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大豆对SMV SC-13株系群的抗性遗传及基因定位的研究

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摘要: 在接种SC-13株系群的情况下, 鉴定了科丰1号×南农1138-2的P₁、P₂、F₁、F₂和184个重组自交系(RIL)的抗性, 结果显示, 科丰1号(P₁)与F₁全部表现抗病, 南农1138-2(P₂)全部表现感病, 表明抗性为显性; F₂群体和184个重组自交系出现抗感分离, 卡方适合性检测表明F₂群体抗感分离符合3:1的比例, 重组自交系抗感分离符合1:1分离比率。表明对SC-13株系群的抗性由一对基因控制, 以Rsc-13表示。利用已建立的遗传连锁图对Rsc-13进行了连锁分析, 结果将抗病基因Rsc-13定位于N8-D1b-W连锁群上, 并与抗性基因 Rn1、Rn3、Rsc-7连锁。

Abstract: The P₁, P₂, F₁, F₂ and 184 RILs of Kefeng No. 1 × Nannong 1138-2 were inoculated with the SMV strain group SC-13 to evaluate their resistance. The results showed that Kefeng No. 1 and F₁ were resistant to SC-13, Nannong 1138-2 was susceptible. F₂ segregated in a 3:1 ratio and the RILs in a 1:1 ratio, which indicated a dominant gene controlled the resistance to SC-13. Through linkage analysis, the resistance gene Rsc-13 was mapped on the linkage group N8-D1b-W and linked with Rn1, Rn3, Rsc-7.

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