Supplementary Table 11. MR-PRESSO results: Genetics of Ischemic Stroke Functional Outcome, Slope-Hunter-corrected

Protein	Exposure MR analysis	Causal estimate	SD	T-stat	*P*	RSSobs	*P*
CASP-8	beta.exposure: Raw	-0.58	0.26	-2.19	0.06	3.69	0.92
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
CCL25	beta.exposure: Raw	-0.12	0.11	-1.02	0.32	21.06	0.76
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
MCP-3	beta.exposure: Raw	0.52	0.32	1.64	0.13	6.69	0.89
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
DNER	beta.exposure: Raw	0.22	0.26	0.84	0.41	33.46	0.15
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
HGF	beta.exposure: Raw	-1.94	0.48	-4.03	0.01	2.43	0.90
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
S100A12	beta.exposure: Raw	1.10	0.42	2.62	0.03	7.86	0.79
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
SIRT2	beta.exposure: Raw	-0.12	0.49	-0.25	0.81	6.88	0.62
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
TRAIL	beta.exposure: Raw	0.17	0.29	0.58	0.58	5.08	0.86
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
TRANCE	beta.exposure: Raw	0.54	0.52	1.04	0.33	8.85	0.54
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
TNFSF14	beta.exposure: Raw	-0.06	0.38	-0.14	0.89	16.88	0.42
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA

MR-PRESSO, Mendelian Randomization Pleiotropy RESidual Sum and Outlier; SD, standard deviation; NA, not applicable (i.e., no outliers identified).