of the head and determine the presence of the Chiari I malformation and syringomyelia. A small blood sample (about 2 tablespoons) will be drawn for DNA studies relating to the Chiari I malformation. CType I Arnold Chiari Malformation C|Official Title:||Genetic Analysis of the Chiari I Malformation | C|Study Start Date: ||February 2000 | CObjectives: The goal of this study is to establish family pedigrees and undertake genetic linkage analysis that will identify gene loci associated with the Chiari I malformation and underdevelopment of the bone forming the posterior cranial fossa. CStudy Population: Patients and family members of patients with the Chiari I malformation. Because the research institutions are located in the United States and Russia, subjects will be recruited predominantly from these countries. CDesign: Human subjects will undergo 1) neurologic examinations. 2) head and cervical MRI scans to evaluate for the Chiari I malformation, syringomyelia, and maldevelopment of the posterior fossa, and 3) isolation and analysis of genomic DNA from whole blood for linkage analysis. COutcome

Measures: Pedigrees will be established based on

(MRI) of the brain and cervical (neck) spinal cord to measure the size