

SUPPLEMENTAL FIGURES

Figure S1. Chromosome 3 Sagittal Synostosis TDT Results using all Trios. *P*-values ($-\log_{10}$) are plotted against the genomic position of each SNV on the chromosome 3 associated, with genes in the region depicted below. The LD (r^2) between the top associated SNV rs3773855 ($P=1.42 \times 10^{-5}$) and other SNVs are indicated.

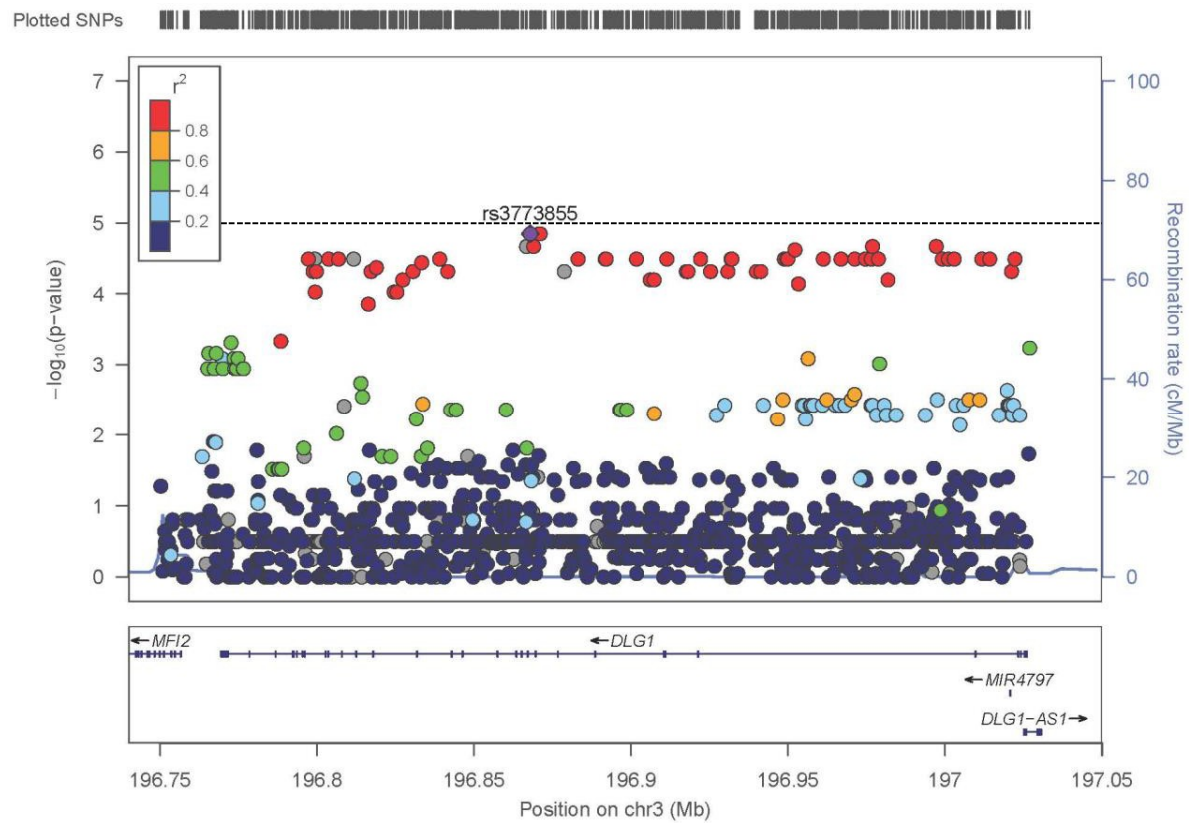


Figure S2. Chromosome 7 Sagittal Synostosis TDT Results using all Trios. P -values ($-\log_{10}$) are plotted against the genomic position of each SNV on the chromosome 7 associated, with genes in the region depicted below. The LD (r^2) between the top associated SNV rs1983511 ($P=1.54 \times 10^{-5}$) and other SNVs are indicated. The dotted line represents the genome-wide suggestive threshold of 1×10^{-5} .

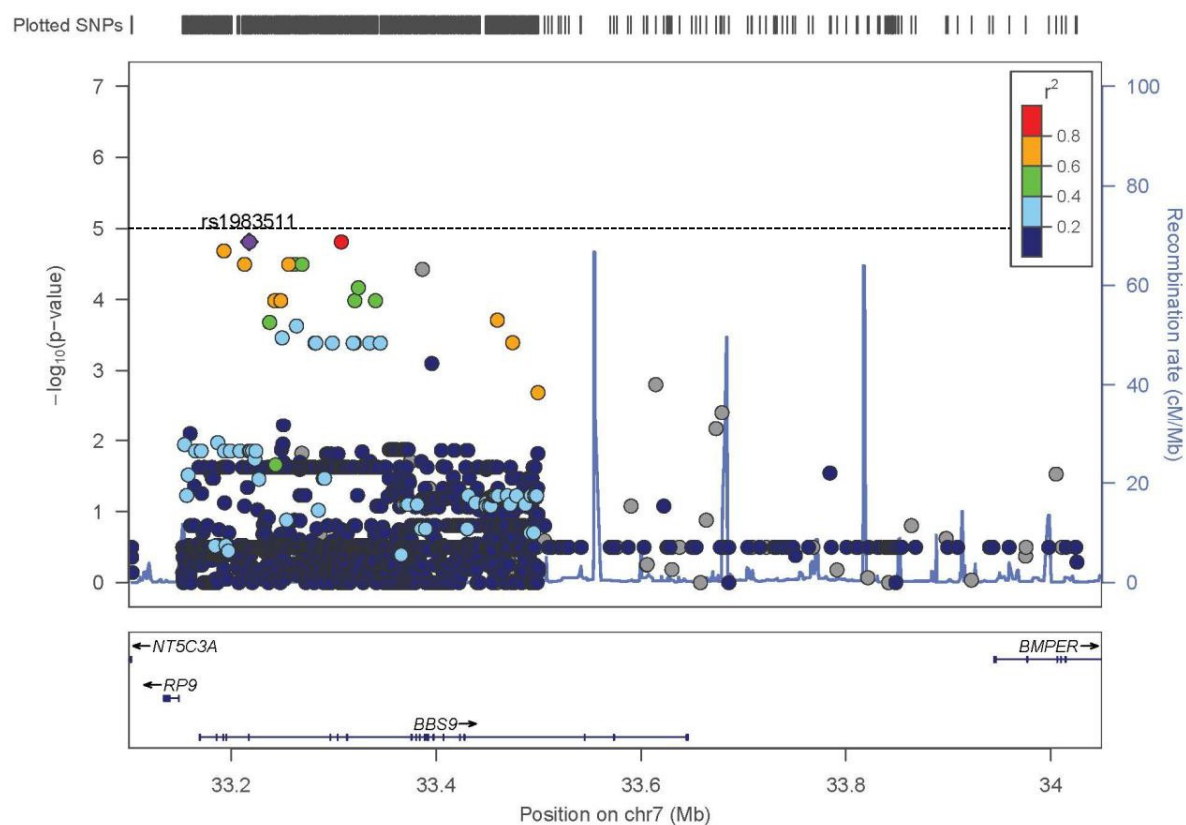


Figure S3. Chromosome 7 Metopic Synostosis TDT Results using all Trios. P -values ($-\log_{10}$) are plotted against the genomic position of each SNV on the chromosome 7 associated, with genes in the region depicted below. The LD (r^2) between the top associated SNV rs12538649 ($P=0.00029$) and other SNVs are indicated. The dotted line represents the genome-wide suggestive threshold of 1×10^{-5} .

