A Comparison of Recombination Rates Between Adapted Soybean Cultivars, and Random Lines. (C01-stefaniak081612-Poster)

Authors:

• T.R.Stefaniak* - *University of Kentucky*

Abstract:

It has been estimated that in the period between 1930 and 1980 soybean yields in America increased 93% due to the introduction of superior cultivars. However, it is unclear what source of genetic variation has been utilized for this improvement. The objective of this study was to investigate the possible contribution of genetic variation from meiotic recombination in the improvement of soybean cultivars. Crossing-over events were detected using 88 SSR markers spanning 13 of soybeans linkage groups. The two populations consisted of 10 adapted high yielding cultivars and 156 Random Recombinant Inbred Lines from the cross Williams x Essex. The calculated recombination rates were standardized for potential crossovers. For total recombination the rates for the 10 cultivars were significantly greater than those of the RRILs. These rates were 0.24 and 0.36 for the RRILs and the cultivars respectively. The population by linkage group interaction was nonsignificant. These results indicate that breeding progress has been accomplished in part utilizing the genetic variation resulting from recombination.

Corresponding Author Information:

Thomas Stefaniak phone: (859)257-1825 University of Kentucky fax: (859)257-2185

S106 Agriculture Science Center N. e-mail: trstef1@pop.uky.edu

Lexington, KY 40546

Presentation Information:

Presentation Date: Tuesday, November 12, 2002

Presentation Time: 4:00-6:00 pm

Poster Board Number: 1014

Keywords:

Recombination, Linkage, Molecular Markers, Simple Sequence Repeats