

APPENDIX FIGURE 1. Pair-wise marker LD between SNPs around the *HNF4A* region.

The lower left and upper right triangles describe pair-wise LD in cases and controls,

respectively. A: $|D'|$. B: r^2 .

APPENDIX FIGURE 1 METHODS

Pair-wise marker LD measures ($|D'|$ and r^2) were estimated using the EM algorithm (1). A pictogram describing pair-wise marker LD was created using a program written in SAS (Cary, NC).

REFERENCE

1. Excoffier L, Slatkin M (1995) Maximum-likelihood estimation of molecular haplotype frequencies in a diploid population. *Mol Biol Evol* 12:921-927