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# the relationship between plasmid and some antibiotics resistance in pseudomonaeruginosa

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*Pseudomonas aeruginosa* has been considered to be one of the causes of serious diseases in man and it infrequently is the cause of infection in normal individuals unless they have suffered from major trauma or burns. It is involved in respiratory, cutaneous and disseminated infections in individuals who have defective host defenses of cutaneous barriers, granulocytes or a immune system. Several plasmids encoding resistance to one or more antibiotics could be identified from clinical isolates of *Pseudomonas aeruginosa*. The present investigation was carried out in order to clarify the 400 and 500 µg/ml respectively and for isolate no 5 were; 12.5, 25, 100, 200 and 100 µg/ml respectively. Single colonies of isolate no. 1 showed great variations in their levels of resistance to the different antibiotics indicating that the resistant genes are located in different plasmids. Isolate no. 2 single colonies were similar in their resistance to gentamicin levels indicating chromosomal mode of inheritance. In isolate no. 3. Km and Sm resistant genes were encoded by the same plasmids. Isolates no. 4 and 5 single colonies showed different levels of resistance to Km, SOI and Gm with great agreement for resistance levels indicating the existence of the genes in the same plasmids. The percentages of *E. coli* K12 transformants which resist 10 µg/ml of each antibiotics ranged from 11.75 when plasmid DNA was used as donor. The highest levels of resistance appeared for *E. coli* K12 transformants to streptomycin followed by those; Km, Gm and Tm respectively. Transformation of the Gram positive *S. aureus* and *B. subtilis* with plasmid DNA isolated from *Pseudomonas aeruginosa* was unsuccessful. *E. coli* K12 transformants indicated that the resistance to fun which can be acquired with a plasmid is probably 10 µg Km/ml medium. The 50 µg Km resistance can be obtained from dose of plasmid as no. 20, 30 or 40 µg Km/ml lowest single level double, so, is also could be obtained. This increase in the double dose had been attributed to gene interactions and cAMP regulatory mechanisms. Three plasmids were responsible for the level of kanamycin resistance reached 100 µg/ml, 4 plasmids for the level of 200 µg/ml. One plasmid gave a resistance level of 1 µg medium. The possibility of gene interactions can be explained. the same could be for tobramycin. Positive correlation between plasmid was estimated to give 50 µg Sm/ml levels, 2 for the level of 100 µg, 3 and 4 for the levels of 150 and 200 µg/ml resistance Gm/ml also can be high bacterial counts and high phage contents and no clear correlation was noticed between phage contents and the resistance of any of the antibiotics studied. In conclusion plasmids play an essential role in antibiotics resistance. To

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solve this problem it is essential to look for plasmid curing. Curing can occur spontaneously during growth and cell divisions or following treatments with some agents such as elevated temperatures, acridine hydrochloride, ascorbic acid and the best therapeutic treatments might be through safe vaccination.