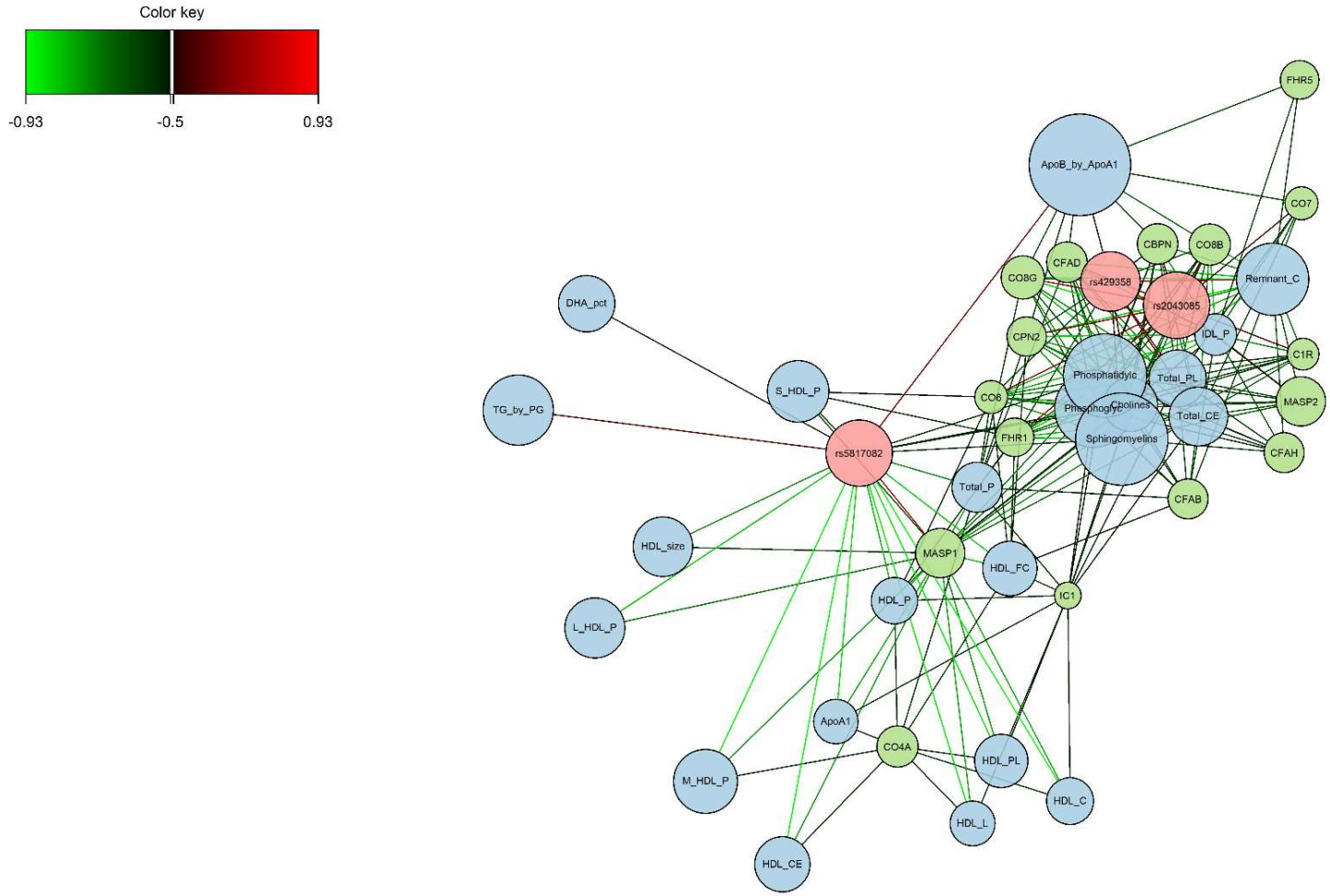Figure S7. correlations between complement proteins (n=16), lipids (n=23) and genetic variants (n=3) selected by the multi-block sPLS-DA model (correlations≥0.5 are considered in this graph).