

04-1-39

Study for Searching the *Anthracnose* Resistance Related Genes from Pepper

Ukjo Kim^{1*}, Soon Ho Choi¹, Nam Han Her¹, Jang Ha Lee¹, Koon Bo Kim², Shin Jae Kim², Jae Bok Yoon³,
Hyo Geun Park³, Seung Geun Yang¹, Chee Hark Harn¹

¹Nong Woo Bio Co., Yeosu, Gyeonggi, Korea; ²F&P Co., Suwon, Gyeonggi, Korea; ³ Dept. of Horticulture,
Seoul National Univ., Seoul, Korea

Objectives

To characterize the *Anthracnose* resistance related genes from Pepper

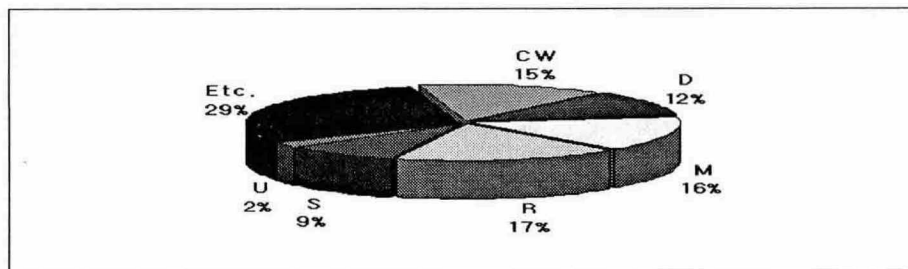
Materials and Methods

1. Materials : *C. annuum* X *C. baccutum*
2. Methods : High-Density cDNA Macroarray, Sequencing

Results and Discussion

A cDNA library was constructed from the transcripts isolated from *Anthracnose* resistant line and susceptible line. A total of 266 EST clones were identified by differential screening of High-Density cDNA Macroarray. From the sequencing data, several defense related genes and transcription factors were found.

Classification of EST Clones



□ R: resistance □ M: metabolism ■ S: signal transduction
 ■ D: development ■ U: unknown ■ CW: cell wall
 ■ Etc: proteins