

研究报告

小麦抗白粉病相关基因的转化

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摘要 利用玉米花青素苷合成调节基因*C1-Lc*作为报告基因, 通过瞬间表达后愈伤组织表面红色斑点的统计分析, 优化了小麦幼胚愈伤组织的基因枪转化参数。小麦*Beclin1*类似基因*TaTBL*和硫代硫酸硫转移酶基因*TaTST*是2个在白粉菌诱导条件下具有增强表达特性的抗病相关基因。本实验进一步利用基因枪将*ubi*强启动子控制下的2个基因导入到小麦品种扬麦158的幼胚愈伤组织细胞中, 使用除草剂经两轮选择培养基上的筛选和再生获得抗性植株, 进一步通过抗性植株的PCR分析获得转*TaTBL*基因植株5株, 转*TaTST*基因植株6株。转基因植株离体叶片的人工接种实验表明, 外源基因的导入不同程度上增强了植株的白粉病抗性, 表现为延缓了白粉菌的发育。利用玉米花青素苷合成调节基因*C1-Lc*作为报告基因, 通过瞬间表达后愈伤组织表面红色斑点的统计分析, 优化了小麦幼胚愈伤组织的基因枪转化参数。小麦*Beclin1*类似基因*TaTBL*和硫代硫酸硫转移酶基因*TaTST*是两个在白粉菌诱导条件下具有增强表达特性的抗病相关基因。本实验进一步利用基因枪将*ubi*强启动子控制下的两个基因导入到小麦品种扬麦158的幼胚愈伤组织细胞中, 使用除草剂经两轮选择培养基上的筛选和再生获得抗性植株, 进一步通过抗性植株的PCR分析获得转*TaTBL*基因植株5株, 转*TaTST*基因植株6株。转基因植株离体叶片的人工接种实验表明, 外源基因的导入不同程度上增强了植株的白粉病抗性, 表现为延缓了白粉菌的发育。

关键词 [小麦](#) [白粉病](#) [抗病相关基因](#) [基因转化](#)

分类号

Transformation of powdery mildew resistance-related genes of wheat

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Abstract

<P>Anthocyanin synthesis regulation gene C1-Lc was used as the reporter gene to optimize the parameters of gene-gun transformation protocol through counting of red spots on wheat calli after transient expression. Wheat *Beclin1* like gene TaTBL and thiosulfate sulfotransferase gene TaTST proved to have an increased expression level after induction of wheat powdery mildew fungus (Erysiphe graminis f.sp. tritici Em. Marchal.). These two resistance-related genes were constructed into expression vectors driven by the strong ubi promoter and used to perform genetic transformation on wheat cv Yangmai158 immature embryo-derived calli through particle bombardment. After two rounds of herbicide bialaphos selection and regeneration, herbicide-resistance plants were obtained, which were subsequently subjected to PCR analysis. Five TaTBL transgenic plants and six TaTST transgenic plants were identified. Pathogen inoculation of detached leaves showed that the introduction of exogenous gene increased wheat resistance level by delaying the development of powdery mildew symptoms.</P>

Key words [wheat](#) [powdery mildew](#) [resistance-related gene](#) [genetic transformation](#)

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