

SUPPLEMENTARY FIGURE LEGENDS

Supplementary Figure 1: Principal Components of whole genome data from the POSH study. The plot shows first two principal component based on post quality control genome wide data.

Supplementary Figure 2: Regional plot of p-values arising from cox-proportional hazard models, 250 kb either side of the rs3884558 variant in stage-1. P-values are from imputed data alone.

Supplementary Figure 3: Regional plot of p-values arising from cox-proportional hazard models, 250 kb either side of the rs3785982 variant in stage-1. P-values are from imputed data alone.

Supplementary Figure 4: Regional plot of p-values arising from cox-proportional hazard models, 250 kb either side of the rs2774307 variant in stage-1. P-values are from imputed data alone.

Supplementary Figure 5: Regional plot of p-values arising from cox-proportional hazard models, 250 kb either side of the rs1387389 variant in stage-1. P-values are from imputed data alone.