# Genetic Variation in Ethanol Yield of Lowland Switchgrass Plant Jessica N. Hentchel and Hem S. Bhandari University of Tennessee, Department of Plant Sciences Image: Colspan="2">Image: Colspan="2" Image: Colspan="2"

### INTRODUCTION

Switchgrass (*Panicum virgatum* L.) has shown high potential for biofuel production. Switchgrass needs improvement in terms of biofuel yield per unit of area for its sustainable use as a bioenergy feedstock. These objectives can be achieved through improvement in biomass yield and feedstock composition



RESULTS

quality. Genetic improvement in biomass yields are in progress. Breeding with

priority on feedstock composition quality is now receiving more focus from

researchers.

# MATERIALS AND METHODS

Initial Genetic Materials: Four-year-old Alamo sward population

Phenotypic selection based on visual vigor

200 plants selected (fall 2011), open pollinated seed harvested from each plant constituted a half-sib (HS) family

Selection based on seed

**Figure 1**. Ethanol yield variation among parent genotypes of half-sib families. Results were significant at p<0.01. As expected Ethanol yield was positively correlated with IVDMD (r = 0.82), and negatively correlated with NDF (r = -0.8).





62 HS evaluated in a replicated trial at Crossville. Each family row in each rep had 9 plants spaced 30cm;

Parents of HS were evaluated at Knoxville in a space plant nursery, with 90cm x 90cm plant spacing **Figure 2**. Ethanol yield variation among Alamo half-sib families. Results were significant only at p<0.15, and the range is narrow as compared to parental genotypes.

# FUTURE RESEARCH

There was no correspondence between ethanol yields of parent genotypes

and their half-sib families. This could be likely due to (i) high genotype x

environment interaction, or (ii) due to different plant spacing, parent

genotypes spaced 90cm x 90cm and HS family were spaced 30cm x 90cm



### (i.e., between plants x within row and between rows). It is also possible

due to poor reliability of feedstock composition data from the

establishment year stands. More detail analysis will be conducted using

biomass samples from a mature stand after at the end of fall 2013.



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