

Supplemental Figure 1. Association between myeloid neoplasm and plasma protein levels stratified by disease subtype. Multivariable cox regression adjusted for age, sex, smoking status, year of sample collection, blood count measurements, and the top 10 principal components of genetic ancestry was used to test for an association between plasma protein expression and myeloid neoplasm (MN) subtype (all MN, myeloproliferative neoplasm (MPN) only, myelodysplastic syndrome (MDS)/acute myeloid leukemia (AML) only. All proteins significantly associated with at least one MN subtype are shown in the circular heatmap. Each box represents a protein-MN association and is colored according to the beta-coefficient (reflects protein expression levels); statistically significant associations (FDR-corrected p <0.05) are labeled with an asterisk (*).