.1 M MES, pH 6.5) from MemGold1[™] and the condition E10 (33% PEG400, 0.23 M sodium chloride, 0.05 M sodium acetate, pH 4.5) from MemGold2TM screening kits. The crystals have a long tetragonal shape with estimated sizes of $300 - 500 \,\mu\text{m}$. The predicted structure of EcChiP in complex with minocycline, gentamycin, and tigecycline indicated that all three antibiotics occupied the constriction zone of the protein pore. Understanding the antibiotics-EcChiP interactions may suggest an effective approach to design for novel anti-microbial agents against infectious strains of Escherichia coli.

Keywords: antibiotic resistance, chitoporin, E. coli, single channel



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