

Molecular Characterization of Efflux Pump Mediated Resistance among Multidrug Resistant Clinical Isolates of *Pseudomonas aeruginosa*

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BACKGROUND

- > Pseudomonas aeruginosa is an opportunistic human pathogen characterized by an innate resistance to multiple antimicrobial agents.
- A major contribution to this intrinsic multidrug resistance(MDR) is provided by a number of broadly-specific multidrug efflux systems, among which members of the Resistance Nodulation cell Division family (RND) viz., MexAB-oprM, MexXY-OprM, MexCD-OprJ & MexEF-OprN are predominant.
- MDR infections are difficult to treat with all available antibiotics and these can be controlled by combining drugs with efflux pump inhibitors (EPIs) in order to prevent treatment failure.
- > Studies on efflux mediated resistance among MDR are lacking.
- Hence, this study is aimed to determine the effect of efflux pump inhibitor and the overexpression of efflux pumps in clinical isolates of *P.aeruginosa*.

| Efflux pump of RND family | Substrates |
|---------------------------|---|
| MexAB-OprM | Fluoroquinolones, tetracycline, chloramphenicol and β-lactam |
| MexCD-OprJ | Fluoroquinolones and the antipseudomonal β-lactams (piperacillin, cefepime and meropenem) |
| MexXY-OprM | Fluoroquinolones, aminoglycosides, trimethoprim and chloramphenicol |
| MexEF-OprN | Fluoroquinolone, aminoglycoside and selected β-lactam (piperacillin, cefepime and meropenem but not carbenicillin, ceftazidime or imipenem) |

MATERIALS AND METHODS

1. Strains

- ✓ A total of 213 *P. aeruginosa* clinical isolates were collected from two tertiary care hospital and these were isolated from pus(67%), tracheal wash(15%), urine(10%), blood(4%), bronchoalveolar lavage(2%), semen (1%), tracheal wash (1%).
- ✓ Identification of *P. aeruginosa* was done by standard biochemical methods.
- ✓ PAO∆*mexR::Gm* (MexAB-OprM), PAONB(MexCD-OprJ),
 PAO↑EF(MexEF-OprN), PAO∆*mexZ::Gm* (MexXY-OprM) were used as reference strains; PAO1 was used as control strain.

2. Antimicrobial Susceptibility Testing(AST)

✓ AST was performed for the following antibiotics and interpreted according to CLSI guidelines 2013.

| Piperacillin (100µg) | Piperacillin/ tazobactam (100μg/10μg) | Ceftazidime (30µg) |
|----------------------|---|-----------------------|
| Cefepime (30µg) | Ceftazidime/ Clavulanic acid (100µg/10µg) | Aztreonam (30µg) |
| Amikacin (30µg) | Gentamicin (10µg) | Netilmicin (30µg) |
| Tobramicin (10µg) | Ciprofloxacin (5µg) | Levofloxacin (5µg) |
| Ofloxacin (5µg) | Imipenem (10µg) | Meropenem (10µg) |

3.Phenotypic analysis

- The following reporter antibiotics were used as phenotypic markers of the Mex efflux pumps: carbenicillin (MexAB-OprM), erythromycin (MexCD-OprJ), gentamicin (MexXY-OprM) and norfloxacin (MexEF-OprN).
- * Minimum Inhibitory Concentration (MICs) was determined by microbroth dilution in the absence or presence of a broad spectrum inhibitor of Mex pumps, namely Phe-Arg β-naphthylamide (PAβN), also known as MC-207,110 at a concentration of 50µg/ml (Narcisa Mesaros *et al.*, 2007).
- * Efflux was considered as possible when there was a reduction in $2\log_2$ dilution of MIC with the addition of PAβN.

4. Genotypic analysis

- ✓ Twenty one clinical isolates of *P. aeruginosa* and reference strains were harvested at the late log phase of growth and total RNA was isolated by Qiagen kit method.
- ✓ Conversion of cDNAs was carried out by Qiagen reverse transcription kit.

Detection of mexA, mexC, mexE and mexX genes by real-time PCR

SYBR green master mix was used with the following conditions: 2 min denaturation at 94°C; 40 cycles (94°C for 10sec, 55°C for 1min and 72°C for 1min); melting curve analysis 60–95°C with continuous fluorescence readings; *rpoD* used as endogenous control.

List of primers used in this study(Quale et al., 2006)

| Gene | Primer Sequence 5' 3' |
|------|--------------------------|
| rpoD | F- GGGCTGTCTCGAATACGTTGA |
| | R- ACCTGCCGGAGGATATTTCC |
| mexA | F-AACCGAACAACGAGCTG |
| | R- ATGGCCTTCTGCTTGACG |
| mexC | F-GGAAGAGCGACAGGAGGC |
| | R-CTGCACCGTCAGGCCCTC |
| mexE | TACTGGTCCTGAGCGCCT |
| | TCAGCGGTTGTTCGATGA |
| mexX | GGCTTGGTGGAAGACGTG |
| | GGCTGATGATCCAGTCGC |

RESULTS

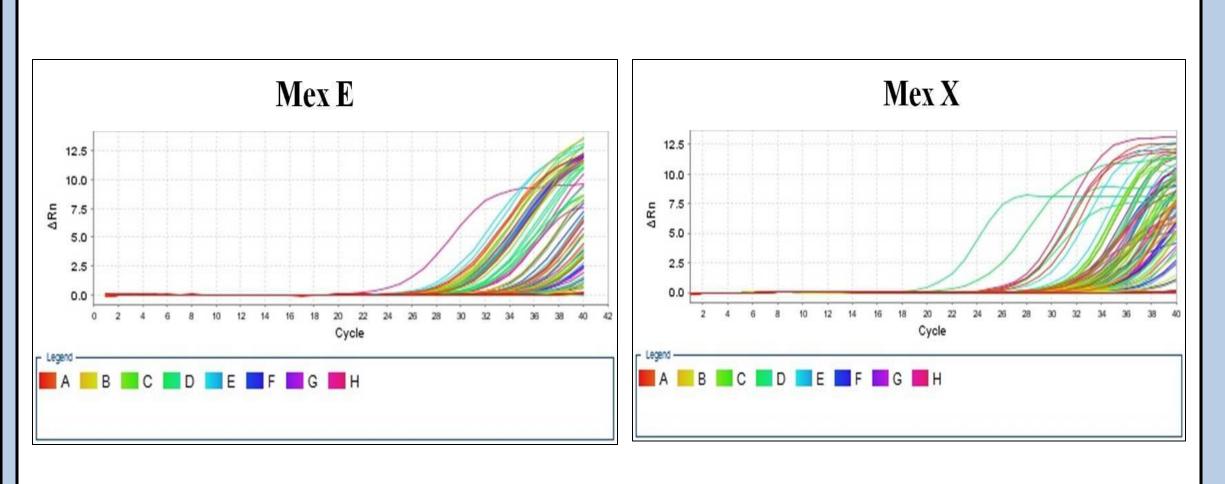
1. Phenotypic results

- ✓ Forty isolates which were resistant to aminoglycosides, fluoroquinolones and cephalosporins by AST were chosen for the phenotypic study of efflux pump.
- ✓ Efflux was considered likely if the MIC of a given strain was at least 2 log2 dilutions higher than the wild-type strain.
- \checkmark For the reference strains overexpressing a mex pump, the addition of PAβN reduced the MIC values for the corresponding antibiotic markers 4 to 64 folds.
- ✓ The lowest-reduction (4 fold) was observed for gentamicin MIC in $PAO\Delta mexZ::Gm$ (MexXY-OprM).
- ✓ For clinical isolates, MICs difference were ranging from 2 to 32 fold for carbenicillin; 2 to 256 fold for erythromycin; 2 to 64 fold for erythromycin; 4 fold in 4, 8 in 1 and 64 in 1 strain for gentamicin.
- ✓ The higher fold difference 128 and 256 were observed in 3 and 1 strain for erythromicin; 64 in 3 and 32 in 2 strains for norfloxacin; 32 in 4 strains for carbenicillin

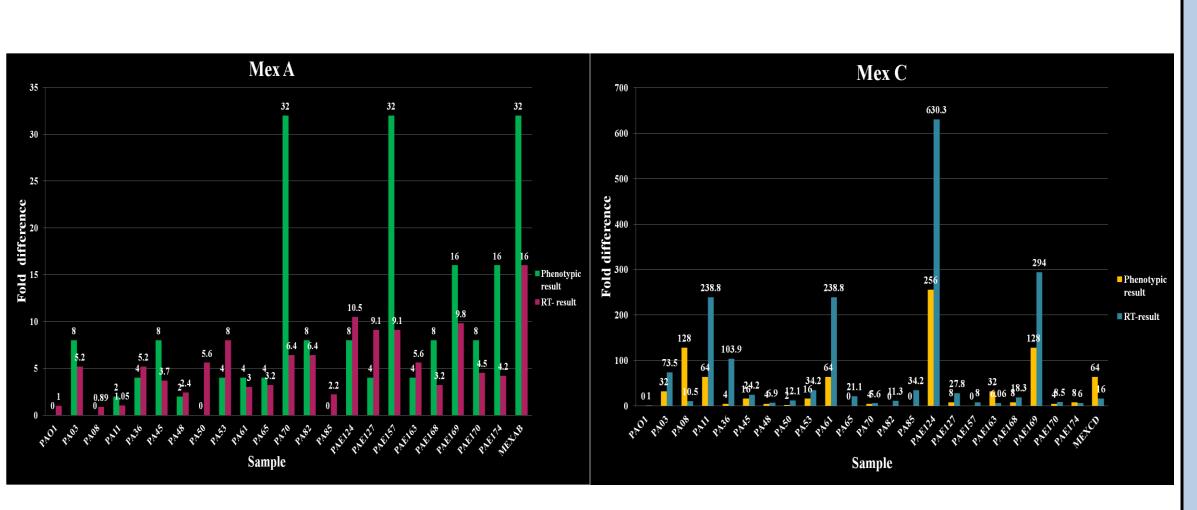
2. Genotypic results

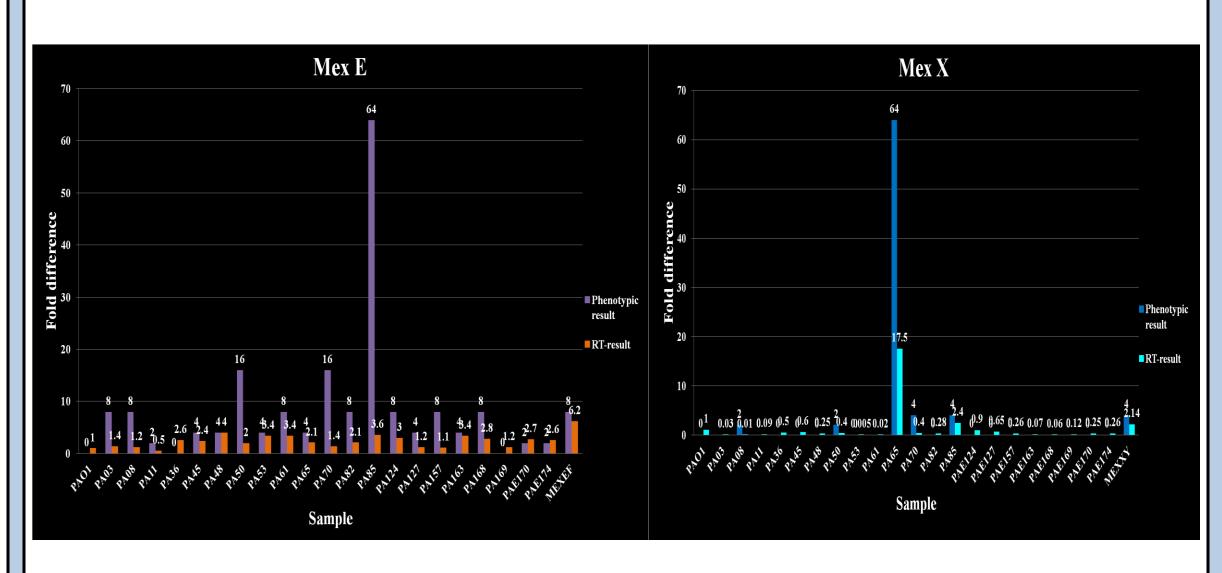
- ✓ MexAB-OprM, MexCD-OprJ, MexEF-OprN and MexXY-OprM efflux was considered likely for 3-fold mexA, mexC, mexE and mexX overexpression than wild type PAO1 strain by QC-RT-PCR.
- ✓ mexA and mexC were overexpressed by 16 times whereas mex E and mexX were overexpressed by 6.2 and 2.14 times in reference strains.
- ✓On comparison with PAO1 strain, expression of *mexA* and *mexE* increased by 3-10 and 2-3fold respectively in all clinical isolates.
- ✓ mexC expression ranged from 5-24 fold increase in 14(67%) isolates, 73-294 fold and 630-fold in 4 and 1 isolate respectively.
- ✓One isolate each showed 2 and 17 fold difference in *mexX* expression whereas no expression was observed for the remaining isolates.
- ✓ For reference strains, a good correlation was observed between the phenotypic and genotypic methods.
- ✓ For the clinical strains, the correlation was good for MexXY-OprM (18 strains with complete convergence), MexAB-OprM and MexCD-OprJ (17 strains with complete convergence) and fair for MexEF-OprN (4 strains with complete convergence).

Amplification plot Mex A Mex C Outline B C D B E F G H



Comparison of phenotypic and genotypic results





CONCLUSION

- ✓ The present study documents the prevalence of efflux pump mediated resistance among MDR isolates of *Pseudomonas aeruginosa*.
- ✓ All MDR isolates over-expressed at least one of the efflux pumps.
- ✓ Among the RND family genes studied, *mexC* showed a wide range as well as high levels of expression indicating its role in multidrug resistance in *P.aeruginosa*.

REFERENCES

- 1. Narcisa Mesaros, Youri Glupczynski, Lae "titia Avrain, Nancy E. Caceres, Paul M. Tulkensm and Francoise Van Bambeke. A combined phenotypic and genotypic method for the detection of Mex efflux pumps in *Pseudomonas aeruginosa. J Antimicrob Chemother* 2007; 59:378–386.
- 2. John Quale, Simona Bratu, Jyoti Gupta, and David Landman. Interplay of Efflux System, *ampC*, *and oprD* Expression in Carbapenem Resistance of *Pseudomonas aeruginosa* Clinical Isolates. *Antimicrob Agents Chemother* 2006; 50: 1633–1641.

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