解淀粉芽孢杆菌赖氨酸-3基因在枯草芽孢杆菌中的克隆

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限制性核酸内切酶EcoRI酶切解淀粉芽孢杆菌AS1.1099染色体DNA和质粒pUB 110 DNA, T4DNA 连接酶连接, 转化枯草芽孢杆菌BR151(trpC2 met B5 1ys3 Neo s)感受态细胞。用选择性培 养基两次筛选,得到Neo r和与 BR151 kys3营养缺陷突变互补的菌落。用快速测定质粒的方 法检查,其中有13株转化子带有大小相同的重组质 粒。分离其中1株转化子BR151-29的重组 策粒pB-L29 DNA,再次转化BR151感受态细胞和原生质体。测定第二次得<mark>▶加入引用管理器</mark> 到的转化子性状, 均与转化子BR151-29相同。琼脂糖凝胶电泳检查这些转化子的质粒,其分子大小也与重组质 粒pB-L29相同。以上试验证明,重组质粒pB-L29是由pUB110和解淀粉芽孢杆菌的1ys3基因片 段构成的。根据琼脂 糖凝胶电泳测定pB-L29的分子量约为5.0kb,推断出1ys3基因片段约为0.5kb。

关键词

分类号

Clonig of Lys3+ Gene of Bacillus amyloliquefaciens in Bacillus subtilis

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Abstract

Chromosomal DNA of Bacillus amyloliquefaciens H AS.1.1099 and plasmid pUB-110 DN A were digested with restriction endonuclease EcoRI and ligated with T4 DNA liga se. Bacillus subtilis BR151 (trpC2 metB5 lys3 Neo s) competent cells were trans-f ormed by ligated mixture. Cells were plated onto complete medium containing 5µg/ ml neomycin. A total of 3,180 Neo r colonies were obtained. These colonies were pi cked picked onto minimal medium containing 5µg/ml meomycin and lacking the tryp tophan, meth-ionine, or lysine separately. 74 resistant-neomycin and only complemen t BR151 lys3 mutation colonies were obtained Agarose gel electrophoretic analysi s of plasmid DNAs from these clonies showed that 13 of them contained inserts of identical size. One of these recombinant plasmid, pB-L29, was studied further. Upon retransformation into BR151 competent cells and protoplasts, pB-L29 was found to confer upon the host resistance to neomycin and complement lys3 mutation indica ting that Lys3+ phenotype is associated with the inserted fragment. Electrophoret ic mobilities of plasmid DNAs isolated from the secondary transformants were ide ntical to pB-L129. It is a reliable proof that recombinant plasmid pB-L29 consist s of plasmid pUB110 with an insert of B amyloliquefaciens H Lya3+ fragment at it s EcoRI site. According to this DNA fragment migration, it was calculated that the MW of pB-L29 is about 5.0 kb, and the MW of Lys3+ fragment is about 0.5 kb.

Key words

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