



THE CLAVULANIC ACID GENE CLUSTERS IN TWO ACTINOMYCETES: HETEROLOGOUS EXPRESSION OF *S. clavuligerus* ATCC 27064 CLAVULANIC ACID CLUSTER IN *S. flavogriseus* ATCC 33331

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Introduction. In a survey of genomes we found the complete cluster for clavulanic acid (CA) biosynthesis in the actinomycete *S. flavogriseus*. This CA cluster contain blocks of genes conserved in the same order as those of *S. clavuligerus* CA cluster but ensambled in a different organization. The <u>Streptomycesactivator regulatory protein</u> (SARP) CcaR, that activates clavulanic acid genes expression is present as a separate unit in *S. flavogriseus*. However, whereas *ccaR* in *S. clavuligerus* is located inside the cephamycin C cluster, in *S. flavogriseus* is in the middle of the clavulanic acid cluster

Methods. Clavulanic acid production was quantified by bioassay using *Klebsiella pneumoniae* ATCC 29665 (Romero *et al.* 1984). In addition, samples derivatized with imidazole were quantified by HPLC as described by Foulstone and Reading (1982). RNA was obtained using the RNeasy kit (Quiagen) and RT-PCR was tested as indicated by Santamarta *et al.* (2011).

Results. S. flavogriseus ATCC3331 was grown in different complex and defined media but clavulanic acid production was undetectable by bioassays or HPLC analysis. RT-PCR analysis showed that ccaR was expressed, but the expression of five genes, cas2, car, orf12, orf14 and orf16 is very low or absent in all the media tested. It was of interest to know whether the S. flavogriseus CcaR protein was functional. Therefore we introduced in S. flavogriseus the clavulanic acid gene cluster from S. clavuligerus located in cosmid SCos-CA. The new strain, S. flavogriseus SCos-CA, produced CA in TBO and MEY media but not in other media tested (SA, TSB, MG, R5. ISP4). Confirmation of the structure of the CA produced by S.

flavogriseus SCos-CA was obtained by LC-MS.

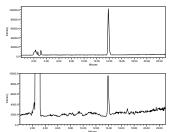


Fig.1 Detecting mass chromatograms at m/z 198. (λ = 220 nm). Up: CA standard. Down: broth of *S.flavogriseus SCos-AC* in TBO medium.

Conclusions. *S. flavogriseus* contains a silent CA cluster and does not produce clavulanic. *S. clavuligerus* CA cluster, present in cosmid SCos-CA is expressed heterologously in *S. flavogriseus* and the genetically modified strain is able to produce clavulanic acid.

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References.

1. Romero, J., Liras, P. and Martín, J.F. (1984) Dissociation of cephamycin and clavulanic acid biosynthesis in *Streptomyces clavuligerus*. *Appl Microbiol Biotechnol* (20): 318-325.

2. Foulstone, M. and Reading, C. (1982) Assay of amoxicillin and clavulanic acid, the components of Augmentin, in biological fluids with high-performance liquid chromatography. Antimicrob Agents Chemoter (22):753–762.

3. Santamarta, I., López-García, M.T., Kurt, A., Nárdiz, N., Álvarez-Álvarez, R., Pérez-Redondo, R., Martín, J.F. and Liras, P, (2011) Characterization of DNA-binding sequences for CcaR in the cephamycin–clavulanic acid supercluster of Streptomyces clavuligerus. Mol Microbiol. 81:968-981.