

SL304

srf Operon and the Role of *comS* gene in Competence
Development of *Bacillus subtilis*

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srf operon of *Bacillus subtilis* encodes surfactin synthetase complex which catalyzes the nonribosomal biosynthesis of the lipopeptide surfactin using the multienzyme thio-template mechanism. The complex is composed, in part, of three subunits SrfA, SrfB, and SrfC, which constitute the seven amino-acid activating domains required for the synthesis of the peptide portion of surfactin. Immunoprecipitation experiments using a hemagglutinin(HA) epitope - tagged SrfA derivative and anti-HA antiserum resulted in co-precipitation of these three subunits, indicating that they are working in a complex form. Single amino acid substitution mutations in the active site or intermodule domain showed no effect on this subunit co-precipitation.

The *comS* gene, which is nested in the *srfB* gene and encodes a small, 46 amino acid protein, is required for competence development at early stationary phase. It has been known that ComS antagonizes ComK-inhibitor MecA, thereby activate ComK, a key protein responsible for the expression of late competence genes. Both SinR and DegU are also known to promote competence development. Through the multicopy suppression experiments with plasmid - amplified *comS*, *sinR*, and *degU* genes, it was found that multicopy *comS* suppresses *sinR* mutation but not *degU* mutation, indicating *sinR* acts at upstream of *comS* and *degU* at downstream, respectively. Experiments testing the efficiencies of *srf-lacZ* and *comS-lacZ* expressions under the $\Delta sinR$ or $\Delta degU$ background showed that only the *comS-lacZ* expression was reduced by *sinR* mutation, while neither *srf-lacZ* nor *comS-lacZ* expression was affected by *degU* mutation. These results suggest that *sinR* is required for optimal *comS* expression but not transcription from the *srf* promoter, and *degU* appears to be in a separate, parallel pathway controlling competence development.