

## **Integrative Plasma Metabolic and Lipidomic Modelling of SARS-CoV-2 Infection in Relation to Clinical Severity and Early Mortality Prediction**

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**Table S1 - Full cohort demographic data for the control samples and SARS-CoV-2 positive patients.**

	Controls (n=89)	SARS-CoV- 2 Positive (+) Group B severity (n=18)	SARS-CoV- 2 Positive (+) Group C severity (n=139)	SARS-CoV- 2 Positive (+) Group D severity (n=92)	SARS-CoV- 2 Positive (+) Group E severity (n=57)
Sex, Male	43 (48.31%)	5 (27.78%)	56 (40.29%)	44 (47.83%)	33 (57.89%)
Age, years [SD]	48.00 [±11.95]	62.00 [±18.13]	68.00 [±18.21]	71.00 [±17.22]	75.00 [±13.46]
BMI, kg/m <sup>2</sup> [SD]	23.81 [±4.57]	-	-	-	-
Pre-diabetic/ diabetic	-	3 (16.67%)	29 (20.86%)	24 (26.09%)	19 (33.33%)
Hypertension	-	5 (27.78%)	48 (34.53%)	50 (54.35%)	33 (57.89%)
Cardiovascular disease	-	3 (16.67%)	33 (23.74%)	29 (31.52%)	21 (36.84%)
Cerebrovascular disease	-	2 (11.11%)	16 (11.51%)	9 (9.78%)	10 (17.54%)
Chronic Renal Disease	-	0	8 (5.76%)	9 (9.78%)	6 (10.53%)
Chronic liver disease	-	0	2 (1.44%)	1 (1.09%)	2 (3.51%)
Patients who died of SARS-CoV-2	-	0	0	0	32 (56.14%)

### Quantification of Plasma Lipoproteins, Lipids and the low molecular weight metabolites

Plasma lipoproteins – A total of 112 lipoprotein parameters were quantified based on 1D NMR experiments as part of Bruker's IVD<sup>r</sup> experiment suite for blood plasma. This approach is termed Bruker's IVD<sup>r</sup> lipoprotein class and subclass Analysis (B.I.-LISA™) and lipid analytes include cholesterol, free cholesterol, phospholipids, triglycerides, apolipoproteins A1/A2/B100 and ratio B100/A1, in total plasma concentration and resolved for main lipoprotein classes and subclasses. Main classes of plasma-lipoproteins were defined as: high-density lipoprotein (HDL, density 1.063–1.210 kg/L), intermediate-density lipoprotein (IDL, density 1.006–1.019 kg/L) low-density lipoprotein (LDL, density 1.09–1.63 kg/L) and very low-density lipoprotein (VLDL, 0.950–1.006 kg/L). The main lipoprotein classes HDL, LDL, VLDL were further divided into different lipoprotein sub-classes: (LDL-1: 1.019–1.031 kg/L, LDL-2: 1.031–1.034 kg/L, LDL-3: 1.034–1.037 kg/L, LDL-4: 1.037–1.040 kg/L, LDL-5: 1.040–1.044 kg/L, LDL-6: 1.044–1.063 kg/L), and

the HDL sub-fractions into four density classes (HDL-1 1.063–1.100 kg/L, HDL-2 1.100–1.125 kg/L, HDL-3 1.125–1.175 kg/L, and HDL-4 1.175–1.210 kg/L), the VLDL sub-fractions divided into 5 density classes.

**Table S2- Annotation of the keys used by the Bruker IVDr Lipoprotein Subclass Analysis (B.I.-LISA™) method.** Abbreviations: LDL – low-density lipoprotein; HDL – high-density lipoprotein; VLDL – very low-density lipoprotein; IDL – intermediate-density lipoprotein.

Key	Class/subclass	Compound	Concentration unit
TPTG	Total Plasma	Triglycerides	mg/dL
TPCH	Total Plasma	Cholesterol	mg/dL
LDCH	LDL	Cholesterol	mg/dL
HDCH	HDL	Cholesterol	mg/dL
TPA1	Total Plasma	Apolipoprotein-A1	mg/dL
TPA2	Total Plasma	Apolipoprotein-A2	mg/dL
TPAB	Total Plasma	Apolipoprotein-B100	mg/dL
LDHD	Ratio LDL and HDL Cholesterol	LDL Cholesterol / HDL Cholesterol	-/-
ABA1	Ratio of Apolipoproteins A1 and Apolipoprotein-A1 B100	/ Apolipoprotein--/-B100	
TBPN	Apolipoprotein-B100 particles	carrying Particle Number	nmol/L
VLPN	VLDL	Particle Number	nmol/L
IDPN	IDL	Particle Number	nmol/L
LDPN	LDL	Particle Number	nmol/L
L1PN	LDL-1	Particle Number	nmol/L
L2PN	LDL-2	Particle Number	nmol/L
L3PN	LDL-3	Particle Number	nmol/L
L4PN	LDL-4	Particle Number	nmol/L
L5PN	LDL-5	Particle Number	nmol/L
L6PN	LDL-6	Particle Number	nmol/L
VLTG	VLDL Class	Triglycerides	mg/dL
IDTG	IDL Class	Triglycerides	mg/dL
LDTG	LDL Class	Triglycerides	mg/dL
HDTG	HDL Class	Triglycerides	mg/dL
VLCH	VLDL Class	Cholesterol	mg/dL
IDCH	IDL Class	Cholesterol	mg/dL
LDCH	LDL Class	Cholesterol	mg/dL
HDCH	HDL Class	Cholesterol	mg/dL
VLFC	VLDL Class	Free Cholesterol	mg/dL
IDFC	IDL Class	Free Cholesterol	mg/dL
LDFC	LDL Class	Free Cholesterol	mg/dL
HDFC	HDL Class	Free Cholesterol	mg/dL
VLPL	VLDL Class	Phospholipids	mg/dL
IDPL	IDL Class	Phospholipids	mg/dL
LDPL	LDL Class	Phospholipids	mg/dL
HDPL	HDL Class	Phospholipids	mg/dL

HDA1	HDL Class	Apolipoprotein-A1	mg/dL
HDA2	HDL Class	Apolipoprotein-A2	mg/dL
VLAB	VLDL Class	Apolipoprotein-B100	mg/dL
IDAB	IDL Class	Apolipoprotein-B100	mg/dL
LDAB	LDL Class	Apolipoprotein-B100	mg/dL
V1TG	VLDL-1 Subclass	Triglycerides	mg/dL
V2TG	VLDL-2 Subclass	Triglycerides	mg/dL
V3TG	VLDL-3 Subclass	Triglycerides	mg/dL
V4TG	VLDL-4 Subclass	Triglycerides	mg/dL
V5TG	VLDL-5 Subclass	Triglycerides	mg/dL
V1CH	VLDL-1 Subclass	Cholesterol	mg/dL
V2CH	VLDL-2 Subclass	Cholesterol	mg/dL
V3CH	VLDL-3 Subclass	Cholesterol	mg/dL
V4CH	VLDL-4 Subclass	Cholesterol	mg/dL
V5CH	VLDL-5 Subclass	Cholesterol	mg/dL
V1FC	VLDL-1 Subclass	Free Cholesterol	mg/dL
V2FC	VLDL-2 Subclass	Free Cholesterol	mg/dL
V3FC	VLDL-3 Subclass	Free Cholesterol	mg/dL
V4FC	VLDL-4 Subclass	Free Cholesterol	mg/dL
V5FC	VLDL-5 Subclass	Free Cholesterol	mg/dL
V1PL	VLDL-1 Subclass	Phospholipids	mg/dL
V2PL	VLDL-2 Subclass	Phospholipids	mg/dL
V3PL	VLDL-3 Subclass	Phospholipids	mg/dL
V4PL	VLDL-4 Subclass	Phospholipids	mg/dL
V5PL	VLDL-5 Subclass	Phospholipids	mg/dL
L1TG	LDL-1 Subclass	Triglycerides	mg/dL
L2TG	LDL-2 Subclass	Triglycerides	mg/dL
L3TG	LDL-3 Subclass	Triglycerides	mg/dL
L4TG	LDL-4 Subclass	Triglycerides	mg/dL
L5TG	LDL-5 Subclass	Triglycerides	mg/dL
L6TG	LDL-6 Subclass	Triglycerides	mg/dL
L1CH	LDL-1 Subclass	Cholesterol	mg/dL
L2CH	LDL-2 Subclass	Cholesterol	mg/dL
L3CH	LDL-3 Subclass	Cholesterol	mg/dL
L4CH	LDL-4 Subclass	Cholesterol	mg/dL
L5CH	LDL-5 Subclass	Cholesterol	mg/dL
L6CH	LDL-6 Subclass	Cholesterol	mg/dL
L1FC	LDL-1 Subclass	Free apoHDA Cholesterol	mg/dL
L2FC	LDL-2 Subclass	Free Cholesterol	mg/dL
L3FC	LDL-3 Subclass	Free Cholesterol	mg/dL
L4FC	LDL-4 Subclass	Free Cholesterol	mg/dL
L5FC	LDL-5 Subclass	Free Cholesterol	mg/dL
L6FC	LDL-6 Subclass	Free Cholesterol	mg/dL
L1PL	LDL-1 Subclass	Phospholipids	mg/dL
L2PL	LDL-2 Subclass	Phospholipids	mg/dL

L3PL	LDL-3 Subclass	Phospholipids	mg/dL
L4PL	LDL-4 Subclass	Phospholipids	mg/dL
L5PL	LDL-5 Subclass	Phospholipids	mg/dL
L6PL	LDL-6 Subclass	Phospholipids	mg/dL
L1AB	LDL-1 Subclass	Apolipoprotein-B100	mg/dL
L2AB	LDL-2 Subclass	Apolipoprotein-B100	mg/dL
L3AB	LDL-3 Subclass	Apolipoprotein-B100	mg/dL
L4AB	LDL-4 Subclass	Apolipoprotein-B100	mg/dL
L5AB	LDL-5 Subclass	Apolipoprotein-B100	mg/dL
L6AB	LDL-6 Subclass	Apolipoprotein-B100	mg/dL
H1TG	HDL-1 Subclass	Triglycerides	mg/dL
H2TG	HDL-2 Subclass	Triglycerides	mg/dL
H3TG	HDL-3 Subclass	Triglycerides	mg/dL
H4TG	HDL-4 Subclass	Triglycerides	mg/dL
H1CH	HDL-1 Subclass	Cholesterol	mg/dL
H2CH	HDL-2 Subclass	Cholesterol	mg/dL
H3CH	HDL-3 Subclass	Cholesterol	mg/dL
H4CH	HDL-4 Subclass	Cholesterol	mg/dL
H1FC	HDL-1 Subclass	Free Cholesterol	mg/dL
H2FC	HDL-2 Subclass	Free Cholesterol	mg/dL
H3FC	HDL-3 Subclass	Free Cholesterol	mg/dL
H4FC	HDL-4 Subclass	Free Cholesterol	mg/dL
H1PL	HDL-1 Subclass	Phospholipids	mg/dL
H2PL	HDL-2 Subclass	Phospholipids	mg/dL
H3PL	HDL-3 Subclass	Phospholipids	mg/dL
H4PL	HDL-4 Subclass	Phospholipids	mg/dL
H1A1	HDL-1 Subclass	Apolipoprotein-A1	mg/dL
H2A1	HDL-2 Subclass	Apolipoprotein-A1	mg/dL
H3A1	HDL-3 Subclass	Apolipoprotein-A1	mg/dL
H4A1	HDL-4 Subclass	Apolipoprotein-A1	mg/dL
H1A2	HDL-1 Subclass	Apolipoprotein-A2	mg/dL
H2A2	HDL-2 Subclass	Apolipoprotein-A2	mg/dL
H3A2	HDL-3 Subclass	Apolipoprotein-A2	mg/dL
H4A2	HDL-4 Subclass	Apolipoprotein-A2	mg/dL

**Table S3 - Annotation of the low molecular weight metabolites. Those in italics were quantified by NMR, while the remaining were quantified by LC-MS.**

Abbreviations in models	Metabolite	Concentration unit
<i>Acetic acid</i>	<i>Acetic acid</i>	mmol/L
<i>Acetoacetic acid</i>	<i>Acetoacetic acid</i>	mmol/L
<i>Acetone</i>	<i>Acetone</i>	mmol/L
<i>Citric acid</i>	<i>Citric acid</i>	mmol/L
<i>Creatine</i>	<i>Creatine</i>	mmol/L
<i>Creatinine</i>	<i>Creatinine</i>	mmol/L
<i>Formic acid</i>	<i>Formic acid</i>	mmol/L

<i>Glucose</i>	<i>Glucose</i>	<i>mmol/L</i>
<i>D-3-hydroxybutyric acid</i>	<i>D-3-hydroxybutyric acid</i>	<i>mmol/L</i>
<i>Lactic acid</i>	<i>Lactic acid</i>	<i>mmol/L</i>
<i>Pyruvic acid</i>	<i>Pyruvic acid</i>	<i>mmol/L</i>
Tryptophan	Tryptophan	$\mu\text{mol/L}$
3-Hydroxykynurenine	3-Hydroxykynurenine	$\mu\text{mol/L}$
3-Hydroxyanthranilic acid	3-Hydroxyanthranilic acid	$\mu\text{mol/L}$
Kynurenic acid	Kynurenic acid	$\mu\text{mol/L}$
Quinolinic acid	Quinolinic acid	$\mu\text{mol/L}$
Picolinic acid	Picolinic acid	$\mu\text{mol/L}$
Xanthurenic acid	Xanthurenic acid	$\mu\text{mol/L}$
Kynurenine	Kynurenine	$\mu\text{mol/L}$
Indole-3-acetic acid	Indole-3-acetic acid	$\mu\text{mol/L}$
5-Hydroxyindole acetic acid	5-Hydroxyindole acetic acid	$\mu\text{mol/L}$
Neopterin	Neopterin	$\mu\text{mol/L}$
Serotonin	Serotonin	$\mu\text{mol/L}$
1-methylhistidine	1-methylhistidine	$\mu\text{mol/L}$
3-methylhistidine	3-methylhistidine	$\mu\text{mol/L}$
Alanine	Alanine	$\mu\text{mol/L}$
Alpha aminobutyric acid	Alpha aminobutyric acid	$\mu\text{mol/L}$
Arginine	Arginine	$\mu\text{mol/L}$
Aspartic acid	Aspartic acid	$\mu\text{mol/L}$
Citrulline	Citrulline	$\mu\text{mol/L}$
Glutamic acid	Glutamic acid	$\mu\text{mol/L}$
Glutamine	Glutamine	$\mu\text{mol/L}$
Histidine	Histidine	$\mu\text{mol/L}$
Isoleucine	Isoleucine	$\mu\text{mol/L}$
Leucine	Leucine	$\mu\text{mol/L}$
Lysine	Lysine	$\mu\text{mol/L}$
Methionine	Methionine	$\mu\text{mol/L}$
Ornithine	Ornithine	$\mu\text{mol/L}$
Phenylalanine	Phenylalanine	$\mu\text{mol/L}$
Proline	Proline	$\mu\text{mol/L}$
Serine	Serine	$\mu\text{mol/L}$
Taurine	Taurine	$\mu\text{mol/L}$
Threonine	Threonine	$\mu\text{mol/L}$
Tyrosine	Tyrosine	$\mu\text{mol/L}$
Valine	Valine	$\mu\text{mol/L}$
<i>Lac.pyr</i>	<i>Lactate/pyruvate</i>	-/-
Kynurenine/tryptophan	Kynurenine/tryptophan	-/-
3-hydroxykynurenine/tryptophan	3-hydroxykynurenine/tryptophan	-/-
Kynurenic acid/tryptophan	Kynurenic acid/tryptophan	-/-
Neopterin/tryptophan	Neopterin/tryptophan	-/-
Serotonin/tryptophan	Serotonin/tryptophan	-/-
Quinolinic acid/tryptophan	Quinolinic acid/tryptophan	-/-
Fisher's ratio	Fisher's ratio	-/-

Branched AA/taurine	Branched chain amino acids/aromatic -/ amino acids	
Asp...glu...asn...gln	Asp:Glu:Asn:Gln	-/-

**Table S4 - Annotation of the lipids**

Abbreviation	Subclasses	Concentration unit
CE	Cholesterol esters	µmol/L
CER	Ceramides	µmol/L
DAG	Diacylglycerides	µmol/L
DCER	Dihydroceramides	µmol/L
FFA	Free fatty acids	µmol/L
HCER	Hexosylceramides	µmol/L
LCER	Lactosylceramides	µmol/L
LPC	Lysophosphocholines	µmol/L
LPE	Lysophosphoethanolamines	µmol/L
LPG	Lysophosphoglycerols	µmol/L
LPI	Lysophosphoinositols	µmol/L
MAG	Monoacylglycerols	µmol/L
PC	Phosphocholines	µmol/L
PE	Phosphoethanolamines	µmol/L
PG	Phosphoglycerols	µmol/L
PI	Phosphoinositols	µmol/L
PS	Phosphoserines	µmol/L
SM	Sphinogomyelins	µmol/L
TAG	Triacylglycerides	µmol/L

### Integrated Spectroscopy Modeling

**Table S5 - OPLS loadings, Cliff's delta and adjusted p-values of the combined model (low molecular weight metabolites, lipoproteins and lipids) for the comparison of the controls and SARS-CoV-2 positive patients.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p value
Formic acid	0.063	1.00	1.18x10 <sup>-43</sup>
PC (18:2/18:2)	0.085	-0.94	4.48x10 <sup>-39</sup>
PE.O (16:0/20:4)	0.073	-0.94	4.48x10 <sup>-39</sup>
Asp:Glu:Asn:Gln	0.039	0.93	1.21x10 <sup>-38</sup>
Pyruvic acid	0.050	0.93	2.67x10 <sup>-38</sup>
PE.P (18:1/20:4)	0.076	-0.93	2.67x10 <sup>-38</sup>
PC (18:1/18:2)	0.085	-0.92	5.41x10 <sup>-38</sup>
H4CH	0.069	-0.92	8.87x10 <sup>-38</sup>
H4PL	0.071	-0.92	1.08x10 <sup>-37</sup>

PE.P (18:1/18:2)	0.082	-0.92	1.21x10 <sup>-37</sup>
PE.P (16:0/18:2)	0.082	-0.91	2.00x10 <sup>-37</sup>
PE.P (16:0/22:5)	0.073	-0.91	2.00x10 <sup>-37</sup>
PE.O (16:0/18:2)	0.074	-0.91	2.60x10 <sup>-37</sup>
PE.P (18:1/20:4)	0.077	-0.91	2.60x10 <sup>-37</sup>
LPC (18:2)	0.077	-0.91	8.25x10 <sup>-37</sup>
PE.P (18:0/18:2)	0.079	-0.90	8.25x10 <sup>-37</sup>
PE.P (18:0/22:5)	0.076	-0.90	2.27x10 <sup>-36</sup>
PE.P (16:0/20:4)	0.074	-0.90	3.76x10 <sup>-36</sup>
PC (18:2/20:3)	0.077	-0.90	3.86x10 <sup>-36</sup>
PC (18:2/20:2)	0.076	-0.89	5.52x10 <sup>-36</sup>
PE.P (18:2/18:2)	0.076	-0.88	4.01x10 <sup>-35</sup>
Lactate/pyruvate	0.046	-0.88	6.10x10 <sup>-35</sup>
PC (20:0/20:3)	0.075	-0.88	6.10x10 <sup>-35</sup>
PE.P (18:0/18:1)	0.077	-0.88	7.95x10 <sup>-35</sup>
HCER.d (18:0/18:0)	0.074	-0.88	8.72x10 <sup>-35</sup>
PE.O (16:0/20:1)	0.073	-0.88	9.46x10 <sup>-35</sup>
H4A1	0.067	-0.87	2.68x10 <sup>-34</sup>
PE.P (18:1/22:5)	0.074	-0.87	2.95x10 <sup>-34</sup>
PE.O (16:0/22:5)	0.072	-0.87	3.00x10 <sup>-34</sup>
PE.P (18:0/20:4)	0.073	-0.87	3.40x10 <sup>-34</sup>
PE.P (16:0/20:1)	0.077	-0.87	4.35x10 <sup>-34</sup>
PC (18:2/20:4)	0.079	-0.87	4.81x10 <sup>-34</sup>
H4A2	0.067	-0.86	6.21x10 <sup>-34</sup>
PE.P (16:0/18:3)	0.066	-0.86	6.21x10 <sup>-34</sup>
PC (18:0/20:1)	0.076	-0.86	6.96x10 <sup>-34</sup>
PE.O (18:0/22:5)	0.070	-0.86	6.96x10 <sup>-34</sup>
LPC (20:0)	0.075	-0.86	1.03x10 <sup>-33</sup>
PE.P (18:0/22:4)	0.069	-0.86	1.22x10 <sup>-33</sup>
PE.O (16:0/18:1)	0.067	-0.86	1.70x10 <sup>-33</sup>
PE.P (18:0/20:1)	0.074	-0.85	4.02x10 <sup>-33</sup>
PE.P (18:0/20:3)	0.073	-0.85	4.59x10 <sup>-33</sup>
PC (18:2/18:3)	0.063	-0.84	2.40x10 <sup>-32</sup>
PE.P (18:0/16:0)	0.075	-0.84	2.71x10 <sup>-32</sup>
PE.P (18:1/22:4)	0.062	-0.84	4.30x10 <sup>-32</sup>
PE.P (18:0/20:2)	0.074	-0.84	5.79x10 <sup>-32</sup>
PE.P (18:1/18:1)	0.063	-0.84	6.28x10 <sup>-32</sup>

HCER (22:0)	0.069	-0.83	8.56x10 <sup>-32</sup>
Glutamic acid	0.045	0.83	1.59x10 <sup>-31</sup>
HCER (26:0)	0.072	-0.83	1.59x10 <sup>-31</sup>
Glutamic acid	0.046	0.83	1.69x10 <sup>-31</sup>
HCER.d (18:0/22:0)	0.069	-0.83	2.01x10 <sup>-31</sup>
PE.P (18:0/18:0)	0.071	-0.83	3.79x10 <sup>-31</sup>
LDTG	0.048	0.83	4.06x10 <sup>-31</sup>
PC (20:0/18:1)	0.068	-0.82	4.58x10 <sup>-31</sup>
PI (18:1/18:1)	0.068	-0.82	7.43x10 <sup>-31</sup>
PI (20:0/18:2)	0.070	-0.82	1.04x10 <sup>-30</sup>
PE.O (18:0/22:4)	0.052	-0.81	2.46x10 <sup>-30</sup>
PC (18:2/22:6)	0.074	-0.81	3.22x10 <sup>-30</sup>
HCER.d (18:0/24:0)	0.069	-0.81	3.25x10 <sup>-30</sup>
SM (20:0)	0.064	-0.81	4.74x10 <sup>-30</sup>
HCER (24:0)	0.069	-0.81	5.28x10 <sup>-30</sup>
PE.P (16:0/22:4)	0.057	-0.81	5.45x10 <sup>-30</sup>
TPA2	0.065	-0.81	7.21x10 <sup>-30</sup>
CER (26:0)	0.072	-0.81	7.64x10 <sup>-30</sup>
PC (18:2/22:5)	0.072	-0.80	1.55x10 <sup>-29</sup>
LPI (20:3)	0.062	-0.80	1.73x10 <sup>-29</sup>
PE.P (18:1/20:1)	0.070	-0.80	3.81x10 <sup>-29</sup>
PC (14:0/18:2)	0.071	-0.79	7.22x10 <sup>-29</sup>
SM (24:0)	0.065	-0.79	8.74x10 <sup>-29</sup>
PE.P (18:1/16:0)	0.034	-0.79	1.98x10 <sup>-28</sup>
LPC (20:3)	0.069	-0.79	2.02x10 <sup>-28</sup>
PC (16:1/18:2)	0.068	-0.78	3.14x10 <sup>-28</sup>
PC (16:0/18:2)	0.071	-0.78	4.28x10 <sup>-28</sup>
PI (18:1/20:4)	0.069	-0.78	6.18x10 <sup>-28</sup>
PI (18:1/18:2)	0.067	-0.78	6.55x10 <sup>-28</sup>
H4FC	0.062	-0.78	6.60x10 <sup>-28</sup>
MAG (20:3)	0.015	0.78	7.44x10 <sup>-28</sup>
LPI (18:1)	0.058	-0.78	8.17x10 <sup>-28</sup>
PC (18:2/20:5)	0.057	-0.78	8.17x10 <sup>-28</sup>
LPC (18:1)	0.069	-0.77	3.10x10 <sup>-27</sup>
SM (26:0)	0.064	-0.77	3.17x10 <sup>-27</sup>
Aspartic acid	0.046	0.77	4.05x10 <sup>-27</sup>
DCER (24:0)	0.072	-0.77	4.09x10 <sup>-27</sup>

PI (18:0/20:3)	0.069	-0.77	4.09x10 <sup>-27</sup>
HDA2	0.063	-0.77	4.99x10 <sup>-27</sup>
PE.P (16:0/20:5)	0.060	-0.76	7.34x10 <sup>-27</sup>
PE.P (16:0/18:1)	0.063	-0.76	8.05x10 <sup>-27</sup>
HDA1	0.066	-0.76	1.29x10 <sup>-26</sup>
PC (18:1/20:3)	0.070	-0.75	3.20x10 <sup>-26</sup>
PE.O (16:0/22:4)	0.056	-0.75	3.59x10 <sup>-26</sup>
PC (14:0/20:4)	0.066	-0.75	4.26x10 <sup>-26</sup>
HCER (20:0)	0.059	-0.75	4.62x10 <sup>-26</sup>
PC (14:0/18:3)	0.050	-0.75	8.38x10 <sup>-26</sup>
LPC (18:3)	0.060	-0.75	8.78x10 <sup>-26</sup>
Neopterin/tryptophan	0.024	0.71	1.49x10 <sup>-25</sup>
MAG (20:4)	0.015	0.74	1.52x10 <sup>-25</sup>
TPA1	0.066	-0.74	1.68x10 <sup>-25</sup>
PC (18:2/16:1)	0.065	-0.74	1.68x10 <sup>-25</sup>
PC (16:0/20:1)	0.068	-0.74	1.69x10 <sup>-25</sup>
Fisher's ratio	0.046	-0.74	2.07x10 <sup>-25</sup>
PI (20:0/20:4)	0.065	-0.74	2.27x10 <sup>-25</sup>
PC (20:0/20:4)	0.067	-0.74	2.38x10 <sup>-25</sup>
PC (18:0/20:2)	0.069	-0.74	4.68x10 <sup>-25</sup>
PI (18:1/20:3)	0.063	-0.73	5.49x10 <sup>-25</sup>
LPI (18:2)	0.049	-0.73	5.56x10 <sup>-25</sup>
PC (14:0/22:6)	0.065	-0.73	1.10x10 <sup>-24</sup>
PC (18:0/18:2)	0.068	-0.73	1.44x10 <sup>-24</sup>
H3FC	0.062	-0.72	2.53x10 <sup>-24</sup>
Quinolinic acid/tryptophan	0.016	0.72	3.12x10 <sup>-24</sup>
PE.P (18:1/20:3)	0.062	-0.72	3.79x10 <sup>-24</sup>
PE.P (18:1/20:5)	0.055	-0.72	6.03x10 <sup>-24</sup>
PS (20:0/18:2)	0.065	-0.72	6.67x10 <sup>-24</sup>
LPC (20:2)	0.067	-0.72	7.30x10 <sup>-24</sup>
LPI (20:4)	0.048	-0.72	8.16x10 <sup>-24</sup>
PG (20:0/18:2)	0.056	-0.72	8.18x10 <sup>-24</sup>
PE.O (16:0/22:6)	0.055	-0.71	8.75x10 <sup>-24</sup>
CER (24:0)	0.066	-0.71	9.16x10 <sup>-24</sup>
DCER (26:0)	0.062	-0.71	1.48x10 <sup>-23</sup>
PI (18:2/18:2)	0.051	-0.71	1.49x10 <sup>-23</sup>
HCER (14:0)	0.057	-0.71	1.61x10 <sup>-23</sup>

HDCH	0.061	-0.71	2.04x10 <sup>-23</sup>
PC (18:1/20:1)	0.061	-0.71	2.39x10 <sup>-23</sup>
PS (18:0/18:0)	0.041	0.71	2.40x10 <sup>-23</sup>
PC (14:0/20:3)	0.062	-0.71	2.71x10 <sup>-23</sup>
LPE (18:2)	0.052	-0.70	4.36x10 <sup>-23</sup>
Phenylalanine	0.046	0.70	6.63x10 <sup>-23</sup>
H2TG	0.038	0.70	6.63x10 <sup>-23</sup>
PC (18:1/18:3)	0.049	-0.70	8.92x10 <sup>-23</sup>
CE (16:0)	0.060	-0.70	9.44x10 <sup>-23</sup>
H3CH	0.061	-0.70	1.18x10 <sup>-22</sup>
L4TG	0.042	0.70	1.27x10 <sup>-22</sup>
PE.P (16:0/20:3)	0.059	-0.69	1.75x10 <sup>-22</sup>
PE.P (16:0/16:1)	0.054	-0.69	1.99x10 <sup>-22</sup>
PE.P (18:1/20:2)	0.063	-0.69	2.55x10 <sup>-22</sup>
MAG (18:1)	0.015	0.69	2.81x10 <sup>-22</sup>
PC (14:0/18:1)	0.058	-0.68	6.84x10 <sup>-22</sup>
SM (14:0)	0.063	-0.68	7.59x10 <sup>-22</sup>
PC (20:0/22:5)	0.057	-0.68	2.20x10 <sup>-21</sup>
DCER (26:1)	0.060	-0.67	2.36x10 <sup>-21</sup>
HCER.d (18:0/20:0)	0.055	-0.67	2.36x10 <sup>-21</sup>
PI (18:0/20:4)	0.061	-0.67	2.36x10 <sup>-21</sup>
PC (14:0/20:5)	0.049	-0.67	3.20x10 <sup>-21</sup>
Histidine	0.046	-0.67	3.34x10 <sup>-21</sup>
CE (18:1)	0.057	-0.67	4.03x10 <sup>-21</sup>
PC (18:0/20:3)	0.063	-0.67	4.09x10 <sup>-21</sup>
PE.P (18:1/18:0)	0.055	-0.67	5.09x10 <sup>-21</sup>
LCER (24:0)	0.054	-0.67	5.37x10 <sup>-21</sup>
L1TG	0.037	0.67	5.50x10 <sup>-21</sup>
L5TG	0.041	0.67	5.69x10 <sup>-21</sup>
Quinolinic acid	0.020	0.67	6.90x10 <sup>-21</sup>
PC (18:1/22:4)	0.056	-0.66	1.00x10 <sup>-20</sup>
CE (18:3)	0.050	-0.66	1.11x10 <sup>-20</sup>
PS (18:0/20:4)	0.047	-0.66	1.29x10 <sup>-20</sup>
Neopterin	0.026	0.66	1.39x10 <sup>-20</sup>
CER (18:0)	0.036	0.66	2.41x10 <sup>-20</sup>
PC (14:0/14:0)	0.040	-0.66	2.90x10 <sup>-20</sup>
CE (14:0)	0.053	-0.65	3.66x10 <sup>-20</sup>

Citrulline	0.042	-0.65	4.91x10 <sup>-20</sup>
L2TG	0.041	0.65	5.56x10 <sup>-20</sup>
PC (14:0/22:5)	0.053	-0.65	5.69x10 <sup>-20</sup>
PC (16:0/18:0)	0.019	-0.65	6.19x10 <sup>-20</sup>
HDPL	0.058	-0.65	7.11x10 <sup>-20</sup>
LPE (20:3)	0.042	-0.65	9.47x10 <sup>-20</sup>
PE (18:2/16:1)	0.055	-0.65	1.24x10 <sup>-19</sup>
LPC (20:4)	0.057	-0.64	2.25x10 <sup>-19</sup>
LPC (22:5)	0.055	-0.64	3.79x10 <sup>-19</sup>
PC (16:0/20:3)	0.063	-0.64	4.08x10 <sup>-19</sup>
H3PL	0.055	-0.64	4.49x10 <sup>-19</sup>
H3A1	0.058	-0.63	7.57x10 <sup>-19</sup>
PG (20:0/20:3)	0.038	-0.63	8.70x10 <sup>-19</sup>
PE.P (18:0/20:5)	0.051	-0.63	1.12x10 <sup>-19</sup>
LCER.d (18:0/22:0)	0.052	-0.63	1.44x10 <sup>-18</sup>
PC (18:0/18:1)	0.049	-0.63	1.44x10 <sup>-18</sup>
CE (22:0)	0.053	-0.63	1.44x10 <sup>-18</sup>
PC (18:0/18:0)	0.050	-0.63	1.51x10 <sup>-18</sup>
PS (20:0/18:3)	0.049	-0.63	1.51x10 <sup>-18</sup>
LPC (18:0)	0.062	-0.62	1.87x10 <sup>-18</sup>
CE (24:0)	0.047	-0.62	2.09x10 <sup>-18</sup>
LPE (20:4)	0.045	-0.62	3.38x10 <sup>-18</sup>
LCER.d (18:0/24:0)	0.051	-0.62	3.52x10 <sup>-18</sup>
PS (20:0/20:4)	0.057	-0.62	4.75x10 <sup>-18</sup>
DCER (22:0)	0.052	-0.62	5.19x10 <sup>-18</sup>
PC (14:0/22:4)	0.049	-0.61	6.72x10 <sup>-18</sup>
PS (20:0/20:5)	0.047	-0.61	6.94x10 <sup>-18</sup>
PE.O (18:0/18:2)	0.039	-0.61	1.07x10 <sup>-17</sup>
VLPN	0.032	0.61	1.35x10 <sup>-17</sup>
VLAB	0.032	0.61	1.35x10 <sup>-17</sup>
ABA1	0.040	0.61	1.51x10 <sup>-17</sup>
LPC (22:4)	0.052	-0.61	1.57x10 <sup>-17</sup>
PG (18:0/18:2)	0.049	-0.60	2.60x10 <sup>-17</sup>
PC (18:1/22:6)	0.057	-0.60	2.78x10 <sup>-17</sup>
PC (18:0/22:4)	0.052	-0.60	3.27x10 <sup>-17</sup>
DCER (22:1)	0.050	-0.60	3.91x10 <sup>-17</sup>
V4TG	0.033	0.60	4.83x10 <sup>-17</sup>

PI (16:0/20:3)	0.053	-0.60	5.15x10 <sup>-17</sup>
PI (18:0/20:2)	0.042	-0.59	6.92x10 <sup>-17</sup>
LPE (18:1)	0.051	-0.59	7.48x10 <sup>-17</sup>
PE (18:2/20:4)	0.054	-0.59	1.03x10 <sup>-16</sup>
PE (18:2/18:2)	0.053	-0.59	1.19x10 <sup>-16</sup>
Glutamine	0.037	-0.59	1.27x10 <sup>-16</sup>
LPC (22:6)	0.050	-0.59	1.40x10 <sup>-16</sup>
PG (18:0/18:1)	0.051	-0.59	1.71x10 <sup>-16</sup>
SM (20:1)	0.050	-0.59	1.91x10 <sup>-16</sup>
PC (16:0/20:4)	0.053	-0.59	1.96x10 <sup>-16</sup>
PC (14:0/20:2)	0.051	-0.59	2.07x10 <sup>-16</sup>
PC (18:1/18:1)	0.050	-0.58	2.26x10 <sup>-16</sup>
LPE (18:3)	0.028	-0.58	2.52x10 <sup>-16</sup>
PI (18:0/18:1)	0.050	-0.58	3.01x10 <sup>-16</sup>
CE (18:0)	0.052	-0.58	4.67x10 <sup>-16</sup>
SM (24:1)	0.051	-0.58	5.75x10 <sup>-16</sup>
HCER.d (18:0/26:0)	0.045	-0.58	5.78x10 <sup>-16</sup>
PG (18:1/16:1)	0.045	0.58	6.63x10 <sup>-16</sup>
MAG (22:5)	0.013	0.58	6.77x10 <sup>-16</sup>
V3TG	0.027	0.57	1.40x10 <sup>-15</sup>
CE (20:2)	0.053	-0.57	1.51x10 <sup>-15</sup>
PE.P (16:0/22:6)	0.050	-0.57	1.51x10 <sup>-15</sup>
LPE (20:1)	0.049	-0.57	1.91x10 <sup>-15</sup>
PI (16:0/20:4)	0.047	-0.56	4.21x10 <sup>-15</sup>
PC (16:0/22:4)	-0.050	-0.56	4.85x10 <sup>-15</sup>
V2FC	0.025	0.56	5.15x10 <sup>-15</sup>
H1TG	0.027	0.55	7.66x10 <sup>-15</sup>
PI (18:0/22:4)	0.049	-0.55	1.04x10 <sup>-14</sup>
PG (20:0/20:5)	0.035	-0.55	1.11x10 <sup>-14</sup>
LPE (20:2)	0.041	-0.55	1.15x10 <sup>-14</sup>
LCER (14:0)	0.042	-0.55	1.66x10 <sup>-14</sup>
CE (20:3)	0.053	-0.54	2.30x10 <sup>-14</sup>
L6TG	0.035	0.54	2.54x10 <sup>-14</sup>
HDFC	0.050	-0.54	2.94x10 <sup>-14</sup>
LPC (20:5)	0.040	-0.54	3.71x10 <sup>-14</sup>
Kynurenine/tryptophan	0.033	0.54	3.84x10 <sup>-14</sup>
PS (20:0/22:6)	0.051	-0.54	4.10x10 <sup>-14</sup>

SM (22:1)	0.044	-0.54	4.70x10 <sup>-14</sup>
PG (18:0/20:0)	0.036	0.54	4.80x10 <sup>-14</sup>
PC (18:2/20:1)	0.042	-0.53	1.08x10 <sup>-13</sup>
PE.O (18:0/20:4)	0.040	-0.53	1.13x10 <sup>-13</sup>
PC (16:0/20:5)	0.040	-0.53	1.18x10 <sup>-13</sup>
V3FC	0.024	0.53	1.42x10 <sup>-13</sup>
MAG (22:4)	0.014	0.53	1.51x10 <sup>-13</sup>
PI (20:0/16:1)	0.050	-0.53	1.52x10 <sup>-13</sup>
PC (16:0/14:0)	0.040	-0.53	1.57x10 <sup>-13</sup>
CE (20:1)	0.037	-0.52	2.48x10 <sup>-13</sup>
MAG (16:0)	0.012	0.52	2.88x10 <sup>-13</sup>
PI (16:0/18:1)	0.045	-0.52	3.00x10 <sup>-13</sup>
PC (16:1/18:1)	0.036	-0.52	3.29x10 <sup>-13</sup>
MAG (18:2)	0.013	0.52	5.34x10 <sup>-13</sup>
H2A1	0.048	-0.51	6.06x10 <sup>-13</sup>
LCER (16:0)	0.038	-0.51	8.86x10 <sup>-13</sup>
PG (18:1/18:2)	0.035	0.51	9.70x10 <sup>-13</sup>
PC (18:0/14:0)	0.044	-0.51	1.05x10 <sup>-12</sup>
PE.P (16:0/18:0)	0.044	-0.51	1.22x10 <sup>-12</sup>
Threonine	0.027	-0.51	1.24x10 <sup>-12</sup>
3-Hydroxykynurenine	0.022	0.47	1.42x10 <sup>-12</sup>
H3TG	0.024	0.51	1.53x10 <sup>-12</sup>
PG (20:0/20:1)	0.047	-0.51	1.58x10 <sup>-12</sup>
PC (16:0/22:6)	0.049	-0.51	1.61x10 <sup>-12</sup>
LPC (14:0)	0.046	-0.50	1.87x10 <sup>-12</sup>
PI (16:0/20:2)	0.047	-0.50	1.95x10 <sup>-12</sup>
Lactic acid	0.030	0.50	2.42x10 <sup>-12</sup>
PE (18:1/18:2)	0.036	-0.50	2.38x10 <sup>-12</sup>
HCER.d (18:0/26:1)	0.032	-0.50	2.45x10 <sup>-12</sup>
V4PL	0.031	0.50	2.96x10 <sup>-12</sup>
HCER (22:1)	-0.037	-0.50	2.95x10 <sup>-12</sup>
LPC (20:1)	0.051	-0.50	3.23x10 <sup>-12</sup>
PC (16:0/18:3)	0.026	-0.50	3.96x10 <sup>-12</sup>
PC (16:0/18:1)	0.040	-0.50	4.32x10 <sup>-12</sup>
H3A2	0.046	-0.49	6.54x10 <sup>-12</sup>
PI (18:0/20:0)	0.039	0.49	6.54x10 <sup>-12</sup>
PC (18:1/20:4)	0.046	-0.49	7.01x10 <sup>-12</sup>

V2TG	0.018	0.49	7.88x10 <sup>-12</sup>
PE.P (18:1/22:6)	0.047	-0.49	1.14x10 <sup>-11</sup>
TPCH	0.048	-0.48	1.48x10 <sup>-11</sup>
CE (22:5)	0.043	-0.48	1.51x10 <sup>-11</sup>
LCER (26:1)	0.041	-0.48	1.85x10 <sup>-11</sup>
PS (20:0/22:5)	0.049	-0.48	3.29x10 <sup>-11</sup>
MAG (16:1)	0.011	0.47	5.62x10 <sup>-11</sup>
TPTG	0.013	0.47	5.75x10 <sup>-11</sup>
Glucose	0.024	0.47	5.94x10 <sup>-11</sup>
LCER (24:1)	0.037	-0.47	5.94x10 <sup>-11</sup>
V3PL	0.025	0.47	6.29x10 <sup>-11</sup>
Methionine	0.016	-0.47	7.06x10 <sup>-11</sup>
HDTG	0.021	0.47	7.51x10 <sup>-11</sup>
PE (18:1/18:3)	0.029	-0.46	1.31x10 <sup>-10</sup>
L3TG	0.031	0.46	1.52x10 <sup>-10</sup>
H1FC	0.043	-0.46	1.71x10 <sup>-10</sup>
PC (16:0/20:2)	0.045	-0.46	1.88x10 <sup>-10</sup>
MAG (22:6)	0.012	0.46	2.04x10 <sup>-10</sup>
HCER (26:1)	0.036	-0.45	2.71x10 <sup>-10</sup>
CE (20:0)	0.035	-0.45	2.74x10 <sup>-10</sup>
PE.P (18:0/22:6)	0.042	-0.45	3.55x10 <sup>-10</sup>
HCER.d (18:0/24:1)	0.034	-0.45	3.99x10 <sup>-10</sup>
V5TG	0.024	0.45	5.21x10 <sup>-10</sup>
HCER (18:0)	0.032	-0.44	6.39x10 <sup>-10</sup>
PC (16:0/22:5)	0.042	-0.44	7.67x10 <sup>-10</sup>
SM (18:1)	0.042	-0.44	9.01x10 <sup>-10</sup>
SM (26:1)	0.035	-0.44	1.00x10 <sup>-9</sup>
L4CH	0.041	-0.44	1.26x10 <sup>-9</sup>
MAG (20:2)	0.011	0.43	1.57x10 <sup>-9</sup>
H2FC	0.039	-0.43	1.62x10 <sup>-9</sup>
L1CH	0.028	-0.43	1.77x10 <sup>-9</sup>
PS (20:0/20:3)	0.040	-0.43	1.82x10 <sup>-9</sup>
CE (20:5)	0.035	-0.43	2.84x10 <sup>-9</sup>
Tryptophan	0.033	-0.42	4.02x10 <sup>-9</sup>
V3CH	0.023	0.42	4.15x10 <sup>-9</sup>
DAG (16:0/20:4)	0.008	0.42	4.43x10 <sup>-9</sup>
LPE (16:1)	0.025	-0.42	6.29x10 <sup>-9</sup>

PE.O (18:0/22:6)	0.037	-0.41	8.67x10 <sup>-9</sup>
CER (22:1)	0.022	0.41	9.89x10 <sup>-9</sup>
V4FC	0.026	0.41	1.21x10 <sup>-8</sup>
VLTG	0.010	0.41	1.34x10 <sup>-8</sup>
PE (18:0/22:6)	0.015	0.41	1.38x10 <sup>-8</sup>
PE (18:2/18:3)	0.020	-0.41	1.39x10 <sup>-8</sup>
LPE (22:4)	0.021	-0.41	1.41x10 <sup>-8</sup>
LPC (16:0)	0.046	-0.41	1.46x10 <sup>-8</sup>
V2PL	0.016	0.41	1.74x10 <sup>-8</sup>
Acetone	0.015	0.40	1.96x10 <sup>-8</sup>
LDCH	0.037	-0.40	2.88x10 <sup>-8</sup>
CER (14:0)	0.040	-0.40	3.99x10 <sup>-8</sup>
DAG (16:0/18:1)	0.007	0.40	4.05x10 <sup>-8</sup>
PE.P (16:0/16:0)	0.033	-0.39	5.80x10 <sup>-8</sup>
3-Hydroxybutyric acid	0.022	0.39	6.03x10 <sup>-8</sup>
FFA (24:0)	0.033	-0.39	6.05x10 <sup>-8</sup>
PI (18:1/16:1)	0.032	-0.39	7.33x10 <sup>-8</sup>
Kynurenic acid	0.011	-0.39	8.31x10 <sup>-8</sup>
LCER (18:1)	0.028	-0.39	8.90x10 <sup>-8</sup>
IDPN	0.025	0.39	8.98x10 <sup>-8</sup>
IDAB	0.025	0.39	9.00x10 <sup>-8</sup>
VLCH	0.016	0.39	9.20x10 <sup>-8</sup>
PC (18:1/20:5)	0.024	-0.39	9.86x10 <sup>-8</sup>
HCER (24:1)	0.032	-0.38	1.08x10 <sup>-7</sup>
PE (16:0/18:1)	0.012	0.38	1.09x10 <sup>-7</sup>
PS (20:0/22:4)	0.027	-0.38	1.16x10 <sup>-7</sup>
DCER (18:0)	0.025	0.38	1.17x10 <sup>-7</sup>
LPC (16:1)	0.035	-0.38	1.26x10 <sup>-7</sup>
LPG (18:1)	0.025	-0.38	1.34x10 <sup>-7</sup>
PE.P (18:0/16:1)	0.037	-0.38	1.84x10 <sup>-7</sup>
L4PL	0.037	-0.38	2.00x10 <sup>-7</sup>
PG (18:1/20:4)	0.028	0.38	2.18x10 <sup>-7</sup>
DAG (18:1/20:4)	0.011	0.37	2.43x10 <sup>-7</sup>
DAG (16:0/22:6)	0.008	0.37	2.46x10 <sup>-7</sup>
PS (18:0/18:1)	0.013	0.37	2.66x10 <sup>-7</sup>
TAG (55:5)_FA (20:4)	0.008	0.37	2.70x10 <sup>-7</sup>
3-Hydroxykynurenine	0.019	0.37	3.60x10 <sup>-7</sup>

V2CH	0.015	0.37	3.84x10 <sup>-7</sup>
L4FC	0.037	-0.37	4.70x10 <sup>-7</sup>
PE (18:0/22:4)	0.025	-0.36	5.13x10 <sup>-7</sup>
PE (18:1/20:4)	0.031	-0.36	5.47x10 <sup>-7</sup>
PI (16:0/22:4)	0.032	-0.36	6.05x10 <sup>-7</sup>
MAG (18:0)	0.017	0.36	7.02x10 <sup>-7</sup>
V4CH	0.025	0.36	7.26x10 <sup>-7</sup>
LPE (22:5)	0.013	-0.36	7.64x10 <sup>-7</sup>
L3CH	0.032	-0.36	8.91x10 <sup>-7</sup>
PI (18:0/18:3)	0.023	-0.36	8.99x10 <sup>-7</sup>
PI (16:0/18:2)	0.034	-0.35	1.10x10 <sup>-6</sup>
PI (18:0/18:2)	0.036	-0.35	1.35x10 <sup>-6</sup>
TAG (54:4)_FA (20:4)	0.010	0.35	1.40x10 <sup>-6</sup>
TAG (54:5)_FA (20:4)	0.013	0.35	1.53x10 <sup>-6</sup>
PC (18:1/22:5)	0.036	-0.35	2.02x10 <sup>-6</sup>
LPE (16:0)	0.029	-0.34	2.90x10 <sup>-6</sup>
PC (18:0/18:3)	0.011	-0.34	2.99x10 <sup>-6</sup>
CER (20:1)	0.019	0.34	3.23x10 <sup>-6</sup>
CER (22:0)	0.032	-0.34	3.43x10 <sup>-6</sup>
DAG (16:0/18:2)	0.002	0.33	4.63x10 <sup>-6</sup>
LCER (20:0)	0.027	-0.33	5.13x10 <sup>-6</sup>
DAG (18:2/22:4)	0.007	0.33	5.14x10 <sup>-6</sup>
HCER (16:0)	0.026	-0.33	5.29x10 <sup>-6</sup>
CE (22:4)	0.026	-0.33	5.39x10 <sup>-6</sup>
PG (20:0/22:5)	0.012	-0.33	5.69x10 <sup>-6</sup>
TAG (54:5)_FA (16:0)	0.007	0.33	5.91x10 <sup>-6</sup>
TAG (52:4)_FA (20:4)	0.006	0.32	1.06x10 <sup>-5</sup>
PI (20:0/18:1)	0.022	-0.32	1.10x10 <sup>-5</sup>
PE (18:1/20:3)	0.028	-0.32	1.18x10 <sup>-5</sup>
TAG (56:6)_FA (18:0)	0.012	0.32	1.23x10 <sup>-5</sup>
TAG (54:6)_FA (20:4)	0.008	0.32	1.25x10 <sup>-5</sup>
TAG (56:5)_FA (20:4)	0.014	0.32	1.30x10 <sup>-5</sup>
VLFC	0.011	0.32	1.44x10 <sup>-5</sup>
TAG (54:6)_FA (16:0)	0.004	0.31	1.98x10 <sup>-5</sup>
IDTG	0.003	0.31	2.54x10 <sup>-5</sup>
CER (20:0)	0.013	0.31	2.56x10 <sup>-5</sup>
DAG (18:1/22:4)	0.007	0.31	2.68x10 <sup>-5</sup>

DAG (16:0/16:0)	0.004	0.31	2.78x10 <sup>-5</sup>
V1TG	0.001	0.31	2.89x10 <sup>-5</sup>
LDPL	0.029	-0.31	2.91x10 <sup>-5</sup>
DAG (18:2/20:4)	0.006	0.31	2.93x10 <sup>-</sup>
TAG (54:4)_FA (22:4)	0.002	0.30	3.57x10 <sup>-5</sup>
TAG (56:5)_FA (18:0)	0.012	0.30	3.77x10 <sup>-5</sup>
PI (16:0/18:0)	0.027	-0.30	4.40x10 <sup>-5</sup>
TAG (58:7)_FA (18:0)	0.006	0.30	4.43x10 <sup>-5</sup>
PG (18:1/20:2)	0.027	-0.30	4.77x10 <sup>-5</sup>
CE (16:1)	0.024	-0.30	5.97x10 <sup>-5</sup>
PC (16:0/16:1)	0.023	-0.29	6.60x10 <sup>-5</sup>
SM (18:0)	0.024	-0.29	6.88x10 <sup>-5</sup>
TAG (56:4)_FA (20:4)	0.009	0.29	6.88x10 <sup>-5</sup>
DAG (18:1/22:6)	0.008	0.29	8.08x10 <sup>-5</sup>
H2A2	0.029	-0.29	8.36x10 <sup>-5</sup>
TAG (58:6)_FA (18:0)	0.003	0.29	8.82x10 <sup>-5</sup>
PI (18:0/20:5)	0.014	-0.29	9.04x10 <sup>-5</sup>
DAG (16:0/20:5)	0.004	0.29	1.05x10 <sup>-4</sup>
CER (24:1)	0.009	0.29	1.09x10 <sup>-4</sup>
Citric acid	0.020	0.28	1.10x10 <sup>-4</sup>
L3PL	0.026	-0.28	1.19x10 <sup>-4</sup>
DAG (16:0/22:5)	0.005	0.28	1.21x10 <sup>-4</sup>
TAG (54:5)_FA (22:5)	0.003	0.28	1.29x10 <sup>-4</sup>
Asparagine	0.010	-0.28	1.34x10 <sup>-4</sup>
MAG (14:0)	0.008	0.28	1.43x10 <sup>-4</sup>
PG (18:2/16:1)	0.028	0.28	1.59x10 <sup>-4</sup>
H2CH	0.031	-0.28	1.59x10 <sup>-4</sup>
FFA (20:4)	0.008	0.28	1.59x10 <sup>-4</sup>
PI (18:0/22:5)	0.030	-0.28	1.61x10 <sup>-4</sup>
V1PL	0.001	0.28	1.72x10 <sup>-4</sup>
VLPL	0.009	0.28	1.92x10 <sup>-4</sup>
LDFC	0.027	-0.27	2.18x10 <sup>-4</sup>
PG (20:0/20:4)	0.011	-0.27	2.24x10 <sup>-4</sup>
PS (20:0/16:1)	0.015	-0.27	2.42x10 <sup>-4</sup>
L1FC	0.018	-0.27	2.53x10 <sup>-4</sup>
TAG (53:5)_FA (20:4)	0.002	0.27	2.74x10 <sup>-4</sup>
DAG (18:0/18:1)	0.004	0.27	2.89x10 <sup>-4</sup>

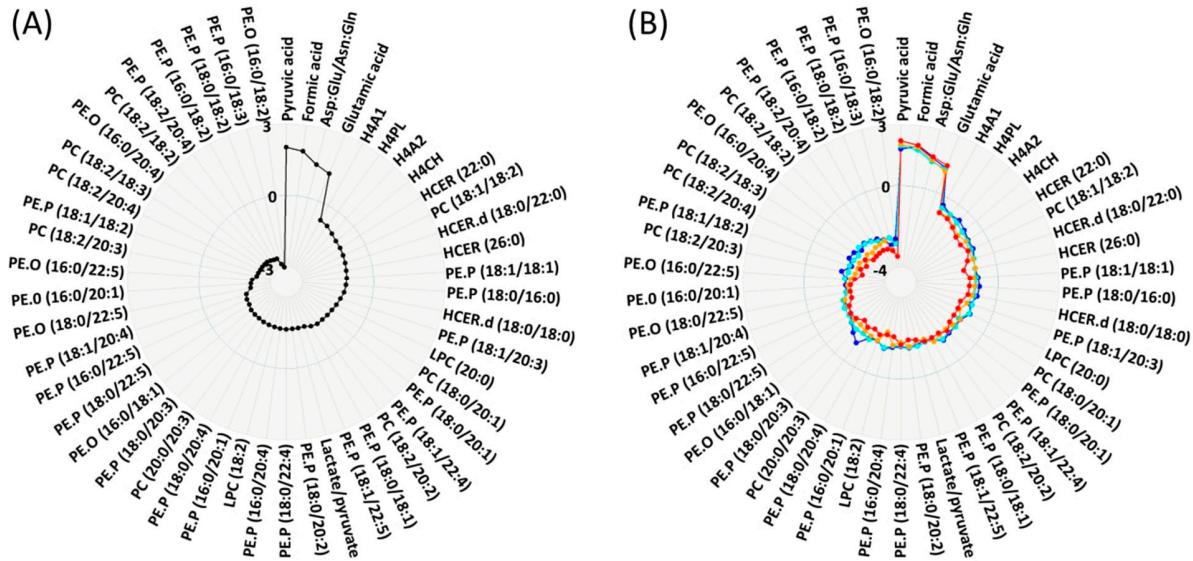
TAG (58:8)_FA (20:3)	0.030	-0.27	3.20x10 <sup>-4</sup>
Indole-3-acetic acid	0.008	-0.26	3.63x10 <sup>-4</sup>
TAG (56:6)_FA (20:4)	0.006	0.26	3.66x10 <sup>-4</sup>
L4PN	0.028	-0.26	4.34x10 <sup>-4</sup>
IDCH	0.014	0.26	4.34x10 <sup>-4</sup>
L1PL	0.013	-0.26	4.34x10 <sup>-4</sup>
L4AB	0.028	-0.26	4.34x10 <sup>-4</sup>
CER (16:0)	0.013	0.26	4.43x10 <sup>-4</sup>
MAG (18:3)	0.010	0.26	4.63x10 <sup>-4</sup>
DAG (16:0/16:1)	0.001	0.26	4.74x10 <sup>-4</sup>
CE (22:2)	0.023	-0.26	5.08x10 <sup>-4</sup>
Kynurenone	0.021	0.25	6.21x10 <sup>-4</sup>
PE (16:0/20:5)	0.020	-0.25	6.29x10 <sup>-4</sup>
CE (22:6)	0.028	-0.25	6.33x10 <sup>-4</sup>
PI (16:0_22:6)	0.025	-0.25	6.34x10 <sup>-4</sup>
TAG (58:6)_FA (22:4)	0.004	0.25	7.31x10 <sup>-4</sup>
Creatine	0.015	0.25	7.31x10 <sup>-4</sup>
TAG (51:1)_FA (18:1)	0.001	0.25	8.20x10 <sup>-4</sup>
DAG (18:1/18:1)	0.004	0.25	8.28x10 <sup>-4</sup>
DAG (16:1/20:4)	0.004	0.25	8.99x10 <sup>-4</sup>
L2CH	0.016	-0.24	1.10x10 <sup>-3</sup>
TAG (50:1)_FA (18:1)	0.006	0.24	1.21x10 <sup>-3</sup>
TAG (51:1)_FA (17:0)	0.000	0.24	1.21x10 <sup>-3</sup>
TAG (54:3)_FA (18:0)	0.005	0.24	1.36x10 <sup>-3</sup>
L3FC	0.024	-0.24	1.43x10 <sup>-3</sup>
DAG (18:0/18:2)	0.000	0.24	1.53x10 <sup>-3</sup>
DAG (16:1/18:2)	0.000	0.24	1.63X10 <sup>-3</sup>
TAG (51:1)_FA (16:0)	0.001	0.23	1.78x10 <sup>-3</sup>
TAG (56:5)_FA (22:4)	0.002	0.23	1.79x10 <sup>-3</sup>
TAG (54:8)_FA (18:3)	0.012	-0.23	1.80x10 <sup>-3</sup>
TAG (51:2)_FA (17:0)	0.001	0.23	1.82x10 <sup>-3</sup>
PE.O (18:0/18:1)	0.015	0.23	1.96x10 <sup>-3</sup>
TAG (50:2)_FA (16:0)	0.004	0.23	2.29x10 <sup>-3</sup>
LPE (18:0)	0.029	-0.23	2.35x10 <sup>-3</sup>
TAG (58:6)_FA (18:1)	0.003	0.23	2.44x10 <sup>-3</sup>
TAG (56:6)_FA (22:4)	0.000	0.23	2.53x10 <sup>-3</sup>
TAG (53:3)_FA (17:0)	0.002	0.23	2.59x10 <sup>-3</sup>

PE (18:0/20:0)	0.031	-0.23	2.69x10 <sup>-3</sup>
TAG (51:4)_FA (20:4)	0.002	0.23	2.70x10 <sup>-3</sup>
PI (18:0/22:6)	0.025	-0.23	2.71x10 <sup>-3</sup>
IDFC	0.011	0.22	2.82x10 <sup>-3</sup>
TAG (50:2)_FA (18:2)	0.000	0.22	2.86x10 <sup>-3</sup>
TAG (58:6)_FA (22:5)	0.001	0.22	2.89x10 <sup>-3</sup>
DCER (20:0)	0.013	-0.22	3.09x10 <sup>-3</sup>
TAG (49:0)_FA (16:0)	0.000	0.22	3.09x10 <sup>-3</sup>
TAG (53:1)_FA (18:1)	0.000	0.22	3.09x10 <sup>-3</sup>
TAG (49:0)_FA (17:0)	0.002	0.22	3.19x10 <sup>-3</sup>
TAG (53:2)_FA (17:0)	0.001	0.22	3.20x10 <sup>-3</sup>
DAG (16:1/18:0)	0.003	0.22	3.39x10 <sup>-3</sup>
TAG (56:4)_FA (22:4)	0.000	0.22	3.44x10 <sup>-3</sup>
TAG (56:6)_FA (18:1)	0.001	0.22	3.73x10 <sup>-3</sup>
TAG (50:1)_FA (16:0)	0.004	0.22	3.75x10 <sup>-3</sup>
PC (18:0/16:1)	0.016	-0.22	4.20x10 <sup>-3</sup>
TAG (52:2)_FA (18:0)	0.002	0.22	4.22x10 <sup>-3</sup>
TAG (54:2)_FA (18:0)	0.004	0.22	4.22x10 <sup>-3</sup>
Isoleucine	0.011	-0.22	4.39x10 <sup>-3</sup>
TAG (53:2)_FA (18:1)	0.000	0.22	4.39x10 <sup>-3</sup>
DAG (18:0/18:3)	0.000	0.22	4.39x10 <sup>-3</sup>
TAG (48:5)_FA (18:3)	0.021	-0.21	4.43x10 <sup>-3</sup>
TAG (52:2)_FA (16:1)	0.001	0.21	4.82x10 <sup>-3</sup>
LCER (18:0)	0.018	-0.21	4.82x10 <sup>-3</sup>
TAG (51:2)_FA (18:2)	0.005	0.21	4.83x10 <sup>-3</sup>
TAG (58:8)_FA (18:1)	0.005	0.21	5.48x10 <sup>-3</sup>
PS (18:0/20:0)	0.019	0.21	5.63x10 <sup>-3</sup>
TAG (52:3)_FA (18:0)	0.000	0.21	6.00x10 <sup>-3</sup>
TAG (48:1)_FA (16:1)	0.001	0.21	6.48x10 <sup>-3</sup>
PG (20:0/16:1)	0.003	-0.21	6.58x10 <sup>-3</sup>
SM (22:0)	0.005	-0.21	6.84x10 <sup>-3</sup>
TAG (53:1)_FA (17:0)	0.001	0.20	7.18x10 <sup>-3</sup>
TAG (56:8)_FA (16:0)	0.003	0.20	7.91x10 <sup>-3</sup>
FFA (22:6)	0.012	-0.20	8.06x10 <sup>-3</sup>
TAG (56:5)_FA (16:0)	0.000	0.20	8.06x10 <sup>-3</sup>
FFA (20:1)	0.011	0.20	8.41x10 <sup>-3</sup>
TAG (50:2)_FA (16:1)	0.003	0.20	9.09x10 <sup>-3</sup>

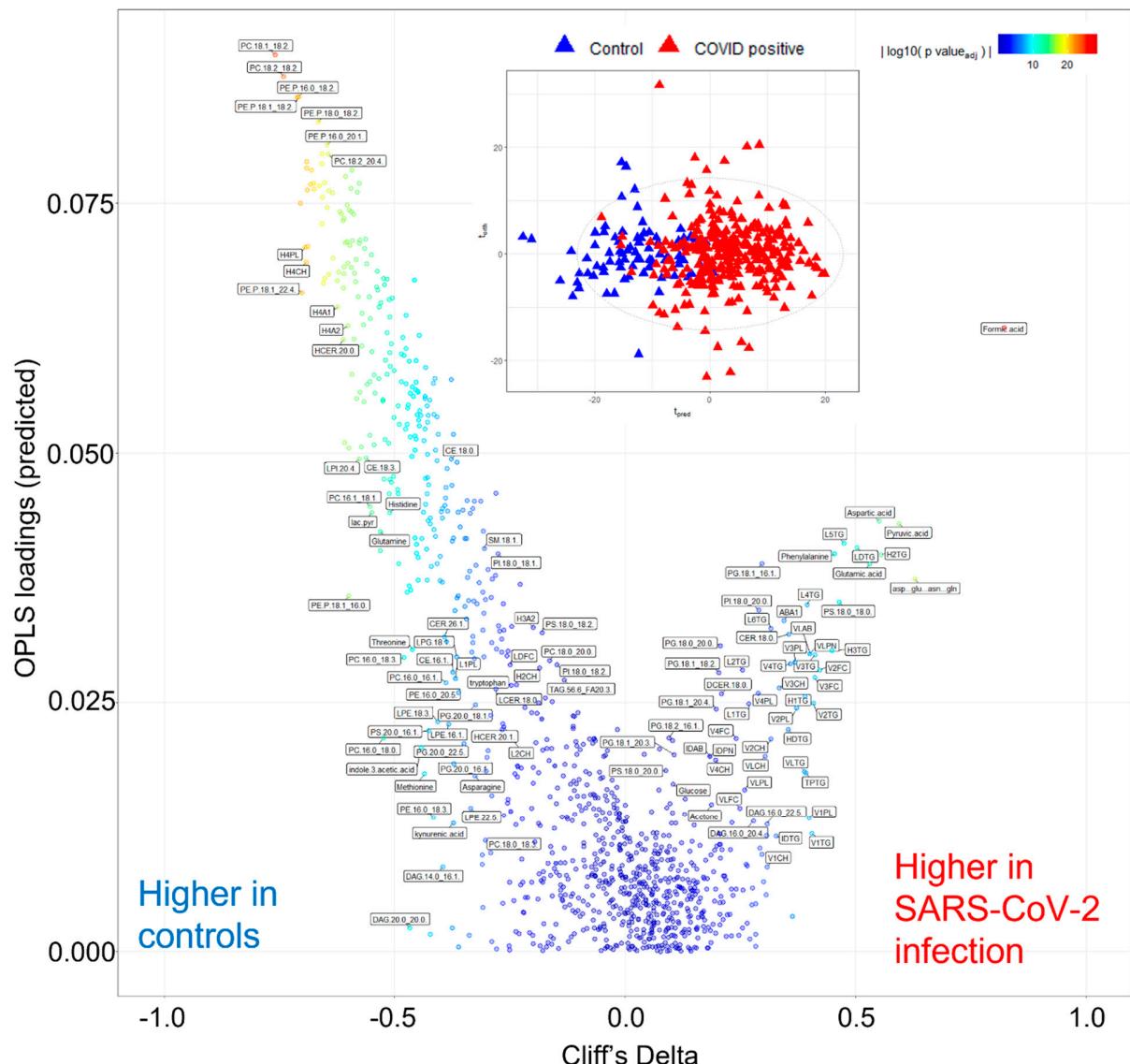
DAG (18:2/22:5)	0.004	0.20	9.37x10 <sup>-3</sup>
PG (18:2/18:2)	0.001	-0.20	9.51x10 <sup>-3</sup>
PG (16:0/18:2)	0.031	-0.20	9.56x10 <sup>-3</sup>
HCER (20:1)	0.015	-0.20	9.93x10 <sup>-3</sup>
TAG (58:7)_FA (18:1)	0.002	0.20	9.93x10 <sup>-3</sup>
TAG (56:7)_FA (20:3)	0.025	-0.20	9.98x10 <sup>-3</sup>
TAG (52:1)_FA (18:1)	0.003	0.20	1.00x10 <sup>-2</sup>
TAG (51:1)_FA (18:0)	0.001	0.20	1.03x10 <sup>-2</sup>
PE (18:0/18:2)	0.020	-0.20	1.05x10 <sup>-2</sup>
TAG (53:1)_FA (18:0)	0.000	0.20	1.05x10 <sup>-2</sup>
TAG (52:3)_FA (20:0)	0.001	0.20	1.05x10 <sup>-2</sup>
PC (18:1/20:2)	0.022	-0.19	1.07x10 <sup>-2</sup>
TAG (56:6)_FA (16:0)	0.002	0.19	1.10x10 <sup>-2</sup>
TAG (56:6)_FA (20:3)	0.025	-0.19	1.10x10 <sup>-2</sup>
TAG (56:7)_FA (20:4)	0.001	0.19	1.15x10 <sup>-2</sup>
TAG (54:3)_FA (18:1)	0.002	0.19	1.21x10 <sup>-2</sup>
DAG (16:1/22:6)	0.003	0.19	1.23x10 <sup>-2</sup>
PI (16:0/22:5)	0.022	-0.19	1.26x10 <sup>-2</sup>
TAG (56:6)_FA (22:6)	0.002	0.19	1.27x10 <sup>-2</sup>
TAG (54:4)_FA (18:0)	0.000	0.19	1.27x10 <sup>-2</sup>
PI (18:0/18:0)	0.015	-0.19	1.31x10 <sup>-2</sup>
TAG (49:1)_FA (18:1)	0.004	0.19	1.32x10 <sup>-2</sup>
PE (18:0/20:1)	0.021	-0.19	1.32x10 <sup>-2</sup>
TAG (56:2)_FA (20:0)	0.016	-0.19	1.32x10 <sup>-2</sup>
TAG (52:1)_FA (16:0)	0.003	0.19	1.36x10 <sup>-2</sup>
PE (18:1/18:1)	0.007	0.19	1.40x10 <sup>-2</sup>
TAG (53:2)_FA (18:2)	0.004	0.19	1.43x10 <sup>-2</sup>
PG (20:0/18:1)	0.007	-0.19	1.43x10 <sup>-2</sup>
TAG (50:4)_FA (20:3)	0.020	-0.19	1.54x10 <sup>-2</sup>
TAG (54:2)_FA (18:1)	0.002	0.18	1.61x10 <sup>-2</sup>
TAG (49:1)_FA (16:1)	0.002	0.18	1.64x10 <sup>-2</sup>
LDHD	0.011	0.18	1.66x10 <sup>-2</sup>
TAG (51:2)_FA (18:1)	0.003	0.18	1.73x10 <sup>-2</sup>
TAG (56:3)_FA (20:0)	0.019	-0.18	1.73x10 <sup>-2</sup>
TAG (48:0)_FA (16:0)	0.000	0.18	1.82x10 <sup>-2</sup>
TAG (48:4)_FA (18:3)	0.019	-0.18	1.82x10 <sup>-2</sup>
PE (18:0/20:4)	0.002	0.18	1.84x10 <sup>-2</sup>

TAG (52:1)_FA (18:0)	0.004	0.18	1.84x10 <sup>-2</sup>
TAG (54:3)_FA (18:2)	0.003	0.18	1.84x10 <sup>-2</sup>
TAG (51:2)_FA (16:0)	0.004	0.18	1.87x10 <sup>-2</sup>
TAG (54:6)_FA (18:3)	0.021	-0.18	1.91x10 <sup>-2</sup>
TAG (56:9)_FA (18:3)	0.013	-0.18	1.91x10 <sup>-2</sup>
TAG (58:7)_FA (22:4)	0.001	0.18	1.91x10 <sup>-2</sup>
V5FC	0.007	-0.18	1.92x10 <sup>-2</sup>
TPAB	0.004	0.18	1.93x10 <sup>-2</sup>
TBPN	0.004	0.18	1.93x10 <sup>-2</sup>
LCER.d (18:0/20:0)	0.015	-0.18	1.95x10 <sup>-2</sup>
TAG (55:3)_FA (18:1)	0.000	0.18	1.97x10 <sup>-2</sup>
TAG (51:2)_FA (16:1)	0.002	0.18	1.97x10 <sup>-2</sup>
TAG (58:9)_FA (18:1)	0.000	0.18	2.00x10 <sup>-2</sup>
TAG (56:7)_FA (22:6)	0.002	0.18	2.03x10 <sup>-2</sup>
TAG (49:1)_FA (16:0)	0.005	0.18	2.05x10 <sup>-2</sup>
TAG (51:0)_FA (16:0)	0.001	0.18	2.06x10 <sup>-2</sup>
PE (16:0/18:3)	0.008	-0.18	2.08x10 <sup>-2</sup>
V5CH	0.009	0.18	2.13x10 <sup>-2</sup>
TAG (53:4)_FA (20:4)	0.001	0.18	2.16x10 <sup>-2</sup>
TAG (53:3)_FA (18:2)	0.006	0.18	2.17x10 <sup>-2</sup>
Valine	0.006	-0.18	2.19x10 <sup>-2</sup>
LPE (22:6)	0.013	-0.18	2.25x10 <sup>-2</sup>
TAG (52:5)_FA (20:5)	0.001	0.17	2.27x10 <sup>-2</sup>
TAG (58:7)_FA (22:6)	0.001	0.17	2.29x10 <sup>-2</sup>
Lysine	0.006	-0.17	2.31x10 <sup>-2</sup>
TAG (54:5)_FA (22:4)	0.003	0.17	2.33x10 <sup>-2</sup>
TAG (52:2)_FA (18:2)	0.003	0.17	2.33x10 <sup>-2</sup>
TAG (53:2)_FA (16:0)	0.002	0.17	2.33x10 <sup>-2</sup>
TAG (47:0)_FA (16:0)	0.003	0.17	2.35x10 <sup>-2</sup>
TAG (51:0)_FA (17:0)	0.002	0.17	2.35x10 <sup>-2</sup>
DAG (16:0/18:0)	0.002	0.17	2.48x10 <sup>-2</sup>
TAG (52:5)_FA (20:4)	0.004	0.17	2.55x10 <sup>-2</sup>
TAG (52:5)_FA (20:3)	0.023	-0.17	2.56x10 <sup>-2</sup>
DAG (16:0/18:3)	0.003	0.17	2.61x10 <sup>-2</sup>
TAG (51:0)_FA (18:0)	0.002	0.17	2.61x10 <sup>-2</sup>
L2PL	0.009	-0.17	2.63x10 <sup>-2</sup>
TAG (58:8)_FA (22:6)	0.002	0.17	2.63x10 <sup>-2</sup>

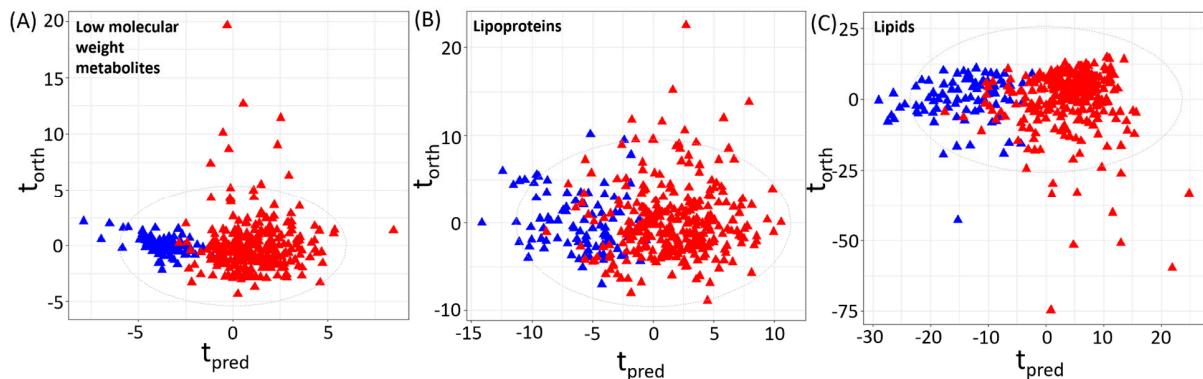
PG (18:1/20:3)	0.019	0.17	$2.65 \times 10^{-2}$
TAG (58:8)_FA (20:4)	0.001	0.17	$2.82 \times 10^{-2}$
TAG (58:7)_FA (18:2)	0.004	0.17	$2.90 \times 10^{-2}$
V5PL	0.008	0.17	$3.05 \times 10^{-2}$
TAG (52:3)_FA (18:1)	0.000	0.17	$3.19 \times 10^{-2}$
TAG (50:5)_FA (18:3)	0.019	-0.16	$3.24 \times 10^{-2}$
TAG (54:7)_FA (18:3)	0.017	-0.16	$3.28 \times 10^{-2}$
TAG (53:4)_FA (17:0)	0.004	0.16	$3.34 \times 10^{-2}$
DAG (18:1/20:1)	0.002	0.16	$3.44 \times 10^{-2}$
TAG (56:7)_FA (18:1)	0.002	0.16	$3.47 \times 10^{-2}$
TAG (52:3)_FA (16:1)	0.002	0.16	$3.60 \times 10^{-2}$
PC (18:0/22:5)	0.002	0.16	$3.63 \times 10^{-2}$
L2FC	0.010	-0.16	$3.65 \times 10^{-2}$
TAG (49:2)_FA (18:2)	0.006	0.16	$3.69 \times 10^{-2}$
TAG (53:6)_FA (20:4)	0.006	0.16	$3.76 \times 10^{-2}$
Creatinine	0.018	0.16	$3.77 \times 10^{-2}$
TAG (50:5)_FA (14:0)	0.019	-0.16	$3.83 \times 10^{-2}$
TAG (50:3)_FA (16:0)	0.003	0.16	$4.01 \times 10^{-2}$
TAG (50:1)_FA (16:1)	0.002	0.16	$4.09 \times 10^{-2}$
V1CH	0.003	0.16	$4.13 \times 10^{-2}$
TAG (58:10)_FA (20:5)	0.006	-0.16	$4.13 \times 10^{-2}$
L5CH	0.018	-0.16	$4.32 \times 10^{-2}$
TAG (49:2)_FA (16:0)	0.006	0.16	$4.38 \times 10^{-2}$
TAG (51:3)_FA (17:0)	0.003	0.16	$4.37 \times 10^{-2}$
TAG (54:4)_FA (18:1)	0.002	0.16	$4.38 \times 10^{-2}$
H1CH	0.023	-0.16	$4.40 \times 10^{-2}$
TAG (54:8)_FA (18:2)	0.009	-0.16	$4.49 \times 10^{-2}$
PE (18:1/22:6)	0.003	0.16	$4.52 \times 10^{-2}$
PE (18:1/20:1)	0.016	-0.16	$4.53 \times 10^{-2}$
DAG (18:0/22:6)	0.002	0.15	$4.61 \times 10^{-2}$
TAG (49:0)_FA (18:0)	0.000	0.15	$4.65 \times 10^{-2}$
TAG (56:7)_FA (16:0)	0.004	0.15	$4.72 \times 10^{-2}$
Ornithine	0.010	0.15	$4.73 \times 10^{-2}$
PS (18:1/18:2)	0.001	0.15	$4.75 \times 10^{-2}$



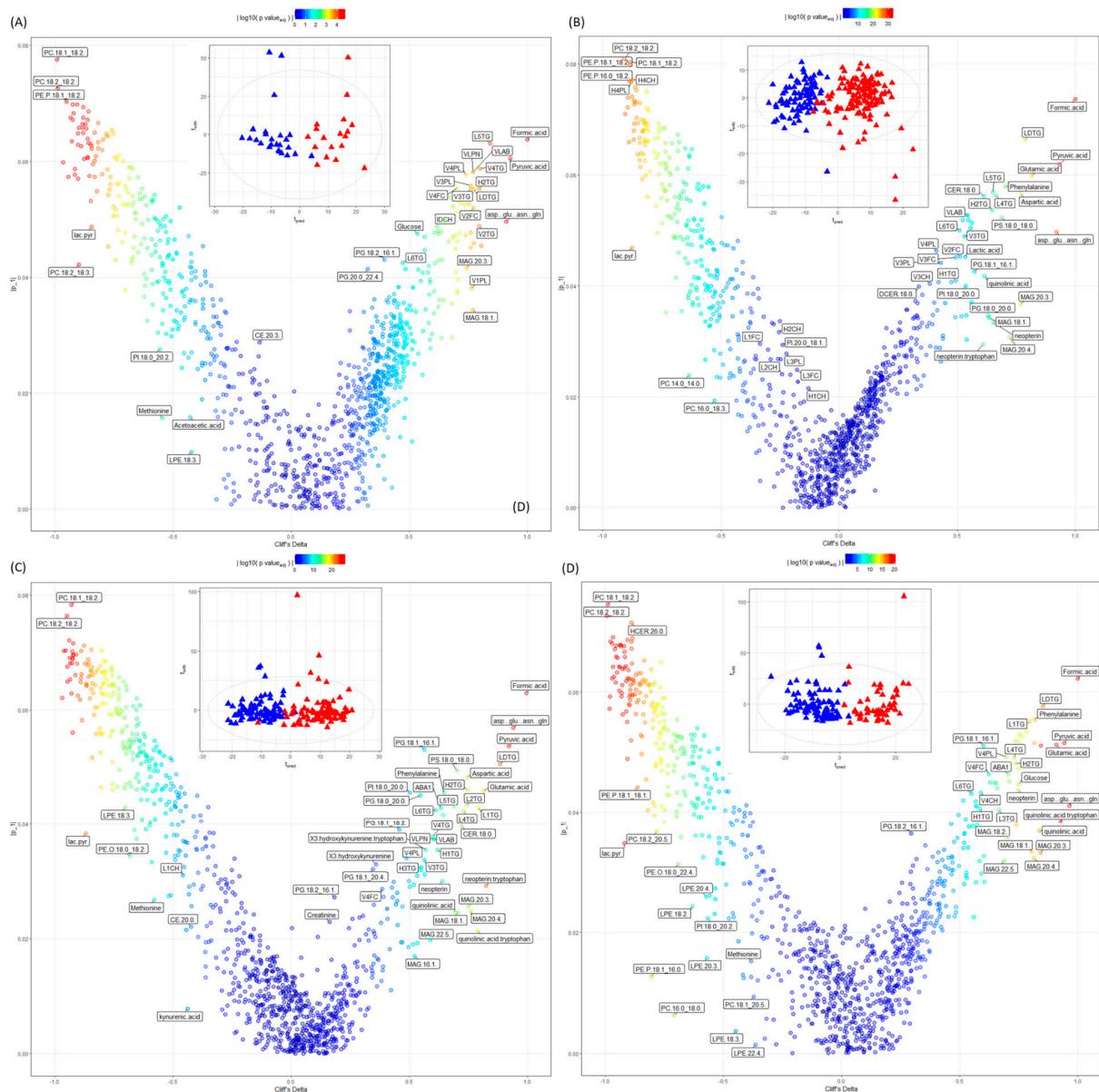
**Figure S1** - (A) Radar plot showing the  $\log_2$  fold change of the top 50 metabolites, lipoproteins and lipids with the most significant p-values for the control versus SARS-CoV-2 positive patients. (B) Radar plot showing the  $\log_2$  fold change of the top 50 metabolites, lipoproteins and lipids with the most significant p-values for the control versus SARS-CoV-2 positive model but coloured by severity class. Control vs group B severity  $\log_2$  fold change (blue), Control vs group C severity  $\log_2$  fold change (cyan), Control vs group D severity  $\log_2$  fold change (orange), Control vs group E severity  $\log_2$  fold change (red). The radar plots are ordered by fold change.



**Figure S2 - Orthogonal Partial Least Squared (O-PLS-DA) and eruption plots between controls and SARS-CoV-2 positive patients of the integrative pan-metabolic model after correction for age.** Eruption plot of the controls vs SARS-CoV-2 positive patients with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 positive patients (red triangles) ( $R^2X=0.10$ , AUROC=0.94) inset.



**Figure S3 – O-PLS-DA scores plots of the low molecular weight metabolites, lipoproteins and lipids.** (A) O-PLS-DA of control samples (blue triangles) and SARS-CoV-2 positive patients (red triangles) for the low molecular weight metabolites,  $R_2X=0.12$ , AUROC=1.00. (B) O-PLS-DA of control samples (blue triangles) and SARS-CoV-2 positive patients (red triangles) for the NMR derived lipoproteins,  $R_2X=0.26$ , AUROC=0.95. (C) O-PLS-DA of control samples (blue triangles) and SARS-CoV-2 positive patients (red triangles) for the MS derived lipids,  $R_2X=0.17$ , AUROC=0.98.



**Figure S4 - Orthogonal Partial Least Squared (O-PLS-DA) and eruption plots between controls and SARS-CoV-2 positive patients of each severity group for the four Integrated assays.** (A) Eruption plot of the controls vs group B severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group B severity patients (red triangles) ( $R_2X=0.21$ , AUROC=0.99) inset. (B) Eruption plot of the controls vs group C severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group B severity patients (red triangles) ( $R_2X=0.12$ , AUROC=0.99) inset. (C) Eruption plot of the controls vs group D severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group B severity patients (red triangles) ( $R_2X=0.20$ , AUROC=0.99) inset. (D) Eruption plot of the controls vs group E severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group B severity patients (red triangles) ( $R_2X=0.22$ , AUROC=1.00) inset.

**Table S6 - OPLS loadings, Cliff's delta and adjusted p-values of all the assays combined for the comparison of the controls and SARS-CoV-2 positive patients in severity class B.** Only the 50 most significant metabolites are shown.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.064	1.00	1.59x10 <sup>-5</sup>
PC (18:1/18:2)	0.078	-0.99	1.59x10 <sup>-5</sup>
PC (18:2/18:2)	0.073	-0.99	1.59x10 <sup>-5</sup>
PE.O (16:0/18:2)	0.060	-0.96	2.37x10 <sup>-5</sup>
PE.P (18:1/18:2)	0.070	-0.95	2.85x10 <sup>-5</sup>
Pyruvic acid	0.061	0.93	2.86x10 <sup>-5</sup>
LPC (18:2)	0.057	-0.94	2.86x10 <sup>-5</sup>
PE.P (16:0/18:2)	0.070	-0.92	2.86x10 <sup>-5</sup>
PE.P (18:0/18:2)	0.068	-0.93	2.86x10 <sup>-5</sup>
PE.P (18:0/20:1)	0.067	-0.92	2.86x10 <sup>-5</sup>
PE.P (18:2/20:4)	0.063	-0.93	2.86x10 <sup>-5</sup>
HCER.d (18:0/18:0)	0.068	-0.92	2.92x10 <sup>-5</sup>
PE.O (16:0/20:4)	0.059	-0.92	2.92x10 <sup>-5</sup>
PE.P (16:0/18:3)	0.058	-0.92	2.92x10 <sup>-5</sup>
Asp:Glu/Asn/Gln	0.050	0.91	3.09x10 <sup>-5</sup>
PC (18:2/20:2)	0.057	-0.90	3.16x10 <sup>-5</sup>
PE.O (18:0/22:5)	0.065	-0.90	3.16x10 <sup>-5</sup>
PE.P (18:1/20:1)	0.070	-0.91	3.16x10 <sup>-5</sup>
PE.P (18:0/20:4)	0.063	-0.90	3.16x10 <sup>-5</sup>
PC (18:2/18:3)	0.042	-0.90	3.25x10 <sup>-5</sup>
PC (18:2/20:4)	0.066	-0.90	3.25x10 <sup>-5</sup>
HCER (22:0)	0.063	-0.89	3.99x10 <sup>-5</sup>
PC (16:0/18:2)	0.066	-0.88	4.15x10 <sup>-5</sup>
PE.O (16:0/20:1)	0.066	-0.88	4.15x10 <sup>-5</sup>
PC (20:0/20:4)	0.065	-0.88	4.18x10 <sup>-5</sup>
PE.P (18:0/18:1)	0.068	-0.88	4.18x10 <sup>-5</sup>
PE.P (18:1/22:5)	0.069	-0.88	4.18x10 <sup>-5</sup>
PE.O (18:0/22:4)	0.060	-0.88	4.56x10 <sup>-5</sup>
H4CH	0.066	-0.87	4.99x10 <sup>-5</sup>
PC (18:0/20:1)	0.066	-0.87	5.44x10 <sup>-5</sup>
PE.O (16:0/18:1)	0.061	-0.86	5.44x10 <sup>-5</sup>
PE.P (16:0/22:5)	0.067	-0.86	5.44x10 <sup>-5</sup>
PE.P (18:0/20:4)	0.061	-0.86	5.44x10 <sup>-5</sup>

PE.P (18:1/22:4)	0.064	-0.86	5.44x10 <sup>-5</sup>
PE.P (18:0/22:5)	0.069	-0.86	5.80x10 <sup>-5</sup>
PE.P (18:2/18:2)	0.063	-0.86	5.80x10 <sup>-5</sup>
PE.O (16:0/22:5)	0.057	-0.85	6.38x10 <sup>-5</sup>
PE.O (16:0/22:6)	0.053	-0.85	6.83x10 <sup>-5</sup>
PE.P (18:0/22:4)	0.063	-0.85	6.83x10 <sup>-5</sup>
Lactate/pyruvate	0.049	-0.84	7.15x10 <sup>-5</sup>
PC (20:0/18:1)	0.062	-0.84	7.15x10 <sup>-5</sup>
PE.P (18:0/18:0)	0.063	-0.84	7.15x10 <sup>-5</sup>
L5TG	0.063	0.84	7.41x10 <sup>-5</sup>
PE.P (16:0/20:4)	0.059	-0.84	7.69x10 <sup>-5</sup>
HCER.d (18:0/22:0)	0.060	-0.83	9.13x10 <sup>-5</sup>
PE.P (16:0/20:1)	0.065	-0.83	9.13x10 <sup>-5</sup>
PE.P (18:1/16:0)	0.057	-0.83	9.13x10 <sup>-5</sup>
PE.P (18:0/16:0)	0.061	-0.83	1.01x10 <sup>-4</sup>
PC (20:0/20:3)	0.055	-0.82	1.11x10 <sup>-4</sup>
H4PL	0.063	-0.82	1.17x10 <sup>-4</sup>

**Table S7 - OPLS loadings, Cliff's delta and adjusted p-values of all the assays combined for the comparison of the controls and SARS-CoV-2 positive patients in severity class C.** Only the 50 most significant metabolites are shown.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.074	1.00	4.47x10 <sup>-34</sup>
Pyruvic acid	0.062	0.93	8.99x10 <sup>-30</sup>
Asp:Glu/Asn:Gln	0.050	0.92	4.05x10 <sup>-29</sup>
PC (18:2/18:2)	0.081	-0.91	1.28x10 <sup>-28</sup>
PE.O (16:0/20:4)	0.071	-0.91	1.66x10 <sup>-28</sup>
PE.O (16:0/18:2)	0.072	-0.89	8.50x10 <sup>-28</sup>
PE.P (18:2/20:4)	0.074	-0.89	8.50x10 <sup>-28</sup>
PE.P (16:0/22:5)	0.066	-0.89	1.31x10 <sup>-27</sup>
PE.P (16:0/18:2)	0.077	-0.89	1.49x10 <sup>-27</sup>
PE.P (18:1/18:2)	0.080	-0.89	1.49x10 <sup>-27</sup>
PC (18:1/18:2)	0.080	-0.88	1.88x10 <sup>-27</sup>
H4PL	0.077	-0.88	3.71x10 <sup>-27</sup>
Lactate/pyruvate	0.047	-0.88	4.94x10 <sup>-27</sup>
PE.P (18:0/18:2)	0.074	-0.88	4.94x10 <sup>-27</sup>

H4CH	0.077	-0.87	$5.88 \times 10^{-27}$
HCER.d (18:0/18:0)	0.073	-0.87	$1.24 \times 10^{-26}$
PE.O (16:0/18:1)	0.066	-0.87	$1.24 \times 10^{-26}$
PE.O (16:0/20:1)	0.068	-0.87	$1.24 \times 10^{-26}$
PE.P (18:0/18:1)	0.073	-0.87	$1.47 \times 10^{-26}$
PE.P (18:0/22:5)	0.069	-0.87	$1.47 \times 10^{-26}$
PE.P (18:0/20:4)	0.073	-0.87	$1.54 \times 10^{-26}$
PC (18:2/20:2)	0.076	-0.86	$2.12 \times 10^{-26}$
PE.P (16:0/20:1)	0.073	-0.86	$3.91 \times 10^{-26}$
LPC (18:2)	0.075	-0.86	$4.11 \times 10^{-26}$
PC (18:2/20:3)	0.070	-0.86	$4.22 \times 10^{-26}$
PE.P (16:0/20:4)	0.069	-0.85	$7.46 \times 10^{-26}$
PE.O (18:0/22:5)	0.064	-0.85	$7.52 \times 10^{-26}$
PC (20:0/20:3)	0.067	-0.84	$2.15 \times 10^{-25}$
PE.P (18:0/22:4)	0.063	-0.84	$2.15 \times 10^{-25}$
PE.P (18:2/18:2)	0.073	-0.84	$2.54 \times 10^{-25}$
PC (18:0/20:1)	0.066	-0.84	$2.68 \times 10^{-25}$
PE.O (18:0/22:4)	0.061	-0.84	$2.72 \times 10^{-25}$
PE.P (18:0/20:1)	0.070	-0.84	$4.69 \times 10^{-25}$
LPI (18:1)	0.064	-0.84	$4.87 \times 10^{-25}$
LPI (18:2)	0.059	-0.83	$6.88 \times 10^{-25}$
PE.P (18:1/22:5)	0.068	-0.83	$9.11 \times 10^{-25}$
HCER (26:0)	0.068	-0.83	$1.15 \times 10^{-24}$
PE.P (18:0/20:3)	0.068	-0.83	$1.20 \times 10^{-24}$
PE.P (16:0/18:3)	0.064	-0.83	$1.36 \times 10^{-24}$
PE.P (18:0/20:4)	0.069	-0.83	$1.97 \times 10^{-24}$
PE.O (16:0/22:5)	0.066	-0.82	$2.14 \times 10^{-24}$
PE.P (18:0/20:2)	0.070	-0.82	$2.14 \times 10^{-24}$
HCER (22:0)	0.066	-0.82	$2.54 \times 10^{-24}$
HCER.d (18:0/22:0)	0.065	-0.82	$2.54 \times 10^{-24}$
LPI (20:3)	0.063	-0.82	$2.65 \times 10^{-24}$
PE.P (18:0/18:0)	0.070	-0.82	$2.65 \times 10^{-24}$
PC (20:0/18:1)	0.060	-0.82	$4.46 \times 10^{-24}$
PE.P (18:1/18:1)	0.069	-0.82	$4.56 \times 10^{-24}$
PC (18:2/20:4)	0.068	-0.82	$5.19 \times 10^{-24}$
LPC (20:0)	0.075	-0.82	$6.05 \times 10^{-24}$

**Table S8 - OPLS loadings, Cliff's delta and adjusted p-values of all the assays combined for the comparison of the controls and SARS-CoV-2 positive patients in severity class D.** Only the 50 most significant metabolites are shown.

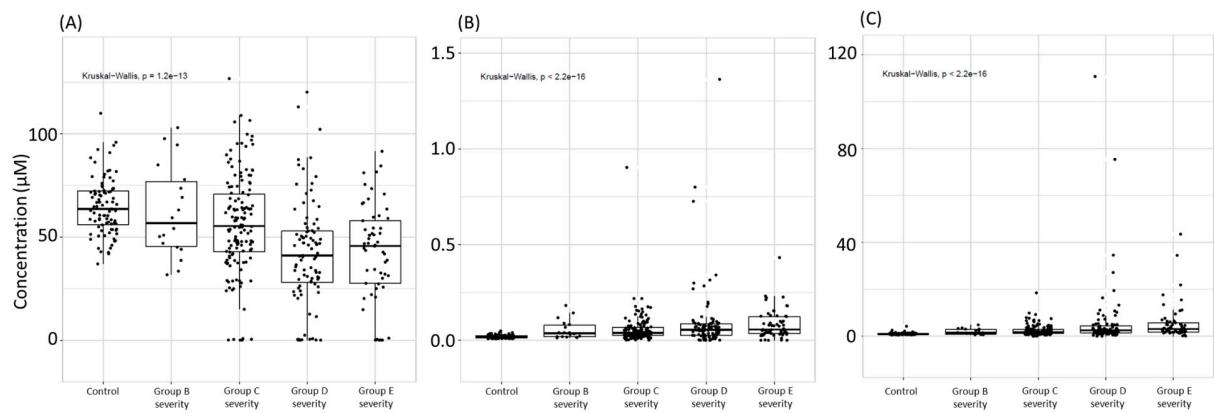
Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.063	1.00	5.46x10 <sup>-28</sup>
PE.O (16:0/20:4)	0.062	-0.97	1.14x10 <sup>-26</sup>
H4CH	0.070	-0.96	1.89x10 <sup>-26</sup>
H4PL	0.070	-0.95	4.13x10 <sup>-26</sup>
PC (18:2/18:2)	0.076	-0.95	5.90x10 <sup>-26</sup>
PE.P (18:1/20:4)	0.068	-0.95	5.90x10 <sup>-26</sup>
PE.P (18:2/20:4)	0.065	-0.95	5.90x10 <sup>-26</sup>
Asp:Glu/Asn:Gln	0.057	0.94	1.04x10 <sup>-25</sup>
PE.P (18:1/18:2)	0.072	-0.94	1.04x10 <sup>-25</sup>
LPC (18:2)	0.069	-0.94	1.27x10 <sup>-25</sup>
PE.P (16:0/20:4)	0.065	-0.94	1.41x10 <sup>-25</sup>
PE.P (16:0/22:5)	0.068	-0.94	1.41x10 <sup>-25</sup>
PC (18:1/18:2)	0.078	-0.93	2.41x10 <sup>-25</sup>
PE.O (16:0/18:2)	0.063	-0.93	2.53x10 <sup>-25</sup>
H4A2	0.068	-0.93	3.02x10 <sup>-25</sup>
PC (18:2/20:2)	0.069	-0.93	3.02x10 <sup>-25</sup>
H4A1	0.070	-0.93	3.12x10 <sup>-25</sup>
PE.P (18:0/18:2)	0.069	-0.92	3.65x10 <sup>-25</sup>
PC (18:2/18:3)	0.056	-0.92	3.91x10 <sup>-25</sup>
PE.P (16:0/18:2)	0.072	-0.92	4.20x10 <sup>-25</sup>
PE.P (18:0/22:5)	0.070	-0.92	4.26x10 <sup>-25</sup>
Pyruvic acid	0.054	0.92	4.96x10 <sup>-25</sup>
PC (18:2/20:3)	0.070	-0.92	4.96x10 <sup>-25</sup>
PE.P (18:2/18:2)	0.066	-0.91	1.08x10 <sup>-24</sup>
LPC (20:0)	0.070	-0.91	1.59x10 <sup>-24</sup>
PE.P (18:1/22:5)	0.068	-0.91	2.88x10 <sup>-24</sup>
PE.O (16:0/22:5)	0.063	-0.90	4.36x10 <sup>-24</sup>
PE.O (16:0/18:1)	0.062	-0.89	1.31x10 <sup>-23</sup>
PE.O (18:0/22:5)	0.065	-0.89	1.55x10 <sup>-23</sup>
PE.P (18:0/20:4)	0.066	-0.89	1.69x10 <sup>-23</sup>
PE.P (18:0/18:1)	0.070	-0.89	1.73x10 <sup>-23</sup>
PC (20:0/20:3)	0.066	-0.89	1.89x10 <sup>-23</sup>
Neopterin/tryptophan	0.029	0.82	1.93x10 <sup>-23</sup>

PE.P (16:0/18:3)	0.058	-0.89	2.32x10 <sup>-23</sup>
LDTG	0.051	0.88	2.94x10 <sup>-23</sup>
PC (18:2/20:4)	0.071	-0.88	7.29x10 <sup>-23</sup>
HCER.d (18:0/18:0)	0.068	-0.87	9.77x10 <sup>-23</sup>
Lactate/pyruvate	0.038	-0.87	1.70x10 <sup>-22</sup>
PE.O (16:0/20:1)	0.065	-0.87	1.91x10 <sup>-22</sup>
PI (20:0/18:2)	0.064	-0.87	2.09x10 <sup>-22</sup>
PE.P (18:1/22:4)	0.064	-0.87	2.23x10 <sup>-22</sup>
LPC (20:3)	0.068	-0.86	3.99x10 <sup>-22</sup>
PE.P (16:0/20:1)	0.069	-0.86	3.99x10 <sup>-22</sup>
PE.P (18:0/16:0)	0.068	-0.86	5.19x10 <sup>-22</sup>
PE.P (18:0/22:4)	0.066	-0.86	5.53x10 <sup>-22</sup>
PI (18:1/18:1)	0.062	-0.85	1.04x10 <sup>-21</sup>
TPA2	0.066	-0.85	1.05x10 <sup>-21</sup>
H4FC	0.067	-0.85	1.41x10 <sup>-21</sup>
PE.P (18:0/20:3)	0.064	-0.85	1.55x10 <sup>-21</sup>
SM (24:0)	0.067	-0.85	1.55x10 <sup>-21</sup>

**Table S9 - OPLS loadings, Cliff's delta and adjusted p-values of all the assays combined for the comparison of the controls and SARS-CoV-2 positive patients in severity class E.** Only the 50 most significant metabolites are shown.

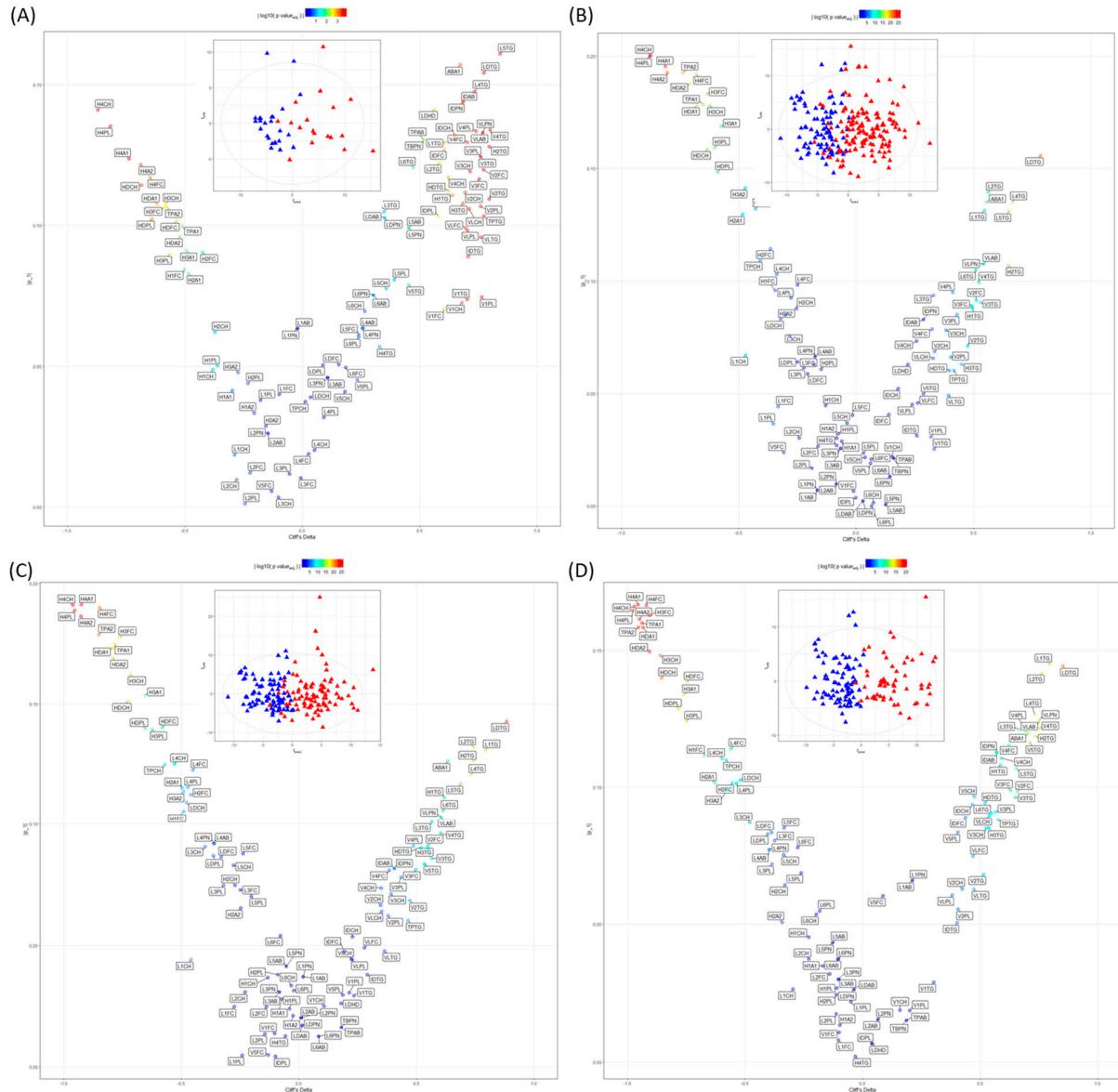
Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.064	1.00	2.50x10 <sup>-21</sup>
PC (18:1/18:2)	0.075	-0.99	2.67x10 <sup>-21</sup>
PC (18:2/18:2)	0.073	-0.99	2.67x10 <sup>-21</sup>
PE.O (16:0/20:4)	0.057	-0.98	3.92x10 <sup>-21</sup>
PE.P (18:2/20:4)	0.059	-0.98	3.92x10 <sup>-21</sup>
H4PL	0.068	-0.98	4.16x10 <sup>-21</sup>
H4CH	0.068	-0.97	7.25x10 <sup>-21</sup>
PC (20:0/20:3)	0.062	-0.97	7.25x10 <sup>-21</sup>
PC (18:2/20:3)	0.064	-0.97	8.31x10 <sup>-21</sup>
PC (18:2/20:4)	0.069	-0.97	8.31x10 <sup>-21</sup>
Asp:Glu/Asn:Gln	0.041	0.97	8.65x10 <sup>-21</sup>
PE.P (18:1/20:4)	0.063	-0.96	2.50x10 <sup>-21</sup>
PE.P (16:0/18:2)	0.067	-0.96	1.26x10 <sup>-20</sup>
LPC (18:2)	0.062	-0.96	1.73x10 <sup>-20</sup>

TPA2	0.066	-0.95	1.81x10 <sup>-20</sup>
H4A1	0.068	-0.95	1.81x10 <sup>-20</sup>
PE.P (16:0/22:5)	0.063	-0.95	2.28x10 <sup>-20</sup>
H4A2	0.066	0.95	2.62x10 <sup>-20</sup>
Neopterin/tryptophan	0.051	0.84	4.83x10 <sup>-20</sup>
Pyruvic acid	0.052	0.94	4.93x10 <sup>-20</sup>
PE.P (18:1/18:2)	0.064	-0.94	4.93x10 <sup>-20</sup>
PE.P (16:0/20:4)	0.060	-0.94	5.06x10 <sup>-20</sup>
TPA1	0.068	-0.94	5.63x10 <sup>-20</sup>
PE.P (18:0/22:5)	0.064	-0.94	5.63x10 <sup>-20</sup>
PC (18:0/20:1)	0.068	-0.94	7.36x10 <sup>-20</sup>
HDA1	0.068	-0.93	8.26x10 <sup>-20</sup>
PE.P (18:2/18:2)	0.059	-0.93	1.00x10 <sup>-20</sup>
PE.P (18:0/20:3)	0.060	-0.93	1.17x10 <sup>-19</sup>
PE.P (18:0/20:4)	0.061	-0.93	1.17x10 <sup>-19</sup>
Quinolinic acid/tryptophan	0.038	0.93	1.19x10 <sup>-19</sup>
PE.P (18:0/18:2)	0.064	-0.93	1.19x10 <sup>-19</sup>
PE.O (16:0/22:5)	0.056	-0.93	1.29x10 <sup>-19</sup>
H4FC	0.068	-0.92	2.37x10 <sup>-19</sup>
PC (18:2/22:5)	0.064	-0.92	2.60x10 <sup>-19</sup>
HCER (22:0)	0.067	-0.92	2.62x10 <sup>-19</sup>
Lactate/pyruvate	0.035	-0.92	2.75x10 <sup>-19</sup>
HCER.d (18:0/22:0)	0.066	-0.92	3.11x10 <sup>-19</sup>
LPC (20:0)	0.064	-0.91	3.80x10 <sup>-19</sup>
PE.P (18:0/22:4)	0.062	-0.91	4.47x10 <sup>-19</sup>
PI (20:0/18:2)	0.057	-0.91	4.70x10 <sup>-19</sup>
Glutamic acid	0.051	0.91	5.54x10 <sup>-19</sup>
PE.O (16:0/18:2)	0.055	-0.91	6.52x10 <sup>-19</sup>
PE.P (16:0/18:3)	0.049	-0.91	6.86x10 <sup>-19</sup>
H3FC	0.069	-0.91	7.32x10 <sup>-19</sup>
HDA2	0.062	-0.91	7.33x10 <sup>-19</sup>
PE.P (16:0/20:1)	0.064	-0.90	9.68x10 <sup>-19</sup>
PC (18:2/22:6)	0.062	-0.90	1.06x10 <sup>-18</sup>
PC (18:2/20:2)	0.058	-0.90	1.12x10 <sup>-18</sup>
PE.O (16:0/20:1)	0.059	-0.90	1.37x10 <sup>-18</sup>
PE.P (18:1/22:4)	0.058	-0.90	1.50x10 <sup>-18</sup>

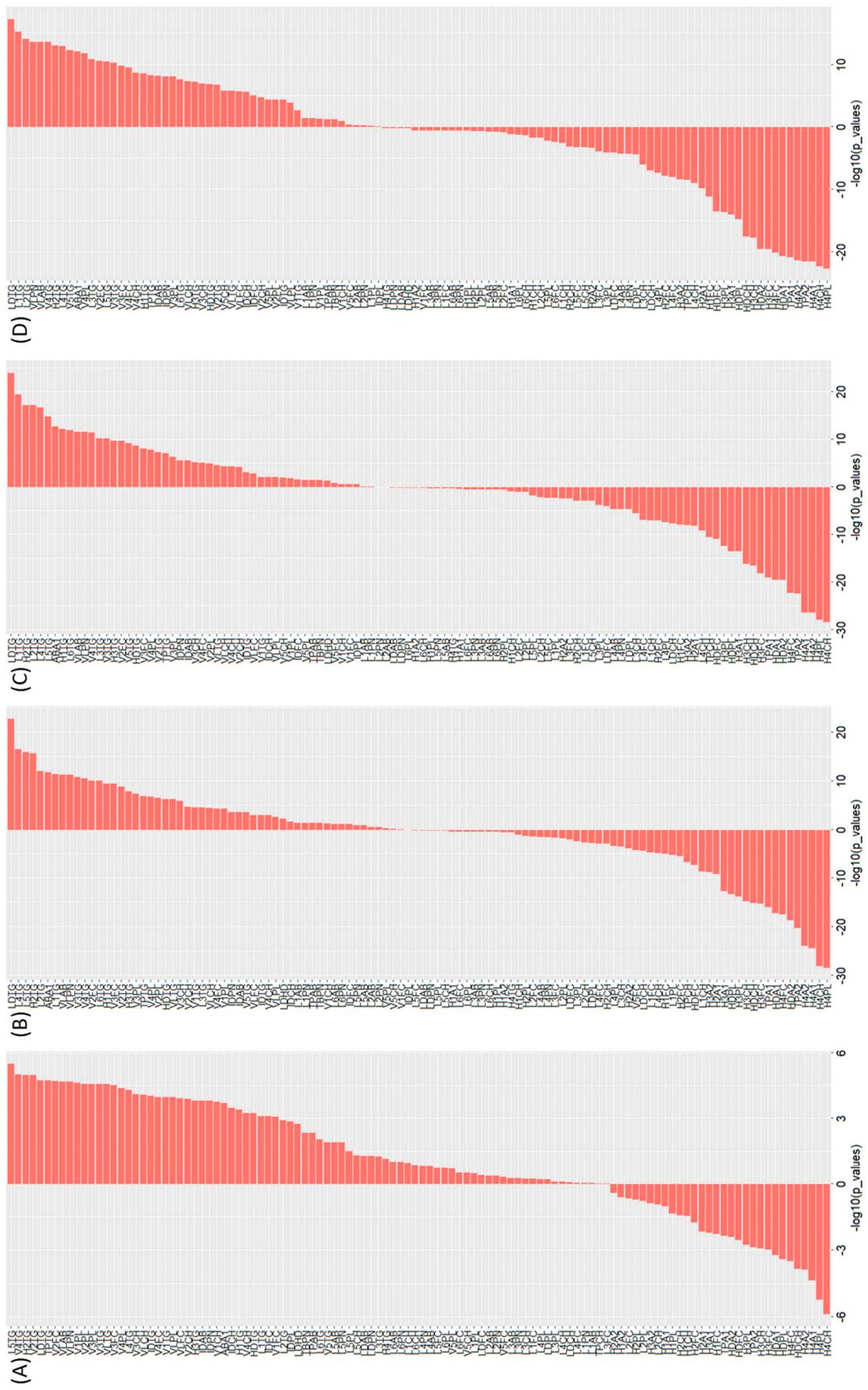


**Figure S5 – Concentrations of metabolites for the controls and each severity class. (A) tryptophan, (B) neopterin and (C) quinolinic acid.**

## Section 4 - Severity group modeling



**Figure S6 - Orthogonal Partial Least Squared (O-PLS-DA) and eruption plots between controls and SARS-CoV-2 positive patients of each severity group for the lipoproteins.** (A) Eruption plot of the controls vs group B severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group B severity patients (red triangles) ( $R_2X=0.39$ , AUROC=0.91) inset. (B) Eruption plot of the controls vs group C severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group C severity patients (red triangles) ( $R_2X=0.27$ , AUROC=0.95) inset. (C) Eruption plot of the controls vs group D severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group D severity patients (red triangles) ( $R_2X=0.29$ , AUROC=0.97) inset. (D) Eruption plot of the controls vs group E severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group E severity patients (red triangles) ( $R_2X=0.36$ , AUROC=1.00) inset. Eruption plots are formed from the Cliff's delta (abscissa) and O-PLS-DA loadings (ordinate). Coloured by significance.



**Figure S6 -  $-\log_{10}$  (p-values) between controls and SARS-CoV-2 positive patients of each severity group for the lipoproteins.** (A) Controls vs group B severity. (B) Controls vs Group C severity. (D) Controls vs Group D severity. (E) Controls vs Group E severity.

**Table S10 - OPLS loadings, Cliff's delta and adjusted p-values of the lipoproteins for the comparison of the controls and SARS-CoV-2 positive patients in severity class B.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
H4CH	0.141	-0.87	$1.57 \times 10^{-4}$
L5TG	0.161	0.84	$1.73 \times 10^{-4}$
TPTG	0.103	0.77	$1.79 \times 10^{-4}$
VLPN	0.133	0.77	$1.79 \times 10^{-4}$
VLTG	0.096	0.76	$1.79 \times 10^{-4}$
LDTG	0.154	0.77	$1.79 \times 10^{-4}$
VLAB	0.133	0.77	$1.79 \times 10^{-4}$
V2TG	0.109	0.80	$1.79 \times 10^{-4}$
V3TG	0.124	0.76	$1.79 \times 10^{-4}$
V4TG	0.134	0.80	$1.79 \times 10^{-4}$
V2FC	0.120	0.77	$1.79 \times 10^{-4}$
V1PL	0.075	0.76	$1.79 \times 10^{-4}$
V2PL	0.107	0.76	$1.79 \times 10^{-4}$
V3PL	0.124	0.76	$1.79 \times 10^{-4}$
H2TG	0.128	0.80	$1.79 \times 10^{-4}$
H4PL	0.135	-0.82	$1.79 \times 10^{-4}$
V3FC	0.117	0.75	$1.97 \times 10^{-4}$
V4PL	0.133	0.74	$2.58 \times 10^{-4}$
H4A1	0.124	-0.74	$2.58 \times 10^{-4}$
L4TG	0.148	0.73	$2.72 \times 10^{-4}$
V3CH	0.119	0.72	$3.93 \times 10^{-4}$
VLCH	0.106	0.71	$4.17 \times 10^{-4}$
IDTG	0.089	0.71	$4.32 \times 10^{-4}$
VLPL	0.098	0.70	$4.32 \times 10^{-4}$
V1TG	0.074	0.70	$4.32 \times 10^{-4}$
V4FC	0.130	0.70	$4.32 \times 10^{-4}$
VLFC	0.099	0.70	$4.84 \times 10^{-4}$
V2CH	0.112	0.69	$4.91 \times 10^{-4}$
HDCH	0.114	-0.68	$5.09 \times 10^{-4}$

IDPN	0.114	0.68	5.09x10 <sup>-4</sup>
IDAB	0.144	0.68	5.09x10 <sup>-4</sup>
H3TG	0.108	0.68	5.09x10 <sup>-4</sup>
H4A2	0.122	-0.69	5.09x10 <sup>-4</sup>
V1CH	0.073	0.68	5.45x10 <sup>-4</sup>
ABA1	0.157	0.67	6.08x10 <sup>-4</sup>
IDCH	0.132	0.65	9.85x10 <sup>-4</sup>
H4FC	0.117	-0.65	9.85x10 <sup>-4</sup>
HDPL	0.102	-0.64	1.13x10 <sup>-3</sup>
H1TG	0.111	0.64	1.13x10 <sup>-3</sup>
HDTG	0.112	0.62	1.55x10 <sup>-3</sup>
V4CH	0.117	0.62	1.55x10 <sup>-3</sup>
HDA1	0.108	-0.62	1.66x10 <sup>-3</sup>
L1TG	0.130	0.61	1.94x10 <sup>-3</sup>
IDFC	0.127	0.61	1.98x10 <sup>-3</sup>
V1FC	0.070	0.60	2.02x10 <sup>-3</sup>
H3FC	0.106	-0.59	2.69x10 <sup>-3</sup>
L2TG	0.122	0.58	2.87x10 <sup>-3</sup>
H3CH	0.108	-0.58	2.94x10 <sup>-3</sup>
TPA2	0.107	-0.58	3.07x10 <sup>-3</sup>
IDPL	0.103	0.58	3.07x10 <sup>-3</sup>
LDHD	0.141	0.56	3.81x10 <sup>-3</sup>
H3PL	0.090	-0.56	3.81x10 <sup>-3</sup>
HDFC	0.102	-0.54	6.36x10 <sup>-3</sup>
HDA2	0.096	-0.52	8.24x10 <sup>-3</sup>
TPAB	0.131	0.52	8.59x10 <sup>-3</sup>
TBPN	0.131	0.52	8.59x10 <sup>-3</sup>
TPA1	0.100	-0.51	9.12x10 <sup>-3</sup>
H1FC	0.084	-0.50	1.04x10 <sup>-2</sup>
H3A1	0.091	-0.49	1.19x10 <sup>-2</sup>
H2A1	0.083	-0.48	1.36x10 <sup>-2</sup>
L6TG	0.121	0.47	1.60x10 <sup>-2</sup>
L5PN	0.099	0.45	2.11x10 <sup>-2</sup>
V5TG	0.079	0.45	2.11x10 <sup>-2</sup>
L5AB	0.099	0.45	2.11x10 <sup>-2</sup>
H2FC	0.090	-0.42	3.22x10 <sup>-2</sup>

**Table S11 - OPLS loadings, Cliff's delta and adjusted p-values of the lipoproteins for the comparison of the controls and SARS-CoV-2 positive patients in severity class C.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
H4CH	0.200	-0.87	4.77x10 <sup>-27</sup>
H4PL	0.195	-0.88	4.77x10 <sup>-27</sup>
H4A1	0.195	-0.81	1.72x10 <sup>-23</sup>
H4A2	0.193	-0.81	3.19x10 <sup>-23</sup>
LDTG	0.156	0.79	3.29x10 <sup>-23</sup>
TPA2	0.193	-0.74	1.24x10 <sup>-19</sup>
HDA2	0.188	-0.71	3.22x10 <sup>-18</sup>
H4FC	0.192	-0.68	4.81x10 <sup>-17</sup>
HDA1	0.178	-0.68	8.30x10 <sup>-17</sup>
L4TG	0.135	0.66	3.07x10 <sup>-16</sup>
TPA1	0.178	-0.65	1.05x10 <sup>-15</sup>
L5TG	0.131	0.65	1.05x10 <sup>-15</sup>
H2TG	0.107	0.65	1.52x10 <sup>-15</sup>
H3FC	0.183	-0.64	4.79x10 <sup>-15</sup>
HDCH	0.159	-0.63	5.62x10 <sup>-15</sup>
H3CH	0.178	-0.62	1.44x10 <sup>-14</sup>
H3PL	0.159	-0.60	1.15x10 <sup>-13</sup>
HDPL	0.148	-0.59	3.56x10 <sup>-13</sup>
H3A1	0.169	-0.58	1.36x10 <sup>-12</sup>
L2TG	0.139	0.56	4.71x10 <sup>-12</sup>
ABA1	0.135	0.56	7.45x10 <sup>-12</sup>
L1TG	0.131	0.55	1.83x10 <sup>-11</sup>
VLPN	0.108	0.54	2.31x10 <sup>-11</sup>
VLAB	0.108	0.54	2.31x10 <sup>-11</sup>
V3TG	0.090	0.53	6.74x10 <sup>-11</sup>
V4TG	0.099	0.52	1.29x10 <sup>-10</sup>
V2FC	0.091	0.51	3.20x10 <sup>-10</sup>
L6TG	0.105	0.51	3.64x10 <sup>-10</sup>
V3FC	0.089	0.49	1.25x10 <sup>-9</sup>
H1TG	0.088	0.49	1.25x10 <sup>-9</sup>
H2A1	0.130	-0.48	2.77x10 <sup>-9</sup>
V2TG	0.071	0.47	5.31x10 <sup>-9</sup>
H3A2	0.141	-0.47	6.35x10 <sup>-9</sup>

L1CH	0.067	-0.47	6.99x10 <sup>-9</sup>
H3TG	0.063	0.45	3.72x10 <sup>-8</sup>
V3PL	0.085	0.43	1.29x10 <sup>-7</sup>
HDFC	0.132	-0.43	1.42x10 <sup>-7</sup>
TPTG	0.060	0.42	3.60x10 <sup>-7</sup>
V4PL	0.095	0.41	4.48x10 <sup>-7</sup>
TPCH	0.109	-0.41	5.74x10 <sup>-7</sup>
V2PL	0.066	0.40	8.18x10 <sup>-7</sup>
HDTG	0.060	0.40	1.26x10 <sup>-6</sup>
VLTG	0.049	0.39	1.39x10 <sup>-6</sup>
V3CH	0.078	0.38	2.77x10 <sup>-6</sup>
H2FC	0.114	-0.36	8.54x10 <sup>-6</sup>
L1PL	0.037	-0.35	1.71x10 <sup>-5</sup>
H1FC	0.096	-0.34	2.75x10 <sup>-5</sup>
L4CH	0.103	-0.34	3.61x10 <sup>-5</sup>
V2CH	0.069	0.33	4.82x10 <sup>-5</sup>
L1FC	0.044	-0.33	4.85x10 <sup>-5</sup>
V1TG	0.025	0.33	5.22x10 <sup>-5</sup>
L3TG	0.094	0.33	6.05x10 <sup>-5</sup>
VLCH	0.066	0.33	6.44x10 <sup>-5</sup>
V4FC	0.079	0.32	8.42x10 <sup>-5</sup>
LDCH	0.083	-0.32	9.33x10 <sup>-5</sup>
V1PL	0.031	0.32	9.91x10 <sup>-5</sup>
V5FC	0.024	-0.31	1.24x10 <sup>-4</sup>
H2A2	0.085	-0.30	2.41x10 <sup>-4</sup>
IDPN	0.083	0.29	5.12x10 <sup>-4</sup>
IDAB	0.083	0.29	5.13x10 <sup>-4</sup>
V5TG	0.050	0.29	5.13x10 <sup>-4</sup>
L3CH	0.076	-0.28	5.40x10 <sup>-4</sup>
L4PL	0.093	-0.28	7.32x10 <sup>-4</sup>
VLFC	0.046	0.26	1.56x10 <sup>-3</sup>
IDTG	0.031	0.26	1.67x10 <sup>-3</sup>
V4CH	0.073	0.26	1.85x10 <sup>-3</sup>
H2CH	0.088	-0.25	2.02x10 <sup>-3</sup>
L4FC	0.098	-0.25	2.38x10 <sup>-3</sup>
LDPL	0.064	-0.25	2.58x10 <sup>-3</sup>
L2CH	0.030	-0.24	3.33x10 <sup>-3</sup>

VLPL	0.045	0.24	3.70x10 <sup>-3</sup>
L3PL	0.062	-0.22	7.06x10 <sup>-3</sup>
LDHD	0.063	0.22	7.59x10 <sup>-3</sup>
LDFC	0.059	-0.21	1.30x10 <sup>-2</sup>
L2PL	0.017	-0.19	2.40x10 <sup>-2</sup>
L3FC	0.063	-0.18	3.35x10 <sup>-2</sup>
IDCH	0.052	0.18	3.37x10 <sup>-2</sup>
L4PN	0.067	-0.17	3.75x10 <sup>-2</sup>
L4AB	0.066	-0.17	3.75x10 <sup>-2</sup>
L1PN	0.007	-0.17	4.79x10 <sup>-2</sup>
L1AB	0.007	-0.17	4.79x10 <sup>-2</sup>

**Table S12 - OPLS loadings, Cliff's delta and adjusted p-values of the lipoproteins for the comparison of the controls and SARS-CoV-2 positive patients in severity class D.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
H4CH	0.191	-0.96	6.16x10 <sup>-27</sup>
H4PL	0.189	-0.95	8.95x10 <sup>-27</sup>
H4A1	0.191	-0.93	1.44x10 <sup>-25</sup>
H4A2	0.186	-0.93	1.44x10 <sup>-25</sup>
LDTG	0.143	0.88	2.23x10 <sup>-23</sup>
TPA2	0.179	-0.85	8.89x10 <sup>-22</sup>
H4FC	0.190	-0.85	1.05x10 <sup>-21</sup>
HDA1	0.173	-0.79	3.96x10 <sup>-19</sup>
HDA2	0.169	-0.79	3.96x10 <sup>-19</sup>
L1TG	0.131	0.79	3.96x10 <sup>-19</sup>
TPA1	0.175	-0.78	1.02x10 <sup>-18</sup>
H3FC	0.178	-0.76	7.79x10 <sup>-18</sup>
H2TG	0.126	0.74	4.40x10 <sup>-17</sup>
L2TG	0.132	0.74	4.46x10 <sup>-17</sup>
L4TG	0.121	0.73	1.53x10 <sup>-16</sup>
HDCH	0.151	-0.73	2.40x10 <sup>-16</sup>
H3CH	0.162	-0.72	5.26x10 <sup>-16</sup>
L5TG	0.117	0.69	7.42x10 <sup>-15</sup>
H3A1	0.154	-0.66	1.83x10 <sup>-13</sup>

HDPL	0.140	-0.65	1.94x10 <sup>-13</sup>
ABA1	0.126	0.63	9.32x10 <sup>-13</sup>
H3PL	0.139	-0.62	2.01x10 <sup>-12</sup>
H1TG	0.111	0.62	3.20x10 <sup>-12</sup>
L6TG	0.106	0.61	5.25x10 <sup>-12</sup>
VLPN	0.103	0.60	1.01x10 <sup>-11</sup>
VLAB	0.103	0.60	1.01x10 <sup>-11</sup>
V4TG	0.096	0.60	1.19x10 <sup>-11</sup>
HDFC	0.140	-0.58	5.71x10 <sup>-11</sup>
TPCH	0.125	-0.57	1.28x10 <sup>-10</sup>
L3TG	0.101	0.57	1.92x10 <sup>-10</sup>
V3TG	0.086	0.56	2.03x10 <sup>-10</sup>
H3TG	0.090	0.55	5.73x10 <sup>-10</sup>
V2FC	0.091	0.55	6.11x10 <sup>-10</sup>
V5TG	0.083	0.53	1.72x10 <sup>-9</sup>
L4CH	0.125	-0.53	2.58x10 <sup>-9</sup>
HDTG	0.090	0.52	5.28x10 <sup>-9</sup>
V3FC	0.081	0.50	2.15x10 <sup>-8</sup>
H2A1	0.115	-0.50	2.20x10 <sup>-8</sup>
H3A2	0.114	-0.49	3.34x10 <sup>-8</sup>
H1FC	0.106	-0.49	3.53x10 <sup>-8</sup>
V4PL	0.091	0.49	4.21x10 <sup>-8</sup>
LDCH	0.109	-0.48	7.00x10 <sup>-8</sup>
L4PL	0.116	-0.47	1.09x10 <sup>-7</sup>
V2TG	0.068	0.47	1.09x10 <sup>-7</sup>
TPTG	0.060	0.46	1.70x10 <sup>-7</sup>
H2FC	0.112	-0.46	2.27x10 <sup>-7</sup>
L1CH	0.044	-0.46	2.48x10 <sup>-7</sup>
L4FC	0.123	-0.45	3.27x10 <sup>-7</sup>
V3PL	0.078	0.43	1.03x10 <sup>-6</sup>
IDPN	0.082	0.41	5.58x10 <sup>-6</sup>
IDAB	0.082	0.40	5.58x10 <sup>-6</sup>
L3CH	0.091	-0.40	7.83x10 <sup>-6</sup>
V3CH	0.071	0.39	1.15x10 <sup>-5</sup>
V4FC	0.081	0.38	1.61x10 <sup>-5</sup>
V2PL	0.062	0.38	2.12x10 <sup>-5</sup>
VLTG	0.048	0.36	4.51x10 <sup>-5</sup>

LDPL	0.087	-0.36	4.81x10 <sup>-5</sup>
L4PN	0.092	-0.36	5.16x10 <sup>-5</sup>
L4AB	0.092	-0.36	5.16x10 <sup>-5</sup>
VLCH	0.064	0.35	8.33x10 <sup>-5</sup>
V4CH	0.074	0.35	8.76x10 <sup>-5</sup>
V2CH	0.067	0.35	9.56x10 <sup>-5</sup>
LDFC	0.086	-0.33	2.08x10 <sup>-4</sup>
L3PL	0.075	-0.32	3.46x10 <sup>-4</sup>
IDTG	0.038	0.29	1.27x10 <sup>-3</sup>
VLFC	0.049	0.28	2.00x10 <sup>-3</sup>
L5CH	0.083	-0.28	2.05x10 <sup>-3</sup>
L1FC	0.025	-0.28	2.27x10 <sup>-3</sup>
H2CH	0.075	-0.27	2.56x10 <sup>-3</sup>
L3FC	0.073	-0.25	6.16x10 <sup>-3</sup>
H2A2	0.065	-0.25	6.30x10 <sup>-3</sup>
L1PL	0.005	-0.24	8.14x10 <sup>-3</sup>
L5FC	0.088	-0.24	8.65x10 <sup>-3</sup>
V1TG	0.029	0.23	1.00x10 <sup>-2</sup>
L2CH	0.031	-0.23	1.16x10 <sup>-2</sup>
IDCH	0.054	0.23	1.17x10 <sup>-2</sup>
VLPL	0.044	0.23	1.21x10 <sup>-2</sup>
V5CH	0.044	0.22	1.40x10 <sup>-2</sup>
V1PL	0.031	0.21	1.82x10 <sup>-2</sup>
L5PL	0.070	-0.20	2.61x10 <sup>-2</sup>
IDFC	0.048	0.19	3.52x10 <sup>-2</sup>
V5PL	0.030	0.19	4.21x10 <sup>-2</sup>
TPAB	0.016	0.18	4.73x10 <sup>-2</sup>
TBPN	0.016	0.18	4.73x10 <sup>-2</sup>

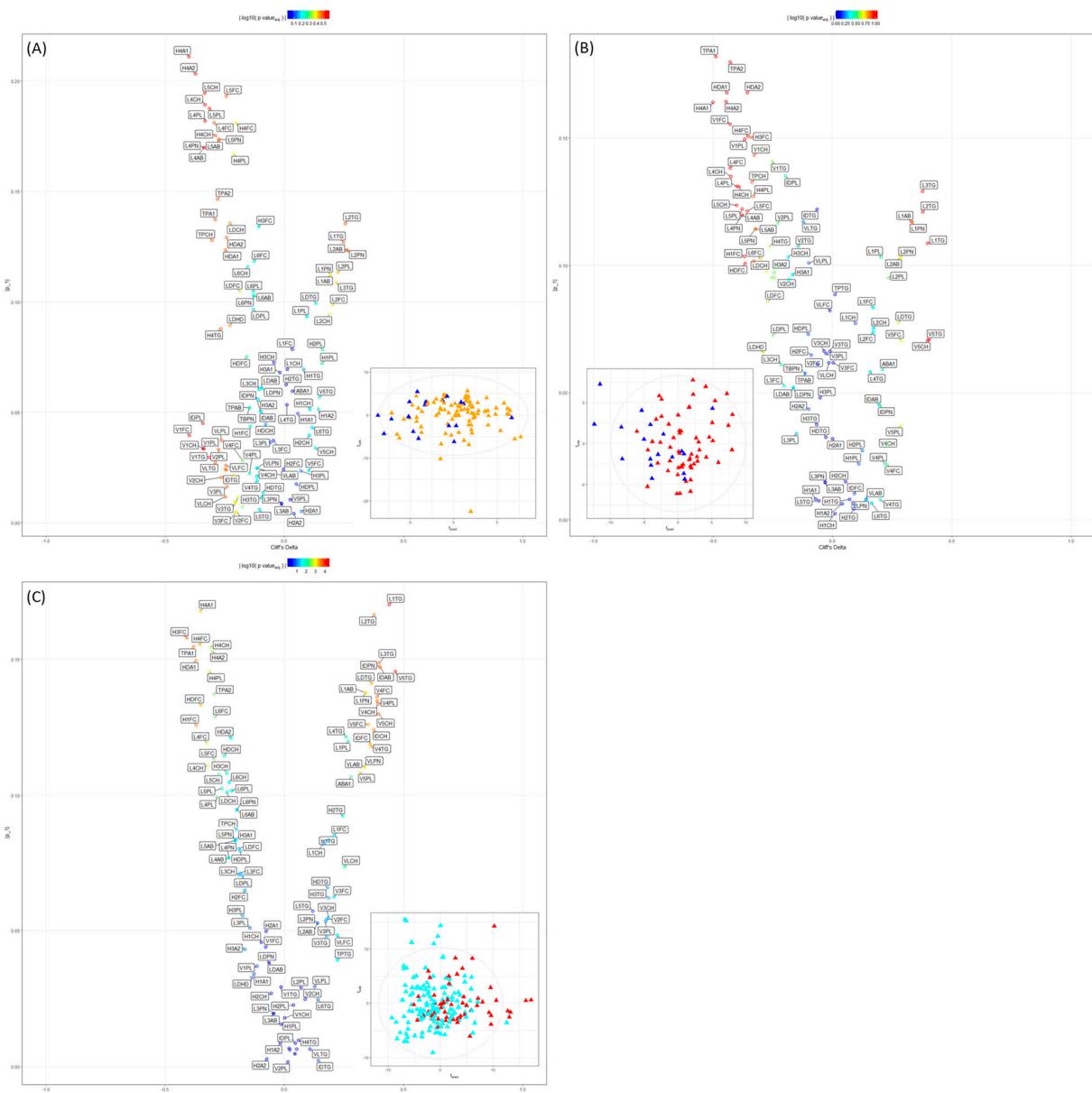
**Table S13 - OPLS loadings, Cliff's delta and adjusted p-values of the lipoproteins for the comparison of the controls and SARS-CoV-2 positive patients in severity class E.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
H4PL	0.165	-0.98	2.70x10 <sup>-21</sup>
H4CH	0.166	-0.97	3.01x10 <sup>-21</sup>
TPA2	0.158	-0.95	7.83x10 <sup>-21</sup>

H4A1	0.167	-0.95	7.83x10 <sup>-21</sup>
H4A2	0.161	-0.95	1.02x10 <sup>-20</sup>
TPA1	0.160	-0.94	2.39x10 <sup>-20</sup>
HDA1	0.158	-0.93	3.32x10 <sup>-20</sup>
H4FC	0.166	-0.92	1.06x10 <sup>-19</sup>
HDA2	0.150	-0.91	3.57x10 <sup>-19</sup>
H3FC	0.163	-0.91	3.57x10 <sup>-19</sup>
H3CH	0.148	-0.86	2.11x10 <sup>-17</sup>
HDCH	0.140	-0.86	3.02x10 <sup>-17</sup>
LDTG	0.144	0.85	4.11x10 <sup>-17</sup>
L1TG	0.145	0.80	4.36x10 <sup>-15</sup>
HDPL	0.129	-0.78	1.36x10 <sup>-14</sup>
L2TG	0.142	0.76	5.23x10 <sup>-14</sup>
H3A1	0.134	-0.76	6.97x10 <sup>-14</sup>
VLPN	0.124	0.75	1.16x10 <sup>-13</sup>
VLAB	0.124	0.75	1.16x10 <sup>-13</sup>
V4TG	0.121	0.75	1.16x10 <sup>-13</sup>
H3PL	0.124	-0.75	1.31x10 <sup>-13</sup>
HDFC	0.138	-0.75	1.47x10 <sup>-13</sup>
H2TG	0.119	0.74	3.52x10 <sup>-13</sup>
L4TG	0.127	0.73	4.65x10 <sup>-13</sup>
V5TG	0.116	0.71	2.18x10 <sup>-12</sup>
ABA1	0.120	0.70	3.23x10 <sup>-12</sup>
V4PL	0.120	0.70	6.23x10 <sup>-12</sup>
H1FC	0.112	-0.67	2.92x10 <sup>-11</sup>
L3TG	0.121	0.67	4.92x10 <sup>-11</sup>
V2FC	0.098	0.66	8.04x10 <sup>-11</sup>
L5TG	0.107	0.65	9.96x10 <sup>-11</sup>
V3TG	0.096	0.65	1.55x10 <sup>-10</sup>
V3FC	0.099	0.63	5.03x10 <sup>-10</sup>
H2A1	0.102	-0.63	5.03x10 <sup>-10</sup>
V4FC	0.115	0.62	7.93x10 <sup>-10</sup>
L4CH	0.110	-0.60	3.78x10 <sup>-9</sup>
V4CH	0.111	0.59	6.08x10 <sup>-9</sup>
H1TG	0.108	0.59	7.39x10 <sup>-9</sup>
TPCH	0.110	-0.58	9.54x10 <sup>-9</sup>
TPTG	0.088	0.58	1.34x10 <sup>-8</sup>

H3A2	0.099	-0.58	1.34x10 <sup>-8</sup>
IDAB	0.113	0.57	1.70x10 <sup>-8</sup>
IDPN	0.113	0.57	1.74x10 <sup>-8</sup>
V3PL	0.091	0.57	2.04x10 <sup>-8</sup>
L4FC	0.114	-0.56	2.94x10 <sup>-8</sup>
H2FC	0.102	-0.55	4.38x10 <sup>-8</sup>
L6TG	0.091	0.55	5.66x10 <sup>-8</sup>
VLCH	0.090	0.54	9.94x10 <sup>-8</sup>
L4PL	0.101	-0.54	1.10x10 <sup>-7</sup>
H3TG	0.085	0.54	1.10x10 <sup>-7</sup>
V3CH	0.085	0.52	2.06x10 <sup>-7</sup>
LDCH	0.103	-0.52	2.59x10 <sup>-7</sup>
HDTG	0.094	0.52	2.60x10 <sup>-7</sup>
V2TG	0.068	0.52	3.31x10 <sup>-7</sup>
L3CH	0.087	-0.48	2.18x10 <sup>-6</sup>
V5CH	0.097	0.48	2.62x10 <sup>-6</sup>
VLTG	0.063	0.47	2.97x10 <sup>-6</sup>
VLFC	0.075	0.47	3.75x10 <sup>-6</sup>
IDCH	0.094	0.47	3.91x10 <sup>-6</sup>
IDFC	0.089	0.44	1.55x10 <sup>-5</sup>
V2CH	0.063	0.42	2.86x10 <sup>-5</sup>
V2PL	0.055	0.41	6.40x10 <sup>-5</sup>
V5PL	0.083	0.41	6.40x10 <sup>-5</sup>
IDTG	0.050	0.40	7.17x10 <sup>-5</sup>
IDPL	0.083	-0.40	7.19x10 <sup>-5</sup>
L4PN	0.077	-0.40	9.02x10 <sup>-5</sup>
L4AB	0.077	-0.40	9.02x10 <sup>-5</sup>
LDFC	0.083	-0.39	1.37x10 <sup>-4</sup>
L3PL	0.072	-0.39	1.41x10 <sup>-4</sup>
VLPL	0.061	0.38	1.69x10 <sup>-4</sup>
L3FC	0.080	-0.37	2.18x10 <sup>-4</sup>
H2A2	0.051	-0.34	7.86x10 <sup>-4</sup>
L5CH	0.075	-0.34	9.84x10 <sup>-4</sup>
L5FC	0.085	-0.33	1.02x10 <sup>-3</sup>
H2CH	0.064	-0.33	1.21x10 <sup>-3</sup>
V1TG	0.029	0.30	3.04x10 <sup>-3</sup>
L1CH	0.026	-0.30	3.84x10 <sup>-3</sup>

L6FC	0.078	-0.28	6.97x10 <sup>-3</sup>
L5PL	0.069	-0.26	1.10x10 <sup>-2</sup>
L2CH	0.037	-0.23	2.76x10 <sup>-2</sup>
H1CH	0.045	-0.23	2.76x10 <sup>-2</sup>
L1PN	0.066	0.21	4.32x10 <sup>-2</sup>
L1AB	0.066	0.21	4.32x10 <sup>-2</sup>



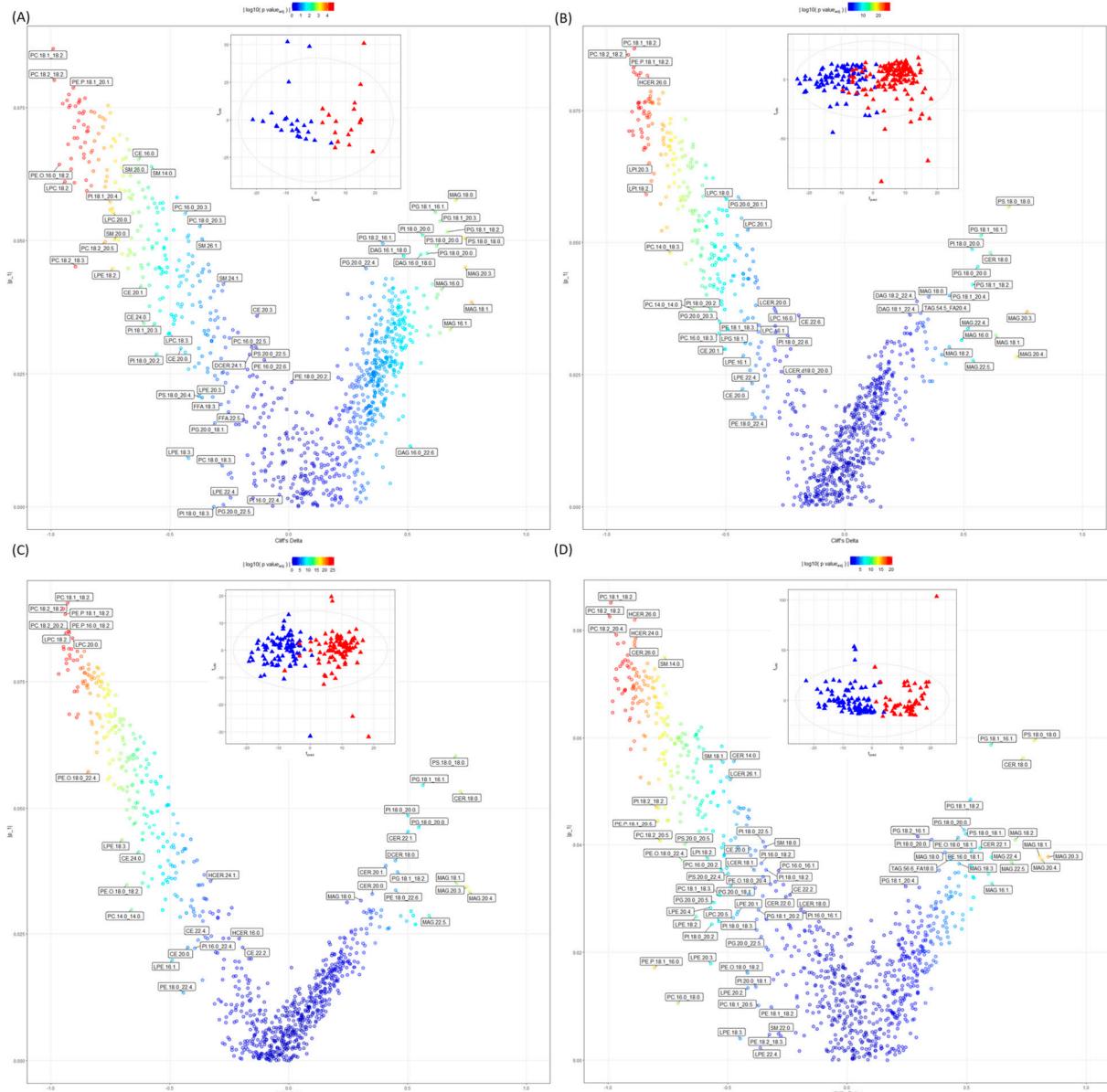
**Figure S8 - Orthogonal Partial Least Squared (O-PLS-DA) and eruption plots between different severity classes of SARS-CoV-2 positive patients for the lipoproteins.** (A) O-PLS-DA SARS-CoV-2 severity group B (blue) and SARS-CoV-2 severity group D (orange) ( $R^2X=0.18$ , AUROC=0.64) and eruption plot. (B) O-PLS-DA SARS-CoV-2 severity group B

(blue) and SARS-CoV-2 severity group E (red) ( $R_2X=0.21$ , AUROC=0.70) and eruption plot. (C) O-PLS-DA SARS-CoV-2 severity group C (cyan) and SARS-CoV-2 severity group E (red) ( $R_2X=0.28$ , AUROC=0.72) and eruption plot. Significant models could not be produced for severity group B vs C, group C vs D or D vs E.

**Table S14 - OPLS loadings, Cliff's delta and adjusted p-values of the lipoproteins for the comparison of severity class C and E.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
V5TG	0.146	0.47	$3.50 \times 10^{-5}$
L1TG	0.170	0.44	$7.69 \times 10^{-5}$
IDPN	0.147	0.40	$2.00 \times 10^{-4}$
IDAB	0.147	0.40	$2.00 \times 10^{-4}$
V5CH	0.130	0.40	$2.00 \times 10^{-4}$
V4PL	0.134	0.40	$2.00 \times 10^{-4}$
L3TG	0.149	0.40	$2.00 \times 10^{-4}$
H3FC	0.158	-0.41	$2.00 \times 10^{-4}$
V4CH	0.135	0.39	$2.20 \times 10^{-4}$
V4FC	0.136	0.39	$2.20 \times 10^{-4}$
TPA1	0.155	-0.38	$3.08 \times 10^{-4}$
L2TG	0.166	0.38	$3.26 \times 10^{-4}$
IDCH	0.124	0.37	$3.40 \times 10^{-4}$
HDA1	0.150	-0.37	$3.58 \times 10^{-4}$
H1FC	0.126	-0.37	$3.65 \times 10^{-4}$
LDTG	0.141	0.37	$4.20 \times 10^{-4}$
V4TG	0.118	0.36	$4.20 \times 10^{-4}$
V5FC	0.126	0.35	$4.57 \times 10^{-4}$
IDFC	0.119	0.36	$4.58 \times 10^{-4}$
H4FC	0.156	-0.35	$5.42 \times 10^{-4}$
H4A1	0.168	-0.35	$6.02 \times 10^{-4}$
HDFC	0.133	-0.35	$6.15 \times 10^{-4}$
L1PN	0.138	0.34	$8.71 \times 10^{-4}$
L1AB	0.138	0.34	$9.64 \times 10^{-4}$
VLPN	0.110	0.34	$9.64 \times 10^{-4}$
VLAB	0.110	0.34	$9.64 \times 10^{-4}$
L4FC	0.119	-0.33	$1.46 \times 10^{-3}$
L4CH	0.111	-0.32	$1.53 \times 10^{-3}$
V5PL	0.109	0.32	$1.91 \times 10^{-3}$
H4PL	0.146	-0.31	$2.08 \times 10^{-3}$
H4CH	0.154	-0.31	$2.39 \times 10^{-3}$

H4A2	0.153	-0.30	$3.07 \times 10^{-3}$
TPA2	0.137	-0.29	$4.04 \times 10^{-3}$
L5FC	0.113	-0.29	$4.04 \times 10^{-3}$
L6FC	0.129	-0.29	$4.70 \times 10^{-3}$
L4PL	0.099	-0.28	$5.81 \times 10^{-3}$
ABA1	0.107	0.28	$6.27 \times 10^{-3}$
L5CH	0.108	-0.27	$7.41 \times 10^{-3}$
L1PL	0.120	0.27	$8.88 \times 10^{-3}$
L5PL	0.103	-0.26	$1.09 \times 10^{-2}$
L4TG	0.121	0.26	$1.26 \times 10^{-2}$
HDCH	0.115	-0.25	$1.49 \times 10^{-2}$
VLCH	0.073	0.25	$1.49 \times 10^{-2}$
H2TG	0.092	0.24	$1.87 \times 10^{-2}$
LDCH	0.101	-0.24	$1.98 \times 10^{-2}$
H3CH	0.108	-0.24	$2.00 \times 10^{-2}$
L4PN	0.077	-0.23	$2.47 \times 10^{-2}$
L4AB	0.077	-0.23	$2.47 \times 10^{-2}$
L6CH	0.105	-0.23	$2.51 \times 10^{-2}$
HDA2	0.121	-0.22	$3.07 \times 10^{-2}$
TPTG	0.039	0.22	$3.14 \times 10^{-2}$
VLFC	0.048	0.22	$3.14 \times 10^{-2}$
L6PL	0.102	-0.22	$3.14 \times 10^{-2}$
L1FC	0.085	0.21	$4.35 \times 10^{-2}$
L5PN	0.083	-0.21	$4.56 \times 10^{-2}$
V3FC	0.062	0.21	$4.56 \times 10^{-2}$
L5AB	0.083	-0.21	$4.56 \times 10^{-2}$
H3A1	0.083	-0.20	$4.81 \times 10^{-2}$



**Figure S9 - Orthogonal Partial Least Squared (O-PLS-DA) and eruption plots between controls and SARS-CoV-2 positive patients of each severity group for the Lipids.** (A) Eruption plot of the controls vs group B severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group B severity patients (red triangles) ( $R_2X=0.22$ , AUROC=0.98) inset. (B) Eruption plot of the controls vs group C severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group C severity patients (red triangles) ( $R_2X=0.18$ , AUROC=0.97) inset. (C) Eruption plot of the controls vs group D severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group D severity patients (red triangles) ( $R_2X=0.12$ , AUROC=0.99) inset. (D) Eruption plot of the controls vs group E severity group with the O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group E severity patients (red triangles) ( $R_2X=0.21$ , AUROC=0.99) inset.

**Table S15 - OPLS loadings, Cliff's delta and adjusted p-values of the lipids for the comparison of the controls and SARS-CoV-2 positive patients in severity class B.** Only the 50 most significant adjusted p-values are shown.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
PC (18:1/18:2)	0.086	-0.99	2.00x10 <sup>-5</sup>
PC (18:2/18:2)	0.080	-0.99	2.00x10 <sup>-5</sup>
PE.O (16:0/18:2)	0.064	-0.96	2.65x10 <sup>-5</sup>
HCER.d (18:0/18:0)	0.075	-0.92	2.85x10 <sup>-5</sup>
LPC (18:2)	0.061	-0.94	2.85x10 <sup>-5</sup>
PE.O (16:0/20:4)	0.063	-0.92	2.85x10 <sup>-5</sup>
PE.P (16:0/18:2)	0.077	-0.92	2.85x10 <sup>-5</sup>
PE.P (16:0/18:3)	0.065	-0.92	2.85x10 <sup>-5</sup>
PE.P (18:0/18:2)	0.076	-0.93	2.85x10 <sup>-5</sup>
PE.P (18:0/20:1)	0.075	-0.92	2.85x10 <sup>-5</sup>
PE.P (18:1/18:2)	0.076	-0.95	2.85x10 <sup>-5</sup>
PE.P (18:2/20:4)	0.068	-0.93	2.85x10 <sup>-5</sup>
PC (18:2/20:2)	0.061	-0.90	3.14x10 <sup>-5</sup>
PE.O (18:0/22:5)	0.071	-0.90	3.14x10 <sup>-5</sup>
PE.P (18:1/20:1)	0.079	-0.91	3.14x10 <sup>-5</sup>
PE.P (18:1/20:4)	0.069	-0.90	3.14x10 <sup>-5</sup>
PC (18:2/18:3)	0.045	-0.90	3.17x10 <sup>-5</sup>
PC (18:2/20:4)	0.075	-0.90	3.17x10 <sup>-5</sup>
HCER (22:0)	0.069	-0.89	3.87x10 <sup>-5</sup>
PC (16:0/18:2)	0.076	-0.88	3.93x10 <sup>-5</sup>
PC (20:0/20:4)	0.074	-0.88	3.93x10 <sup>-5</sup>
PE.O (16:0/20:1)	0.073	-0.88	3.93x10 <sup>-5</sup>
PE.P (18:0/18:1)	0.077	-0.88	3.93x10 <sup>-5</sup>
PE.P (18:1/22:5)	0.076	-0.88	3.93x10 <sup>-5</sup>
PE.O (18:0/22:4)	0.068	-0.88	4.28x10 <sup>-5</sup>
PC (18:0/20:1)	0.074	-0.87	5.16x10 <sup>-5</sup>
PE.O (16:0/18:1)	0.066	-0.86	5.16x10 <sup>-5</sup>
PE.P (16:0/22:5)	0.074	-0.86	5.16x10 <sup>-5</sup>
PE.P (18:0/20:4)	0.066	-0.86	5.16x10 <sup>-5</sup>
PE.P (18:1/22:4)	0.072	-0.86	5.16x10 <sup>-5</sup>
PE.P (18:0/22:5)	0.068	-0.86	5.46x10 <sup>-5</sup>
PE.P (18:2/18:2)	0.068	-0.86	5.46x10 <sup>-5</sup>
PE.O (16:0/22:5)	0.061	-0.85	5.98x10 <sup>-5</sup>

PE.O (16:0/22:6)	0.059	-0.85	6.36x10 <sup>-5</sup>
PE.P (18:0/22:4)	0.071	-0.85	6.36x10 <sup>-5</sup>
PC (20:0/18:1)	0.071	-0.84	6.79x10 <sup>-5</sup>
PE.P (18:0/18:0)	0.071	-0.84	6.79x10 <sup>-5</sup>
PE.P (16:0/20:4)	0.065	-0.84	7.45x10 <sup>-5</sup>
HCER.d (18:0/22:0)	0.067	-0.83	8.76x10 <sup>-5</sup>
PE.P (16:0/20:1)	0.074	-0.83	8.76x10 <sup>-5</sup>
PE.P (18:1/16:0)	0.060	-0.83	8.76x10 <sup>-5</sup>
PE.P (18:0/16:0)	0.067	-0.83	9.62x10 <sup>-5</sup>
PC (20:0/20:3)	0.063	-0.82	1.06x10 <sup>-4</sup>
PC (18:2/20:3)	0.069	-0.82	1.14x10 <sup>-4</sup>
PE.P (18:1/18:1)	0.074	-0.82	1.14x10 <sup>-4</sup>
PC (18:2/22:6)	0.065	-0.80	1.72x10 <sup>-4</sup>
PE.P (18:0/20:2)	0.070	-0.80	1.72x10 <sup>-4</sup>
PC (16:0/20:1)	0.073	-0.80	1.89x10 <sup>-4</sup>
HCER.d (18:0/24:0)	0.073	-0.78	2.55x10 <sup>-4</sup>
PE.P (16:0/18:1)	0.071	-0.78	2.55x10 <sup>-4</sup>

**Table S16 - OPLS loadings, Cliff's delta and adjusted p-values of the lipids for the comparison of the controls and SARS-CoV-2 positive patients in severity class C.** Only the 50 most significant adjusted p-values are shown.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
PC (18:2/18:2)	0.086	-0.91	3.47x10 <sup>-28</sup>
PE.O (16:0/20:4)	0.071	-0.91	3.47x10 <sup>-28</sup>
PE.O (16:0/18:2)	0.075	-0.89	1.24x10 <sup>-27</sup>
PE.P (18:2/20:4)	0.071	-0.89	1.24x10 <sup>-27</sup>
PE.P (16:0/22:5)	0.070	-0.89	1.75x10 <sup>-27</sup>
PE.P (16:0/18:2)	0.082	-0.89	1.78x10 <sup>-27</sup>
PE.P (18:1/18:2)	0.083	-0.89	1.78x10 <sup>-27</sup>
PC (18:1/18:2)	0.087	-0.88	2.16x10 <sup>-27</sup>
PE.P (18:0/18:2)	0.082	-0.88	6.43x10 <sup>-27</sup>
HCER.d (18:0/18:0)	0.076	-0.87	1.56x10 <sup>-26</sup>
PE.O (16:0/18:1)	0.073	-0.87	1.56x10 <sup>-26</sup>
PE.O (16:0/20:1)	0.076	-0.87	1.56x10 <sup>-26</sup>
PE.P (18:0/18:1)	0.082	-0.87	1.76x10 <sup>-26</sup>

PE.P (18:0/22:5)	0.076	-0.87	1.76x10 <sup>-26</sup>
PE.P (18:1/20:4)	0.075	-0.87	1.80x10 <sup>-26</sup>
PC (18:2/20:2)	0.076	-0.86	2.43x10 <sup>-26</sup>
PE.P (16:0/20:1)	0.082	-0.86	4.43x10 <sup>-26</sup>
LPC (18:2)	0.076	-0.86	4.58x10 <sup>-26</sup>
PC (18:2/20:3)	0.071	-0.86	4.64x10 <sup>-26</sup>
PE.O (18:0/22:5)	0.070	-0.85	8.09x10 <sup>-26</sup>
PE.P (16:0/20:4)	0.071	-0.85	8.09x10 <sup>-26</sup>
PC (20:0/20:3)	0.074	-0.84	2.26x10 <sup>-25</sup>
PE.P (18:0/22:4)	0.068	-0.84	2.26x10 <sup>-25</sup>
PE.P (18:2/18:2)	0.075	-0.84	2.65x10 <sup>-25</sup>
PC (18:0/20:1)	0.076	-0.84	2.78x10 <sup>-25</sup>
PE.O (18:0/22:4)	0.068	-0.84	2.80x10 <sup>-25</sup>
PE.P (18:0/20:1)	0.077	-0.84	4.80x10 <sup>-25</sup>
LPI (18:1)	0.068	-0.84	4.94x10 <sup>-25</sup>
LPI (18:2)	0.059	-0.83	6.95x10 <sup>-25</sup>
PE.P (18:1/22:5)	0.074	-0.83	9.14x10 <sup>-25</sup>
HCER (26:0)	0.082	-0.83	1.15x10 <sup>-24</sup>
PE.P (18:0/20:3)	0.074	-0.83	1.19x10 <sup>-24</sup>
PE.P (16:0/18:3)	0.070	-0.83	1.35x10 <sup>-24</sup>
PE.P (18:0/20:4)	0.072	-0.83	1.94x10 <sup>-24</sup>
PE.O (16:0/22:5)	0.069	-0.82	2.09x10 <sup>-24</sup>
PE.P (18:0/20:2)	0.077	-0.82	2.09x10 <sup>-24</sup>
HCER (22:0)	0.075	-0.82	2.46x10 <sup>-24</sup>
HCER.d (18:0/22:0)	0.074	-0.82	2.46x10 <sup>-24</sup>
LPI (20:3)	0.065	-0.82	2.55x10 <sup>-24</sup>
PE.P (18:0/18:0)	0.077	-0.82	2.55x10 <sup>-24</sup>
PC (20:0/18:1)	0.074	-0.82	4.27x10 <sup>-24</sup>
PE.P (18:1/18:1)	0.077	-0.82	4.36x10 <sup>-24</sup>
PC (18:2/20:4)	0.078	-0.82	4.95x10 <sup>-24</sup>
LPC (20:0)	0.079	-0.82	5.75x10 <sup>-24</sup>
HCER.d (18:0/24:0)	0.077	-0.81	7.44x10 <sup>-24</sup>
PE.P (18:1/22:4)	0.062	-0.81	8.83x10 <sup>-24</sup>
PI (18:1/18:1)	0.072	-0.81	1.22x10 <sup>-23</sup>
PE.P (18:0/16:0)	0.076	-0.81	1.33x10 <sup>-23</sup>
HCER (24:0)	0.077	-0.81	1.48x10 <sup>-23</sup>
PC (18:2/18:3)	0.063	-0.80	3.23x10 <sup>-23</sup>

**Table S17 - OPLS loadings, Cliff's delta and adjusted p-values of the lipids for the comparison of the controls and SARS-CoV-2 positive patients in severity class D.** Only the 50 most significant adjusted p-values are shown.

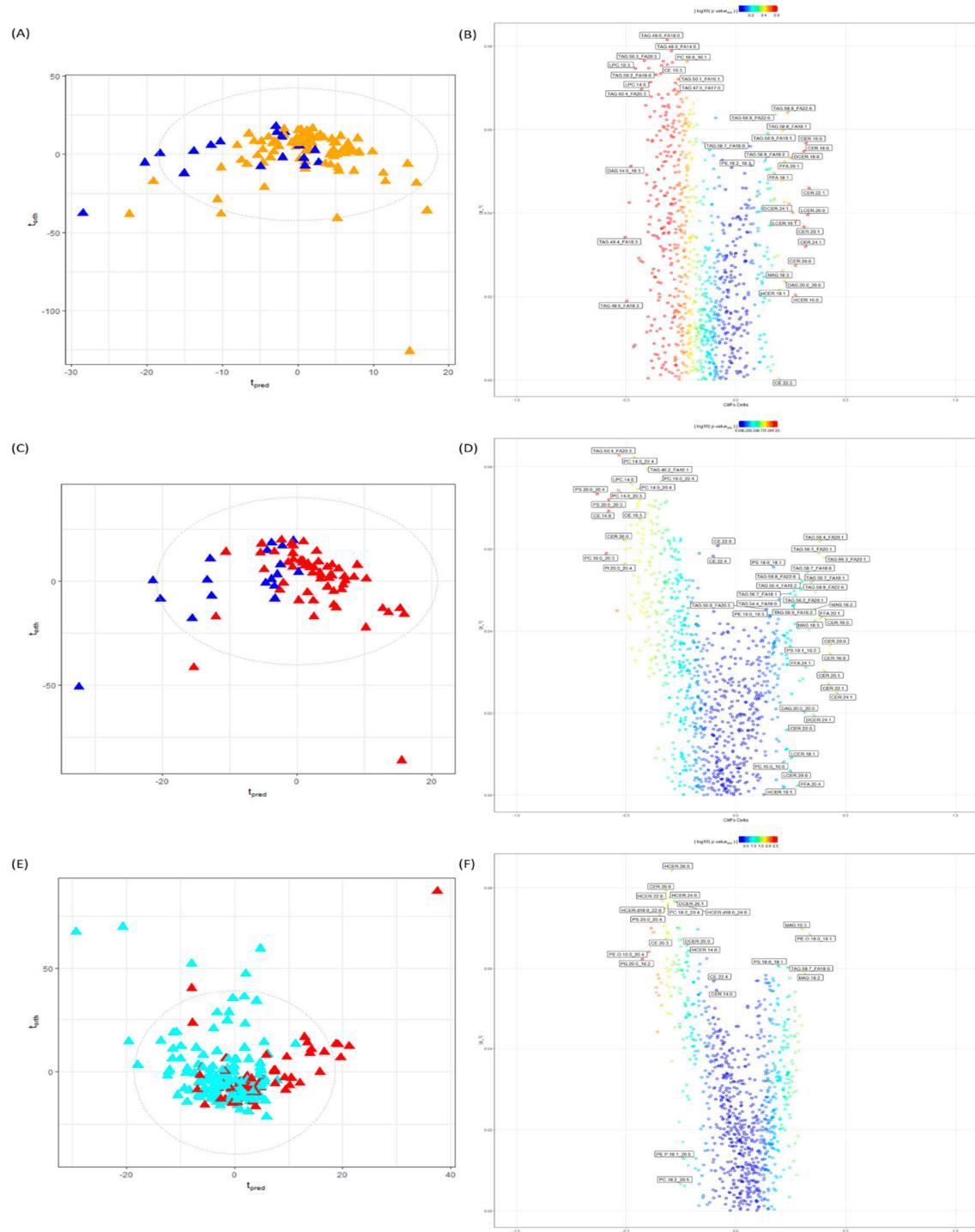
Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
PE.O (16:0/20:4)	0.077	-0.97	1.91x10 <sup>-26</sup>
PC (18:2/18:2)	0.089	-0.95	8.63x10 <sup>-26</sup>
PE.P (18:1/20:4)	0.082	-0.95	8.63x10 <sup>-26</sup>
PE.P (18:2/20:4)	0.081	-0.95	8.63x10 <sup>-26</sup>
PE.P (18:1/18:2)	0.088	-0.94	1.56x10 <sup>-25</sup>
LPC (18:2)	0.085	-0.94	1.76x10 <sup>-25</sup>
PE.P (16:0/20:4)	0.079	-0.94	1.76x10 <sup>-25</sup>
PE.P (16:0/22:5)	0.079	-0.94	1.76x10 <sup>-25</sup>
PC (18:1/18:2)	0.090	-0.93	2.91x10 <sup>-25</sup>
PE.O (16:0/18:2)	0.077	-0.93	2.97x10 <sup>-25</sup>
PC (18:1/20:2)	0.085	-0.93	3.67x10 <sup>-25</sup>
PE.P (18:0/18:2)	0.081	-0.92	4.58x10 <sup>-25</sup>
PC (18:2/18:3)	0.069	-0.92	4.78x10 <sup>-25</sup>
PE.P (16:0/18:2)	0.085	-0.92	4.98x10 <sup>-25</sup>
PE.P (18:0/22:5)	0.080	-0.92	4.98x10 <sup>-25</sup>
PC (18:2/20:3)	0.080	-0.92	5.97x10 <sup>-25</sup>
PE.P (18:2/18:2)	0.081	-0.91	1.28x10 <sup>-24</sup>
LPC (20:0)	0.084	-0.91	1.85x10 <sup>-24</sup>
PE.P (18:1/22:5)	0.079	-0.91	3.30x10 <sup>-24</sup>
PE.O (16:0/22:5)	0.073	-0.90	4.92x10 <sup>-24</sup>
PE.O (16:0/18:1)	0.068	-0.89	1.46x10 <sup>-23</sup>
PE.O (18:0/22:5)	0.073	-0.89	1.71x10 <sup>-23</sup>
PE.P (18:0/20:4)	0.079	-0.89	1.84x10 <sup>-23</sup>
PE.P (18:0/18:1)	0.081	-0.89	1.87x10 <sup>-23</sup>
PC (20:0/20:3)	0.074	-0.89	2.02x10 <sup>-23</sup>
PE.P (16:0/18:3)	0.069	-0.89	2.54x10 <sup>-23</sup>
PC (18:2/20:4)	0.078	-0.88	8.13x10 <sup>-23</sup>
HCER.d (18:0/18:0)	0.080	-0.87	1.08x10 <sup>-22</sup>
PE.O (16:0/20:1)	0.075	-0.87	2.15x10 <sup>-22</sup>
PI (20:0/18:2)	0.078	-0.87	2.33x10 <sup>-22</sup>
PE.P (18:1/22:4)	0.075	-0.87	2.46x10 <sup>-22</sup>
LPC (20:3)	0.077	-0.86	4.35x10 <sup>-22</sup>
PE.P (16:0/20:1)	0.081	-0.86	4.35x10 <sup>-22</sup>

PE.P (18:0/16:0)	0.077	-0.86	5.62x10 <sup>-22</sup>
PE.P (18:0/22:4)	0.075	-0.86	5.95x10 <sup>-22</sup>
PI (18:1/18:1)	0.074	-0.85	1.11x10 <sup>-21</sup>
PE.P (18:0/20:3)	0.074	-0.85	1.70x10 <sup>-21</sup>
SM (24:0)	0.076	-0.85	1.70x10 <sup>-21</sup>
PC (18:0/20:1)	0.074	-0.85	1.91x10 <sup>-21</sup>
PE.O (18:0/22:4)	0.057	-0.84	2.61x10 <sup>-21</sup>
PE.P (18:1/18:1)	0.078	-0.84	2.77x10 <sup>-21</sup>
PE.P (18:0/20:1)	0.077	-0.84	3.03x10 <sup>-21</sup>
PE.P (18:0/18:0)	0.076	-0.84	4.02x10 <sup>-21</sup>
PE.P (16:0/22:4)	0.073	-0.84	5.05x10 <sup>-21</sup>
LPI (20:3)	0.069	-0.83	6.35x10 <sup>-21</sup>
PE.P (18:0/20:2)	0.075	-0.83	1.05x10 <sup>-20</sup>
LPC (18:3)	0.067	-0.83	1.09x10 <sup>-20</sup>
LPC (18:1)	0.076	-0.83	1.16x10 <sup>-20</sup>
SM (20:0)	0.067	-0.83	1.34x10 <sup>-20</sup>
PC (18:2/22:6)	0.077	-0.82	3.24x10 <sup>-20</sup>

**Table S18 - OPLS loadings, Cliff's delta and adjusted p-values of the lipids for the comparison of the controls and SARS-CoV-2 positive patients in severity class E.** Only the 50 most significant adjusted p values are shown.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
PC (18:1/18:2)	0.085	-0.99	3.35x10 <sup>-21</sup>
PC (18:2/18:2)	0.082	-0.99	3.35x10 <sup>-21</sup>
PE.O (16:0/20:4)	0.065	-0.98	4.10x10 <sup>-21</sup>
PE.P (18:2/20:4)	0.067	-0.98	4.10x10 <sup>-21</sup>
PC (20:0/20:3)	0.071	-0.97	9.70x10 <sup>-21</sup>
PC (18:2/20:3)	0.073	-0.97	9.93x10 <sup>-21</sup>
PC (18:2/20:4)	0.079	-0.97	9.93x10 <sup>-21</sup>
PE.P (18:1/20:4)	0.071	-0.96	1.19x10 <sup>-20</sup>
PE.P (16:0/18:2)	0.077	-0.96	1.52x10 <sup>-20</sup>
LPC (18:2)	0.069	-0.96	2.03x10 <sup>-20</sup>
PE.P (16:0/22:5)	0.073	-0.95	2.95x10 <sup>-20</sup>
PE.P (18:1/18:2)	0.074	-0.94	6.91x10 <sup>-20</sup>
PE.P (16:0/20:4)	0.068	-0.94	7.16x10 <sup>-20</sup>

PE.P (18:0/22:5)	0.074	-0.94	8.07x10 <sup>-20</sup>
PC (18:0/20:1)	0.078	-0.94	1.03x10 <sup>-19</sup>
PE.P (18:2/18:2)	0.069	-0.93	1.41x10 <sup>-19</sup>
PE.P (18:0/20:3)	0.069	-0.93	1.58x10 <sup>-19</sup>
PE.P (18:0/20:4)	0.069	-0.93	1.58x10 <sup>-19</sup>
PE.P (18:0/18:2)	0.074	-0.93	1.62x10 <sup>-19</sup>
PE.O (16:0/22:5)	0.065	-0.93	1.73x10 <sup>-19</sup>
HCER (22:0)	0.075	-0.92	3.49x10 <sup>-19</sup>
PC (18:2/22:5)	0.073	-0.92	3.49x10 <sup>-19</sup>
HCER.d (18:0/22:0)	0.074	-0.92	4.19x10 <sup>-19</sup>
LPC (20:0)	0.073	-0.91	5.04x10 <sup>-19</sup>
PE.P (18:0/22:4)	0.071	-0.91	5.84x10 <sup>-19</sup>
PI (20:0/18:2)	0.063	-0.91	6.05x10 <sup>-19</sup>
PE.O (16:0/18:2)	0.064	-0.91	8.49x10 <sup>-19</sup>
PE.P (16:0/18:3)	0.058	-0.91	8.82x10 <sup>-19</sup>
PE.P (16:0/20:1)	0.074	-0.90	1.28x10 <sup>-18</sup>
PC (18:2/22:6)	0.071	-0.90	1.39x10 <sup>-18</sup>
PC (18:2/20:2)	0.065	-0.90	1.45x10 <sup>-18</sup>
PE.O (16:0/20:1)	0.069	-0.90	1.75x10 <sup>-18</sup>
PE.P (18:1/22:4)	0.066	-0.90	1.90x10 <sup>-18</sup>
PE.P (18:0/16:0)	0.073	-0.90	2.14x10 <sup>-18</sup>
PE.P (18:1/22:5)	0.069	-0.89	2.15x10 <sup>-18</sup>
PE.P (16:0/22:4)	0.067	-0.89	2.17x10 <sup>-18</sup>
PE.P (18:0/20:1)	0.070	-0.89	2.19x10 <sup>-18</sup>
HCER.d (18:0/18:0)	0.070	-0.89	2.47x10 <sup>-18</sup>
SM (20:0)	0.059	-0.89	2.90x10 <sup>-18</sup>
CER (26:0)	0.077	-0.89	3.00x10 <sup>-18</sup>
HCER (26:0)	0.082	-0.89	3.00x10 <sup>-18</sup>
HCER.d (18:0/24:0)	0.078	-0.89	3.00x10 <sup>-18</sup>
SM (24:0)	0.073	-0.89	3.28x10 <sup>-18</sup>
HCER (24:0)	0.078	-0.89	3.99x10 <sup>-18</sup>
PE.P (18:0/20:2)	0.069	-0.89	4.05x10 <sup>-18</sup>
PC (14:0/20:4)	0.068	-0.88	4.26x10 <sup>-18</sup>
PI (18:1/20:4)	0.072	-0.88	4.65x10 <sup>-18</sup>
PC (16:1/18:2)	0.070	-0.88	6.09x10 <sup>-18</sup>
PE.P (18:0/18:1)	0.072	-0.88	6.42x10 <sup>-18</sup>
PI (20:0/20:4)	0.066	-0.87	9.72x10 <sup>-18</sup>



**Figure S10 - Orthogonal Partial Least Squared (O-PLS-DA) and eruption plots between different severity classes of SARS-CoV-2 positive patients for the lipids.** (A) O-PLS-DA SARS-CoV-2 severity group B (blue) and SARS-CoV-2 severity group D (orange) ( $R^2X=0.16$ , AUROC=0.64). (B) Eruption plot of severity group B vs severity group D. (C) O-

PLS-DA SARS-CoV-2 severity group B (blue) and SARS-CoV-2 severity group E (red) ( $R^2X=0.15$ , AUROC=0.79). (D) Eruption plot of severity group B vs severity group E. (E) O-PLS-DA SARS-CoV-2 severity group C (cyan) and SARS-CoV-2 severity group E (red) ( $R^2X=0.13$ , AUROC=0.72). Significant models could not be produced for severity group B vs C, group C vs D or D vs E.

**Table S19 - OPLS loadings, Cliff's delta and adjusted p-values of the lipids for the comparison of severity class B and E.** Only metabolites with significant p-values are shown in the table.

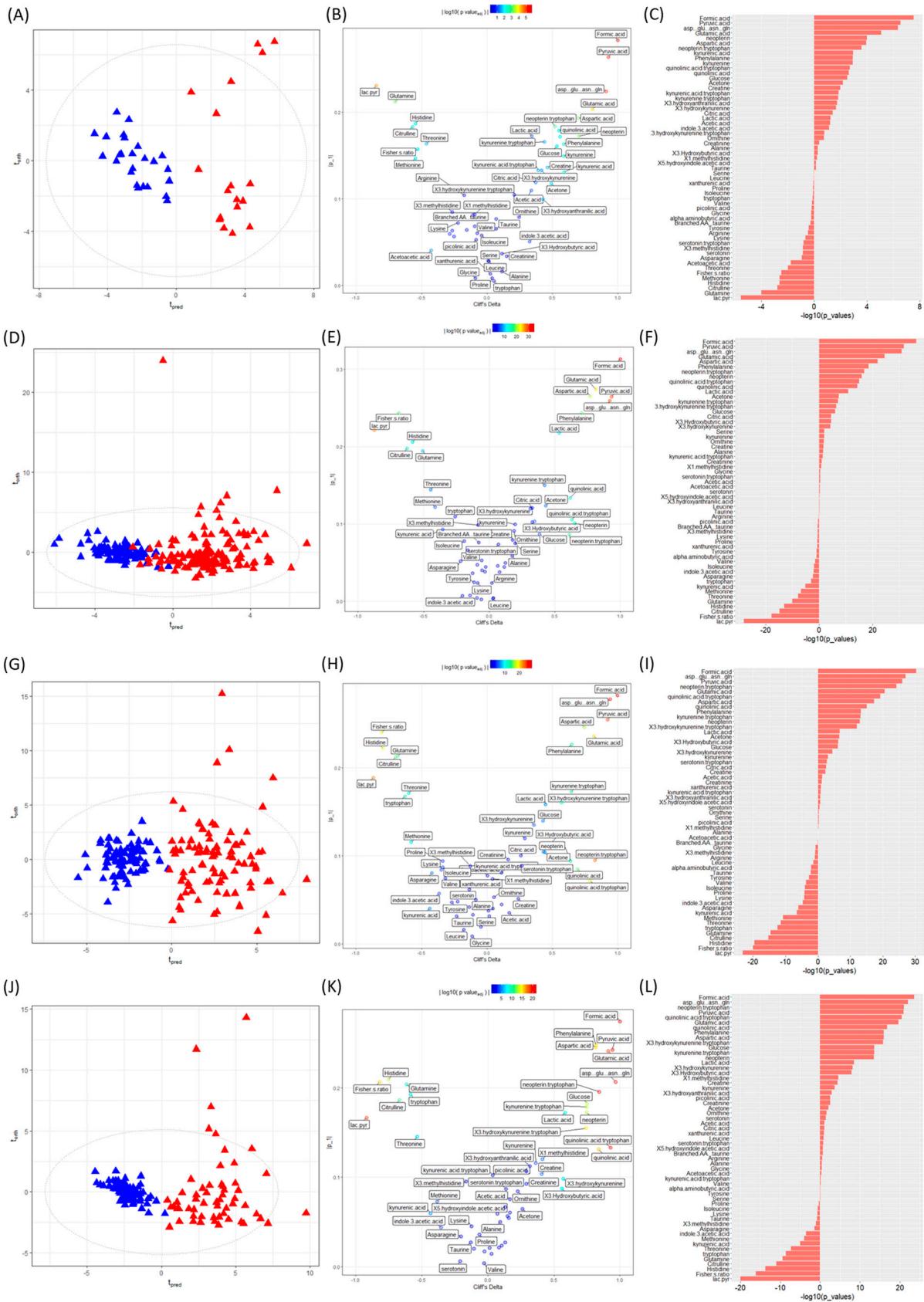
Lipid	OPLS loadings	Cliff's delta	Adjusted p-value
CE (14:0)	0.069	-0.58	4.96x10 <sup>-2</sup>
PC (18:0/20:3)	0.059	-0.59	4.96x10 <sup>-2</sup>
PS (20:0/20:3)	0.072	-0.58	4.96x10 <sup>-2</sup>
PC (20:0/20:4)	0.073	-0.63	4.96x10 <sup>-2</sup>

**Table S20 - OPLS loadings, Cliff's delta and adjusted p-values of the lipids for the comparison of severity class C and E.** Only metabolites with significant p-values are shown in the table.

Lipid	OPLS loadings	Cliff's delta	Adjusted p-value
PG (20:0/18:2)	0.062	-0.43	2.39x10 <sup>-3</sup>
PE.O (16:0/20:4)	0.064	-0.40	6.14x10 <sup>-3</sup>
PE.P (18:0/20:3)	0.070	-0.39	6.18x10 <sup>-3</sup>
PI (20:0/20:4)	0.069	-0.38	7.94x10 <sup>-3</sup>
PC (18:0/20:3)	0.054	-0.36	9.36x10 <sup>-3</sup>
PE.P (16:0/20:3)	0.055	-0.36	9.36x10 <sup>-3</sup>
PG (20:0/20:3)	0.044	-0.36	9.83x10 <sup>-3</sup>
PS (20:0/22:6)	0.058	-0.36	9.83x10 <sup>-3</sup>
PI (20:0/16:1)	0.058	-0.35	9.99x10 <sup>-3</sup>
PI (20:0/18:2)	0.053	-0.35	9.99x10 <sup>-3</sup>
PS (20:0/20:4)	0.073	-0.34	1.21x10 <sup>-2</sup>
PE.P (18:1/20:3)	0.056	-0.34	1.30x10 <sup>-2</sup>
PE (18:2/16:1)	0.051	-0.34	1.33x10 <sup>-2</sup>
HCER (22:0)	0.077	-0.33	1.46x10 <sup>-2</sup>
PE.O (18:0/18:1)	0.069	0.33	1.46x10 <sup>-2</sup>
LPC (20:0)	0.071	-0.33	1.68x10 <sup>-2</sup>
CE (16:0)	0.062	-0.31	2.00x10 <sup>-2</sup>
CE (20:3)	0.067	-0.32	2.00x10 <sup>-2</sup>
CER (26:0)	0.079	-0.32	2.00x10 <sup>-2</sup>
HCER (24:0)	0.077	-0.31	2.00x10 <sup>-2</sup>

HCER.d (18:0/22:0)	0.075	-0.32	2.00x10 <sup>-2</sup>
LPC (14:0)	0.069	-0.31	2.00x10 <sup>-2</sup>
LPC (16:0)	0.059	-0.31	2.00x10 <sup>-2</sup>
LPC (18:0)	0.069	-0.31	2.00x10 <sup>-2</sup>
MAG (18:2)	0.059	0.31	2.00x10 <sup>-2</sup>
PC (18:2/18:2)	0.069	-0.31	2.00x10 <sup>-2</sup>
PC (18:2/20:4)	0.076	-0.31	2.00x10 <sup>-2</sup>
PC (18:2/22:6)	0.060	-0.32	2.00x10 <sup>-2</sup>
PE.P (18:0/22:6)	0.049	-0.31	2.00x10 <sup>-2</sup>
SM (22:1)	0.056	-0.32	2.00x10 <sup>-2</sup>
MAG (18:3)	0.070	0.30	2.15x10 <sup>-2</sup>
PC (18:2/20:3)	0.054	-0.31	2.15x10 <sup>-2</sup>
PE.P (18:2/20:4)	0.058	-0.31	2.15x10 <sup>-2</sup>
PS (20:0/20:3)	0.055	-0.30	2.15x10 <sup>-2</sup>
PC (18:0/18:2)	0.060	-0.30	2.45x10 <sup>-2</sup>
TAG (56:3)_FA (20:1)	0.046	0.30	2.45x10 <sup>-2</sup>
PI (18:0/20:4)	0.049	-0.30	2.54x10 <sup>-2</sup>
SM (20:1)	0.053	-0.30	2.45x10 <sup>-2</sup>
PG (20:0/20:1)	0.053	-0.30	2.58x10 <sup>-2</sup>
HCER (26:0)	0.084	-0.29	2.77x10 <sup>-2</sup>
PC (14:0/22:6)	0.061	-0.29	2.83x10 <sup>-2</sup>
PG (20:0/18:1)	0.037	-0.29	2.94x10 <sup>-2</sup>
TAG (56:4)_FA (20:1)	0.048	0.29	2.94x10 <sup>-2</sup>
TAG (56:2)_FA (18:1)	0.042	0.28	3.37x10 <sup>-2</sup>
SM (24:0)	0.062	-0.28	3.42x10 <sup>-2</sup>
PE.P (16:0/22:6)	0.055	-0.28	3.45x10 <sup>-2</sup>
CER (16:0)	0.018	0.28	3.50x10 <sup>-2</sup>
HCER.d (18:0/24:0)	0.077	-0.28	3.50x10 <sup>-2</sup>
PC (16:0/22:6)	0.045	-0.28	3.55x10 <sup>-2</sup>
TAG (56:4)_FA (18:2)	0.044	0.28	3.55x10 <sup>-2</sup>
CER (24:1)	0.009	0.28	3.93x10 <sup>-2</sup>
DCER (26:1)	0.077	-0.28	3.93x10 <sup>-2</sup>
PE.P (18:0/20:4)	0.061	-0.28	3.93x10 <sup>-2</sup>
TAG (56:2)_FA (18:0)	0.035	0.28	3.93x10 <sup>-2</sup>
TAG (55:4)_FA (18:1)	0.030	0.27	4.15x10 <sup>-2</sup>
CER (20:0)	0.017	0.27	4.18x10 <sup>-2</sup>
FFA (20:1)	0.050	0.27	4.18x10 <sup>-2</sup>

HCER (20:0)	0.063	-0.27	$4.18 \times 10^{-2}$
MAG (18:1)	0.054	0.27	$4.18 \times 10^{-2}$
PC (18:1/20:4)	0.059	-0.27	$4.18 \times 10^{-2}$
PE (16:0/18:1)	0.047	0.27	$4.18 \times 10^{-2}$
TAG (54:3)_FA (18:0)	0.039	0.27	$4.18 \times 10^{-2}$
TAG (56:3)_FA (18:0)	0.043	0.27	$4.18 \times 10^{-2}$
TAG (56:7)_FA (18:1)	0.054	0.27	$4.18 \times 10^{-2}$
TAG (58:7)_FA (18:1)	0.049	0.27	$4.18 \times 10^{-2}$
PS (18:1/18:2)	0.050	0.27	$4.19 \times 10^{-2}$
PC (18:0/22:4)	0.070	-0.27	$4.28 \times 10^{-2}$
CE (22:5)	0.070	-0.26	$4.47 \times 10^{-2}$
TAG (58:3)_FA (18:1)	0.043	0.27	$4.47 \times 10^{-2}$
HCER.d (18:0/20:0)	0.060	-0.26	$4.48 \times 10^{-2}$
CER (22:1)	0.012	0.26	$4.54 \times 10^{-2}$
PE.P (18:1/20:4)	0.062	-0.26	$4.74 \times 10^{-2}$
TAG (55:3)_FA (18:1)	0.029	0.26	$4.74 \times 10^{-2}$
LPC (20:4)	0.054	-0.26	$4.84 \times 10^{-2}$
PE.P (16:0/20:4)	0.061	-0.26	$4.84 \times 10^{-3}$
PE.P (16:0/20:5)	0.022	-0.26	$4.84 \times 10^{-3}$
PC (14:0/20:4)	0.070	-0.30	$4.89 \times 10^{-3}$
PC (20:0/20:4)	0.067	-0.26	$4.89 \times 10^{-3}$
PC (18:0/20:2)	0.036	0.26	$4.89 \times 10^{-3}$
PE.P (18:2/18:2)	0.045	-0.26	$4.89 \times 10^{-3}$



**Figure S11 - Orthogonal Partial Least Squared (O-PLS-DA), eruption plots and  $-\log_{10}$  (p-values) between controls and SARS-CoV-2 positive patients of each severity group for the low molecular weight metabolites.** (A) O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group B severity patients (red triangles) ( $R^2X=0.22$ , AUROC=0.98). (B) Eruption plot of the controls vs group B severity group. (C)  $-\log_{10}$  (p-values) of the controls vs group B severity group. (D) O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group C severity patients (red triangles) ( $R^2X=0.15$ , AUROC=0.99). (E) Eruption plot of the controls vs group C severity group. (F)  $-\log_{10}$  (p-values) of the controls vs group C severity group. (G) O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group D severity patients (red triangles) ( $R^2X=0.19$ , AUROC=1.00). (H) Eruption plot of the controls vs group D severity group. (I)  $-\log_{10}$  (p-values) of the controls vs group D severity group. (J) O-PLS-DA of controls (blue triangles) and SARS-CoV-2 group E severity patients (red triangles) ( $R^2X=0.21$ , AUROC=1.00). (K) Eruption plot of the controls vs group E severity group. (L)  $-\log_{10}$  (p-values) of the controls vs group E severity group. Eruption plots are formed from the Cliff's delta (abscissa) and O-PLS-DA loadings (ordinate). Coloured by significance.

**Table S21 - OPLS loadings, Cliff's delta and adjusted p-values of the low molecular weight metabolites for the comparison of the controls and SARS-CoV-2 positive patients in severity class B.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.283	1.00	$1.68 \times 10^{-6}$
Pyruvic acid	0.264	0.93	$8.01 \times 10^{-6}$
Asp/Glu:Asn/Gln	0.224	0.91	$8.50 \times 10^{-6}$
Lactate/pyruvate	0.231	-0.84	$4.14 \times 10^{-5}$
Glutamic acid	0.204	0.80	$9.52 \times 10^{-5}$
Neopterin	0.172	0.70	$7.91 \times 10^{-4}$
Aspartic acid	0.193	0.70	$7.91 \times 10^{-4}$
Glutamine	0.212	-0.70	$7.91 \times 10^{-4}$
Neopterin/tryptophan	0.184	0.52	$1.66 \times 10^{-3}$
Kynurenic acid	0.131	0.59	$5.49 \times 10^{-3}$
Kynurenine	0.149	0.47	$1.66 \times 10^{-3}$
Phenylalanine	0.163	0.59	$5.49 \times 10^{-3}$
Citrulline	0.183	-0.57	$7.13 \times 10^{-3}$
Quinolinic acid/tryptophan	0.172	0.56	$7.80 \times 10^{-3}$
Quinolinic acid	0.176	0.55	$8.25 \times 10^{-3}$
Histidine	0.187	-0.55	$8.25 \times 10^{-3}$
Methionine	0.147	-0.55	$8.25 \times 10^{-3}$
Glucose	0.161	0.53	$9.92 \times 10^{-3}$
Fisher's ratio	0.157	-0.53	$1.02 \times 10^{-2}$
Acetone	0.118	0.49	$1.84 \times 10^{-2}$
Creatine	0.135	0.46	$2.71 \times 10^{-2}$
Threonine	0.164	-0.46	$2.71 \times 10^{-2}$

Kynurenic acid/tryptophan	0.133	0.39	$3.44 \times 10^{-2}$
Kynurenine/tryptophan	0.166	0.44	$3.50 \times 10^{-2}$
Acetoacetic acid	0.040	-0.43	$3.88 \times 10^{-2}$
3-hydroxyanthranilic acid	0.100	0.43	$3.88 \times 10^{-2}$
3-hydroxykynurenine	0.133	0.42	$4.49 \times 10^{-2}$

**Table S22 - OPLS loadings, Cliff's delta and adjusted p-values of the low molecular weight metabolites for the comparison of the controls and SARS-CoV-2 positive patients in severity class C.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.313	1.00	$2.46 \times 10^{-35}$
Pyruvic acid	0.265	0.93	$4.95 \times 10^{-31}$
Asp/Glu:Asn/Gln	0.259	0.92	$2.23 \times 10^{-30}$
Lactate/pyruvate	0.221	-0.88	$9.32 \times 10^{-28}$
Glutamic acid	0.274	0.81	$3.79 \times 10^{-24}$
Aspartic acid	0.265	0.77	$1.24 \times 10^{-21}$
Phenylalanine	0.243	0.70	$2.64 \times 10^{-18}$
Fisher's ratio	0.243	-0.69	$1.08 \times 10^{-17}$
Neopterin/tryptophan	0.086	0.61	$8.43 \times 10^{-17}$
Neopterin	0.101	0.65	$6.01 \times 10^{-16}$
Quinolinic acid/tryptophan	0.106	0.63	$4.95 \times 10^{-15}$
Citrulline	0.197	-0.63	$7.53 \times 10^{-15}$
Quinolinic acid	0.134	0.62	$2.02 \times 10^{-14}$
Histidine	0.206	-0.59	$3.70 \times 10^{-13}$
Lactic acid	0.218	0.53	$3.92 \times 10^{-11}$
Glutamine	0.194	-0.51	$3.95 \times 10^{-10}$
Threonine	0.144	-0.45	$4.12 \times 10^{-8}$
Acetone	0.123	0.43	$1.35 \times 10^{-7}$
Kynurenine/tryptophan	0.150	0.42	$2.42 \times 10^{-7}$
Methionine	0.121	-0.41	$4.40 \times 10^{-7}$
3-hydroxykynurenine/tryptophan	0.103	0.35	$8.87 \times 10^{-7}$
Glucose	0.087	0.38	$2.50 \times 10^{-6}$
Kynurenic acid	0.093	-0.35	$1.55 \times 10^{-5}$
Citric acid	0.120	0.33	$6.50 \times 10^{-5}$
3-Hydroxybutyric acid	0.102	0.33	$6.85 \times 10^{-5}$
3-Hydroxykynurenine	0.120	0.31	$1.45 \times 10^{-4}$
Tryptophan	0.109	-0.26	$2.07 \times 10^{-3}$

Asparagine	0.052	-0.22	1.05x10 <sup>-2</sup>
Indole-3-acetic acid	0.007	-0.21	1.71x10 <sup>-2</sup>
Serine	0.075	0.20	2.17x10 <sup>-2</sup>
Kynurenone	0.099	0.20	2.28x10 <sup>-2</sup>
Isoleucine	0.078	-0.19	2.71x10 <sup>-2</sup>
Ornithine	0.091	0.19	2.89x10 <sup>-2</sup>
Creatine	0.079	0.18	4.08x10 <sup>-2</sup>
Alanine	0.058	0.18	4.22x10 <sup>-2</sup>

**Table S23 - OPLS loadings, Cliff's delta and adjusted p-values of the low molecular weight metabolites for the comparison of the controls and SARS-CoV-2 positive patients in severity class D.** Only metabolites with significant p-values are shown in the table.

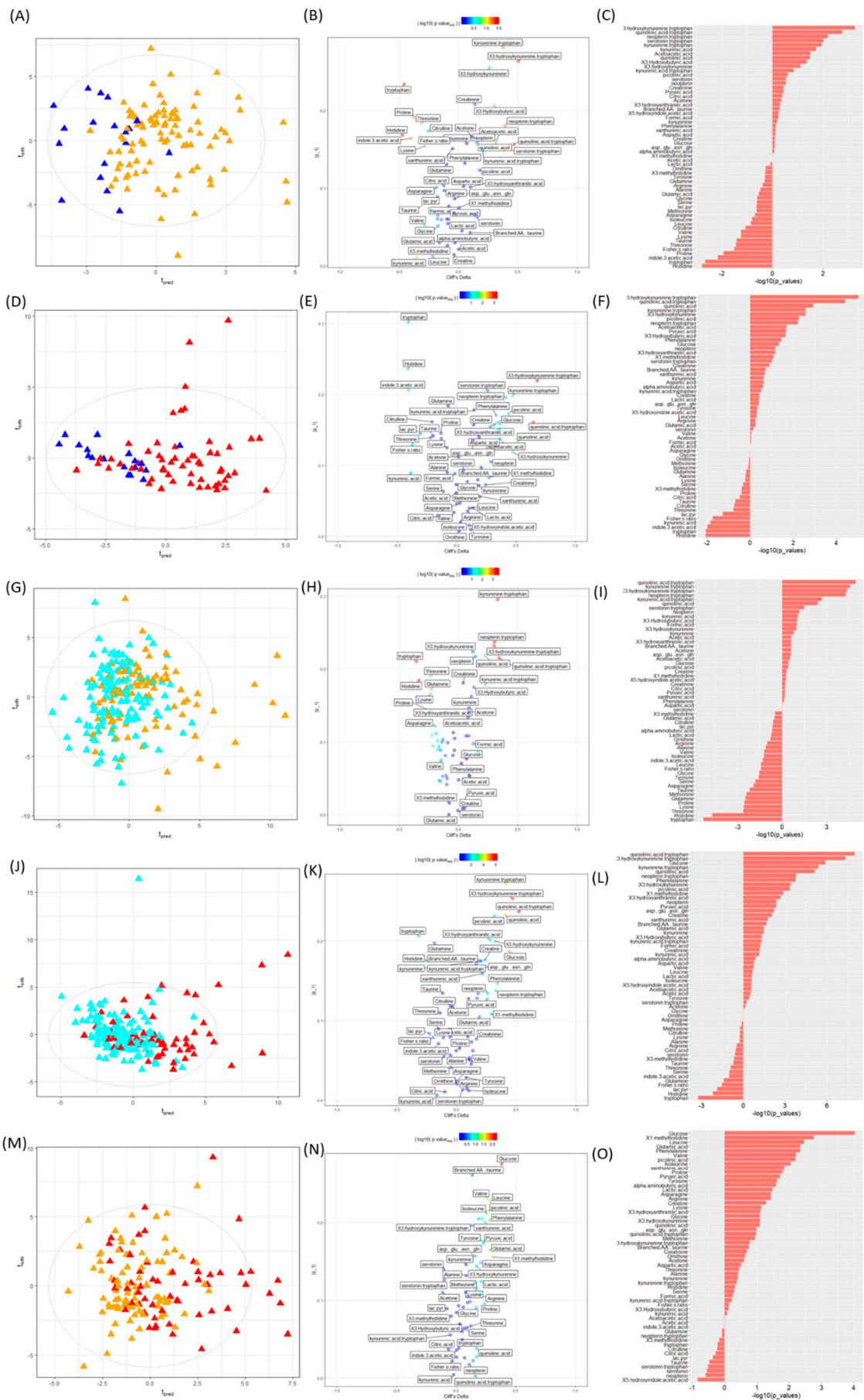
Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.282	1.00	3.01x10 <sup>-29</sup>
Asp/Glu:Asn/Gln	0.278	0.94	2.41x10 <sup>-26</sup>
Pyruvic acid	0.254	0.92	2.01x10 <sup>-25</sup>
Neopterin/tryptophan	0.095	0.82	8.77x10 <sup>-24</sup>
Lactate/pyruvate	0.189	-0.87	7.12x10 <sup>-23</sup>
Glutamic acid	0.236	0.82	2.67x10 <sup>-20</sup>
Fisher's ratio	0.240	-0.80	7.12x10 <sup>-20</sup>
Histidine	0.222	-0.80	1.72x10 <sup>-19</sup>
Quinolinic acid/tryptophan	0.072	0.79	3.62x10 <sup>-19</sup>
Aspartic acid	0.246	0.74	2.95x10 <sup>-17</sup>
Citrulline	0.212	-0.70	2.68x10 <sup>-15</sup>
Quinolinic acid	0.085	0.69	4.10x10 <sup>-15</sup>
Glutamine	0.213	-0.68	9.50x10 <sup>-15</sup>
Phenylalanine	0.227	0.65	2.36x10 <sup>-13</sup>
Kynurenone/tryptophan	0.173	0.64	2.82x10 <sup>-13</sup>
Neopterin	0.094	0.64	4.01x10 <sup>-13</sup>
Tryptophan	0.168	-0.63	1.04x10 <sup>-12</sup>
3-hydroxykynurenone/tryptophan	0.161	0.57	3.21x10 <sup>-12</sup>
Threonine	0.171	-0.60	1.08x10 <sup>-11</sup>
Methionine	0.116	-0.58	4.99x10 <sup>-11</sup>
Lactic acid	0.158	0.45	5.41x10 <sup>-7</sup>
Acetone	0.104	0.44	7.24x10 <sup>-7</sup>
Kynurenic acid	0.040	-0.44	8.95x10 <sup>-7</sup>
3-hydroxybutyric acid	0.105	0.43	1.33x10 <sup>-6</sup>

Glucose	0.140	0.43	1.59x10 <sup>-6</sup>
Asparagine	0.080	-0.42	2.19x10 <sup>-6</sup>
Indole-3-acetic acid	0.057	-0.36	4.73x10 <sup>-5</sup>
3-hydroxykynurenine	0.135	0.36	6.30x10 <sup>-5</sup>
Lysine	0.084	-0.35	1.03x10 <sup>-4</sup>
Proline	0.094	-0.34	1.54x10 <sup>-4</sup>
Isoleucine	0.087	-0.34	1.60x10 <sup>-4</sup>
Valine	0.075	-0.32	4.01x10 <sup>-4</sup>
Kynurenine	0.120	0.29	1.40x10 <sup>-3</sup>
Tyrosine	0.047	-0.27	3.02x10 <sup>-3</sup>
Serotonin/tryptophan	0.089	0.27	3.02x10 <sup>-3</sup>
Citric acid	0.100	0.26	4.09x10 <sup>-3</sup>
Creatine	0.052	0.24	8.30x10 <sup>-3</sup>
Taurine	0.032	-0.23	1.00x10 <sup>-2</sup>
Alpha-aminobutyric acid	0.048	-0.23	1.29x10 <sup>-2</sup>

**Table S24 - OPLS loadings, Cliff's delta and adjusted p-values of the low molecular weight metabolites for the comparison of the controls and SARS-CoV-2 positive patients in severity class E.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Formic acid	0.274	1.00	1.38x10 <sup>-22</sup>
Asp/Glu:Asn/Gln	0.207	0.97	2.62x10 <sup>-21</sup>
Pyruvic acid	0.243	0.94	1.43x10 <sup>-20</sup>
Neopterin/tryptophan	0.196	0.84	1.43x10 <sup>-20</sup>
Quinolinic acid/tryptophan	0.133	0.93	3.93x10 <sup>-20</sup>
Lactate/pyruvate	0.167	-0.92	9.10x10 <sup>-20</sup>
Glutamic acid	0.242	0.91	1.79x10 <sup>-19</sup>
Quinolinic acid	0.131	0.84	1.04x10 <sup>-16</sup>
Fisher's ratio	0.207	-0.82	4.71x10 <sup>-16</sup>
Phenylalanine	0.248	0.82	5.29x10 <sup>-16</sup>
Aspartic acid	0.246	0.82	5.32x10 <sup>-16</sup>
3-hydroxykynurenine/tryptophan	0.155	0.74	6.96x10 <sup>-16</sup>
Histidine	0.210	-0.75	8.42x10 <sup>-14</sup>
Glucose	0.183	0.75	8.35x10 <sup>-12</sup>
Neopterin	0.173	0.75	1.01x10 <sup>-13</sup>
Kynurenine/tryptophan	0.179	0.75	1.01x10 <sup>-13</sup>
Citrulline	0.187	-0.67	3.78x10 <sup>-11</sup>

Glutamine	0.204	-0.61	$1.25 \times 10^{-9}$
Lactic acid	0.172	0.58	$7.98 \times 10^{-9}$
Tryptophan	0.192	-0.58	$7.98 \times 10^{-9}$
3-hydroxykynurenine	0.098	0.57	$1.82 \times 10^{-8}$
3-Hydroxybutyric acid	0.087	0.56	$3.21 \times 10^{-8}$
Threonine	0.145	-0.53	$1.31 \times 10^{-7}$
Kynurenic acid	0.060	-0.44	$2.81 \times 10^{-5}$
1-methylhistidine	0.120	0.41	$5.82 \times 10^{-5}$
Creatine	0.104	0.41	$8.15 \times 10^{-5}$
Methionine	0.073	-0.38	$2.02 \times 10^{-4}$
Kynurenine	0.116	0.36	$4.63 \times 10^{-4}$
Indole-3-acetic acid	0.044	-0.35	$6.06 \times 10^{-4}$
3-hydroxyanthranilic acid	0.112	0.32	$2.35 \times 10^{-3}$
Creatinine	0.093	0.29	$5.29 \times 10^{-3}$
Picolinic acid	0.107	0.29	$5.29 \times 10^{-3}$
Acetone	0.065	0.26	$1.32 \times 10^{-2}$
Ornithine	0.084	0.23	$3.55 \times 10^{-2}$



**Figure S12 - Orthogonal Partial Least Squared (O-PLS-DA), eruption plots and  $-\log_{10}$  (p-values) between different severity classes of SARS-CoV-2 positive patients for the low molecular weight metabolites.** (A) O-PLS-DA of SARS-CoV-2 severity group B (blue triangles) and SARS-CoV-2 severity group D (orange triangles) ( $R_2X=0.11$ , AUROC=0.76). (B) Eruption plot of group B vs group D. (C)  $-\log_{10}$  (p-values) of group B vs group D. (D) O-PLS-DA of SARS-CoV-2 severity group B (blue) and SARS-CoV-2 severity group E (red) ( $R_2X=0.09$ , AUROC=0.78). (E) Eruption plot of group B vs group E. (F)  $-\log_{10}$  (p-values) of group B vs group E. (G) O-PLS-DA of SARS-CoV-2 severity group C (cyan) and SARS-CoV-2 severity group D (orange) ( $R_2X=0.15$ , AUROC=0.65). (H) Eruption plot of group C vs group D. (I)  $-\log_{10}$  (p-values) of group C vs group D. (J) O-PLS-DA of SARS-CoV-2 severity group C (cyan) and SARS-CoV-2 severity group E (red) ( $R_2X=0.13$ , AUROC=0.74). (K) Eruption plot of group C vs group E. (L)  $-\log_{10}$  (p-values) of group C vs group E. (M) O-PLS-DA of SARS-CoV-2 severity group D (orange) and SARS-CoV-2 severity group E (red) ( $R_2X=0.12$ , AUROC=0.62). (N) Eruption plot of group D vs group E. (O)  $-\log_{10}$  (p-values) of group D vs group E. Comparison of severity group B vs severity group C produced no model (AUROC=0.58). Eruption plots are formed from the Cliff's delta (abscissa) and O-PLS-DA loadings (ordinate). Coloured by significance.

**Table S25 - OPLS loadings, Cliff's delta and adjusted p-values for the low molecular weight metabolites for the comparison of severity class B and D.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
3-hydroxykynurenine/tryptophan	0.263	0.50	$3.14 \times 10^{-2}$
Tryptophan	0.234	-0.46	$3.22 \times 10^{-2}$
Histidine	0.168	-0.47	$3.22 \times 10^{-2}$
Quinolinic acid/tryptophan	0.156	0.47	$3.22 \times 10^{-2}$

**Table S26 - OPLS loadings, Cliff's delta and adjusted p-values for the low molecular weight metabolites for the comparison of severity class B and E.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
3-hydroxykynurenine/tryptophan	0.220	0.68	$5.40 \times 10^{-4}$
Quinolinic acid/tryptophan	0.161	0.65	$1.14 \times 10^{-3}$
Quinolinic acid	0.133	0.51	$2.29 \times 10^{-2}$
3-hydroxykynurenine	0.119	0.47	$3.05 \times 10^{-2}$
Kynurenine/tryptophan	0.199	0.47	$3.05 \times 10^{-2}$

**Table S27 - OPLS loadings, Cliff's delta and adjusted p-values for the low molecular weight metabolites for the comparison of severity class C and D.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value

Tryptophan	0.232	-0.36	2.68x10 <sup>-4</sup>
Quinolinic acid/tryptophan	0.215	0.34	3.14x10 <sup>-4</sup>
Histidine	0.185	-0.33	3.46x10 <sup>-4</sup>
Kynurenine/tryptophan	0.296	0.33	3.73x10 <sup>-4</sup>
3-hydroxykynurenine/tryptophan	0.232	0.30	4.80x10 <sup>-4</sup>
Neopterin/tryptophan	0.237	0.30	4.80x10 <sup>-4</sup>
Lysine	0.166	-0.23	1.58x10 <sup>-2</sup>
Proline	0.142	-0.23	1.58x10 <sup>-2</sup>
Threonine	0.190	-0.24	1.58x10 <sup>-2</sup>
Kynurenic acid/tryptophan	0.179	0.22	1.58x10 <sup>-2</sup>
Glutamine	0.173	-0.23	1.68x10 <sup>-2</sup>
Quinolinic acid	0.219	0.22	1.78x10 <sup>-2</sup>
Methionine	0.109	-0.23	1.78x10 <sup>-2</sup>
Taurine	0.094	-0.21	2.60x10 <sup>-2</sup>
Asparagine	0.122	-0.20	4.35x10 <sup>-2</sup>

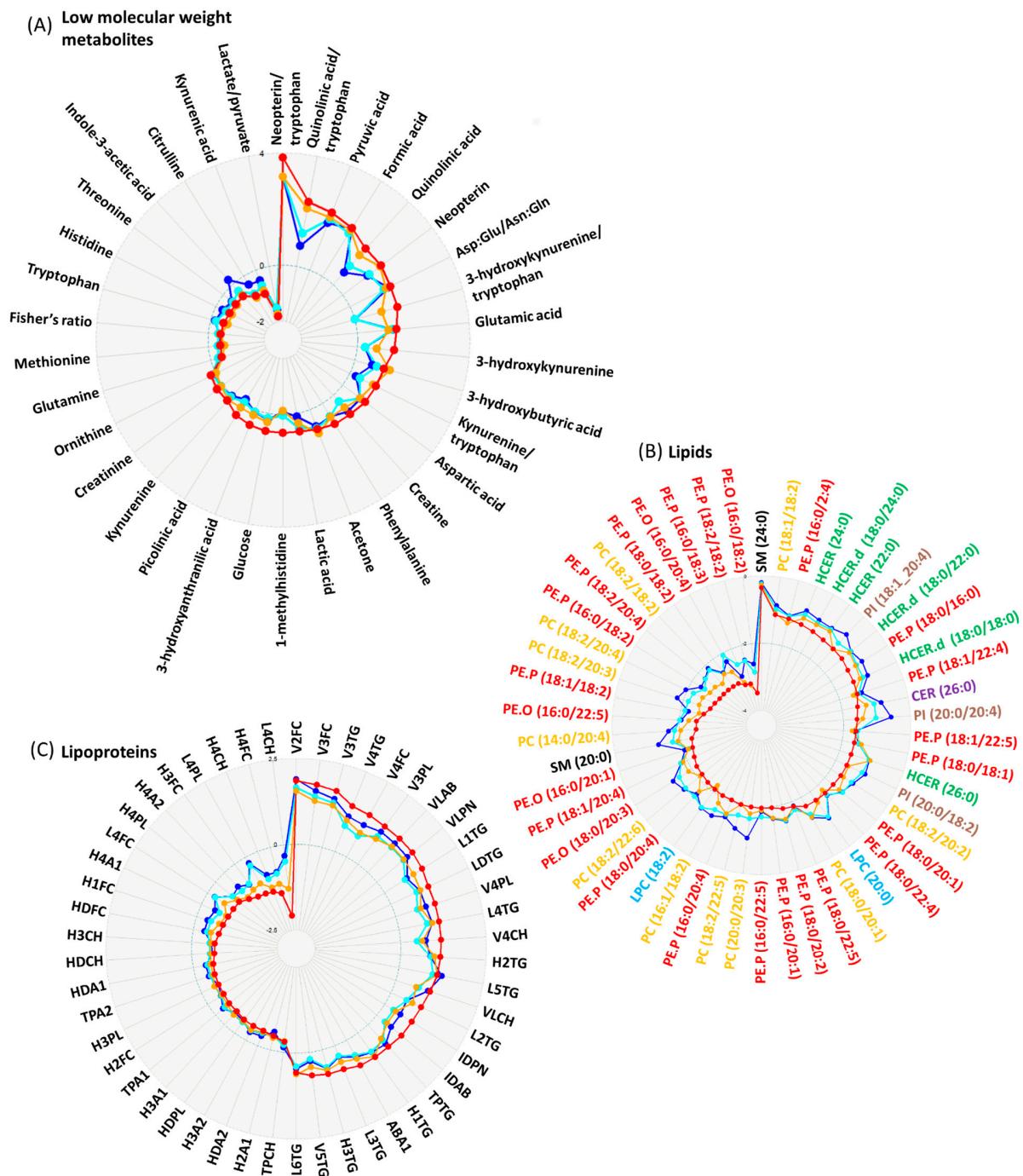
**Table S28 - OPLS loadings, Cliff's delta and adjusted p-values for the low molecular weight metabolites for the comparison of severity class C and E.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Quinolinic acid/tryptophan	0.237	0.52	5.79x10 <sup>-7</sup>
3-hydroxykynurenine/tryptophan	0.253	0.47	1.34x10 <sup>-6</sup>
Glucose	0.187	0.44	2.52x10 <sup>-5</sup>
Kynurenine/tryptophan	0.271	0.42	5.88x10 <sup>-5</sup>
Quinolinic acid	0.234	0.41	8.53x10 <sup>-5</sup>
Phenylalanine	0.147	0.34	1.46x10 <sup>-3</sup>
Neopterin/tryptophan	0.138	0.33	1.46x10 <sup>-3</sup>
3-hydroxykynurenine	0.200	0.32	2.77x10 <sup>-3</sup>
Picolinic acid	0.232	0.32	2.77x10 <sup>-3</sup>
Tryptophan	0.206	-0.31	2.77x10 <sup>-3</sup>
1-methylhistidine	0.108	0.29	6.86x10 <sup>-3</sup>
3-hydroxyanthranilic acid	0.201	0.28	1.01x10 <sup>-2</sup>
Pyruvic acid	0.127	0.26	1.75x10 <sup>-2</sup>
Neopterin	0.135	0.26	1.75x10 <sup>-2</sup>
Asp/Glu:Asn/Gln	0.168	0.25	2.28x10 <sup>-2</sup>
Histidine	0.175	-0.25	2.33x10 <sup>-2</sup>
Creatine	0.176	0.24	2.82x10 <sup>-2</sup>

Lactate/pyruvate	0.084	-0.22	4.32x10 <sup>-2</sup>
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**Table S29 - OPLS loadings, Cliff's delta and adjusted p-values for the low molecular weight metabolites for the comparison of severity class D and E.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
Glucose	0.276	0.38	5.40x10 <sup>-3</sup>



**Figure S13—Radar plots showing the  $\log_2$  fold change between controls and SARS-CoV-2 severity group E patients of the most significant ( $p < 0.05$ ) metabolites, lipids and lipoproteins ordered by  $\log_2$  fold change.** All radar plots are coloured in relation to fold change of each severity class in comparison to the controls. Controls vs group B (blue), controls vs group C severity (cyan), controls vs group D severity (orange) and controls vs group E severity (red). (A) Radar plot of the metabolites with a significant p-value from the controls vs group E severity model for the low molecular weight metabolites. (B) Radar plot of the top 50 most significant lipids from the controls vs group E severity model. (C) Radar plot of the top 50 most significant lipoproteins from the controls vs group E severity model.

**Table S30 - Demographics for the severity group E patients who survived and who did not.**

Patient	Sex	Age	Blood collection date	Date of death	Days between blood collection and death
1	Male	73	01/04/2020	NA	
2	Female	75	01/04/2020	NA	
3	Male	65	01/04/2020	NA	
4	Female	74	01/04/2020	NA	
5	Female	68	01/04/2020	NA	
6	Male	77	01/04/2020	NA	
7	Male	77	01/04/2020	NA	
8	Male	73	01/04/2020	NA	
9	Male	73	01/04/2020	NA	
10	Male	72	01/04/2020	NA	
11	Female	72	01/04/2020	NA	
12	Female	66	01/04/2020	NA	
13	Male	76	01/04/2020	NA	
14	Female	70	01/04/2020	17/04/2020	16
15	Male	77	01/04/2020	14/04/2020	13
16	Male	77	01/04/2020	21/05/2020	50
17	Male	73	01/04/2020	28/04/2020	27
18	Male	73	01/04/2020	10/04/2020	9
19	Female	79	01/04/2020	02/05/2020	31
20	Female	75	01/04/2020	02/05/2020	31
21	Male	70	01/04/2020	01/06/2020	61
22	Female	68	07/04/2020	21/04/2020	14
23	Female	80	09/04/2020	unknown	-
24	Male	66	09/04/2020	17/04/2020	8
25	Female	79	09/04/2020	08/05/2020	29
26	Male	79	09/04/2020	03/05/2020	24

**Table S31 - OPLS loadings, Cliff's delta and adjusted p-values of the combined model (low molecular weight metabolites, lipoproteins and lipids) for the comparison group E severity patients comparing who survived and who died, samples were age matched.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
HCER (16:0)	0.057	0.92	9.15x10 <sup>-3</sup>
HCER (20:0)	0.055	0.92	9.15x10 <sup>-3</sup>
HCER (24:0)	0.063	0.91	9.15x10 <sup>-3</sup>
HCER (24:1)	0.062	0.91	9.15x10 <sup>-3</sup>
HCER (26:0)	0.061	0.95	9.15x10 <sup>-3</sup>
HCER (26:1)	0.063	0.92	9.15x10 <sup>-3</sup>
HCER.d (18:0/24:0)	0.063	0.91	9.15x10 <sup>-3</sup>
HCER.d (18:0/24:1)	0.062	0.94	9.15x10 <sup>-3</sup>
PE.O (18:0/18:1)	0.056	-0.98	9.15x10 <sup>-3</sup>
PE.P (16:0/18:0)	0.053	-0.93	9.15x10 <sup>-3</sup>
HCER (22:0)	0.063	0.89	9.41x10 <sup>-3</sup>
HCER.d (18:0/22:0)	0.063	0.89	9.41x10 <sup>-3</sup>
PE (16:0/18:1)	0.061	-0.88	1.07x10 <sup>-2</sup>
TAG (54:1)_FA (16:0)	0.051	-0.87	1.07x10 <sup>-2</sup>
TAG (54:1)_FA (20:0)	0.053	-0.87	1.07x10 <sup>-2</sup>
TAG (56:1)_FA (16:0)	0.055	-0.87	1.07x10 <sup>-2</sup>
H3FC	0.057	0.83	1.19x10 <sup>-2</sup>
HCER (18:0)	0.052	0.83	1.19x10 <sup>-2</sup>
PE (18:0/18:0)	0.059	-0.85	1.19x10 <sup>-2</sup>
PE (18:0/18:1)	0.058	-0.83	1.19x10 <sup>-2</sup>
PE (18:1/18:1)	0.059	-0.82	1.19x10 <sup>-2</sup>
PE (18:1/18:3)	0.044	-0.82	1.19x10 <sup>-2</sup>
PI (16:0/20:2)	0.040	-0.82	1.19x10 <sup>-2</sup>
SM (26:0)	0.060	0.86	1.19x10 <sup>-2</sup>
TAG (50:1)_FA (16:0)	0.053	-0.82	1.19x10 <sup>-2</sup>
TAG (50:2)_FA (16:0)	0.048	-0.82	1.19x10 <sup>-2</sup>
TAG (52:2)_FA (18:1)	0.055	-0.82	1.19x10 <sup>-2</sup>
TAG (54:2)_FA (20:0)	0.036	-0.82	1.19x10 <sup>-2</sup>
TAG (56:1)_FA (18:1)	0.053	-0.82	1.19x10 <sup>-2</sup>
TAG (56:2)_FA (16:0)	0.054	-0.83	1.19x10 <sup>-2</sup>
TAG (56:2)_FA (20:0)	0.053	-0.83	1.19x10 <sup>-2</sup>
TAG (58:2)_FA (18:1)	0.038	-0.83	1.19x10 <sup>-2</sup>

HCER.d (18:0/20:0)	0.050	0.81	1.25x10 <sup>-2</sup>
PE (16:0/16:1)	0.050	-0.81	1.25x10 <sup>-2</sup>
PE (16:0/18:2)	0.056	-0.81	1.25x10 <sup>-2</sup>
PE (18:0/18:3)	0.048	-0.81	1.25x10 <sup>-2</sup>
PG (18:1/18:1)	0.047	-0.81	1.25x10 <sup>-2</sup>
HDCH	0.050	0.80	1.30x10 <sup>-2</sup>
VLPN	0.050	-0.79	1.30x10 <sup>-2</sup>
VLAB	0.050	-0.79	1.30x10 <sup>-2</sup>
DAG (16:0/16:0)	0.040	-0.79	1.30x10 <sup>-2</sup>
DAG (18:0/18:3)	0.038	-0.79	1.30x10 <sup>-2</sup>
HCER (14:0)	0.049	0.79	1.30x10 <sup>-2</sup>
MAG (18:3)	0.040	-0.79	1.30x10 <sup>-2</sup>
PC (18:1/18:3)	0.048	-0.79	1.30x10 <sup>-2</sup>
PI (18:0/18:3)	0.025	-0.80	1.30x10 <sup>-2</sup>
TAG (50:1)_FA (18:1)	0.056	-0.79	1.30x10 <sup>-2</sup>
TAG (51:2)_FA (16:0)	0.037	-0.80	1.30x10 <sup>-2</sup>
TAG (52:1)_FA (18:1)	0.060	-0.79	1.30x10 <sup>-2</sup>
TAG (55:1)_FA (16:0)	0.048	-0.80	1.30x10 <sup>-2</sup>
TAG (55:1)_FA (18:1)	0.048	-0.80	1.30x10 <sup>-2</sup>
TAG (56:3)_FA (16:0)	0.043	-0.80	1.30x10 <sup>-2</sup>
MAG (18:1)	0.041	-0.78	1.36x10 <sup>-2</sup>
PC (16:0/18:3)	0.044	-0.78	1.36x10 <sup>-2</sup>
PC (18:0/20:4)	0.053	0.78	1.36x10 <sup>-2</sup>
PE (16:0/18:3)	0.048	-0.78	1.36x10 <sup>-2</sup>
TAG (51:1)_FA (16:0)	0.042	-0.78	1.36x10 <sup>-2</sup>
TAG (52:4)_FA (16:0)	0.046	-0.78	1.36x10 <sup>-2</sup>
TAG (53:0)_FA (16:0)	0.039	-0.78	1.36x10 <sup>-2</sup>
TAG (58:3)_FA (18:1)	0.050	-0.78	1.36x10 <sup>-2</sup>
L6FC	0.055	0.75	1.51x10 <sup>-2</sup>
FFA (24:0)	0.041	-0.76	1.51x10 <sup>-2</sup>
TAG (50:5)_FA (18:1)	0.035	-0.76	1.54x10 <sup>-2</sup>
TAG (52:0)_FA (20:0)	0.042	-0.76	1.54x10 <sup>-2</sup>
LPE (18:3)	0.041	-0.75	1.66x10 <sup>-2</sup>
MAG (20:3)	0.040	-0.75	1.66x10 <sup>-2</sup>
PC (18:1/18:1)	0.040	-0.75	1.66x10 <sup>-2</sup>
SM (26:1)	0.043	0.75	1.66x10 <sup>-2</sup>
TAG (52:3)_FA (18:3)	0.031	-0.75	1.66x10 <sup>-2</sup>

TAG (54:0)_FA (16:0)	0.050	-0.75	1.66x10 <sup>-2</sup>
TAG (54:4)_FA (18:1)	0.053	-0.75	1.66x10 <sup>-2</sup>
DAG (16:0/18:0)	0.037	-0.74	1.72x10 <sup>-2</sup>
DAG (16:0/18:1)	0.047	-0.74	1.72x10 <sup>-2</sup>
LPE (18:1)	0.051	-0.74	1.72x10 <sup>-2</sup>
PC (20:0/20:4)	0.052	0.74	1.72x10 <sup>-2</sup>
PE (18:0/20:1)	0.047	-0.74	1.72x10 <sup>-2</sup>
TAG (51:1)_FA (18:1)	0.044	-0.74	1.72x10 <sup>-2</sup>
TAG (52:1)_FA (16:0)	0.057	-0.74	1.72x10 <sup>-2</sup>
TAG (52:2)_FA (20:0)	0.047	-0.74	1.72x10 <sup>-2</sup>
TAG (53:1)_FA (16:0)	0.038	-0.74	1.72x10 <sup>-2</sup>
TAG (54:2)_FA (16:0)	0.046	-0.74	1.72x10 <sup>-2</sup>
TAG (54:5)_FA (18:1)	0.051	-0.74	1.72x10 <sup>-2</sup>
V5TG	0.048	-0.73	1.79x10 <sup>-2</sup>
H1FC	0.054	0.73	1.79x10 <sup>-2</sup>
DCER (26:1)	0.055	0.73	1.79x10 <sup>-2</sup>
MAG (18:0)	0.036	-0.73	1.79x10 <sup>-2</sup>
PC (18:0/18:3)	0.035	-0.73	1.79x10 <sup>-2</sup>
PS (20:0/18:1)	0.041	-0.73	1.79x10 <sup>-2</sup>
TAG (51:1)_FA (17:0)	0.039	-0.73	1.79x10 <sup>-2</sup>
TAG (52:4)_FA (18:3)	0.039	-0.73	1.79x10 <sup>-2</sup>
TAG (54:4)_FA (18:3)	0.038	-0.73	1.79x10 <sup>-2</sup>
TAG (54:5)_FA (18:0)	0.035	-0.73	1.79x10 <sup>-2</sup>
TAG (54:6)_FA (18:1)	0.045	-0.73	1.79x10 <sup>-2</sup>
TAG (57:10)_FA (22:6)	0.034	-0.73	1.79x10 <sup>-2</sup>
TPTG	0.046	-0.72	1.84x10 <sup>-2</sup>
VLCH	0.043	-0.72	1.84x10 <sup>-2</sup>
V4FC	0.050	-0.72	1.84x10 <sup>-2</sup>
CE (16:0)	0.051	0.72	1.84x10 <sup>-2</sup>
DAG (16:0/18:3)	0.036	-0.72	1.84x10 <sup>-2</sup>
HCER.d (18:0/26:1)	0.048	0.72	1.84x10 <sup>-2</sup>
PE (18:1/20:1)	0.044	-0.72	1.84x10 <sup>-2</sup>
PS (18:0/18:1)	0.057	-0.72	1.84x10 <sup>-2</sup>
TAG (49:1)_FA (18:1)	0.032	-0.72	1.84x10 <sup>-2</sup>
TAG (50:3)_FA (18:3)	0.035	-0.72	1.84x10 <sup>-2</sup>
TAG (52:3)_FA (18:1)	0.052	-0.72	1.84x10 <sup>-2</sup>
TAG (52:7)_FA (18:1)	0.034	-0.72	1.84x10 <sup>-2</sup>

TAG (54:2)_FA (18:1)	0.054	-0.72	1.84x10 <sup>-2</sup>
TAG (55:5)_FA (18:1)	0.042	-0.72	1.84x10 <sup>-2</sup>
TAG (57:2)_FA (18:1)	0.051	-0.72	1.84x10 <sup>-2</sup>
Branched AA/Taurine	0.036	-0.70	1.94x10 <sup>-2</sup>
HDFC	0.053	0.70	1.94x10 <sup>-2</sup>
DAG (18:1/20:1)	0.051	-0.70	1.94x10 <sup>-2</sup>
SM (20:1)	0.048	0.70	1.94x10 <sup>-2</sup>
TAG (48:1)_FA (18:1)	0.040	-0.70	1.94x10 <sup>-2</sup>
TAG (50:4)_FA (16:0)	0.031	-0.70	1.94x10 <sup>-2</sup>
TAG (50:5)_FA (16:0)	0.033	-0.70	1.94x10 <sup>-2</sup>
TAG (52:6)_FA (16:0)	0.033	-0.70	1.94x10 <sup>-2</sup>
TAG (52:6)_FA (18:1)	0.035	-0.70	1.94x10 <sup>-2</sup>
TAG (53:1)_FA (18:1)	0.040	-0.70	1.94x10 <sup>-2</sup>
TAG (53:2)_FA (16:0)	0.044	-0.70	1.94x10 <sup>-2</sup>
TAG (54:2)_FA (20:1)	0.048	-0.70	1.94x10 <sup>-2</sup>
TAG (54:3)_FA (18:3)	0.032	-0.70	1.94x10 <sup>-2</sup>
TAG (54:7)_FA (18:1)	0.038	-0.70	1.94x10 <sup>-2</sup>
LDCH	0.054	0.69	2.04x10 <sup>-2</sup>
V4TG	0.043	-0.69	2.04x10 <sup>-2</sup>
MAG (20:2)	0.029	-0.69	2.04x10 <sup>-2</sup>
PE (16:0/22:5)	0.049	-0.69	2.04x10 <sup>-2</sup>
PE (18:0/18:2)	0.050	-0.69	2.04x10 <sup>-2</sup>
PE (18:1/18:2)	0.046	-0.69	2.04x10 <sup>-2</sup>
PI (16:0/20:3)	0.035	-0.69	2.04x10 <sup>-2</sup>
TAG (48:4)_FA (18:1)	0.042	-0.69	2.04x10 <sup>-2</sup>
TAG (49:0)_FA (16:0)	0.031	-0.69	2.04x10 <sup>-2</sup>
TAG (49:0)_FA (17:0)	0.030	-0.69	2.04x10 <sup>-2</sup>
TAG (49:1)_FA (16:0)	0.033	-0.69	2.04x10 <sup>-2</sup>
TAG (49:3)_FA (16:0)	0.023	-0.69	2.04x10 <sup>-2</sup>
TAG (51:2)_FA (17:0)	0.026	-0.69	2.04x10 <sup>-2</sup>
TAG (52:1)_FA (18:0)	0.055	-0.69	2.04x10 <sup>-2</sup>
TAG (52:3)_FA (18:2)	0.040	-0.69	2.04x10 <sup>-2</sup>
Taurine	0.033	0.68	2.18x10 <sup>-2</sup>
L6PN	0.046	0.68	2.18x10 <sup>-2</sup>
V2FC	0.041	-0.68	2.18x10 <sup>-2</sup>
V4PL	0.046	-0.68	2.18x10 <sup>-2</sup>
L6AB	0.046	0.68	2.18x10 <sup>-2</sup>

MAG (22:6)	0.032	-0.68	2.18x10 <sup>-2</sup>
PG (20:0/18:2)	0.052	0.68	2.18x10 <sup>-2</sup>
TAG (51:0)_FA (16:0)	0.034	-0.68	2.18x10 <sup>-2</sup>
TAG (52:1)_FA (20:0)	0.033	-0.68	2.18x10 <sup>-2</sup>
TAG (52:5)_FA (16:0)	0.037	-0.68	2.18x10 <sup>-2</sup>
TAG (53:2)_FA (17:0)	0.046	-0.68	2.18x10 <sup>-2</sup>
TAG (53:2)_FA (18:1)	0.050	-0.68	2.18x10 <sup>-2</sup>
TAG (54:5)_FA (18:3)	0.039	-0.68	2.18x10 <sup>-2</sup>
TAG (56:3)_FA (20:0)	0.047	-0.68	2.18x10 <sup>-2</sup>
H4FC	0.052	0.67	2.22x10 <sup>-2</sup>
IDTG	0.039	-0.67	2.31x10 <sup>-2</sup>
V3FC	0.040	-0.67	2.31x10 <sup>-2</sup>
CE (22:4)	0.049	0.67	2.31x10 <sup>-2</sup>
DAG (16:0/20:5)	0.024	-0.67	2.31x10 <sup>-2</sup>
LPE (18:2)	0.038	-0.67	2.31x10 <sup>-2</sup>
MAG (16:0)	0.037	-0.67	2.31x10 <sup>-2</sup>
MAG (18:2)	0.040	-0.67	2.31x10 <sup>-2</sup>
MAG (22:4)	0.037	-0.67	2.31x10 <sup>-2</sup>
PI (16:0/20:4)	0.036	-0.67	2.31x10 <sup>-2</sup>
TAG (48:4)_FA (16:0)	0.029	-0.67	2.31x10 <sup>-2</sup>
TAG (51:1)_FA (18:0)	0.029	-0.67	2.31x10 <sup>-2</sup>
TAG (51:2)_FA (16:1)	0.032	-0.67	2.31x10 <sup>-2</sup>
TAG (51:4)_FA (18:3)	0.025	-0.67	2.31x10 <sup>-2</sup>
TAG (53:4)_FA (16:0)	0.029	-0.67	2.31x10 <sup>-2</sup>
TAG (54:2)_FA (18:0)	0.055	-0.67	2.31x10 <sup>-2</sup>
L6CH	0.048	0.66	2.46x10 <sup>-2</sup>
H1CH	0.040	0.66	2.46x10 <sup>-2</sup>
DAG (18:0/18:1)	0.051	-0.66	2.46x10 <sup>-2</sup>
MAG (20:4)	0.036	-0.66	2.46x10 <sup>-2</sup>
PC (16:1/18:1)	0.037	-0.66	2.46x10 <sup>-2</sup>
PE (16:0/20:1)	0.050	-0.66	2.46x10 <sup>-2</sup>
PE (18:0/16:0)	0.045	-0.66	2.46x10 <sup>-2</sup>
PE (18:1/20:3)	0.046	-0.66	2.46x10 <sup>-2</sup>
PG (20:0/20:1)	0.047	0.66	2.46x10 <sup>-2</sup>
TAG (48:1)_FA (16:0)	0.039	-0.66	2.46x10 <sup>-2</sup>
TAG (49:1)_FA (16:1)	0.025	-0.66	2.46x10 <sup>-2</sup>
TAG (49:2)_FA (18:1)	0.027	-0.66	2.46x10 <sup>-2</sup>

TAG (50:2)_FA (18:1)	0.047	-0.66	2.46x10 <sup>-2</sup>
TAG (51:2)_FA (18:1)	0.043	-0.66	2.46x10 <sup>-2</sup>
TAG (52:5)_FA (22:5)	0.029	-0.66	2.46x10 <sup>-2</sup>
TAG (53:1)_FA (17:0)	0.036	-0.66	2.46x10 <sup>-2</sup>
TAG (54:4)_FA (16:0)	0.037	-0.66	2.46x10 <sup>-2</sup>
TAG (56:3)_FA (18:2)	0.040	-0.66	2.46x10 <sup>-2</sup>
$\alpha$ -aminobutyric acid	0.029	-0.64	2.46x10 <sup>-2</sup>
Isoleucine	0.023	-0.64	2.60x10 <sup>-2</sup>
DAG (18:1/20:5)	0.026	-0.64	2.60x10 <sup>-2</sup>
MAG (22:5)	0.037	-0.64	2.60x10 <sup>-2</sup>
PC (16:0/18:1)	0.037	-0.64	2.60x10 <sup>-2</sup>
PI (18:0/20:5)	0.023	-0.64	2.60x10 <sup>-2</sup>
PI (18:1/18:1)	0.031	-0.64	2.60x10 <sup>-2</sup>
TAG (46:3)_FA (16:0)	0.030	-0.64	2.60x10 <sup>-2</sup>
TAG (48:1)_FA (14:0)	0.036	-0.64	2.60x10 <sup>-2</sup>
TAG (48:2)_FA (16:0)	0.022	-0.64	2.60x10 <sup>-2</sup>
TAG (49:2)_FA (16:0)	0.022	-0.64	2.60x10 <sup>-2</sup>
TAG (50:0)_FA (18:0)	0.035	-0.64	2.60x10 <sup>-2</sup>
TAG (50:2)_FA (16:1)	0.043	-0.64	2.60x10 <sup>-2</sup>
TAG (52:0)_FA (16:0)	0.045	-0.64	2.60x10 <sup>-2</sup>
TAG (52:1)_FA (20:1)	0.039	-0.64	2.60x10 <sup>-2</sup>
TAG (52:2)_FA (20:2)	0.034	-0.64	2.60x10 <sup>-2</sup>
TAG (52:3)_FA (20:2)	0.025	-0.64	2.60x10 <sup>-2</sup>
TAG (53:1)_FA (18:0)	0.036	-0.64	2.60x10 <sup>-2</sup>
TAG (54:1)_FA (18:0)	0.053	-0.64	2.60x10 <sup>-2</sup>
TAG (54:1)_FA (18:1)	0.053	-0.64	2.60x10 <sup>-2</sup>
TAG (56:7)_FA (20:5)	0.032	-0.64	2.60x10 <sup>-2</sup>
VLFC	0.038	-0.63	2.70x10 <sup>-2</sup>
V5FC	0.042	-0.63	2.70x10 <sup>-2</sup>
H2TG	0.049	-0.63	2.70x10 <sup>-2</sup>
CE (20:3)	0.039	0.63	2.70x10 <sup>-2</sup>
DAG (14:0/18:3)	0.029	-0.63	2.70x10 <sup>-2</sup>
DAG (16:0/20:4)	0.033	-0.63	2.70x10 <sup>-2</sup>
DAG (18:0/18:2)	0.034	-0.63	2.70x10 <sup>-2</sup>
DAG (18:1/18:1)	0.043	-0.63	2.70x10 <sup>-2</sup>
PE (18:0/20:2)	0.047	-0.63	2.70x10 <sup>-2</sup>
PE.P (16:0/22:6)	0.034	0.63	2.70x10 <sup>-2</sup>

PE.P (18:0/20:4)	0.040	0.63	2.70x10 <sup>-2</sup>
PI (16:0/18:1)	0.041	-0.63	2.70x10 <sup>-2</sup>
TAG (47:1)_FA (16:0)	0.025	-0.63	2.70x10 <sup>-2</sup>
TAG (49:0)_FA (18:0)	0.024	-0.63	2.70x10 <sup>-2</sup>
TAG (49:3)_FA (18:3)	0.021	-0.63	2.70x10 <sup>-2</sup>
TAG (50:0)_FA (16:0)	0.041	-0.63	2.70x10 <sup>-2</sup>
TAG (50:3)_FA (16:0)	0.029	-0.63	2.70x10 <sup>-2</sup>
TAG (51:3)_FA (18:3)	0.023	-0.63	2.70x10 <sup>-2</sup>
TAG (52:7)_FA (16:0)	0.030	-0.63	2.70x10 <sup>-2</sup>
TAG (54:3)_FA (16:0)	0.036	-0.63	2.70x10 <sup>-2</sup>
TAG (54:3)_FA (18:1)	0.051	-0.63	2.70x10 <sup>-2</sup>
TAG (54:3)_FA (20:2)	0.038	-0.63	2.70x10 <sup>-2</sup>
TAG (54:8)_FA (18:2)	0.031	-0.63	2.70x10 <sup>-2</sup>
TAG (56:2)_FA (18:0)	0.044	-0.63	2.70x10 <sup>-2</sup>
TAG (56:4)_FA (16:0)	0.036	-0.63	2.70x10 <sup>-2</sup>
TAG (56:7)_FA (18:1)	0.041	-0.63	2.70x10 <sup>-2</sup>
Tyrosine	0.032	-0.62	2.90x10 <sup>-2</sup>
V5CH	0.042	-0.62	2.90x10 <sup>-2</sup>
L3FC	0.047	0.62	2.90x10 <sup>-2</sup>
CE (20:0)	0.046	0.62	2.90x10 <sup>-2</sup>
FFA (20:0)	0.038	-0.62	2.90x10 <sup>-2</sup>
LPE (20:1)	0.046	-0.62	2.90x10 <sup>-2</sup>
PC (14:0/18:1)	0.038	-0.62	2.90x10 <sup>-2</sup>
PC (18:1/20:2)	0.037	-0.62	2.90x10 <sup>-2</sup>
PC (18:2/18:3)	0.042	-0.62	2.90x10 <sup>-2</sup>
TAG (45:1)_FA (16:0)	0.025	-0.62	2.90x10 <sup>-2</sup>
TAG (48:1)_FA (16:1)	0.035	-0.62	2.90x10 <sup>-2</sup>
TAG (52:2)_FA (14:0)	0.028	-0.62	2.90x10 <sup>-2</sup>
TAG (52:3)_FA (20:3)	0.035	-0.62	2.90x10 <sup>-2</sup>
TAG (52:5)_FA (20:5)	0.030	-0.62	2.90x10 <sup>-2</sup>
TAG (53:4)_FA (18:3)	0.026	-0.62	2.90x10 <sup>-2</sup>
TAG (54:1)_FA (20:1)	0.041	-0.62	2.90x10 <sup>-2</sup>
TAG (54:2)_FA (20:2)	0.032	-0.62	2.90x10 <sup>-2</sup>
TAG (54:6)_FA (20:5)	0.031	-0.62	2.90x10 <sup>-2</sup>
TAG (55:2)_FA (18:2)	0.028	-0.62	2.90x10 <sup>-2</sup>
TAG (56:5)_FA (18:1)	0.044	-0.62	2.90x10 <sup>-2</sup>
LDFC	0.051	0.61	3.08x10 <sup>-2</sup>

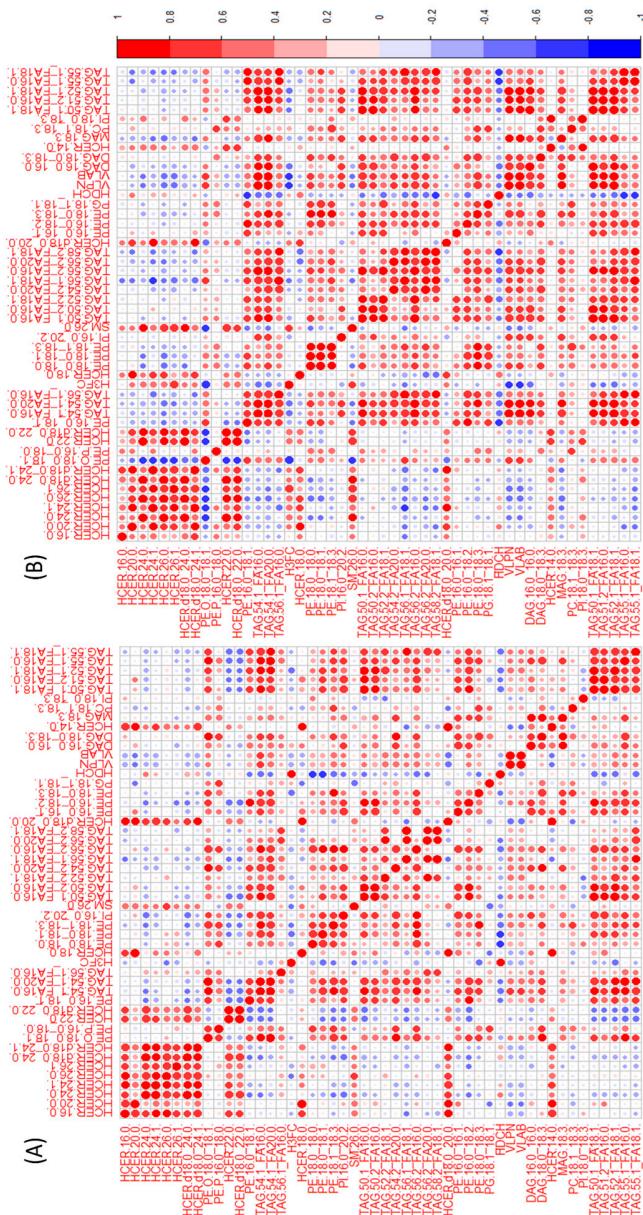
HDA1	0.045	0.61	3.08x10 <sup>-2</sup>
V5PL	0.038	-0.61	3.08x10 <sup>-2</sup>
L6PL	0.047	0.61	3.08x10 <sup>-2</sup>
H4A1	0.045	0.61	3.08x10 <sup>-2</sup>
DAG (16:1/18:0)	0.035	-0.61	3.08x10 <sup>-2</sup>
DAG (18:2/18:3)	0.031	-0.61	3.08x10 <sup>-2</sup>
PE (18:1/22:6)	0.047	-0.61	3.08x10 <sup>-2</sup>
PE.P (16:0/16:1)	0.034	-0.61	3.08x10 <sup>-2</sup>
PE.P (16:0/20:4)	0.042	0.61	3.08x10 <sup>-2</sup>
PI (16:0/22:4)	0.019	-0.61	3.08x10 <sup>-2</sup>
TAG (44:2)_FA (16:0)	0.020	-0.61	3.08x10 <sup>-2</sup>
TAG (44:2)_FA (18:1)	0.025	-0.61	3.08x10 <sup>-2</sup>
TAG (46:3)_FA (18:1)	0.024	-0.61	3.08x10 <sup>-2</sup>
TAG (47:0)_FA (16:0)	0.025	-0.61	3.08x10 <sup>-2</sup>
TAG (52:3)_FA (14:0)	0.024	-0.61	3.08x10 <sup>-2</sup>
TAG (54:4)_FA (20:3)	0.036	-0.61	3.08x10 <sup>-2</sup>
TAG (54:5)_FA (16:0)	0.041	-0.61	3.08x10 <sup>-2</sup>
TAG (54:5)_FA (20:5)	0.029	-0.61	3.08x10 <sup>-2</sup>
TAG (54:5)_FA (22:5)	0.033	-0.61	3.08x10 <sup>-2</sup>
TAG (54:8)_FA (18:3)	0.035	-0.61	3.08x10 <sup>-2</sup>
TAG (56:2)_FA (20:1)	0.047	-0.61	3.08x10 <sup>-2</sup>
TAG (56:3)_FA (18:1)	0.049	-0.61	3.08x10 <sup>-2</sup>
TAG (56:5)_FA (16:0)	0.033	-0.61	3.08x10 <sup>-2</sup>
TPA1	0.047	0.60	3.30x10 <sup>-2</sup>
LDPL	0.049	0.60	3.30x10 <sup>-2</sup>
DAG (16:0/22:5)	0.029	-0.60	3.30x10 <sup>-2</sup>
FFA (20:5)	0.024	-0.60	3.30x10 <sup>-2</sup>
HCER.d (18:0/26:0)	0.036	0.60	3.30x10 <sup>-2</sup>
PC (16:0/16:1)	0.035	-0.60	3.30x10 <sup>-2</sup>
PE (16:0/20:3)	0.045	-0.60	3.30x10 <sup>-2</sup>
PE (16:0/20:4)	0.048	-0.60	3.30x10 <sup>-2</sup>
PE (18:0/22:5)	0.046	-0.60	3.30x10 <sup>-2</sup>
PG (18:2/18:2)	0.028	-0.60	3.30x10 <sup>-2</sup>
PI (16:0/22:5)	0.041	-0.60	3.30x10 <sup>-2</sup>
PI (18:1/18:2)	0.030	-0.60	3.30x10 <sup>-2</sup>
PS (20:0/18:3)	0.034	-0.60	3.30x10 <sup>-2</sup>
SM (18:0)	0.032	0.60	3.30x10 <sup>-2</sup>

SM (22:1)	0.046	0.60	3.30x10 <sup>-2</sup>
TAG (48:0)_FA (16:0)	0.036	-0.60	3.30x10 <sup>-2</sup>
TAG (48:2)_FA (14:0)	0.021	-0.60	3.30x10 <sup>-2</sup>
TAG (48:3)_FA (18:3)	0.025	-0.60	3.30x10 <sup>-2</sup>
TAG (51:0)_FA (18:0)	0.031	-0.60	3.30x10 <sup>-2</sup>
TAG (51:3)_FA (17:0)	0.020	-0.60	3.30x10 <sup>-2</sup>
TAG (54:3)_FA (20:3)	0.037	-0.60	3.30x10 <sup>-2</sup>
TAG (56:3)_FA (18:0)	0.039	-0.60	3.30x10 <sup>-2</sup>
TAG (56:3)_FA (20:2)	0.043	-0.60	3.30x10 <sup>-2</sup>
TAG (56:4)_FA (22:4)	0.030	-0.60	3.30x10 <sup>-2</sup>
CE (22:6)	0.048	0.59	3.59x10 <sup>-2</sup>
DAG (18:1/20:2)	0.039	-0.59	3.59x10 <sup>-2</sup>
DAG (18:2/20:5)	0.024	-0.59	3.59x10 <sup>-2</sup>
PE.O (18:0/18:2)	0.038	-0.59	3.59x10 <sup>-2</sup>
PE.P (18:1/20:4)	0.043	0.59	3.59x10 <sup>-2</sup>
TAG (42:1)_FA (16:0)	0.021	-0.59	3.59x10 <sup>-2</sup>
TAG (47:1)_FA (16:1)	0.023	-0.59	3.59x10 <sup>-2</sup>
TAG (48:3)_FA (16:0)	0.018	-0.59	3.59x10 <sup>-2</sup>
TAG (49:1)_FA (17:0)	0.026	-0.59	3.59x10 <sup>-2</sup>
TAG (49:2)_FA (14:0)	0.023	-0.59	3.59x10 <sup>-2</sup>
TAG (49:2)_FA (16:1)	0.023	-0.59	3.59x10 <sup>-2</sup>
TAG (50:1)_FA (16:1)	0.031	-0.59	3.59x10 <sup>-2</sup>
TAG (50:4)_FA (18:1)	0.032	-0.59	3.59x10 <sup>-2</sup>
TAG (52:1)_FA (16:1)	0.034	-0.59	3.59x10 <sup>-2</sup>
TAG (52:5)_FA (18:1)	0.035	-0.59	3.59x10 <sup>-2</sup>
TAG (54:3)_FA (18:0)	0.047	-0.59	3.59x10 <sup>-2</sup>
TAG (54:5)_FA (20:4)	0.038	-0.59	3.59x10 <sup>-2</sup>
TAG (57:3)_FA (18:2)	0.034	-0.59	3.59x10 <sup>-2</sup>
TAG (58:7)_FA (18:1)	0.038	-0.59	3.59x10 <sup>-2</sup>
H3TG	0.047	-0.58	3.83x10 <sup>-2</sup>
IDPN	0.044	-0.57	3.90x10 <sup>-2</sup>
IDAB	0.044	-0.57	3.90x10 <sup>-2</sup>
V3PL	0.036	-0.57	3.90x10 <sup>-2</sup>
L1TG	0.044	-0.57	3.90x10 <sup>-2</sup>
DAG (18:1/20:4)	0.031	-0.57	3.90x10 <sup>-2</sup>
LPE (16:1)	0.038	-0.57	3.90x10 <sup>-2</sup>
LPE (20:3)	0.037	-0.57	3.90x10 <sup>-2</sup>

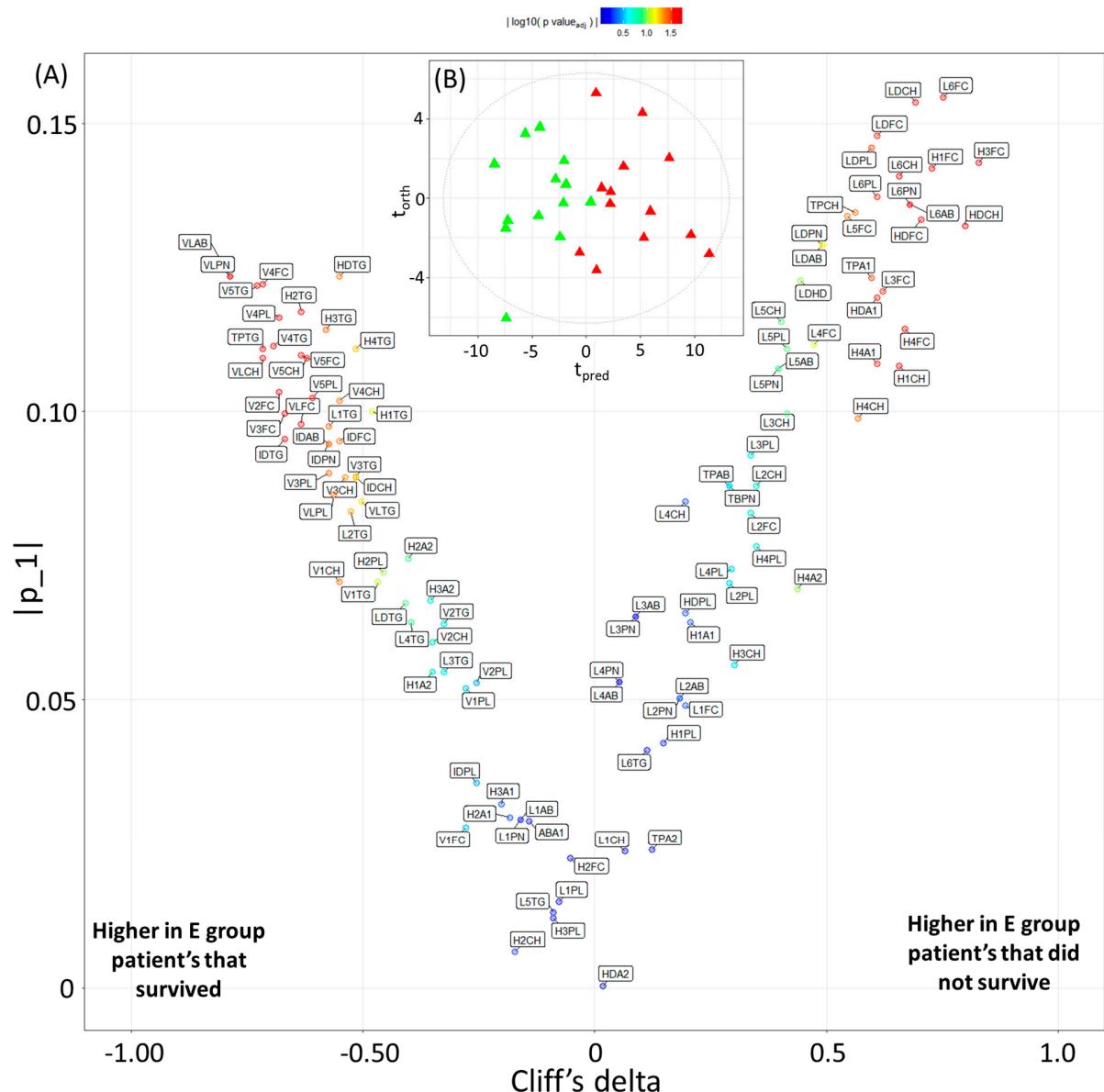
LPE (22:5)	0.043	-0.57	3.90x10 <sup>-2</sup>
MAG (16:1)	0.038	-0.57	3.90x10 <sup>-2</sup>
PE.P (18:1/22:6)	0.040	0.57	3.90x10 <sup>-2</sup>
PI (18:0/18:1)	0.031	-0.57	3.90x10 <sup>-2</sup>
PI (18:1/16:1)	0.036	-0.57	3.90x10 <sup>-2</sup>
PI (18:1/20:3)	0.037	-0.57	3.90x10 <sup>-2</sup>
TAG (46:0)_FA (14:0)	0.028	-0.57	3.90x10 <sup>-2</sup>
TAG (50:1)_FA (20:1)	0.024	-0.57	3.90x10 <sup>-2</sup>
TAG (50:2)_FA (18:2)	0.035	-0.57	3.90x10 <sup>-2</sup>
TAG (51:0)_FA (17:0)	0.032	-0.57	3.90x10 <sup>-2</sup>
TAG (53:3)_FA (17:0)	0.031	-0.57	3.90x10 <sup>-2</sup>
TAG (54:4)_FA (20:4)	0.035	-0.57	3.90x10 <sup>-2</sup>
H4CH	0.043	0.57	4.15x10 <sup>-2</sup>
TPCH	0.046	0.56	4.17x10 <sup>-2</sup>
VLPL	0.034	-0.56	4.17x10 <sup>-2</sup>
DAG (18:0/22:6)	0.028	-0.56	4.17x10 <sup>-2</sup>
DAG (18:1/22:4)	0.036	-0.56	4.17x10 <sup>-2</sup>
DAG (18:1/22:6)	0.030	-0.56	4.17x10 <sup>-2</sup>
LPC (18:3)	0.028	-0.56	4.17x10 <sup>-2</sup>
PE (18:2/18:3)	0.037	-0.56	4.17x10 <sup>-2</sup>
TAG (44:1)_FA (16:0)	0.022	-0.56	4.17x10 <sup>-2</sup>
TAG (46:0)_FA (16:0)	0.028	-0.56	4.17x10 <sup>-2</sup>
TAG (46:1)_FA (16:0)	0.022	-0.56	4.17x10 <sup>-2</sup>
TAG (49:1)_FA (14:0)	0.025	-0.56	4.17x10 <sup>-2</sup>
TAG (50:2)_FA (20:2)	0.023	-0.56	4.17x10 <sup>-2</sup>
TAG (51:2)_FA (18:2)	0.022	-0.56	4.17x10 <sup>-2</sup>
TAG (52:0)_FA (18:0)	0.042	-0.56	4.17x10 <sup>-2</sup>
TAG (52:2)_FA (18:0)	0.042	-0.56	4.17x10 <sup>-2</sup>
TAG (52:4)_FA (18:0)	0.028	-0.56	4.17x10 <sup>-2</sup>
TAG (52:4)_FA (20:4)	0.035	-0.56	4.17x10 <sup>-2</sup>
TAG (52:8)_FA (18:2)	0.032	-0.56	4.17x10 <sup>-2</sup>
TAG (54:2)_FA (18:2)	0.034	-0.56	4.17x10 <sup>-2</sup>
TAG (54:4)_FA (22:4)	0.030	-0.56	4.17x10 <sup>-2</sup>
TAG (54:6)_FA (16:0)	0.034	-0.56	4.17x10 <sup>-2</sup>
TAG (54:6)_FA (18:3)	0.027	-0.56	4.17x10 <sup>-2</sup>
TAG (55:3)_FA (18:1)	0.047	-0.56	4.17x10 <sup>-2</sup>
TAG (56:3)_FA (20:1)	0.047	-0.56	4.17x10 <sup>-2</sup>

TAG (56:5)_FA (22:4)	0.032	-0.56	4.17x10 <sup>-2</sup>
TAG (56:8)_FA (18:1)	0.029	-0.56	4.17x10 <sup>-2</sup>
3-hydroxyanthranilic acid	0.031	-0.55	4.43x10 <sup>-2</sup>
HDTG	0.049	-0.55	4.43x10 <sup>-2</sup>
IDFC	0.043	-0.55	4.43x10 <sup>-2</sup>
V1CH	0.029	-0.55	4.43x10 <sup>-2</sup>
V4CH	0.041	-0.55	4.43x10 <sup>-2</sup>
CE (18:0)	0.039	0.55	4.43x10 <sup>-2</sup>
CE (20:1)	0.035	0.55	4.43x10 <sup>-2</sup>
CE (22:5)	0.044	0.55	4.43x10 <sup>-2</sup>
DAG (16:0/16:1)	0.034	-0.55	4.43x10 <sup>-2</sup>
DAG (18:1/22:5)	0.035	-0.55	4.43x10 <sup>-2</sup>
DCER (22:1)	0.024	-0.55	4.43x10 <sup>-2</sup>
LPE (16:0)	0.035	-0.55	4.43x10 <sup>-2</sup>
PE.O (16:0/20:4)	0.034	0.55	4.43x10 <sup>-2</sup>
PE.P (18:0/22:6)	0.037	0.55	4.43x10 <sup>-2</sup>
PE.P (18:1/18:0)	0.035	-0.55	4.43x10 <sup>-2</sup>
PI (18:1/20:4)	0.030	-0.55	4.43x10 <sup>-2</sup>
SM (24:1)	0.043	0.55	4.43x10 <sup>-2</sup>
TAG (46:2)_FA (16:0)	0.021	-0.55	4.43x10 <sup>-2</sup>
TAG (46:2)_FA (18:1)	0.025	-0.55	4.43x10 <sup>-2</sup>
TAG (47:2)_FA (18:1)	0.024	-0.55	4.43x10 <sup>-2</sup>
TAG (50:4)_FA (18:3)	0.025	-0.55	4.43x10 <sup>-2</sup>
TAG (52:2)_FA (20:1)	0.023	-0.55	4.43x10 <sup>-2</sup>
TAG (52:3)_FA (18:0)	0.032	-0.55	4.43x10 <sup>-2</sup>
TAG (53:2)_FA (18:2)	0.023	-0.55	4.43x10 <sup>-2</sup>
TAG (53:3)_FA (16:0)	0.031	-0.55	4.43x10 <sup>-2</sup>
TAG (55:2)_FA (18:1)	0.048	-0.55	4.43x10 <sup>-2</sup>
TAG (55:4)_FA (18:1)	0.046	-0.55	4.43x10 <sup>-2</sup>
TAG (56:4)_FA (18:0)	0.039	-0.55	4.43x10 <sup>-2</sup>
TAG (56:6)_FA (16:0)	0.033	-0.55	4.43x10 <sup>-2</sup>
TAG (56:6)_FA (18:1)	0.041	-0.55	4.43x10 <sup>-2</sup>
L5FC	0.046	0.54	4.56x10 <sup>-2</sup>
Acetone	0.008	0.53	4.66x10 <sup>-2</sup>
Glucose	0.031	-0.54	4.88x10 <sup>-2</sup>
Xanturenic acid	0.029	-0.54	4.88x10 <sup>-2</sup>
V3CH	0.035	-0.54	4.88x10 <sup>-2</sup>

DAG (16:0/20:3)	0.032	-0.54	$4.88 \times 10^{-2}$
DCER (26:0)	0.044	0.54	$4.88 \times 10^{-2}$
LPE (22:4)	0.036	-0.54	$4.88 \times 10^{-2}$
PC (18:0/18:1)	0.036	-0.54	$4.88 \times 10^{-2}$
PE (18:0/20:3)	0.040	-0.54	$4.88 \times 10^{-2}$
PE (18:0/20:5)	0.031	-0.54	$4.88 \times 10^{-2}$
TAG (45:0)_FA (16:0)	0.023	-0.54	$4.88 \times 10^{-2}$
TAG (48:5)_FA (18:2)	0.031	-0.54	$4.88 \times 10^{-2}$
TAG (50:2)_FA (14:0)	0.040	-0.54	$4.88 \times 10^{-2}$
TAG (50:5)_FA (20:5)	0.028	-0.54	$4.88 \times 10^{-2}$
TAG (52:2)_FA (16:1)	0.041	-0.54	$4.80 \times 10^{-3}$
TAG (54:4)_FA (18:2)	0.036	-0.54	$4.81 \times 10^{-3}$
TAG (54:7)_FA (18:3)	0.028	-0.54	$4.89 \times 10^{-2}$



**Figure S14 - Correlation plot of the (A) survived model and (B) Did not survive model.** The top 50 metabolites are shown according to adjusted p-values.



**Figure S15 – Modeling COVID-19 mortality prediction using only the lipoproteins** (A) Eruption plot of the severity group E patients who survived vs the severity group E patients who died. (B) O-PLS-DA of severity group E patients who survived (green triangles) and severity group E patients who died (red triangles),  $R^2X=0.32$ , AUROC=0.81. Cliff's delta, OPLS loadings values and the adjusted p values for this model can be found in table S30.

**Table S32 - OPLS loadings, Cliff's delta and adjusted p-values of lipoprotein model for the comparison group E severity patients comparing who survived and who died, samples were age matched.** Only metabolites with significant p-values are shown in the table.

Metabolite	OPLS loadings	Cliff's delta	Adjusted p-value
HDCH	0.132	0.80	1.82x10 <sup>-2</sup>
VLPN	0.123	-0.79	1.82x10 <sup>-2</sup>
VLAB	0.123	-0.79	1.82x10 <sup>-2</sup>
H3FC	0.143	0.83	1.82x10 <sup>-2</sup>
L6FC	0.154	0.75	1.98x10 <sup>-2</sup>
TPTG	0.111	-0.72	2.04x10 <sup>-2</sup>
LDCH	0.154	0.69	2.04x10 <sup>-2</sup>
L6PN	0.136	0.68	2.04x10 <sup>-2</sup>
VLCH	0.109	-0.72	2.04x10 <sup>-2</sup>
HDFC	0.133	0.70	2.04x10 <sup>-2</sup>
V4TG	0.111	-0.69	2.04x10 <sup>-2</sup>
V5TG	0.122	-0.73	2.04x10 <sup>-2</sup>
V2FC	0.103	-0.68	2.04x10 <sup>-2</sup>
V4FC	0.122	-0.72	2.04x10 <sup>-2</sup>
V4PL	0.116	-0.68	2.04x10 <sup>-2</sup>
L6AB	0.136	0.68	2.04x10 <sup>-2</sup>
H1FC	0.412	0.73	2.04x10 <sup>-2</sup>
H4FC	0.114	0.67	2.04x10 <sup>-2</sup>
IDTG	0.095	-0.67	2.11x10 <sup>-2</sup>
V3FC	0.100	-0.67	2.11x10 <sup>-2</sup>
L6CH	0.141	0.66	2.16x10 <sup>-2</sup>
H1CH	0.108	0.66	2.25x10 <sup>-2</sup>
VLFC	0.098	-0.63	2.72x10 <sup>-2</sup>
V5FC	0.110	-0.63	2.72x10 <sup>-2</sup>
H2TG	0.117	-0.63	2.72x10 <sup>-2</sup>
LDFC	0.148	0.61	2.89x10 <sup>-2</sup>
HDA1	0.120	0.61	2.89x10 <sup>-2</sup>
V5CH	0.109	-0.62	2.89x10 <sup>-2</sup>
V5PL	0.102	-0.61	2.89x10 <sup>-2</sup>
L3FC	0.121	0.62	2.89x10 <sup>-2</sup>
L6PL	0.137	0.61	2.89x10 <sup>-2</sup>
H4A1	0.108	0.61	2.89x10 <sup>-2</sup>
TPA1	0.123	0.60	3.16x10 <sup>-2</sup>

LDPL	0.146	0.60	$3.16 \times 10^{-2}$
IDPN	0.094	-0.57	$3.70 \times 10^{-2}$
IDAB	0.094	-0.57	$3.70 \times 10^{-2}$
V3PL	0.089	-0.57	$3.70 \times 10^{-2}$
L1TG	0.097	-0.57	$3.70 \times 10^{-2}$
H3TG	0.114	-0.58	$3.70 \times 10^{-2}$
H4CH	0.099	0.57	$3.85 \times 10^{-2}$
TPCH	0.134	0.56	$3.96 \times 10^{-2}$
VLPL	0.086	-0.56	$3.96 \times 10^{-2}$
HDTG	0.123	-0.55	$4.16 \times 10^{-2}$
IDFC	0.095	-0.55	$4.16 \times 10^{-2}$
V1CH	0.071	-0.55	$4.16 \times 10^{-2}$
V4CH	0.102	-0.55	$4.16 \times 10^{-2}$
L5FC	0.135	0.54	$4.20 \times 10^{-2}$
V3CH	0.089	-0.54	$4.58 \times 10^{-2}$

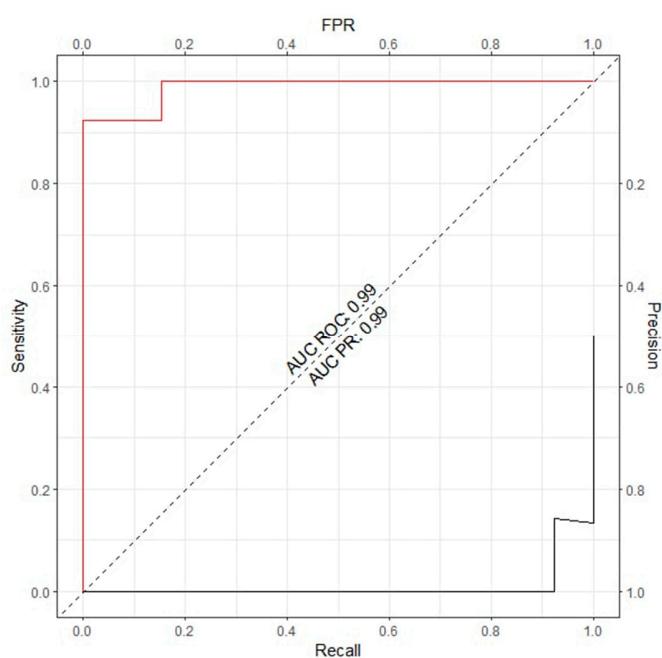


Figure S16 - AUROC showing prediction of mortality (AUCROC = 0.99) built using HCer 16:0 and PEO 18:0\_18:1