

**The Function of *orf13* and *orf16*, Genes Involved
in the Biosynthesis of Rifamycin B by
Amycolatopsis mediterranei S699**

Chang-Joon Kim^{1,2}, Feng Xu Ma¹, Taifo mahmud^{2,3}, Heinz G Floss²,
and Sung Bae Kim¹

¹Department of Chemical & Biological Engineering,
Gyeongsang National University, Jinju 660-701, Korea,

²Department of Chemistry, University of Washington, Box 351700,
Seattle, WA 98195-1700, USA,

³Department of Pharmaceutical Sciences, College of Pharmacy,
Oregon State University, Corvallis, OR 97331-3507, USA

Tel:+82-55-751-5391, FAX: +82-55-753-1806

Synthetically modified derivatives from rifamycin, such as rifampicin, rifabutin, and rifapentine are principal chemotherapeutic agents used for combating tuberculosis, leprosy and AIDS-related mycobacterial infections. Recently, as with many other antibiotics, the incidence of resistance of *Mycobacterium tuberculosis*, the causative agent of tuberculosis, to rifamycin is continuing increase over time, due largely to mutational alterations of the target molecule. This issued the search for improved new drug candidates. Despite the preparation of a large number of rifamycin derivatives by semi-synthetic approaches, the structural modifications have been limited primarily to one of the region of the molecule, the C3- or C4- positions of the aromatic core unit. Hence there is a need for alternative, biological method of structural modifications. To this end, we investigated the reactions involved in rifamycin B biosynthesis in *A. mediterranei* S699 by inactivation of *orf13* or *orf16*, showing strong similarity to bacterial cytochrom P450, in rifamycin biosynthetic gene cluster. The conversion of rifamycin SV to rifamycin B was almost stopped by the *orf16*-disrupted mutants. Rifamycin B was still produced by the *orf13*-disrupted mutants with lowered productivity compared with wild-type cells.

References

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