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The Putative Sigma (YlaC) and Anti-Sigma Factors (YlaD) from Bacillus subtilis

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Two open reading frame designated as ylaC and ylaD in the Bacillus subtilis genomic sequencing project, were cloned using pRB374 vector which is shuttle vector in E. coli and B. subtilis. YlaC and YlaD have the sequence homology to SigX and SigW to YbbM, respectively, which are known to be ECF sigmaand anti-sigma factor, respectively.

The YlaD have CXXC motif and transmembrane domain in C-termanal region. CXXC motif may be concerned in active site to sigma and anti-sigma factor interaction and C-terminal region act as anchor and intermediate of signal transduction. The Overexpressed strains of *ylaC* and *ylaD* genes showed the retardation of initiation of exponential growth stage with the reduced sporulation rate and cell density at stationary stage. The promoter::*gusA* fusion protein for *ylaC* and *ylaD* genes was expressed under condition that was LB broth containing 0.1 mM MnCl₂ or 1 mM paraquat and 5 mM H₂O₂. From this result, we propose that YlaC and YlaD may expert physiology effects on the growth, especially on the sporulation of *B*. subtilis