Supplementary Table 4. Mendelian randomization random-effects inverse-variance weighted method results: Genetics of Ischemic Stroke Functional Outcome, National Institutes of Health Stroke Scale-adjusted and Slope-Hunter-corrected

Protein	Estimate	Lower CI	Upper CI	*P*	FDR	OR	OR lower CI	OR upper CI
CASP-8	-0.929	-1.582	-0.277	0.005	0.017	0.395	0.206	0.758
CCL25	-0.167	-0.464	0.131	0.272	0.389	0.846	0.629	1.140
MCP-3	0.830	0.069	1.591	0.033	0.083	2.293	1.071	4.908
DNER	0.206	-0.270	0.682	0.396	0.495	1.229	0.763	1.979
HGF	-2.411	-4.076	-0.746	0.005	0.017	0.090	0.017	0.474
S100A12	1.735	0.646	2.825	0.002	0.017	5.672	1.907	16.864
SIRT2	0.160	-1.080	1.399	0.801	0.801	1.173	0.340	4.051
TRAIL	0.705	-0.073	1.482	0.076	0.152	2.024	0.930	4.403
TRANCE	0.633	-0.113	1.380	0.096	0.160	1.884	0.893	3.973
TNFSF14	0.181	-0.840	1.202	0.728	0.801	1.198	0.432	3.325

CI, confidence interval; FDR, false discovery rate; OR, odds ratio.