

## **Loblolly Pine Karyotype Using FISH and DAPI Positive Banding**

M. N. Islam-Faridi<sup>1,2</sup>, C. D. Nelson<sup>2</sup>, T. L. Kubisiak<sup>2</sup>, M. V. Gullirno<sup>1</sup>,  
V. H. McNamara<sup>1</sup>, H. J. Price<sup>1</sup> and D. M. Stelly<sup>1</sup>

### **Poster Abstract**

A loblolly pine (*Pinus taeda* L.) karyotype has been developed based on fluorescent *insitu* hybridization (FISH) using cyto-molecular landmarks including plant telomere repeat, 18S-28S rDNA and 5S rDNA probes and DAPI positive bands. Somatic chromosome spreads of loblolly pine root tips were prepared using a modified enzymatic digestion technique. We observed ten pairs of long metacentric, one pair of long sub-metacentric and one pair of short sub-metacentric chromosomes. All the chromosomes showed characteristic DAPI positive bands (A-T rich regions) near and/or around the centromeres. At least one DAPI positive band was also observed in intercalary positions on all chromosome arms. Plant telomere FISH signals were observed towards the end of each chromosomal arm as expected. In addition, most of the chromosomes showed telomeric sites near and/or around the centromeres except for one or possibly two chromosomes. A total of seventeen 18S-28S rDNA sites were identified per haploid genome. Eight of these were located near and/or around the centromeres and seven were at intercalary positions. One major 5S rDNA site was observed in an intercalary region of a metacentric chromosome that lacked 18S-28S rDNA sites. One or possibly two minor 5S rDNA sites were observed near the ends of two different chromosomes. We are also developing a slash pine karyotype for direct comparison with loblolly as well as a comparison with a previously published slash karyotype (Doudrick et al. 1995, Journal of Heredity 86:289-296). Finally, we will provide an update on our progress toward using BAC clones as FISH probes on pine chromosomes.

---

<sup>1</sup> Department of Soil and Crop Sciences, Texas A&M University, College Station, Texas 77843;

<sup>2</sup> USDA Forest Service, Southern Institute of Forest Genetics, Saucier, Mississippi 39574.

**Note: Also presented as a contributed paper.**