

Figure S1. Consort diagram outlining the summary of subject exclusion and inclusion. A total of 468 subjects were analyzed in the present study.


Figure S2. Percent difference (with $95 \% \mathrm{CI}$ ) from GEE models showing the associations between Adipo-IR and longitudinal change in glucose measures. Model 1 adjusted for time, baseline age, sex, family history of diabetes, ethnicity, physical activity, smoking status, and BMI. Model 2: Adjusted for Model 1 + TG: HDL.


Figure S3. GEE model showing the OR per SD increase (with $95 \% \mathrm{CI}$ ) of baseline Adipo-IR and incident dysglycemia over 9-year follow-up in PROMISE. Model 1: adjusted for time, baseline age, sex, family history of diabetes, ethnicity, physical activity, smoking status, and BMI. Model 2: Adjusted for Model 1 + TG: HDL.

## A) Adipose Tissue Function


B) Circulating Lipids

C) Systemic Inflammation


Figure S4. Percent difference (with $95 \% \mathrm{CI}$ ) from GEE models showing the associations between Adipo-IR and longitudinal change in biomarkers of adipose tissue function ((A) adiponectin and sCD163), circulating lipids ((B) total cholesterol, HDL, LDL, TG), and systemic inflammation ((C) TNF- $\alpha$, and II-6). Model 1: adjusted for time, baseline age, sex, family history of diabetes, ethnicity, physical activity, smoking status, and BMI. Model 2: Adjusted for Model 1 + TG: HDL.


Figure S5. GEE model showing the OR per SD increase (with $95 \% \mathrm{CI}$ ) of baseline measures of fasting insulin, HOMA2-IR, 1/ISI, and Adipo-IR with incident dysglycemia over 9 -year follow-up in PROMISE. Adjusted for time, baseline age, sex, family history of diabetes, ethnicity, physical activity, smoking status, and waist circumference.

