

Figure 1 displays six scatter plots showing the relationship between various blood count variables and a genomic region (approximately 80,000 to 81,000). The variables are:

- blood_MONOCYTE_COUNT
- blood_WHITE_COUNT
- blood_HIGH_LIGHT_SCATTER_RETICULOCYTE_COUNT
- blood_EOSINOPHIL_COUNT
- blood_LYMPHOCYTE_COUNT
- blood_RED_COUNT

Each plot shows data points (black dots) and a fitted curve (yellow line). Red circles highlight specific data points, and vertical green lines indicate genomic features. The y-axis scales vary for each variable.

[illegible]