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Supplemental Information

Whole-Genome Analysis Reveals that Mutations in Inositol Polyphosphate Phosphatase-like 1 Cause Opsismodysplasia

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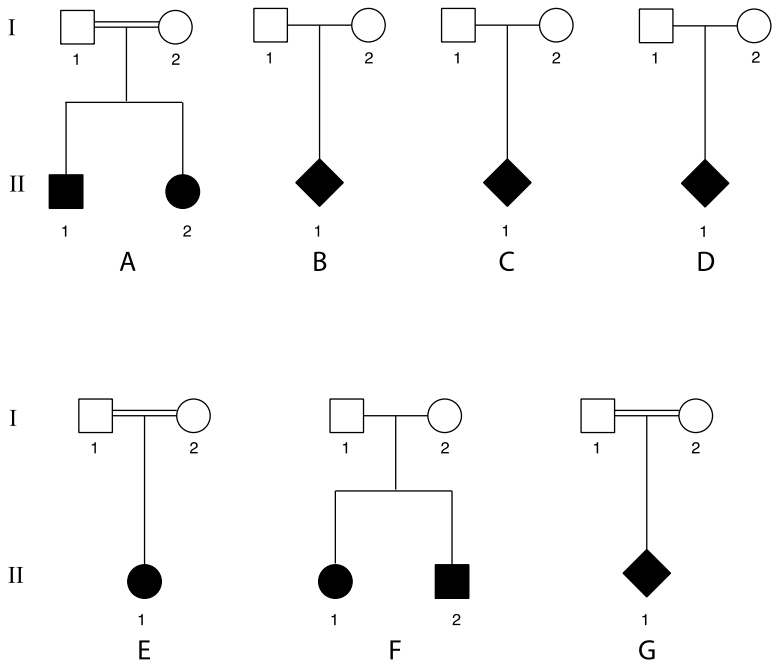

Figure S1. Pedigrees of the seven families in which mutations in *INPPL1* that cause opsismodysplasia were identified.

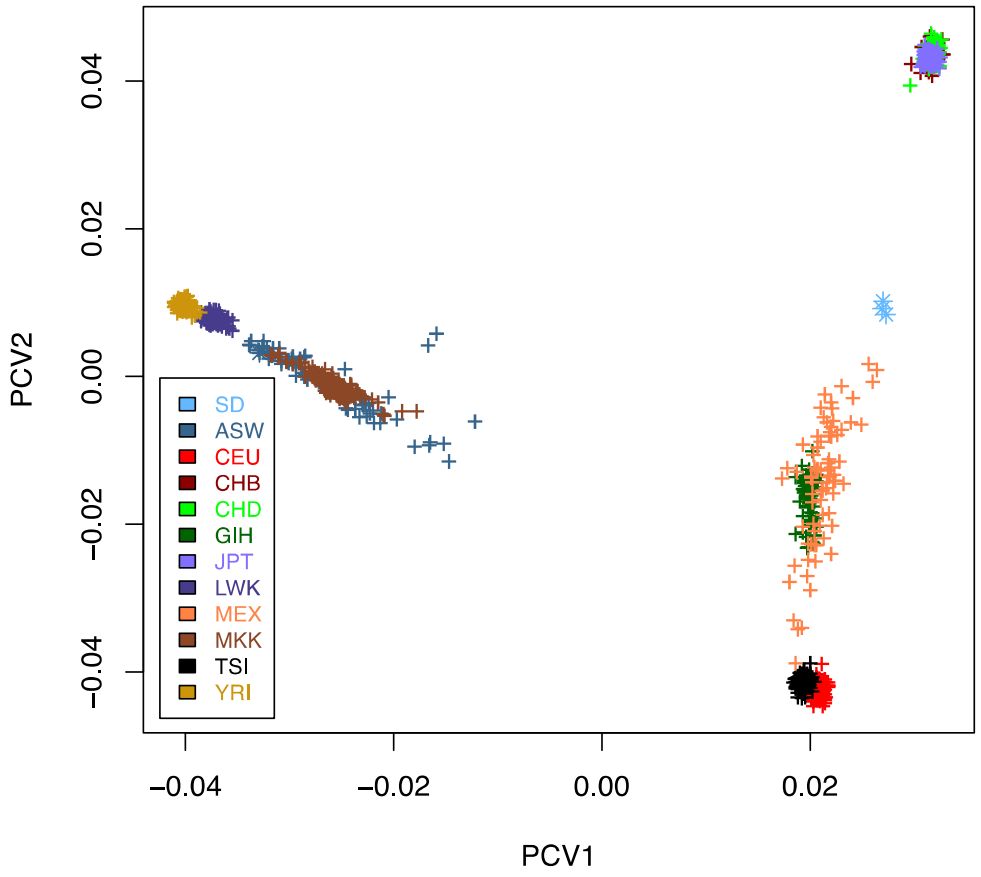

Figure S2. Principal component plot of Family A (light blue, SD – skeletal dysplasia) using genotypes from HumanCytoSNP-12 BeadChips showing clustering is consistent with self-identified ancestry.

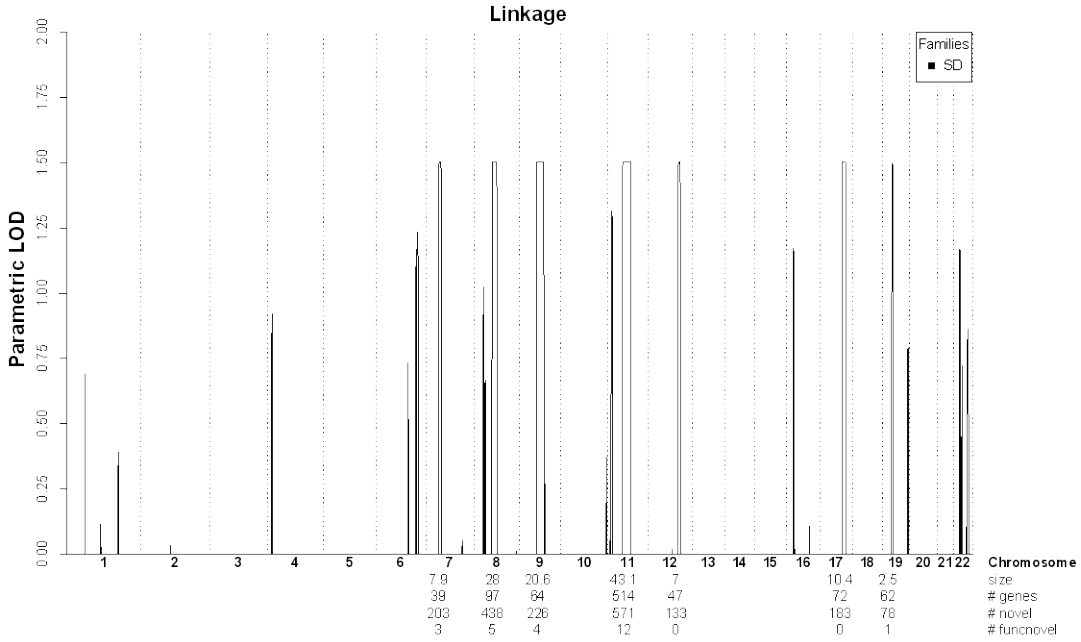

Figure S3. Results of parametric linkage analysis using a fully penetrant rare recessive model (f_2 = 1; q = 0.0001) and allele frequencies estimated from unrelated members of the HapMap CEPH European, Chinese, Japanese, and Mexican American populations performed using ALLEGRO) on an approximately 0.2 cM SNP map. SD – skeletal dysplasia.