

具感染力之蕃茄黃化捲葉病毒之構築與含其複製? 鈹穢]菸草之抗病分析

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摘要

蕃茄黃化捲葉病毒(Tomato yellow leaf curl virus; TYLCV) 屬於雙生科(Geminiviridae)中的Begomovirus屬, 基因體包含單一種環狀DNA核酸(monopartite genomes), 由銀葉粉蝨(*Besisia argentifoli*)傳播, 對蕃茄收成造成嚴重之危害。本實驗利用病原誘導抗病機制(parasite-derived resistance, PDR)以台灣品系之TYLCV的部分複製蛋白?基因(C1), 依不同排列方式, 分別構築正股(sense)、反意股(antisense)及兩段反意股相連(double antisenses)之轉基因質體, 再以農桿菌方式轉殖到植株上, 並挑戰接種含全長度TYLCV質體之農桿菌。本研究先以煙草為模式, 進行抗病分析, 以作為將來轉殖到蕃茄的參考。實驗首先在彰化縣, 包括溪湖鎮、永靖鄉、大村鄉及員林鎮八個蕃茄園區, 採集疑似受病毒感染之蕃茄葉片, 再以聚合?連鎖反應(polymerase chain reaction; PCR)針對病毒之複製?基因(C1), 進行蕃茄黃化捲葉病毒之偵測, 在溪湖鎮40個樣本中, 共偵測到5個染病材料, 其罹病率為12.5%; 在永靖鄉、大村鄉及員林鎮各採集8個樣本, 亦各偵測到1個染病植株, 罹病率為12.5%。將針對TYLCV的PCR產物進行選殖與解序後, 發現無論是C1基因或是全長度基因體, 溪湖品系的TYLCV與泰國品系TYLCV最為相近, 可達98%。所轉殖含C1基因的擬轉基因煙草, 則先利用針對病毒C1基因及NPTII專一性引子進行PCR增幅, 可獲得預期DNA片段, 證明轉基因確實有送入煙草中, 總共每種構築分別得10-15個擬轉基因煙草品系。由於TYLCV無法以機械接種方式感染植物, 因此為了進行轉基因煙草的抗病分析, 本實驗另外再構築具感染力的TYLCV質體。首先以溪湖採集之TYLCV為材料, 以PCR分別放大三段DNA片段, 涵蓋全長度病毒基因體, 以適當酵素進行剪接後, 得到一個含病毒全長度基因體1.27倍TYLCV質體(TI0380AE5), 其在5'端多出346核?酸, 在3'端多出416核?酸, 此質體藉由農桿菌, 接種到非轉基因煙草及蕃茄後, 取接種植物之上位葉進行PCR及解序, 確定此病毒質體(TI0380AE5)確實具有感染力, 比正常感染TYLCV的煙草病徵較弱。此具感染力的TYLCV質體即可用來挑戰接種於轉基因煙草。由本實驗在煙草上的研究經驗, 將可應用到未來蕃茄的轉殖上, 以達抗蕃茄黃化捲葉病毒之目的。

關鍵詞: 蕃茄黃化捲葉病毒; 基因轉殖; 農桿菌; 聚合?連鎖反應

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