Supplementary Table 6. MR-Pleiotropy RESidual Sum and Outlier results: Genetics of Ischemic Stroke Functional Outcome, National Institutes of Health Stroke Scale-adjusted and Slope-Hunter-corrected

Protein	Exposure MR analysis	Causal estimate	SD	T-stat	*P*	RSSobs	*P*
CASP-8	beta.exposure: Raw	-0.93	0.33	-2.79	0.024	4.25	0.93
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
CCL25	beta.exposure: Raw	-0.17	0.15	-1.10	0.283	24.50	0.55
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
MCP-3	beta.exposure: Raw	0.83	0.39	2.14	0.058	6.48	0.87
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
DNER	beta.exposure: Raw	0.21	0.24	0.85	0.405	20.23	0.78
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
HGF	beta.exposure: Raw	-2.41	0.85	-2.84	0.036	5.53	0.62
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
S100A12	beta.exposure: Raw	1.74	0.56	3.12	0.011	10.33	0.65
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
SIRT2	beta.exposure: Raw	0.16	0.63	0.25	0.808	7.77	0.57
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
TRAIL	beta.exposure: Raw	0.70	0.40	1.78	0.113	4.74	0.88
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
TRANCE	beta.exposure: Raw	0.63	0.38	1.66	0.140	2.59	0.97
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA
TNFSF14	beta.exposure: Raw	0.18	0.52	0.35	0.733	20.87	0.21
	beta.exposure: Outlier-corrected	NA	NA	NA	NA	NA	NA

MR, Mendelian randomization; SD, standard deviation; NA, not applicable (i.e., no outliers identified).