

Supplementary Figure S3. SNP associations in the *FOXP2* region. Genome-wide association results in the *FOXP2* region, for BMI[11], waist-hip ratio (adjusted for BMI)[12] and insulin resistance[23], were plotted using LocusZoom (<http://csg.sph.umich.edu/locuszoom>). Linkage disequilibrium with relation to the top SNP (purple diamond) is according to 1000 Genomes (CEU, June 2010 release). The shading in the bottom panel corresponds to the region shown in Figure 3.

