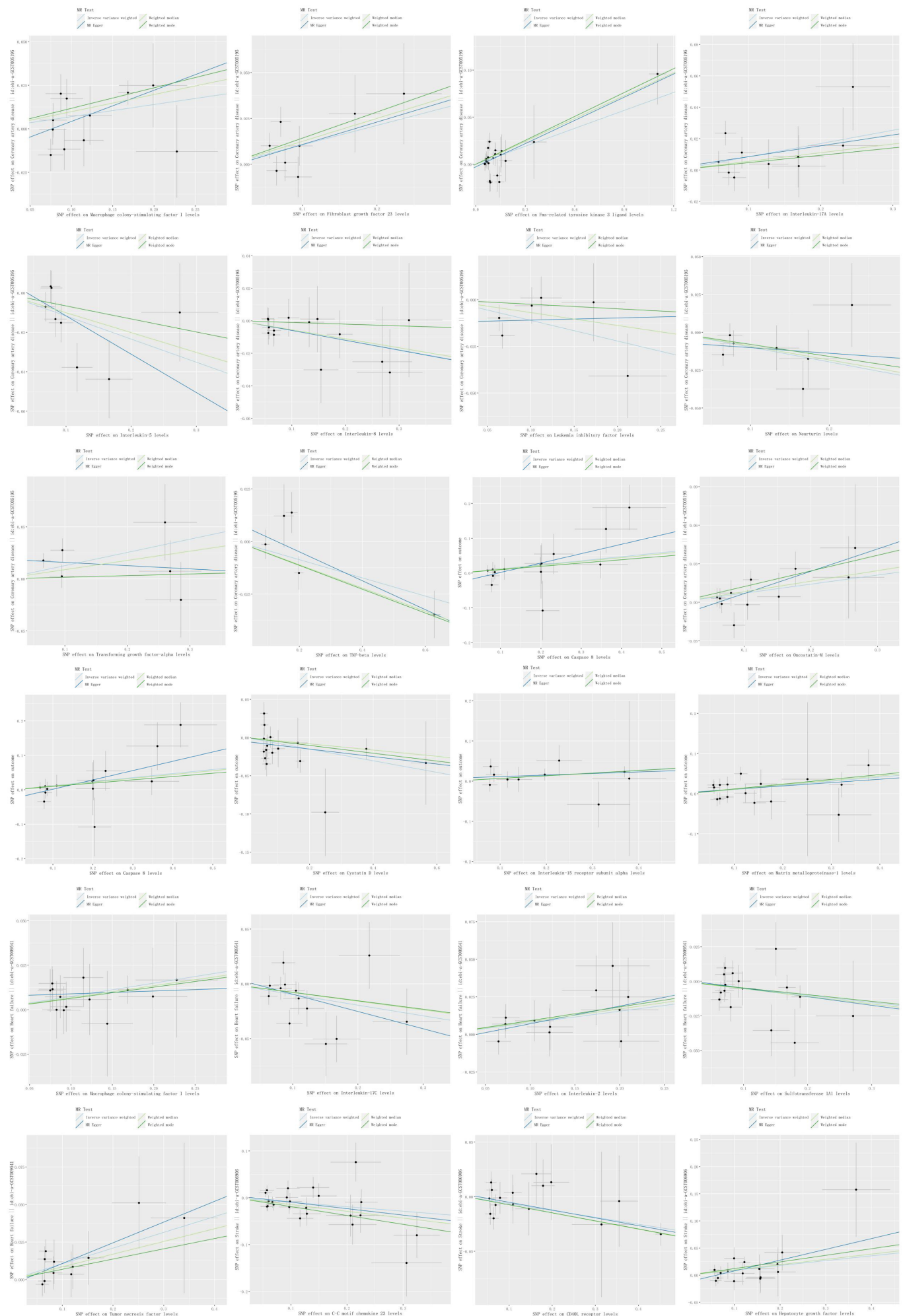
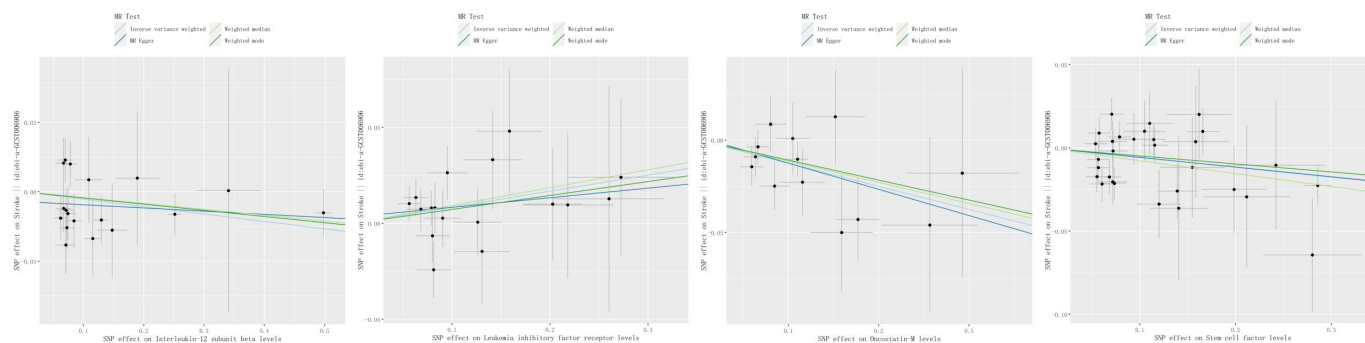






FigureS1. Scatterplot for the significant Mendelian randomization (MR) association (P value < 0.05) between circulating inflammatory proteins and 4 types of cardiovascular diseases phenotypes (Coronary artery disease, Peripheral artery disease, Heart failure and Stroke).





Figure S2. Forest plots for the Mendelian randomization (MR) leave-one-out analysis of the significant inverse variance weighted (IVW) estimates.