

Effect of variant coding on *APOL1* rs73885319 × HDLC interaction¹

| Variable | β (SE) | P-value |
|----------------------|--------------|---------|
| logHDL | 3.5 (4.4) | 0.43 |
| rs73885319 AG | 0.05 (2.6) | 0.98 |
| rs73885319 GG | -1.8 (1.7) | 0.31 |
| rs73885319 AG × HDLC | -4.0 (6.6) | 0.54 |
| rs73885319 GG × HDLC | -46.1 (13.1) | 0.0005 |

¹Results are for a model of eGFR in HUF5, NHAAN, and ARIC individuals including terms for rs73885319 (coded as # of variant alleles), HDLC, rs73885319 AG × HDLC, rs73885319 GG × HDLC, age, BMI, sex, study, and genome-wide proportion African ancestry and a random term for family.