



**Supplemental Figure 7. Prediction of MN risk using clinical, CH, and proteomics data.** (A) Bar chart of the most frequent features that predict MN risk in LASSO regression analysis ordered by the number of times they were included in the model during 100-fold cross validation. Features used in the analysis include demographics, blood counts, CH, and proteomics. (B) Forest plot illustrating the beta coefficients and their confidence intervals (95%) for the top 25 features from (A).