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9th International Conference on

COPD AND LUNGS

August 19-20, 2019 Tokyo, Japan



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Serotype and genetic variations in COPD infected with Streptococcus pneumoniae in Tehran

A total of 100 *Streptococcus pneumoniae* were collected in Tehran, Iran. The strains were tested for antimicrobial susceptibility and Minimum Inhibitory Concentrations (MIC), serotyped, and genotyped by Multilocus Sequence Typing (MLST). The most frequent serotypes amongst the isolates of *S. pneumoniae* (PNSP) were 14 (24%), 23F (18%) and 19F (17%). MLST indicated a high degree of genetic diversity amongst the 93 PNSP with 36 different sequence types. Six internationally known penicillin resistant clones were identified in our isolates amongst which Spain^{23F}-1 (ST81), Spain^{6B}-2 (ST90), Spain^{9V}-3 (ST156) were the predominant clones. The results indicated international identifiable clones of *S. pneumoniae*, especially Spain^{23F}-1 with high penicillin resistance, could play a major role in spread of antimicrobial resistant in Iran. The extensive sequence variation in PBP2x, PBP2b, and PBP1a in resistant strains was suggestive of a widespread homologous recombination within *S. pneumoniae populations*.

Biography

Mohammad Reza Pourshafie is professor of Microbiology Department and had done his Ph.D in Bacteriology at Institute Pasteur of Iran, Tehrat.

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