Supplementary material

Supplementary Table 1: Protein Abbreviation List for Figure 1

Protein.Names First.Protein.Description
VGF HUMAN Neurosecretory protein VGF

TYRO3 HUMAN Tyrosine-protein kinase receptor TYRO3

TIMP2_HUMAN Metalloproteinase inhibitor 2
TIMP1 HUMAN Metalloproteinase inhibitor 1

TICN1 HUMAN Testican-1

SPRL1 HUMAN SPARC-like protein 1

SPRC HUMAN SPARC

SLIK1_HUMAN SLIT and NTRK-like protein 1

SHPS1 HUMAN Tyrosine-protein phosphatase non-receptor type substrate 1

SGCE_HUMAN Epsilon-sarcoglycan SE6L2_HUMAN Seizure 6-like protein 2 IC1_HUMAN Plasma protease C1 inhibitor

A2AP_HUMAN Alpha-2-antiplasmin
AACT_HUMAN Alpha-1-antichymotrypsin
A1AT_HUMAN Alpha-1-antitrypsin
SEM7A_HUMAN Semaphorin-7A
LYAM1 HUMAN L-selectin

7B2 HUMAN Neuroendocrine protein 7B2

SCG2_HUMAN Secretogranin-2 RNAS4 HUMAN Ribonuclease 4

RNAS1_HUMAN Ribonuclease pancreatic

RGMB_HUMAN RGM domain family member B
RET4_HUMAN Retinol-binding protein 4
EPCR HUMAN Endothelial protein C receptor

PRIO_HUMAN Major prion protein KPYM_HUMAN Pyruvate kinase PKM

PGRP2_HUMAN N-acetylmuramoyl-L-alanine amidase PEBP4_HUMAN Phosphatidylethanolamine-binding protein 4

PCSK1 HUMAN ProSAAS

A1AG2_HUMAN Alpha-1-acid glycoprotein 2 A1AG1_HUMAN Alpha-1-acid glycoprotein 1

OMGP_HUMAN Oligodendrocyte-myelin glycoprotein

NRX2A_HUMAN Neurexin-2

NRCAM_HUMAN
Neuronal cell adhesion molecule
NPTXR_HUMAN
Neuronal pentraxin receptor

NPTX1_HUMAN Neuronal pentraxin-1

NEGR1 HUMAN Neuronal growth regulator 1

LYSC HUMAN Lysozyme C

LYVE1 HUMAN Lymphatic vessel endothelial hyaluronic acid receptor 1

LYNX1_HUMAN Ly-6/neurotoxin-like protein 1 LY6H HUMAN Lymphocyte antigen 6H

A2GL HUMAN Leucine-rich alpha-2-glycoprotein

LAMP2 HUMAN Lysosome-associated membrane glycoprotein 2

K2C1 HUMAN Keratin, type II cytoskeletal 1

KLKB1 HUMAN Plasma kallikrein

KLK6 HUMAN Kallikrein-6 Immunoglobulin J chain **IGJ HUMAN** Inter-alpha-trypsin inhibitor heavy chain H4 ITIH4 HUMAN Inter-alpha-trypsin inhibitor heavy chain H2 ITIH2 HUMAN Inter-alpha-trypsin inhibitor heavy chain H1 ITIH1 HUMAN Immunoglobulin lambda variable 3-19 LV319 HUMAN Immunoglobulin lambda variable 1-47 LV147 HUMAN Immunoglobulin lambda-like polypeptide 5 IGLL5 HUMAN Immunoglobulin lambda constant 3 IGLC3 HUMAN Immunoglobulin kappa variable 4-1 KV401 HUMAN Immunoglobulin kappa variable 3-20 KV320 HUMAN Immunoglobulin kappa variable 1-5 KV105 HUMAN Immunoglobulin kappa constant IGKC HUMAN Immunoglobulin heavy variable 3/OR16-9 (non-functional) S4R460 HUMAN Immunoglobulin heavy constant mu IGHM HUMAN IGHG3 HUMAN Immunoglobulin heavy constant gamma 3 IGHG1 HUMAN Immunoglobulin heavy constant gamma 1 Immunoglobulin heavy constant alpha 2 IGHA2 HUMAN IGHA1 HUMAN Immunoglobulin heavy constant alpha 1 Insulin-like growth factor-binding protein 7 **IBP7 HUMAN** IBP6 HUMAN Insulin-like growth factor-binding protein 6 Insulin-like growth factor-binding protein 4 IBP4 HUMAN IBP2_HUMAN Insulin-like growth factor-binding protein 2 Basement membrane-specific heparan sulfate proteoglycan core **PGBM HUMAN** protein Ganglioside GM2 activator SAP3 HUMAN FIBG HUMAN Fibrinogen gamma chain Fibrinogen beta chain FIBB HUMAN Fibrinogen alpha chain FIBA HUMAN IgGFc-binding protein FCGBP HUMAN FA5 HUMAN Coagulation factor V Cathepsin D CATD HUMAN CATB HUMAN Cathepsin B CBPQ HUMAN Carboxypeptidase Q CBPE HUMAN Carboxypeptidase E Ceruloplasmin CERU HUMAN CNTN2 HUMAN Contactin-2 **CLUS HUMAN** Clusterin CSTN3 HUMAN Calsyntenin-3 CSTN1 HUMAN Calsyntenin-1 SCG1 HUMAN Secretogranin-1 CFAI HUMAN Complement factor I CFAD HUMAN Complement factor D CFAB HUMAN Complement factor B CADH2 HUMAN Cadherin-2

Monocyte differentiation antigen CD14 CD14 HUMAN

CADM2 HUMAN Cell adhesion molecule 2

Voltage-dependent calcium channel subunit alpha-2/delta-1 CA2D1 HUMAN

CO9 HUMAN Complement component C9 Complement component C7 CO7 HUMAN Complement component C6 CO6 HUMAN

CO5_HUMAN	Complement C5
CO4A_HUMAN	Complement C4-A
CO2_HUMAN	Complement C2
C1R_HUMAN	Complement C1r subcomponent
C1QC_HUMAN	Complement C1q subcomponent subunit C
C1QB_HUMAN	Complement C1q subcomponent subunit B
B4GA1_HUMAN	Beta-1,4-glucuronyltransferase 1
B2MG_HUMAN	Beta-2-microglobulin
ZA2G_HUMAN	Zinc-alpha-2-glycoprotein
VAS1_HUMAN	V-type proton ATPase subunit S1
A4_HUMAN	Amyloid-beta A4 protein
APOC1_HUMAN	Apolipoprotein C-I
APOA4_HUMAN	Apolipoprotein A-IV
APOA2_HUMAN	Apolipoprotein A-II
APOA1_HUMAN	Apolipoprotein A-I
APLP2_HUMAN	Amyloid-like protein 2
APLP1_HUMAN	Amyloid-like protein 1
AMBP_HUMAN	Protein AMBP
ADA22_HUMAN	Disintegrin and metalloproteinase domain-containing protein 22
A2MG_HUMAN	Alpha-2-macroglobulin

Supplementary Table 2: Serum autoantibody findings and associated clinical characteristics in 10 patients with COVID-19.

Patient	Antibody	Syndrome
#1	NMDAR 1:100	Nystagmus, orofacial myoclonus (first present after
		resuscitation),
		MRI:hypoxic brain damage
#2	Yo 1:100	Brachiofacial myclonia, EEG normal
#3	Caspr2 1:32	Unsteady gait and delir
#4	Myelin 1:100	Protracted waking after mechanical ventilation
#5	Myelin 1:100	Hyperactive delir
#6	Myelin 1:100	Delir
#7	Myelin 1:100	Delir
#8	Caspr2 1:10 and	Very severe delir

NMDAR 1:10

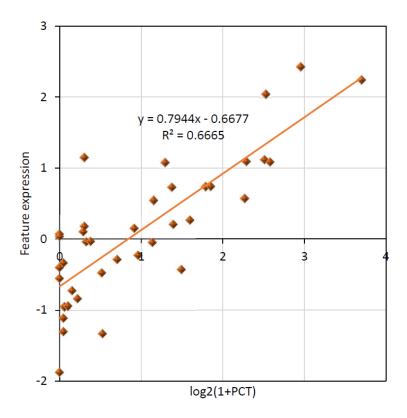
#9	Glycin 1:10	Mnestic difficulties (known AD)
#10	Glycin 1:10	Myalgia, delir

Supplementary Table 3. Change of the data matrix size after pre-processing steps.

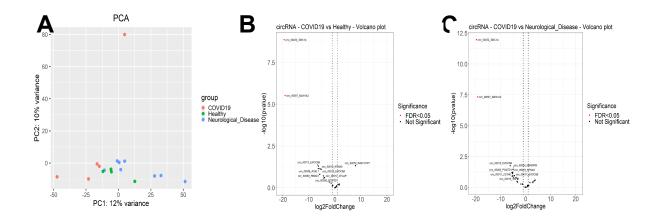
Step	N features	M samples
Input data matrix size	5486	74
After sample removal	5486	72
After features with NA>35% removal	2230	72
After filtering out nonproteotypic peptides	1992	72
After peptides summarization	271	72
Final protein matrix size	271	72

Supplementary Table 4. Clinical groups used for analysis along with some demographic information. COVID-19 Patients were divided into cohort with elevated Procalcitonin levels (PCT>1) indicating a confounding superinfection (C19 high PCT) and a cohort C19 low PCT with normal Procalcitonin levels (PCT \leq 1). All C19 high PCT patients had WHO severity grade 4, C19_low_PCT patients on average had WHO severity grade 3. Two other groups were control and HSV Meningitis (HSVE).

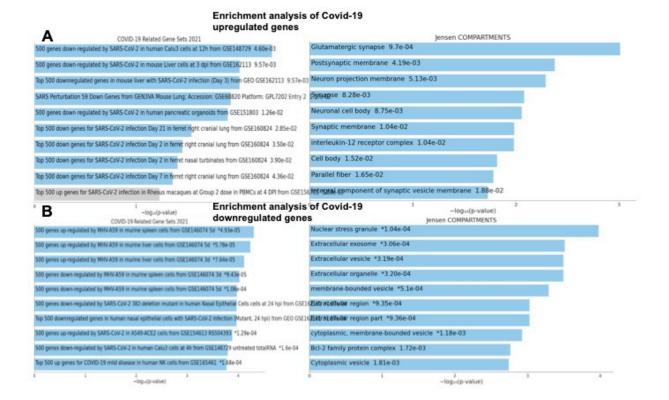
	Female			Male			Total		
	Avg	Avg	NN	Avg	Avg	NN	Avg	Avg	NN
	Age	PCT	pat	Age	PCT	pat	Age	PCT	pat
C19_high_PCT	63.7	1.8	3	71.8	4.0	13	70.3	3.6	16
C19_low_PCT	69.2	0.1	5	68.8	0.3	17	68.9	0.2	22
control	57.6		17	58.8		11	58.1		28
HSVE	53.2		6				53.2		6
Total	59.2		31	67.0		41	63.7		72



Supplementary Figure 1. Linear dependence of combined log2 expression of three **proteins** y = (C4A + CD14)/2 - NRCAM on transformed PCT level x = log2(1 + PCT). Proteins C4A and CD14 show increase of expression with increase of x and have nearly opposite dependence on sex and WHO grade, while NRCAM show decrease with increase of x and is almost independent on sex and WHO grade. Linear fit showed that approximately 67% of the variation in combined proteins expression can be explained by BSI ($R^2 = 0.67$). The fit was conducted on all COVID samples (as shown in the Figure) and on WHO grade 4 only samples with practically the same outcome.



Supplementary Figure 2. CircRNA data analysis. (A) Principal component analysis (PCA) of circRNA profile in CSF of Covid-19 patients, healthy controls and patients with neurological disease. Differential expression analysis of circRNAs in (B) CSF of Covid-19 patients and healthy controls and (C) CSF of Covid-19 patients and patients with neurological disease. Red dots: FDR < 0.05.



Supplementary Figure 3 Supplementary Figure. Enrichment analysis of deregulated genes in COVID-19 CSF. Top 10 enriched items in the COVID-19 related gene sets 2021 and cellular compartment localization in the analysis of (A) COVID-19 up-regulated genes and (B) COVID-19 down-regulated genes. The p-value is listed after each item; the items with pvalue < 0.05 are significantly enriched; An asterisk (*) next to a p-value indicates the term also has a adjusted p-value < 0.05.