

**Table 5.** Results of single SNPs association analyses for  $\beta$ -CTX and P1NP levels

| Dependent<br>Covariates |            | $\beta$ -CTX |              |                |              | P1NP   |              |                |              |
|-------------------------|------------|--------------|--------------|----------------|--------------|--------|--------------|----------------|--------------|
|                         |            | r            | <i>P1</i>    | $\beta$ (SE)   | <i>P2</i>    | r      | <i>P1</i>    | $\beta$ (SE)   | <i>P2</i>    |
| Gene                    | SNP        |              |              |                |              |        |              |                |              |
| GC                      | rs222020   | 0.010        | 0.349        |                |              | 0.008  | 0.379        |                |              |
| GC                      | rs2298849  | -0.004       | 0.436        |                |              | -0.003 | 0.457        |                |              |
| CYP2R1                  | rs12794714 | -0.053       | <b>0.021</b> | -0.056 (0.018) | <b>0.002</b> | -0.064 | <b>0.007</b> | -0.017 (0.007) | <b>0.014</b> |
| CYP2R1                  | rs10741657 | 0.036        | 0.084        |                |              | 0.046  | <b>0.036</b> |                |              |
| CYP2R1                  | rs1562902  | 0.025        | 0.163        |                |              | 0.036  | 0.085        |                |              |
| CYP2R1                  | rs10766197 | -0.018       | 0.246        | 0.044 (0.018)  | <b>0.012</b> | -0.043 | 0.050        |                |              |
| Age                     |            | 0.029        | 0.128        |                |              | 0.030  | 0.121        |                |              |
| BMI                     |            | -0.052       | <b>0.022</b> | -0.003 (0.001) | 0.051        | 0.010  | 0.351        |                |              |

P1NP levels were log-transformed to approximate normality. Bold numbers represent significant *P* values.

r: Pearson correlation coefficient; *P1*: *P* value of Pearson correlation;  $\beta$ : regression coefficient; SE: standard error; *P2*: *P* value of general linear regression.