

**Figure S3: A** - Linkage Disequilibrium (LD) plot of HapMap phase II for YRI, CEU and CHB+JPT data centered on *SERPINB11* region of chromosome 18. The image was built using *Haploview* 4.1 software. The triangular units designate LD blocks. The degree of LD between pairs of markers is indicated by the |D'| statistic (|D'| =1 bright red; |D'| >1 shades of red) LD blocks overlapping between populations are indicted. **B** – Haplotype structure of LD blocks reconstructed by median-joining networks (<a href="http://www.fluxus-engineering.com/sharenet.htm">http://www.fluxus-engineering.com/sharenet.htm</a>).