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| Table S2. Heterogeneity and pleiotropy analysis between six proteins (exposures) and MASLD (outcome). |
|  | MR-Egger regression | IVW |  | MR-Egger |
|  | Intercept | *P* |  | Q | Q\_ *P* value |  | Q | Q\_ *P* value |
| APOE | -0.014  | 0.332  |  | 30.224  | 0.088  |  | 28.800  | 0.092  |
| CNPY4 | -0.026  | 0.068  |  | 25.891  | 0.211  |  | 21.838  | 0.349  |
| ENTPD6 | -0.013  | 0.079  |  | 32.518  | 0.588  |  | 29.230  | 0.701  |
| HLA-A | 0.007  | 0.330  |  | 96.408  | 0.583  |  | 95.448  | 0.582  |
| SCG3 | 0.002  | 0.898  |  | 5.542  | 1.000  |  | 5.525  | 1.000  |
| TOR1AIP1 | 0.014  | 0.169  |  | 42.483  | 0.450  |  | 40.526  | 0.492  |