

Figure 1 displays six vertically stacked Manhattan plots showing GWAS results for various traits. The x-axis for all plots represents the genomic position (chromosome 10), ranging from 55,500 to 56,500. The y-axis represents the association statistic, ranging from -5 to 5. Green vertical lines indicate significant SNPs, and red circles highlight specific SNPs.

- blood_WHITE_COUNT**: Shows a significant association peak around 55,500.
- blood_MONOCYTE_COUNT**: Shows a significant association peak around 55,500.
- body_BMIz**: Shows a significant association peak around 55,500.
- impedance_BASAL_METABOLIC_RATEz**: Shows a significant association peak around 55,500.
- blood_HIGH_LIGHT_SCATTER_RETICULOCYTE_COUNT**: Shows a significant association peak around 55,500.
- blood_EOSINOPHIL_COUNT**: Shows a significant association peak around 55,500.

[illegible]