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**摘要:** 建立转乙肝表面抗原(*HbsAg*)基因樱桃番茄组培苗叶片的蛋白质组双向电泳体系,为转基因番茄的蛋白质组学研究提供技术参数。比较转*HbsAg*基因的樱桃番茄与野生型在不同蛋白裂解液、不同蛋白上样量及不同等电聚焦程序下,叶片总蛋白的双向电泳结果差异。以裂解液III制备叶片总蛋白,采用一向13cm、pH3-10L IPG胶条,二向12.5%的SDS-PAGE凝胶,单次上样120μg叶片总蛋白,聚丙烯酰胺凝胶电泳时,可以得到最理想的双向电泳分析结果。在此操作程序下,野生型樱桃番茄叶片蛋白双向电泳最多可识别699个蛋白点,而转基因型樱桃番茄叶片蛋白双向电泳最多可识别545个蛋白点,其中有368个点匹配,选取丰度百分比差异2倍以上蛋白点25个进行质谱鉴定,16个蛋白点成功鉴定。本研究建立的转*HbsAg*基因番茄植株叶片总蛋白的双向电泳体系可为以番茄试管苗为材料进行的蛋白质组学研究提供技术支持。

**关键词:** 双向电泳 转基因植物 口服植物疫苗 蛋白组 组培苗

### ESTABLISHMENT OF TWO-DIMENSIONAL GEL ELECTROPHORESIS SYSTEM FOR LEAF PROTEINS OF *HBsAg* TRANSGENIC CHERRY TOMATO

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**Abstract:** Two-dimensional gel electrophoresis (2-DE) system was established for detection of the leaf proteins of *HBsAg* transgenic cherry tomato, which provided the experiment data for proteomics study of transgenic cherry tomato. This study compared the effects of different protein lysis buffers, protein loading amounts and isoelectric focusing conditions on the 2-DE maps of the transgenic plants. An optimum system was obtained as follows: dissolving total proteins of samples with the lysis buffer III, separating the proteins with 13cm pH3-10L IPG strips, loading protein samples of 120μg followed by isoelectric focusing program III, and staining the gels with silver staining after 12.5% SDS-PAGE. Under the above conditions, 545 protein spots were identified from the transgenic plants and 699 protein spots from control plants, of which 368 protein spots were matched. A total of 25 protein spots were analyzed by MALDI-TOF MS and 16 proteins were identified by PMF. The 2-DE system for leaf proteins of *HBsAg* transgenic cherry tomato established in this work provides technical base for the studies of tomato proteomics.

**Keywords:** two-dimensional electrophoresis transgenic plants plant-derived oral vaccine proteomics tissue culture plantlet

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