

S3 Table. Changes in the serum levels of the significantly-regulated proteins between COPD patients from Cluster 1 and Cluster 2

Protein ID	Fold changes	p value *	References
<i>Up-regulated in Cluster 2 versus Cluster 1</i>			
Tropomyosin 4	3.093	8.13E-29	S1-S3
Carbonic anhydrase XIII	3.005	1.73E-29	S4,S5
BTK	2.824	3.08E-33	S6,S7
Cyclophilin F	2.694	7.51E-29	S8
CSK	2.549	3.06E-36	S9-S11
FER	2.537	1.26E-28	S12,S13
PKC-B-II	2.535	1.19E-42	S14
SP-D	2.532	2.39E-24	S15
VAV	2.439	5.27E-27	S16-S18
Histone H1.2	2.420	9.64E-20	S19-S22
GRB2 adapter protein	2.417	1.97E-26	S23,S24
H2A3	2.344	2.10E-19	S19,S22,S25
IMB1	2.316	6.15E-27	S26
DRG-1	2.284	1.16E-32	S27-S29
SRCN1	2.264	9.30E-21	S30,S31
LYN	2.242	1.72E-29	S32,S33
SMAD2	2.206	3.20E-33	S34, S39, S124, S137
NSF1C	2.196	1.83E-26	S35
LYNB	2.182	4.98E-25	S32,S33
NDP kinase B	2.157	6.04E-29	S36-S38
Caspase-3	2.157	3.50E-34	S39
SBDS	2.112	1.97E-30	S40-S42
PDPK1	2.106	4.36E-31	S43,S44
CK2-A1:B	2.097	2.88E-27	S45,S46
RAC1	2.077	3.08E-33	S47-S50
IF4G2	2.064	5.73E-27	S51,S52
eIF-4H	2.051	1.54E-25	S53
Aflatoxin B1 aldehyde reductase	2.045	6.17E-23	S54
Sphingosine kinase 1	2.045	1.28E-27	S55
GAPDH, liver	2.042	8.88E-18	S56,S57
UBE2N	2.034	4.11E-24	S58,S61
α -synuclein	2.011	8.92E-19	S62-S63
SGTA	2.006	3.64E-20	S64
SNAA	1.991	3.45E-23	S65-S68
Haemoglobin	1.966	1.81E-08	S69, S70
METAP1	1.958	3.20E-33	S71,S72
IMDH1	1.957	1.70E-17	S73,S74
PRKACA	1.956	2.18E-30	S75
Annexin I	1.921	2.88E-32	S76,S77

S3 Table (continued)

PKC-A	1.921	2.44E-19	S78
RPS6KA3	1.916	1.57E-33	S79,S80
Histone H2A.z	1.914	5.43E-20	S19,S21,S22,S81,S82
hnRNP A2/B1	1.909	1.84E-29	S83-S85
UFM1	1.888	8.39E-23	S58,S86-S88
FYN	1.886	9.28E-13	S89,S90
RAN	1.861	1.85E-14	S91,S92
Myokinase	1.859	4.43E-11	S93,S94
H2B2E	1.841	4.02E-18	S19,S21,S22,S95
41	1.837	6.00E-16	S96,S97
BARK1	1.835	6.95E-28	S98,S99
M2-PK	1.825	4.82E-15	S100
14-3-3 protein zeta/delta	1.785	1.68E-33	S101-S104
PPID	1.781	2.29E-13	S105-S107
6-phosphogluconate dehydrogenase	1.776	7.82E-8	S108
HXK2	1.775	3.32E-13	S109,S110
PTP-1C	1.756	5.02E-20	S111,S112
DUSP3	1.731	2.39E-25	S113,S114
TCTP	1.723	1.83E-27	S115-S117
Triosephosphate isomerase	1.711	1.39E-23	S118,S119
DLRB1	1.681	8.07E-19	S120,S121
SUMO3	1.680	3.88E-26	S58,S122,S123
PA2G4	1.671	3.51E-15	S124,S125
14-3-3 protein β/α	1.669	3.42E-28	S101-S103,S126
PPAC	1.667	1.62E-08	S127,S128
Transketolase	1.660	1.28E-24	S129,S130
UBC9	1.651	2.16E-31	S58,S86,S131
FGF16	1.650	3.70E-18	S132,S133
EDAR	1.610	1.56E-10	S134,S135
SHC1	1.603	2.71E-25	S136,S137
ARGI1	1.601	2.58E-12	S138-S140
GPVI	1.599	3.60E-13	S141,S142
Cyclophilin A	1.592	3.21E-33	S143,S144
BAD	1.587	3.87E-13	S101,S145
ERK-1	1.566	2.59E-28	S146,S147
PLPP	1.560	9.71E-15	S148,S149
CPNE1	1.560	8.80E-18	S150,S151
BPI	1.554	1.79E-06	S152-S154
NCC27	1.552	1.81E-27	S155,S156
MK01	1.548	2.46E-16	S93,S94
eIF-5A-1	1.544	2.47E-16	S157-S160
NACA	1.539	5.58E-09	S161,S162
AREG	1.538	1.51E-13	S163-S165

S3 Table (continued)

ATPO	1.536	1.71E-13	S166
CD40 ligand	1.524	2.49E-11	S167-S169
Prostatic binding protein	1.519	1.51E-13	S98,S170-S172
Ubiquitin+1	1.512	3.62E-11	S58
Lactoferrin	1.510	9.58E-09	S173
Azurocidin	1.507	5.01E-10	S174,S175
Sorting nexin 4	1.502	1.39E-13	S176
STAT3	1.502	2.02E-26	S46, S50
<i>Up-regulated in Cluster 1 versus Cluster 2</i>			
Glucagon	0.666	1.36E-07	S177
MMP-12	0.652	2.55E-09	S178,S179
Renin	0.650	6.58E-06	S180
Lactadherin (MFGM)	0.640	8.42E-14	S181-S183
C3b	0.637	3.26E-05	S184
Midkine	0.505	1.63E-17	S185-S187

* False discover rate from group comparisons

Abbreviations are listed in the Table 1 of the main manuscript

Bold indicates the 15 proteins belonging to the short signature

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