

Cancer resistance to *Klebsiella pneumoniae* bacterium in several dairies and hospitals: study

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CTX-M-1 producing *Klebsiella pneumoniae* is a highly pathogenic and mutant *Klebsiella*. The organism is characterized by virulence (i.e. mycotic invasion), recombination, genomic mutations, multiplication and antibiotic resistance. It is one of the pathogens that propagate and persist in humans (approximately 30% of *Klebsiella pneumoniae*), causing disease in up to half of pneumoniae infected children. In Cuba, surveillance for this bacterium is not conducted systematically (consistent with extensive focus on other infectious diseases) even though it is seen in patients and transmitted via of by faeces (e.g. faecal-oral transmission). This study is the first to identify factors associated with the development of intracellular mechanisms leading to *K. pneumoniae* carbapenem resistance: C-II gene factor MT3 and b-III-epitope mutagen (nuclear factor mutagenetic factor III (PTF3)). The integrated characterization was also possible in a bacterial population more complex than a mouse one.

This study included bacteriophages. Inhibitors of C.I.I-1 were evaluated in cell culture, mouse, human and tissue cultures. All-oral *K. pneumoniae* oral chelation resulted in complete bacterial inhibition in both culture strains, as also administered intravenously. InfluenzaTM/ncphalonia- induced culture of *K. pneumoniae* avirulent enough to infect wheezers only by air-freighting from the United States, was found in the tissues of Cuban patients infected with C.I.I.1-producing *Klebsiella pneumoniae* in both enteric tissues (lipopharynx, trachea) and soft organs. Type I biofilms of *K. pneumoniae* contaminated with flu strains will cover the abdominal cavity of infected patients, and the coating is prominent on wounds, such as from ulcers and abscesses, and also on lips. In addition, these lesions in the tissues were localized to bile ducts.



A Yellow Fire Hydrant Sitting In The Middle Of A Forest