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# A novel streptomycin and spectinomycin resistance Gene cassette occurrence in *E. cloacae* isolated from Zhenjiang

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**Abstracts:** The *aadA* genes, encoding resistance to streptomycin and spectinomycin, have been found as gene cassettes in different gram-negative and gram-positive bacterial species. The present report has revealed that the sequence of a new gene, *aadA5*, combining with the trimethoprim resistance gene *dfr17* occurred in a class I integron. The integron was identified in a nosocomial pathogen *Enterobacter cloacae* isolates, which indicated that integrons are specialized genetic elements that were capable of capturing, integrating and mobilizing gene-cassette by site-specific recombination and integron was a dangerous factor to lead to bacterial resistance.

**Keywords:** *E. Cloacae*, Class I Integron, Resistance

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In recent years, *Enterobacter cloacae* (*E. cloacae*) has become a well-recognized nosocomial pathogen. *E. cloacae* isolates from clinical specimens may be resistant to multiple antibiotics. For example, *E. cloacae* is intrinsically resistant to ampicillin and narrow-spectrum cephalosporins [1]. And a substantial proportion of multiresistant *E. cloacae* isolates carry integrons. Integrons are specialized genetic elements that was capable of capturing, integrating and mobilizing gene cassettes by site-specific recombination. Gene cassettes mainly encoded antibiotics genes. There are five classes integrons involved in bacterial resistance to date, While class I integrons are most frequently found in antibiotic-resistant gram negative bacteria [2]. The structure of the class I integron includes 5' and 3' conserved segments and a variable region [3].

The *aadA* genes which were first reported by Hollingshead and Vapnek [4], are the only characterized genes that encode both streptomycin and spectinomycin resistance, and many of these genes are found as gene cassettes in class I integron [5]. The present report characterizes the nucleotide sequence and expression of a novel streptomycin and spectinomycin resistance gene located as a gene cassette. In a two-year period (2005–2006), 15 multiresistant *E. cloacae* strains were isolated in our laboratory, which carried a *aadA5*-containing integron. To the best of our knowledge, *E. cloacae* strains carrying aminoglycoside-3'-adenylyltransferase genes have not been reported before. Therefore, we undertook this study to

analyse the structure of the *aadA5*-containing integron.

## 1. Materials and Methods

### 1.1. Bacterial Strains and Susceptibility Testing

The *E. cloacae* isolates were identified by the conventional method, and with the VITEK system card (bioMérieux, Hazelwood, MO, USA). Antibiotic-containing discs (Oxoid Ltd., England) were used for routine antibiograms by disc diffusion assay. MICs of antimicrobial agents were determined by agar dilution in a Mueller–Hinton Media (Oxoid Ltd., Basingstoke, Hampshire, England) according to the guidelines of the NCCLS [6]. *Escherichia coli* ATCC 25922 was used as MIC reference strain.

### 1.2. *E. Cloacae* DNA Extraction

The *E. cloacae* DNA was extracted using the boiling lysated method [7]. The template was prepared by suspending a loopful of each isolate in 200mL of sterile water, followed by boiling for 10 min and centrifuging for 3 min.

### 1.3. Amplification and Sequencing of Class I Integrons

The gene cassette regions for the class I integrons were performed by PCR amplification with the following sets of primers: for the class I integron, 5'CS-F (5'-GGC ATC

CAA GCA GCA AG-3') and 3'CS-R (5'-AAG CAG ACT TGA CCT GAT-3') (8). PCRs were performed in volumes of 15µl under the following conditions: 1.5µl 10×PCR Buffer (Mg<sub>2</sub><sup>+</sup> plus), 0.2 mM of each deoxynucleoside triphosphate, 0.1 µM of each primer, and 1.5 units of Taq DNA or LA Taq DNA polymerase (Takara Biotechnology (Dalian) Co., Ltd.) for 10 min at 94°C; 35 cycles, with 1 cycle consisting of 45 seconds at 94°C, 1 to 2 min at 55 to 60°C, and 1 to 2 min at 72°C, and a final step of 10 min at 72°C. Sequencing reactions were performed by Shanghai GeneCore BioTechnologies Co.,Ltd. Database similarity searches for nucleotide and deduced amino acid sequences were carried out at the NCBI website (<http://www.ncbi.nlm.nih.gov>).

#### 1.4. Nucleotide Sequence Accession Number

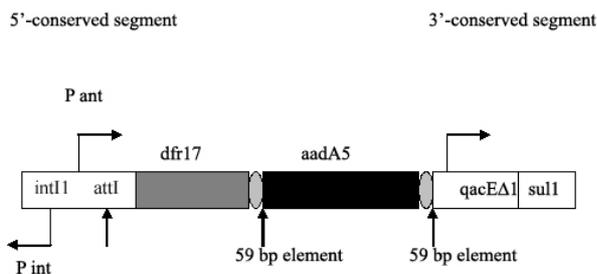
The nucleotide sequences of the *aadA5* gene of *E. cloacae* isolate have been assigned to the GenBank nucleotide sequence database (GenBank accession no EF571855).

## 2. Results and Discussion

### 2.1. Properties of 15 *E. Cloacae* Strains

MIC testing revealed that *E. cloacae* isolates was resistant to most β-lactams, including ampicillin, ampicillin-sulbactam, piperacillin, piperacillin-tazobactam, cefalothin, cefoxitin, cefotaxime, ceftazidime, aztreonam, and *E. cloacae* isolates was high-level resistant to streptomycin and spectinomycin. MICs of aztreonam for the isolate was 64 mg/L. MICs of ampicillin, ampicillin-sulbactam, piperacillin, piperacillin-tazobactam, cefalothin, cefoxitin, cefotaxime and ceftazidime were >128 mg/L.

### 2.2. Sequence Analysis of the *aadA5*-Containing Integron



**Fig 1.** Schematic map of the novel class 1 integron containing the *aadA5* gene cassette from the *E. cloacae* isolate. Designations are as follows: *intI1*, integrase; *attI*, recombination site; *dfr17*, dihydrofolate reductase (trimethoprim resistance) gene; *aadA5*, streptomycin and spectinomycin resistance gene; *qacEA1*, quaternary ammonium compound resistance gene; *sulI*.

PCR-based experiments showed that the *aadA5* gene occurred 15 multiresistant *E. cloacae* strains and also found the 15 isolates had the identical amplicon. Sequence analysis of the 1664 bp amplicon revealed the structure of the class 1 integron, such as the 5'-CS element containing an *IntI1* integrase gene with its own promoter region and the 3'-CS

element containing *qacEA1*. The integron contained insert gene cassettes *dfr17*, *aadA5*, and two putative 59 base element. The *aadA5* gene was located immediately downstream of the first putative 59 base element. The *aadA5* gene cassettes had a 59 bp element, and the *qacEA1* gene also had a core site (Fig. 1).

The aminoglycoside-3'-adenylyltransferase genes have spread among gram-negative bacilli, including *Pseudomonas aeruginosa*, *Acinetobacter baumannii* and *Escherichia coli* [8,9]. However, this enzyme was rarely found in *E. cloacae*. The presence of the mobile *aadA5* gene cassette integron in *E. cloacae* suggests that the resistance can spread to other members of the family Enterobacteriaceae [10-12]. In the new class 1 integron in this study also contained a dihydrofolate reductase gene (*dfr17*) besides *aadA5* cassette, suggesting that the gene responsible for the trimethoprim resistance.

The present study has characterized a novel streptomycin and spectinomycin resistance gene cassette found in a class 1 integron. The distribution of this gene is still to be investigated. The novel gene cassette is present in a class 1 integron, which is mobile. These factors enhance the mobilizing possibilities for the *aadA5* gene, and the conditions of transfer and the molecular epidemiology of this gene cassette will need further attention in the future.

## 3. Conclusion

The present work indicated that integrons are specialized genetic elements that were capable of capturing, integrating and mobilizing gene-cassette by site-specific recombination and integron was a dangerous factor to lead to bacterial resistance.

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