ESM Figure 7: Q-Q plots of the observed versus expected quantiles of \(-\log_{10}(pvalues)\) from meta-analysis of 715,467 SNPs in DCCT/EDIC and LifeLines analysis of SF (M3). A, Plotting all 715,467 SNPs; B, Plotting results after excluding 1MB region (874 SNPs) around NAT2. Genomic control lambda for both A and B were 1.02 [1].