Figure S5. contributions to AMD, expressed odds ratios with their 95% confidence intervals using a logistic regression model, of the complement proteins (n=19) (A), lipids (n=32) (B), and genetic variants (n=3) (C) selected by the multi-block sPLS-DA model. Each variable was tested separately in a different model adjusted on age, sex, hypertension, diabetes, BMI, smoking status and lipid-lowering medications. Red triangles correspond to significant ORs (p<0.05)