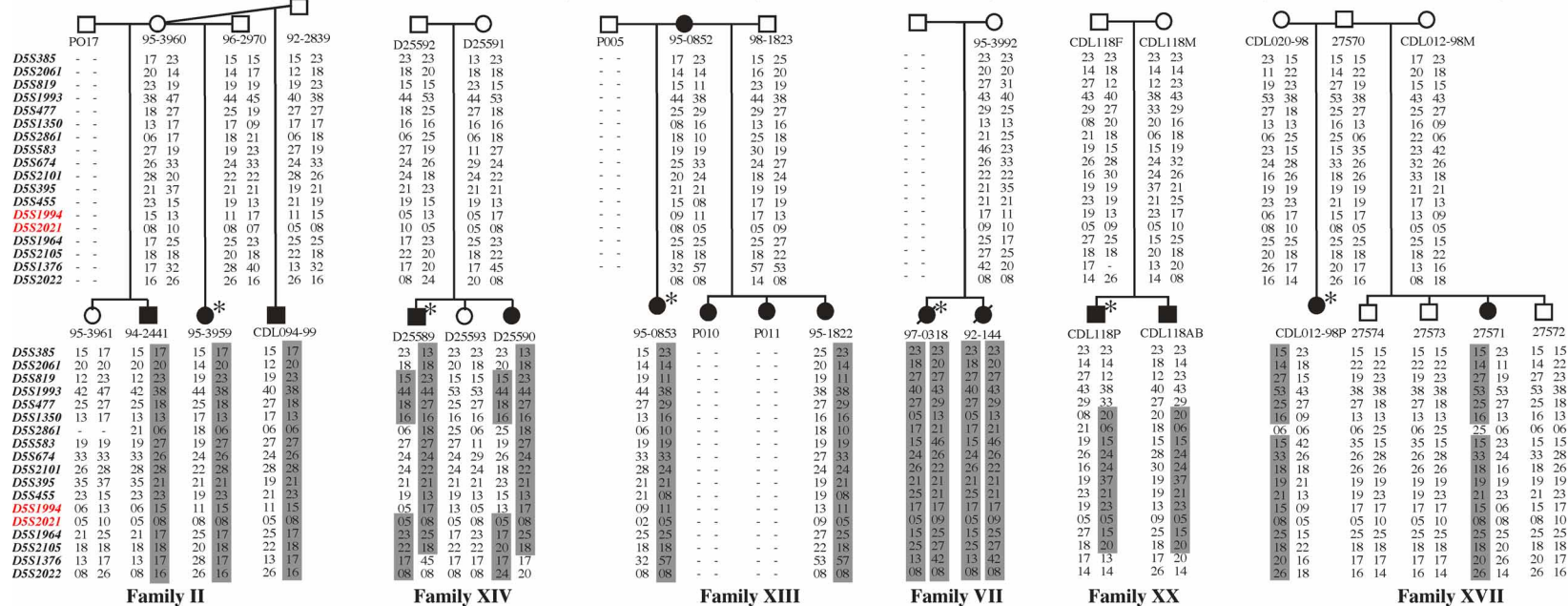
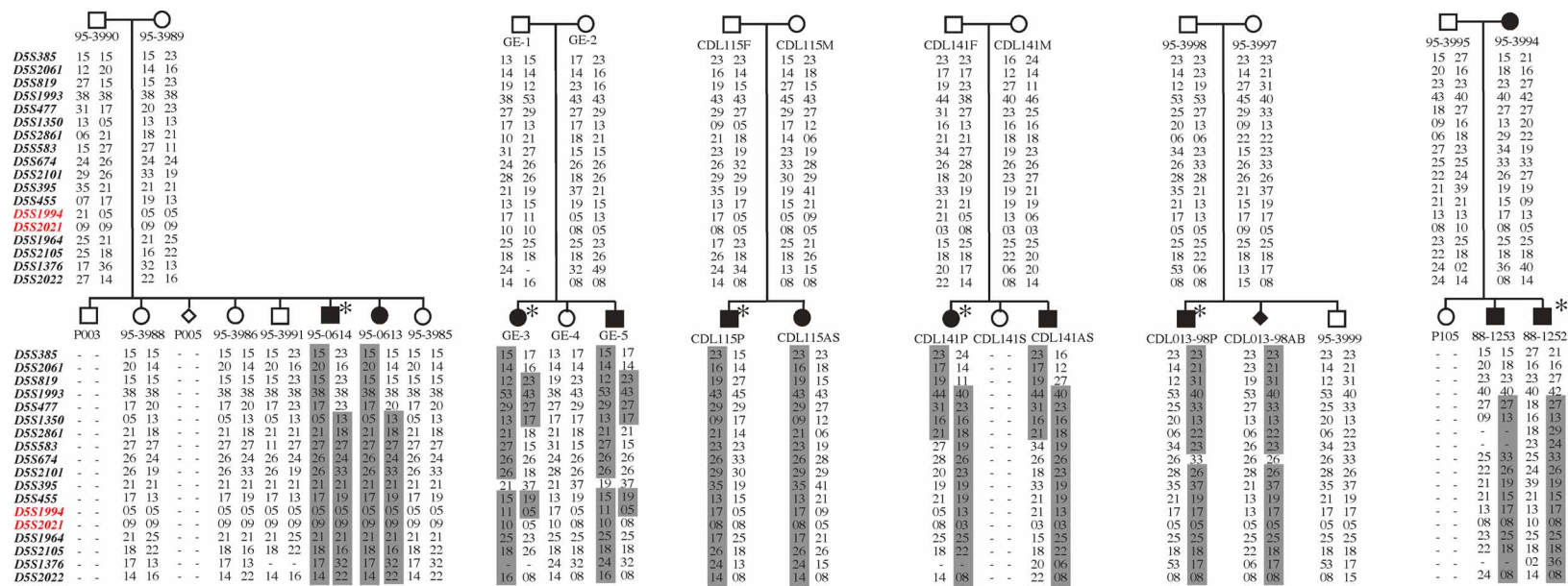




Supplementary Figure 1. High-resolution linkage analysis of 5p13 in the 12 CdLS families.

All microsatellite markers used are indicated on the left (distances between markers are indicated in Figure 3). Haplotypes for each individual are indicated below the individual identification numbers. Paternal alleles are on the left and maternal alleles are on the right. Alleles shared by affected individuals are indicated by gray shading. Family XX allowed for refinement of the distal boundary to marker D5S477 and of the proximal boundary to marker D5S1376. Marker D5S426 which gave the initial positive LOD score in the genome-wide scan lies between markers D5S674 and D5S2101. The gene lies between markers D5S1994 and D5S 2021 (indicated in red).