Draft Genome Sequence of Streptomyces iranensis.

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Projects

Prediction of gene regulatory networks involved in the differentiation, secondary metabolism and cross talk of *Aspergillus nidulans*Details

Abstract

Streptomyces iranensis HM 35 has been shown to exhibit 72.7% DNA-DNA similarity to the important drug rapamycin (sirolimus)-producing Streptomyces rapamycinicus NRRL5491. Here, we report the genome sequence of HM 35, which represents a partially overlapping repertoire of secondary metabolite gene clusters with S. rapamycinicus, including the gene cluster for rapamycin biosynthesis.

Identifier

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