AN AUTOMATED APPROACH TO GENETIC MAPPING WITH RANDOM AMPLIFIED POLYMORPHIC DNA MARKERS

L.S. Nelson, G.N. Johnson, M.L. Crawford, W.L. Nance, C.D. Nelson, R.L. Doudrick¹

<u>Abstract.--</u> At least 10000 Random Amplified Polymorphic DNA (RAPD) reactions are required to complete a modest RAPD-based genetic mapping effort. Most molecular genetics laboratories are not equipped to handle this level of production in a time-efficient manner; and efforts to scale up are costly when considering equipment acquisition and personnel needs. We describe a RAPD laboratory of our own design that is capable of completing 2304 RAPD reactions per 2.0 technician-work-days and minimal space requirements (approximately 500 square feet). The laboratory features a robotic pipettor, custom designed electrophoresis rigs, and a customized database management system. Exhibits of several recently completed RAPD-based maps are included, along with estimates of the times required to complete each map. Future plans for enhancing the throughput of the laboratory are also briefly described.

<u>Ke^{Σ}words:</u> linkage mapping, genetic markers, Random Amplified Polymorphic DNA (RAPD), Polymerase Chain Reaction (PCR), automation.

¹ Biological Laboratory Technician, Biological Laboratory Technician, Biological Laboratory Technician, Project Leader, Research Geneticist, and Research Pathologist, respectively, USDA Forest Service, Southern Forest Experiment Station, Gulfport, MS 39505.