

Supplementary Table 4. The top EV proteins (MSA vs. HC) through BOLD Selector from 1272 overlapping EV proteins between Orbitrap Fusion Lumos and Orbitrap Fusion Lumos FAIMS platform analysis.

MSA vs. HC				
Number	Protein ID	Gene symbol	ranking	Missing rate
v1	P04180	LCAT	1	0
v2	P29622	SERPINA4	1	0
v3	P55060	CSE1L	1	0.00666667
v4	P46109	CRKL	4	0.04666667
v5	P02745	C1QA	5	0
v6	P06310	IGKV2-30	5	0
v7	P09871	C1S	5	0
v8	P12429	ANXA3	5	0
v9	P30038	ALDH4A1	5	0.00666667
v10	Q13586	STIM1	5	0.04666667
v11	Q7LBR1	CHMP1B	5	0.03333333
v12	Q86UD1	OAF	5	0.01333333
v13	A0A075B6K5	IGLV3-9	13	0.00666667
v14	O00602	FCN1	13	0
v15	P01591	JCHAIN	13	0
v16	P01859	IGHG2	13	0
v17	P02671	FGA	13	0
v18	P05543	SERPINA7	13	0
v19	P06239	LCK	13	0
v20	P06681	C2	13	0
v21	P08185	SERPINA6	13	0
v22	P08195	SLC3A2	13	0
v23	P20701	ITGAL	13	0
v24	P35613	BSG	13	0
v25	P49908	SELENOP	13	0
v26	P58166	INHBE	13	0
v27	P62330	ARF6	13	0.08
v28	Q14766	LTBP1	13	0