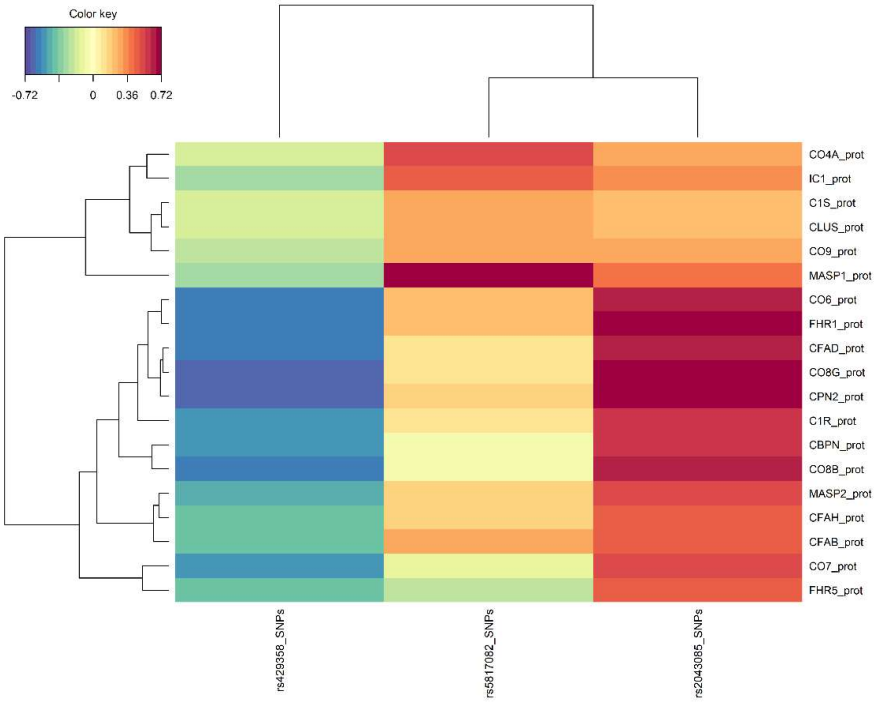Figure S12. correlations between genetic variants (n=3) and complement proteins (n=19) selected by the multi-block sPLS-DA model.