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Abstract (poster session)

First description of the metallo-beta-lactamase GIM-1 in *Acinetobacter pittii* (formerly *Acinetobacter* genomospecies 3)

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Objectives: Multidrug-resistance in species of the *Acinetobacter baumannii-calcoaceticus* complex is an increasing problem since respective strains might cause difficult to control outbreaks in hospitals. Multidrug-resistance in those species is mainly observed in *A. baumannii*, whereas *A. pittii* (formerly *Acinetobacter* genomospecies 3) tends to be more susceptible. Carbapenem resistance in *Acinetobacter* spp. is mainly caused by class D carbapenemases like OXA-23 and in some strains by metallo-beta-lactamases like IMP, SIM or NDM. The metallo-beta-lactamase GIM-1 has never been described in *Acinetobacter* spp.. **Methods:** *A. pittii* strains with reduced susceptibility to imipenem were referred to the German reference laboratory for multidrug-resistant gram-negative bacteria. Species identification was checked by MALDI-TOF analysis and *rpoB* sequencing. Carbapenemases were tested for by a combined test using EDTA, a bioassay based on cell-free extracts as well as PCRs for OXA carbapenemases and metallo-beta-lactamases. The integron structure was sequenced with specific primers. Strain typing was done by pulsed-field gel electrophoresis (PFGE) and identification of a GIM-1 harbouring plasmid was performed by S1 nuclease restriction followed by Southern Blot hybridization. **Results:** GIM-1 was found in four *A. pittii* strains all of which displayed EDTA synergy both in the combined disk test and the bioassay based on cell-free extracts. No other carbapenemase gene was found in those strains. GIM-1 was always part of a class 1 integron. Although strains showed no relatedness when tested by PFGE, all of them harboured GIM-1 encoding plasmids of the same size (approximately 60 kb). **Conclusion:** The GIM-1 carbapenemase has spread to the species *A. pittii* (formerly *Acinetobacter* genomospecies 3) and is obviously transferred by an identical plasmid. The further spread of carbapenemases in this species needs to be monitored carefully.