

Figure 1.

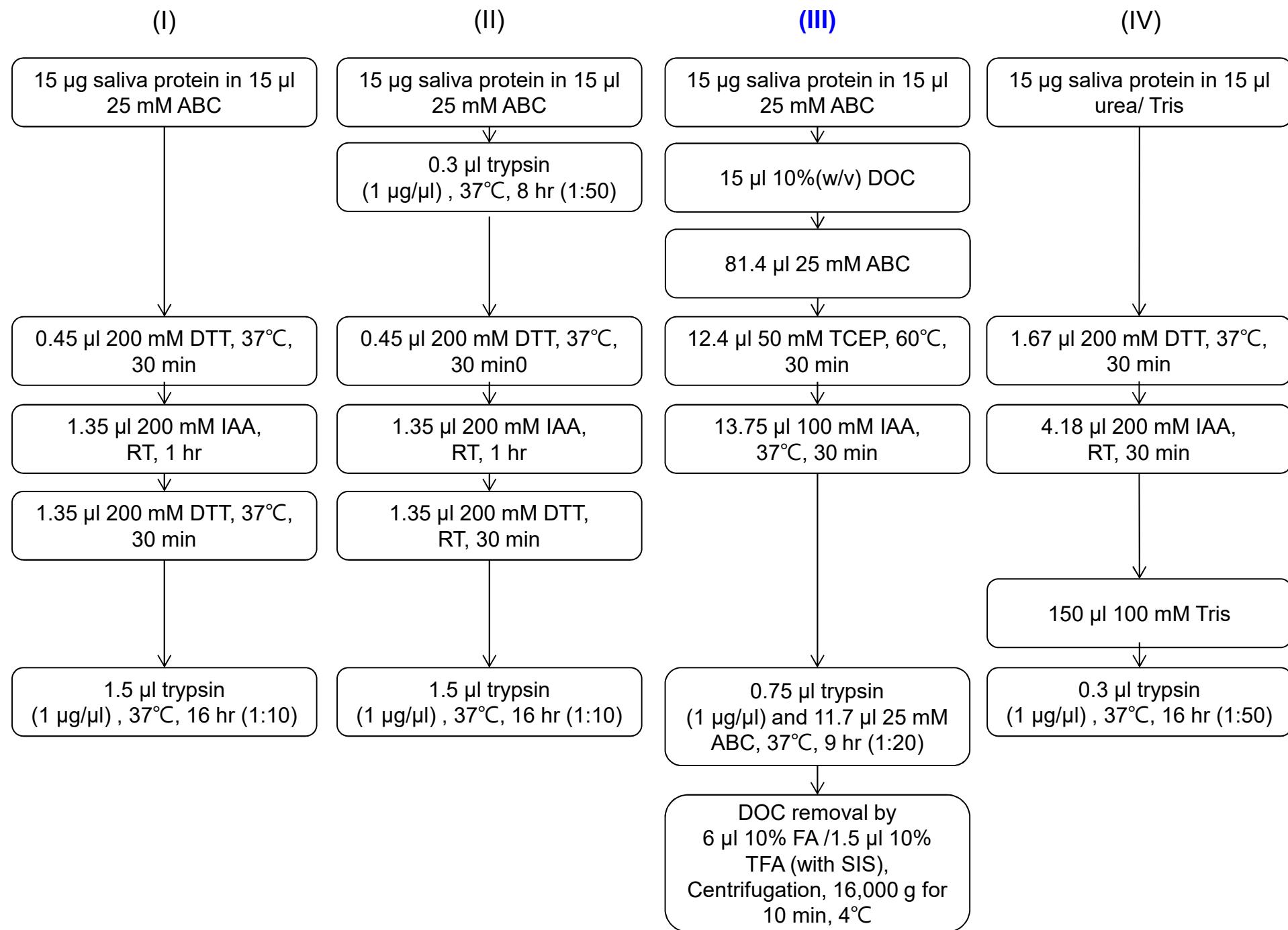


Figure 2.

(A)

Protocol	I	II	III	IV
Best Digestion				
Protein no.	2	8	32	14
CV(%) under 20	17 (30.4%)	17 (30.4%)	39 (69.6%)	22 (39.3%)
CV(%) under 30	24 (42.9%)	22 (39.3%)	43 (76.8%)	37 (66.1%)
Detectable Protein no.	53	54	54	52
Total Protein no.		56		
Hydrophobicity index				
Average	-1.19	-0.67	-0.35	-0.26
SD	0.38	0.35	0.51	0.76
Median	-1.19	-0.56	-0.34	-0.43

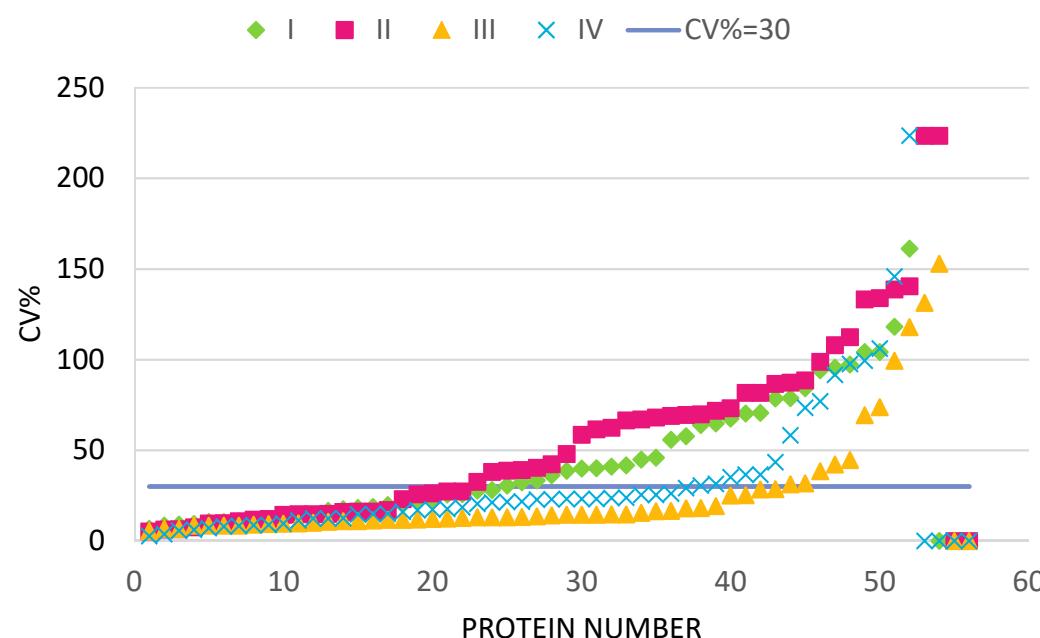


Figure 2.

(B)

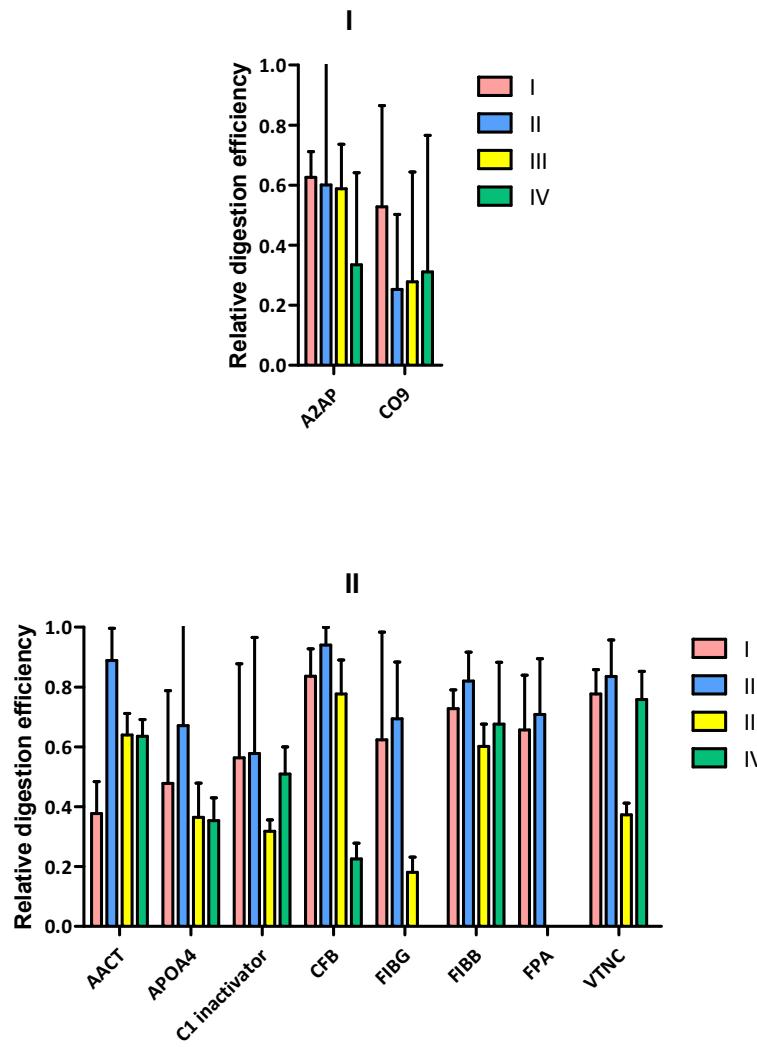


Figure 2.

(C)

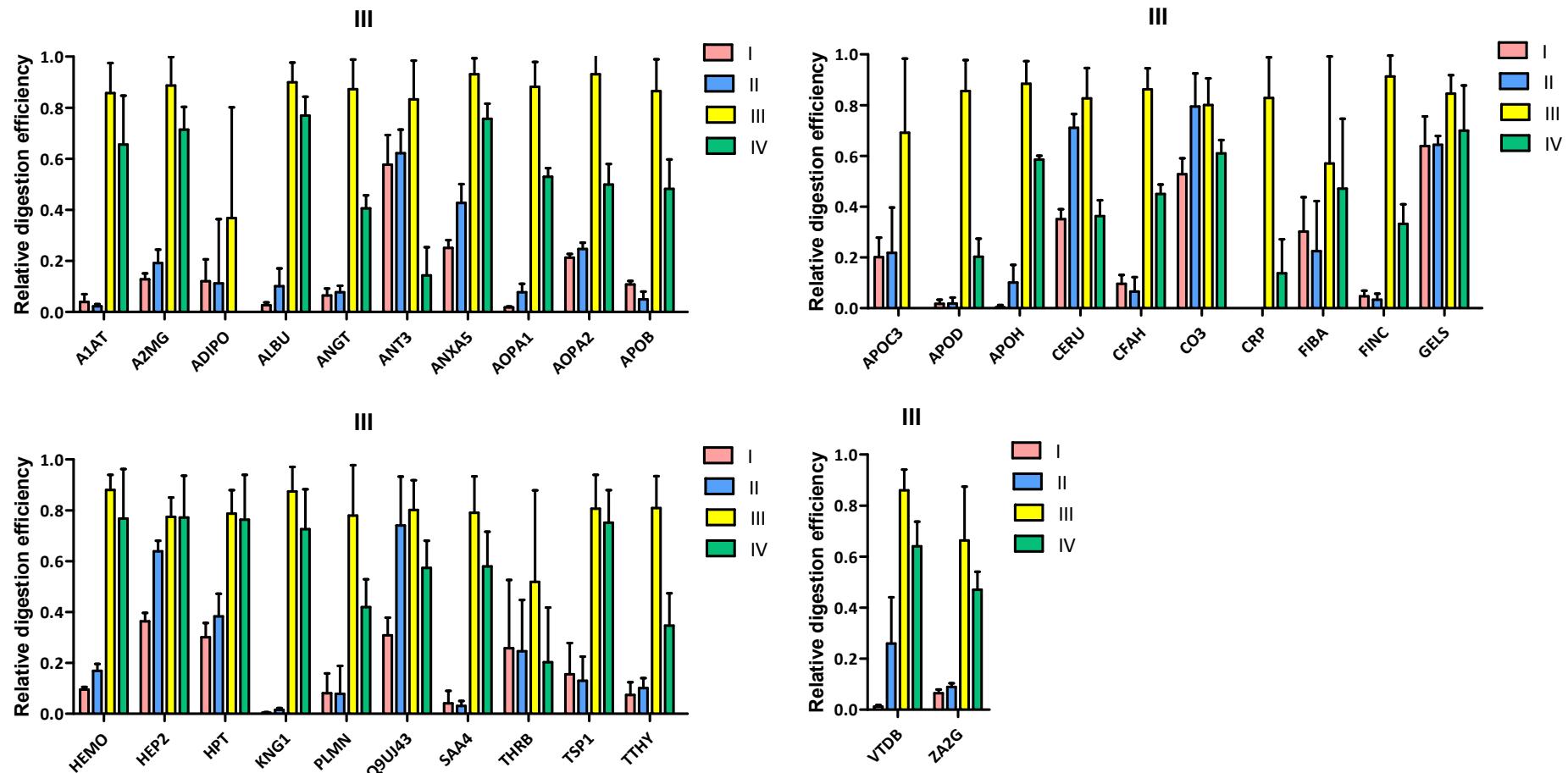


Figure 2.

(D)

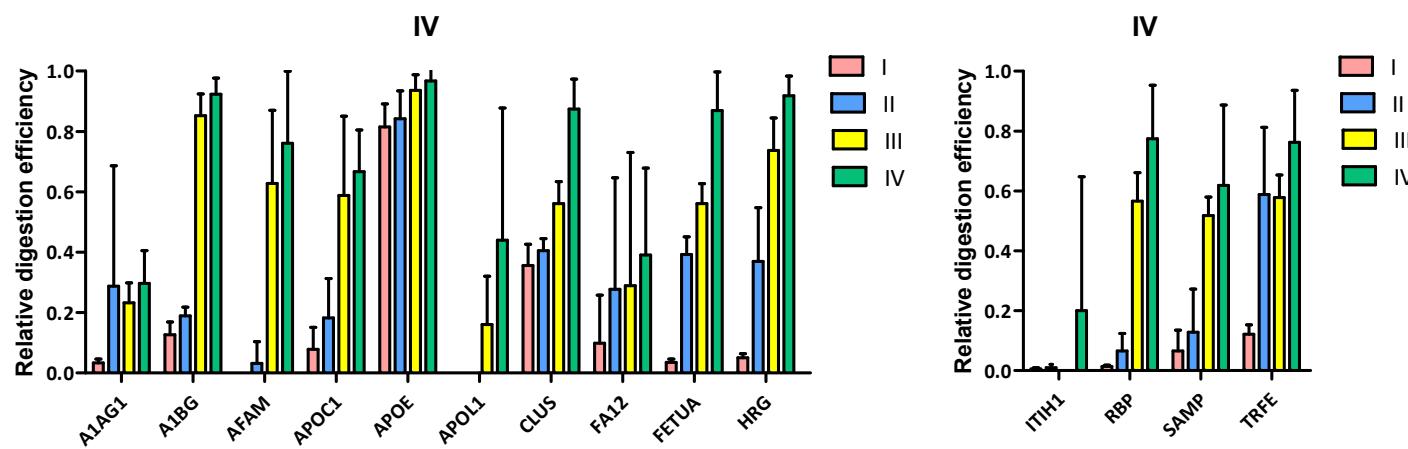


Figure 3(A)

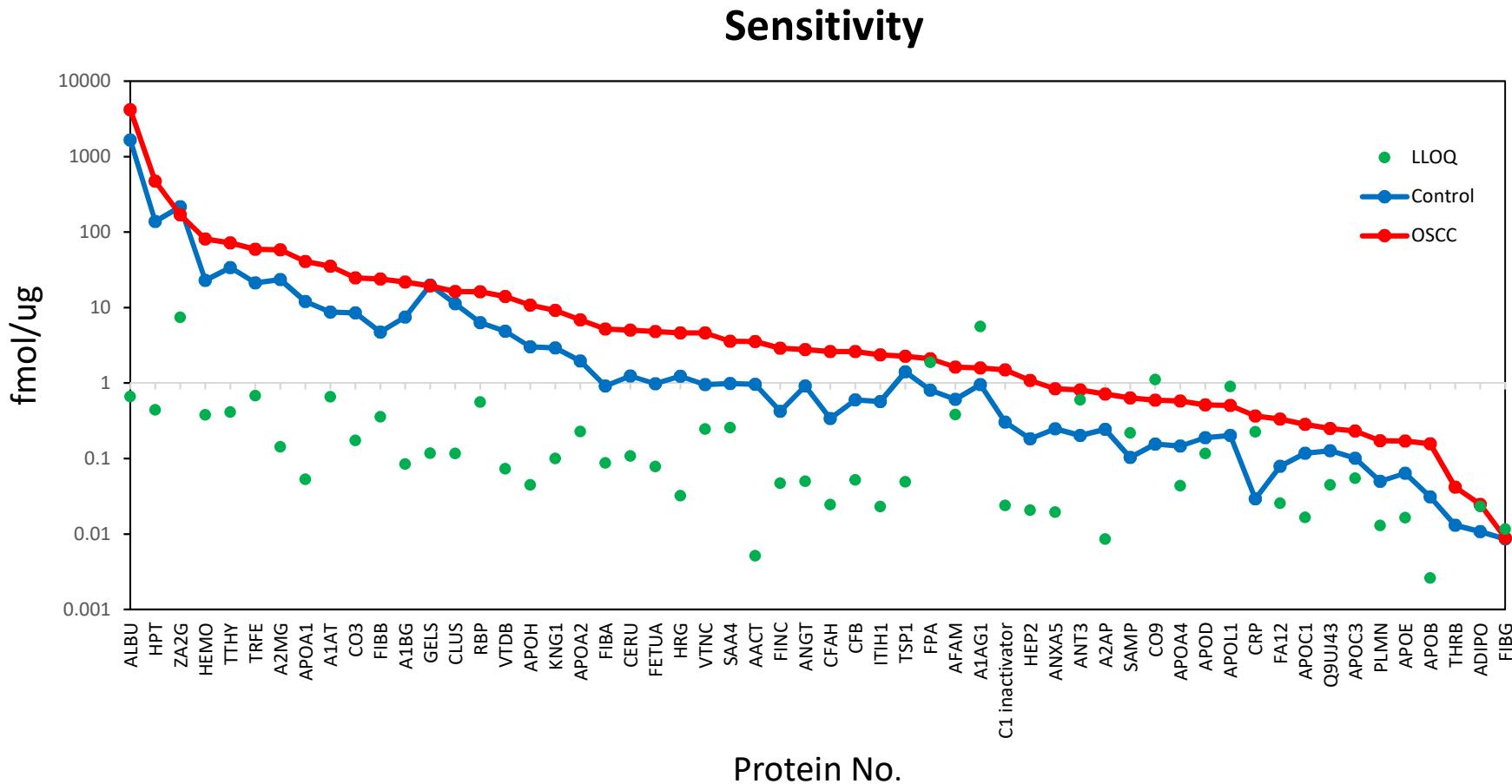
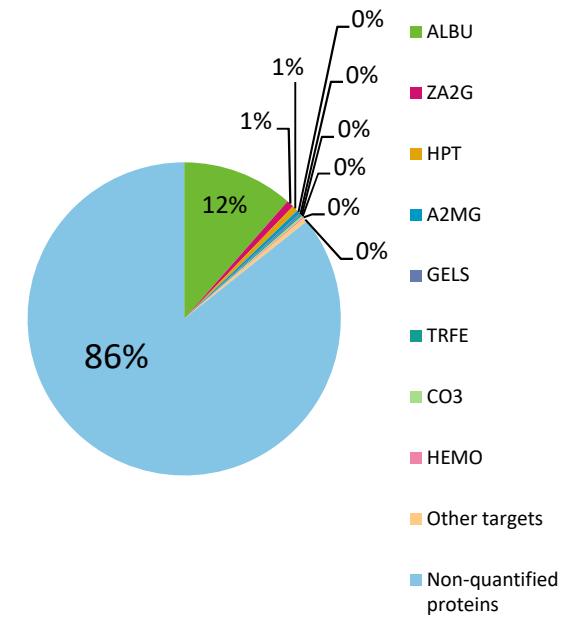
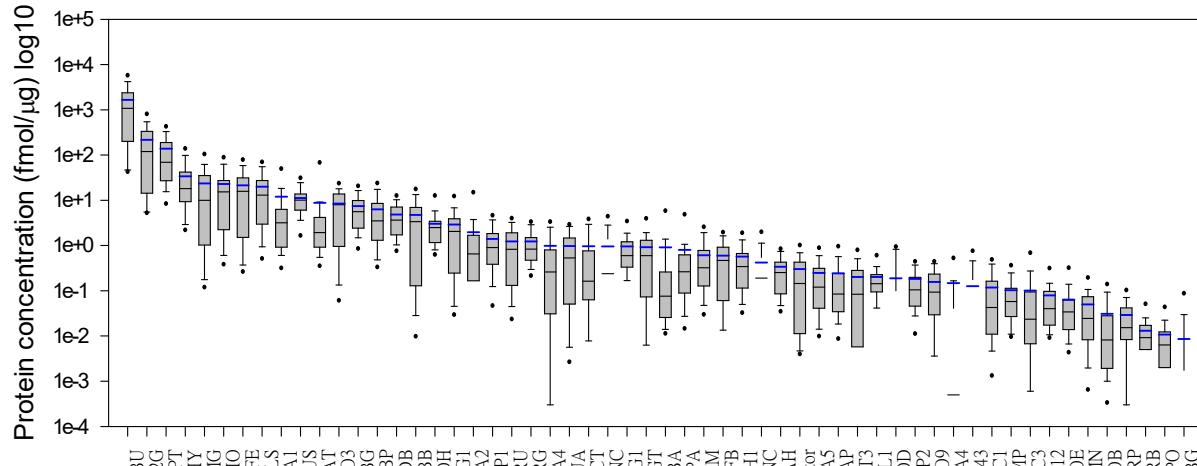


Figure 3(B)

Control (N=58)



OSCC (N=61)

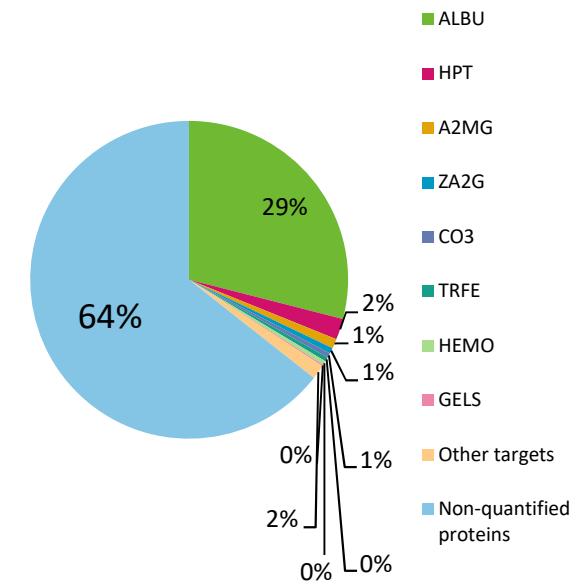
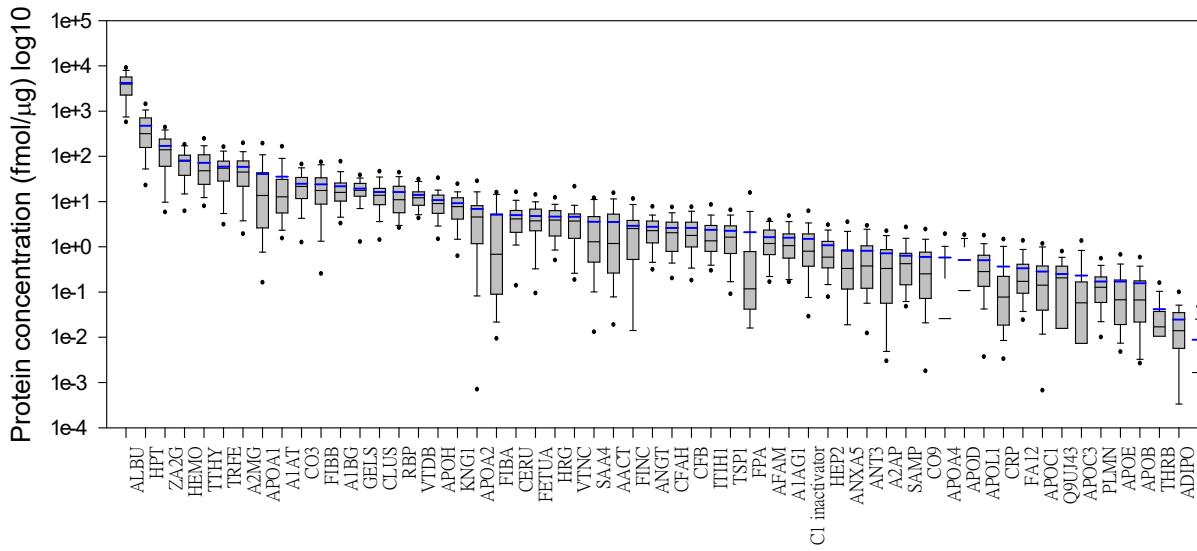


Figure 3(C)

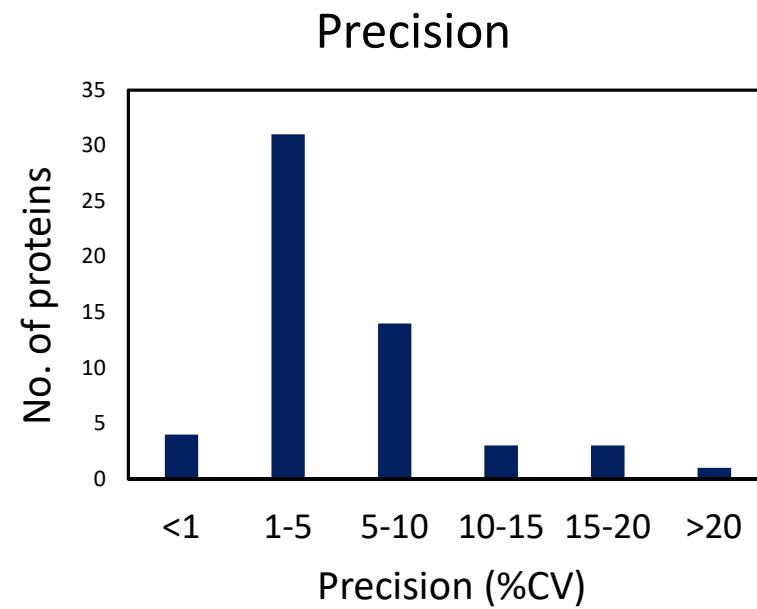


Figure 4(A)

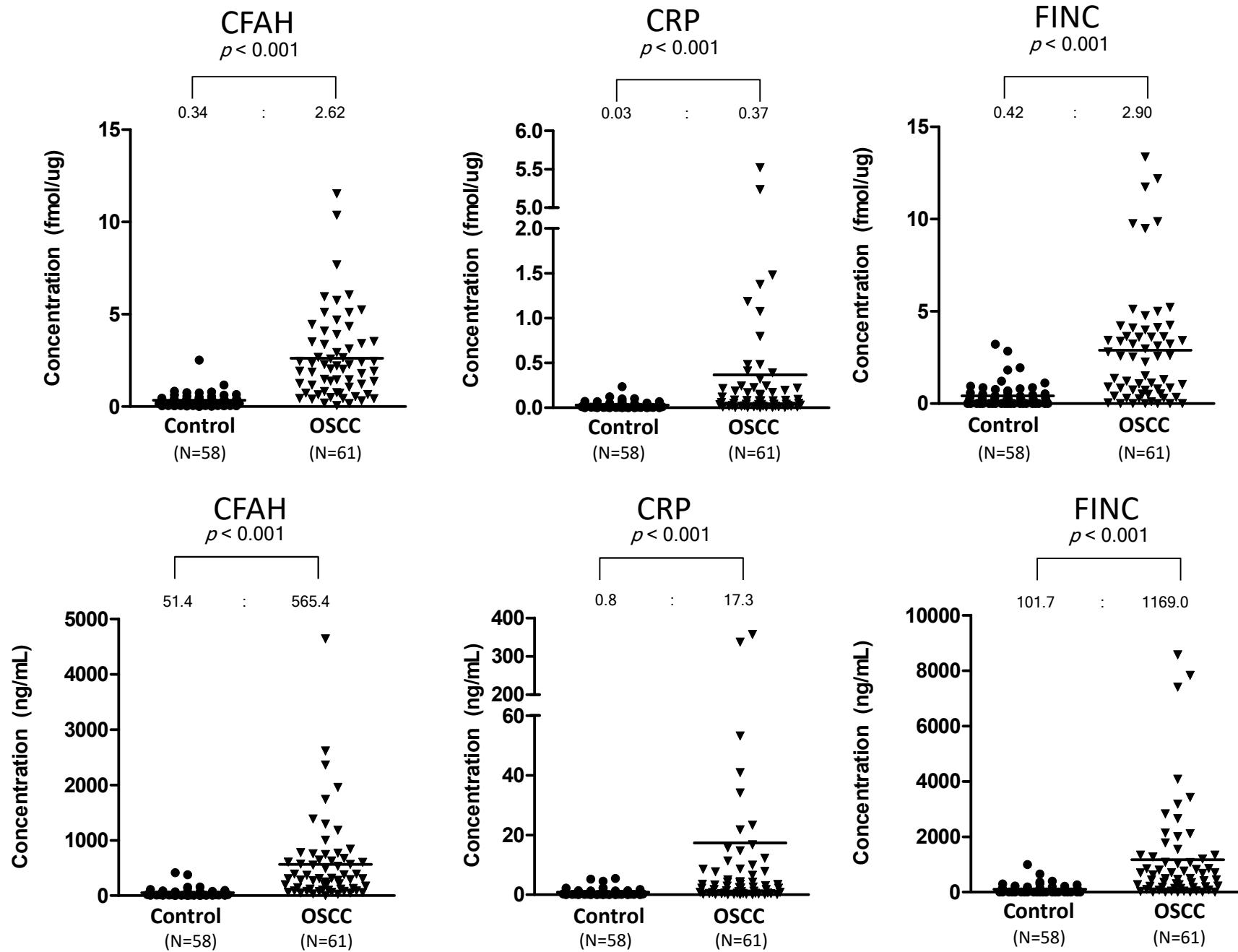


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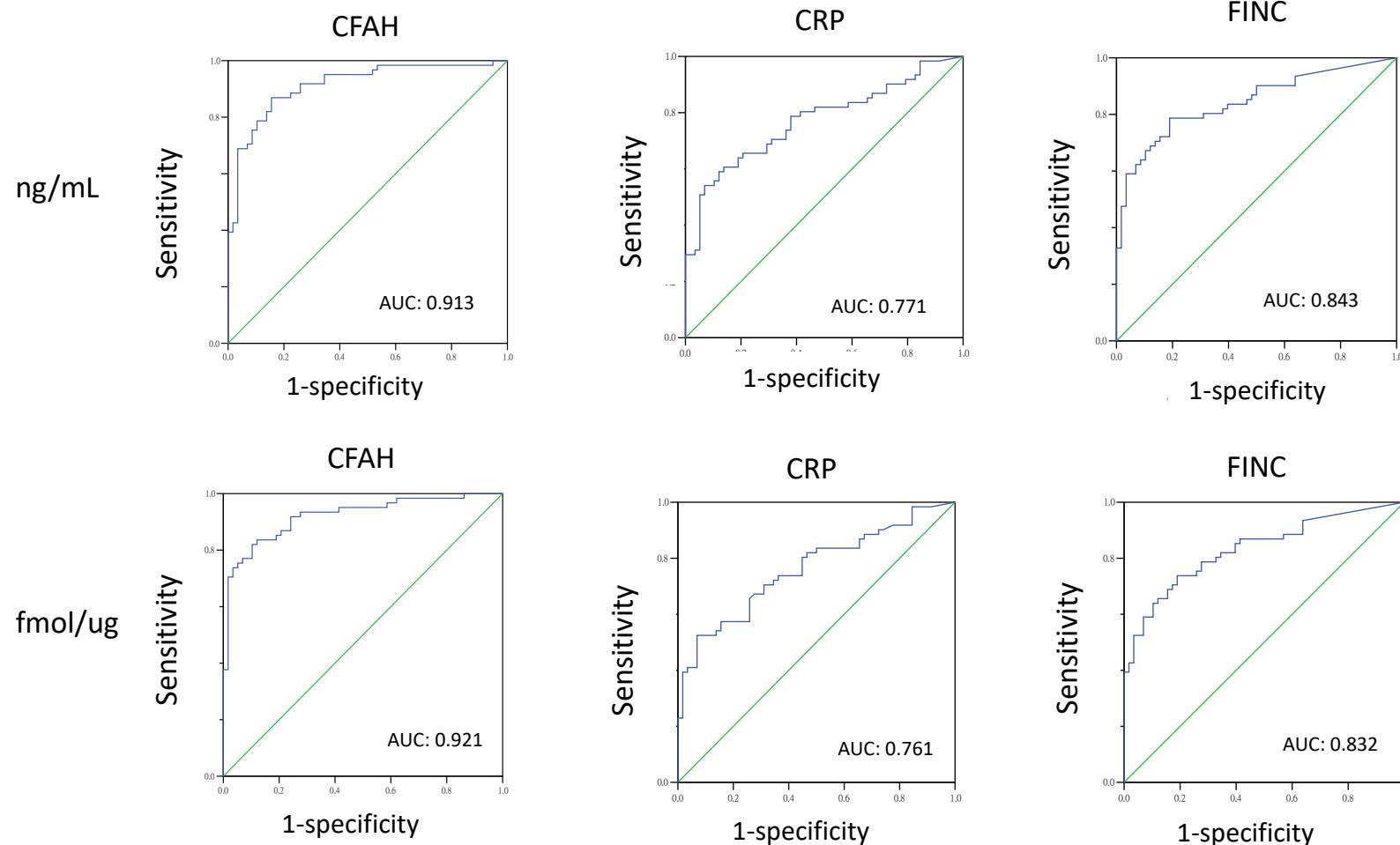
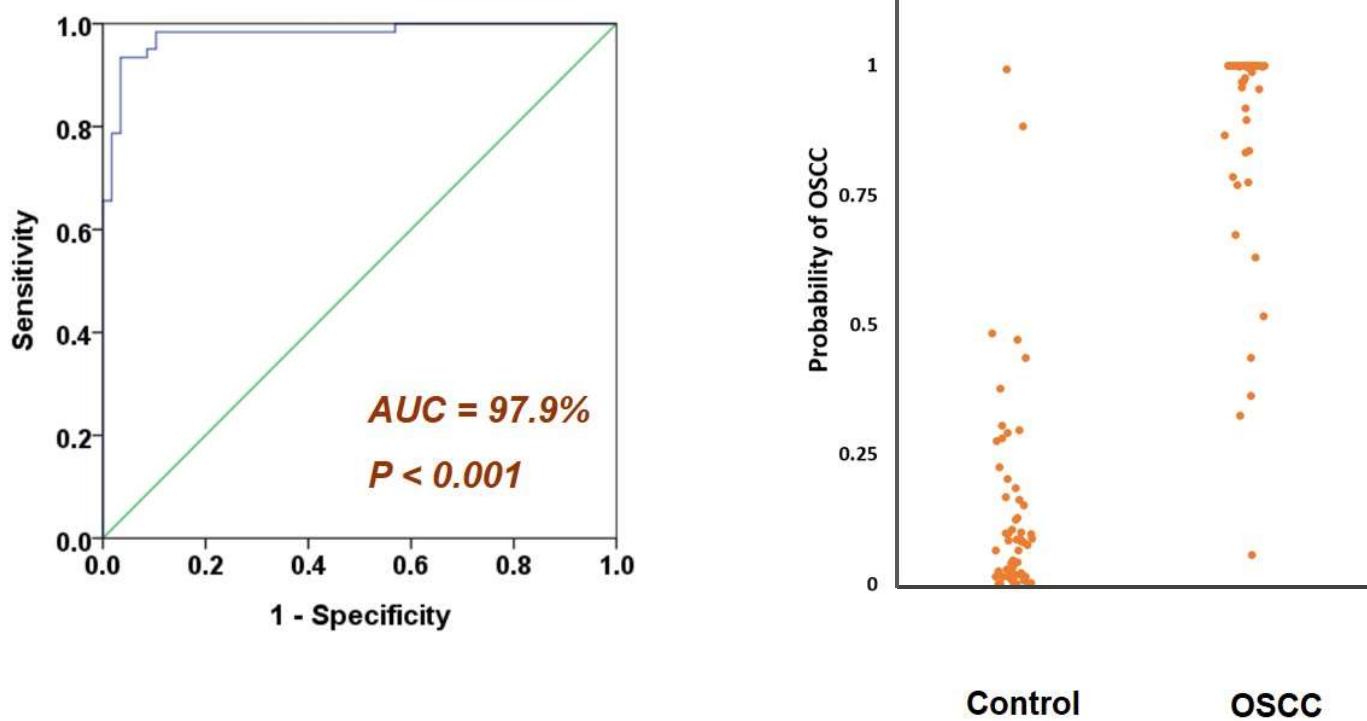
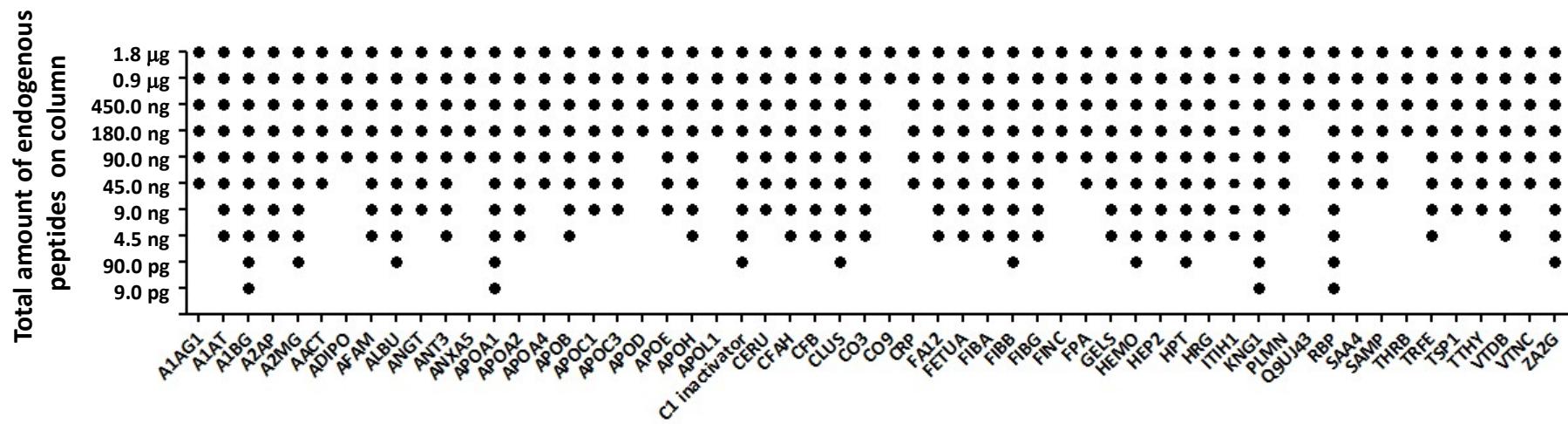


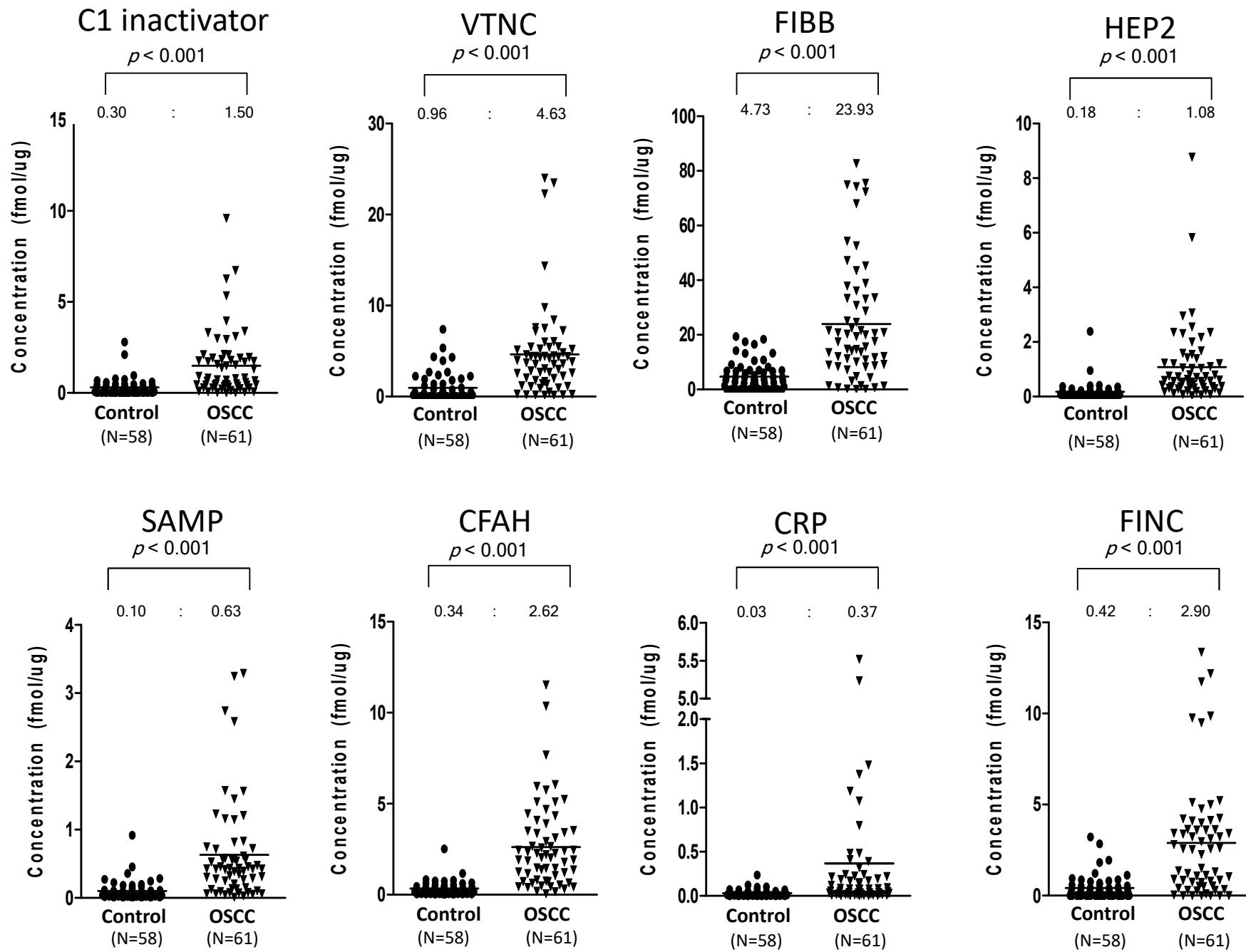
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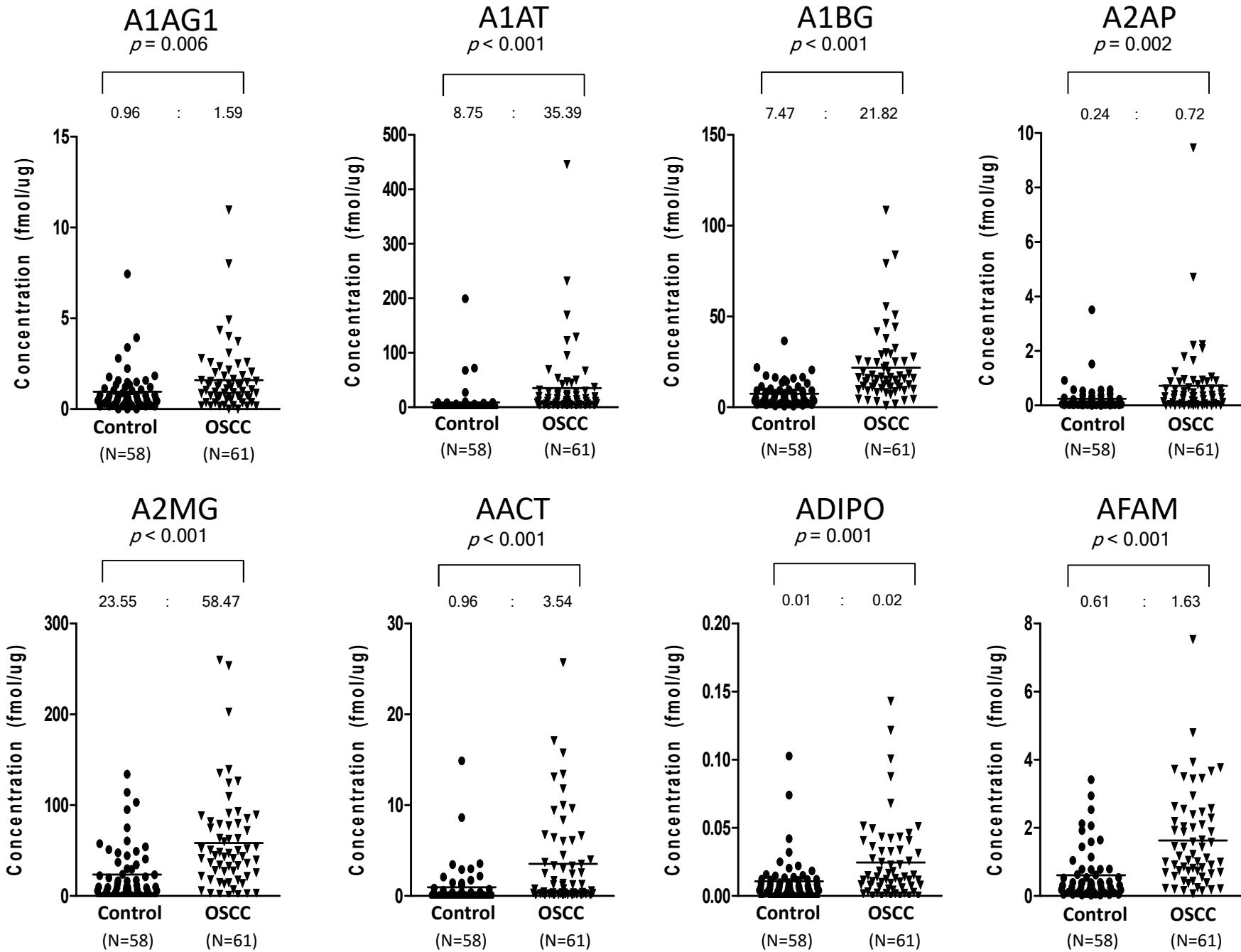


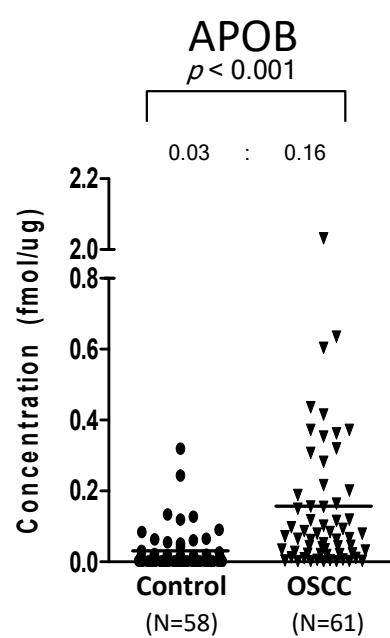
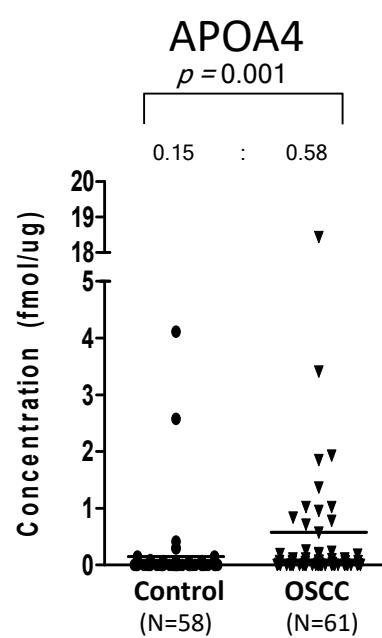
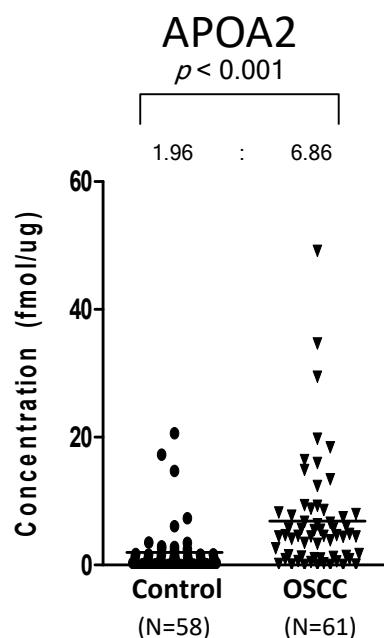
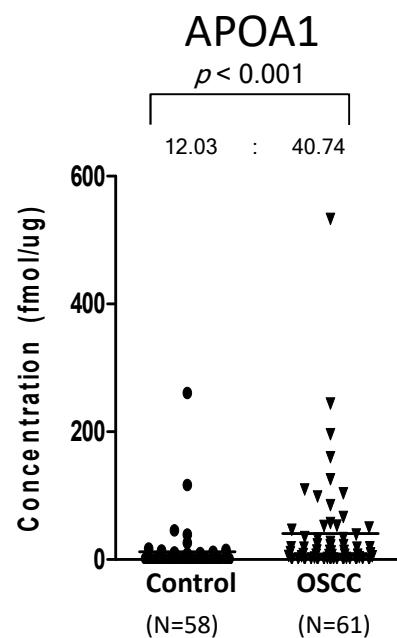
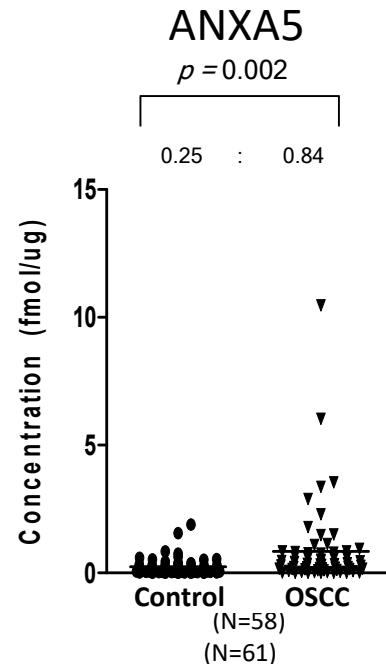
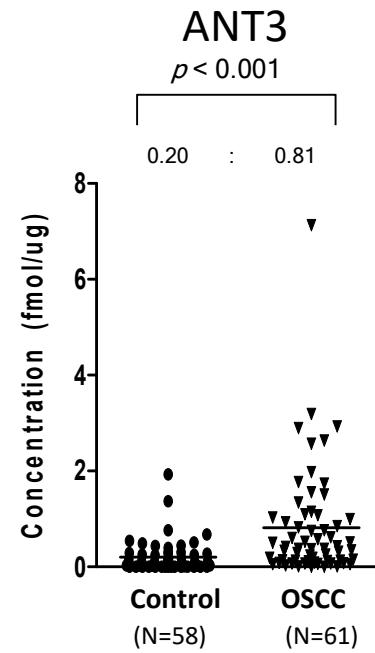
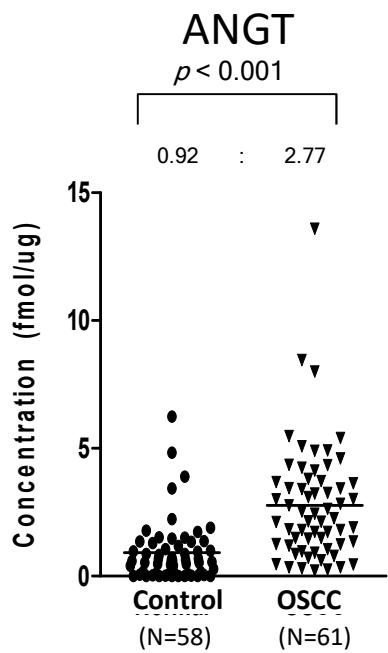
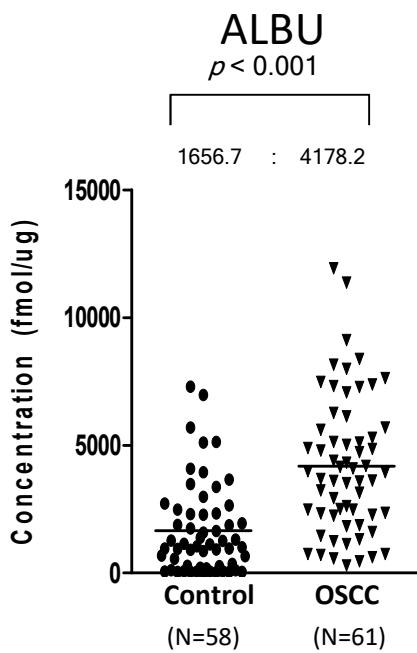
Supplemental Figure S1B

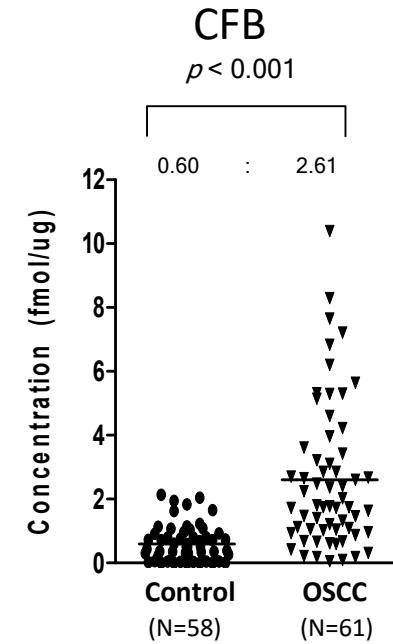
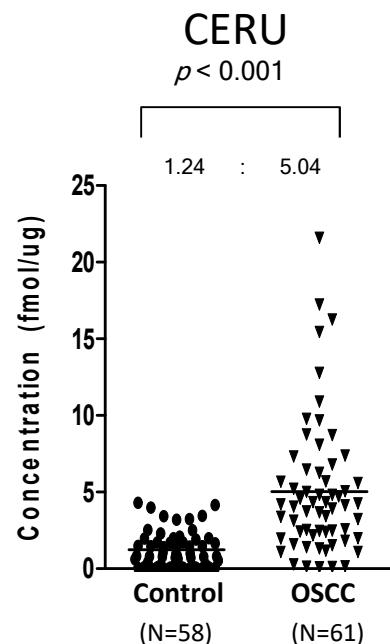
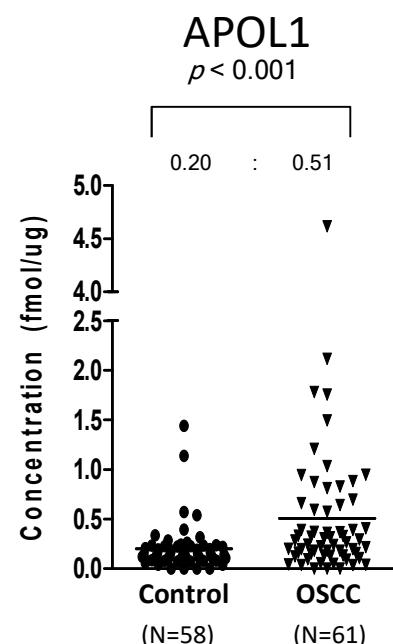
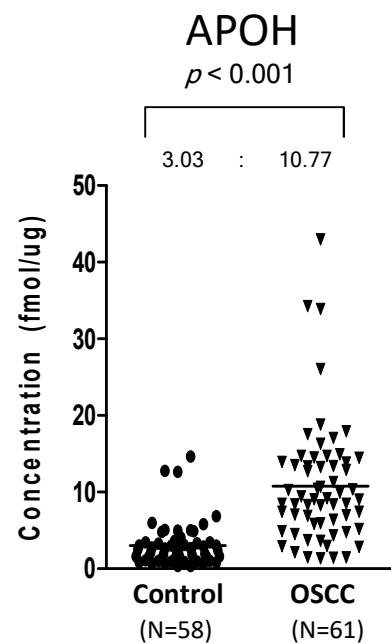
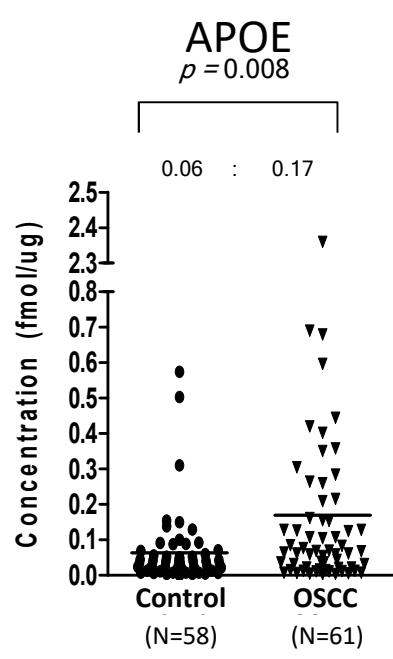
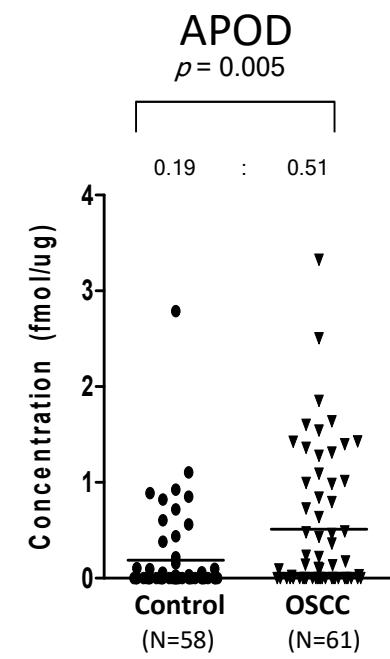
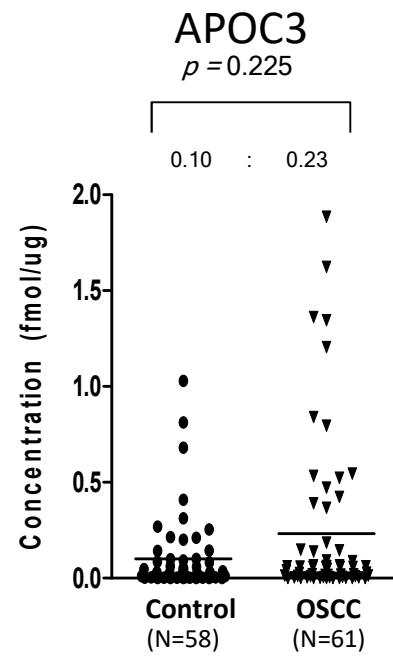
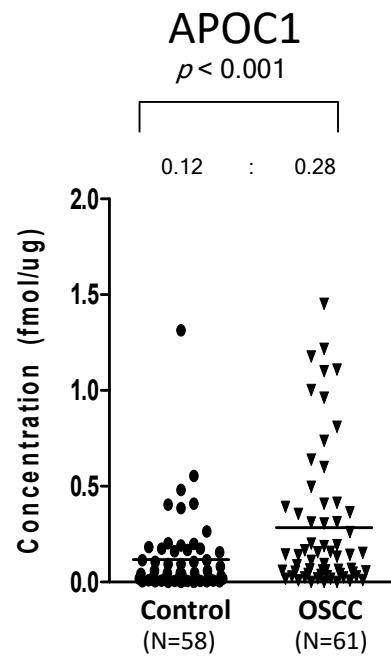


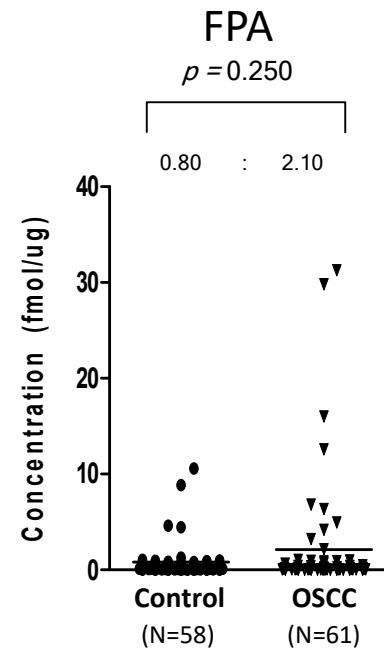
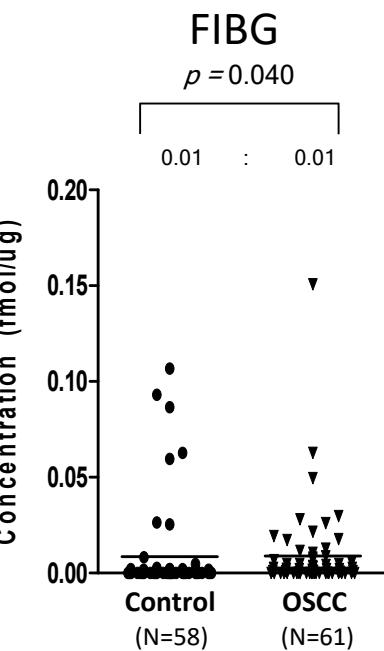
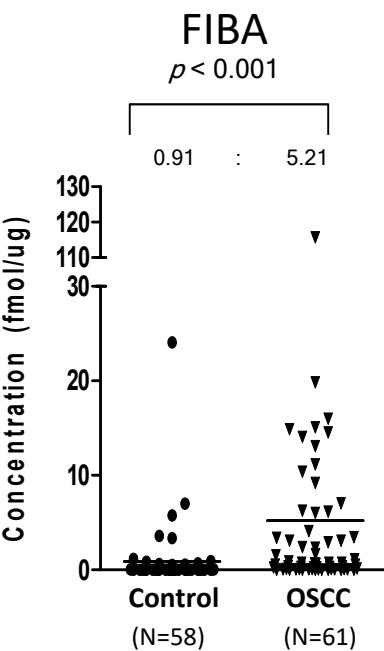
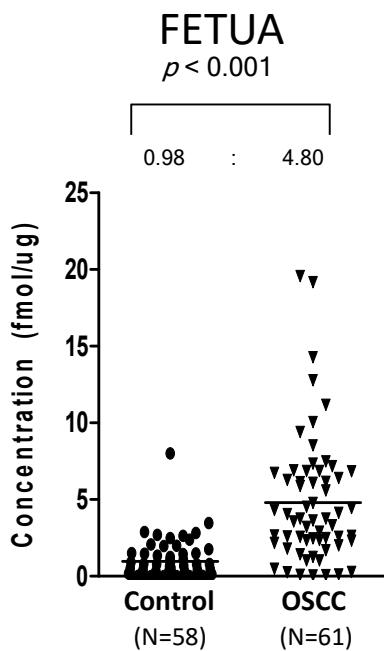
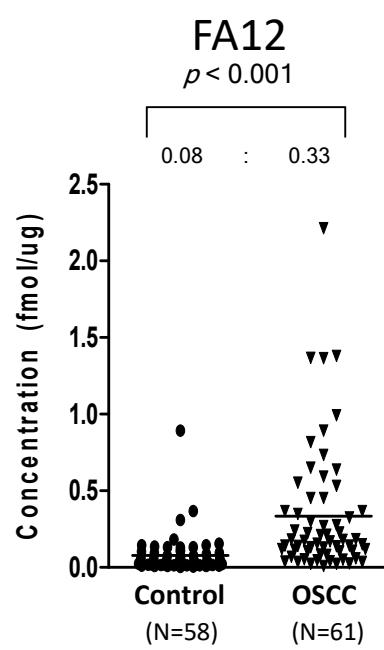
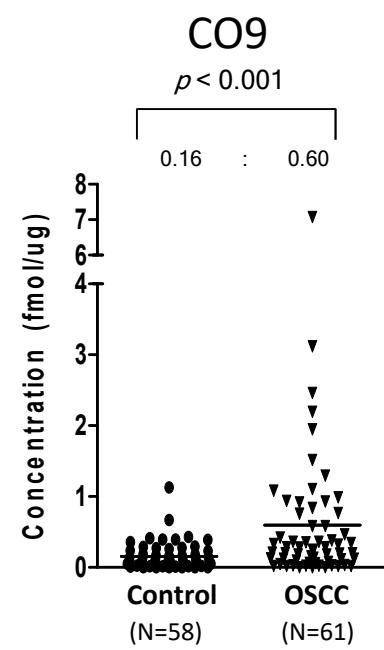
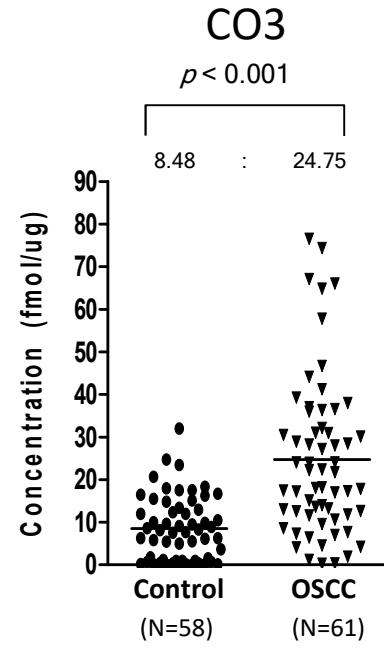
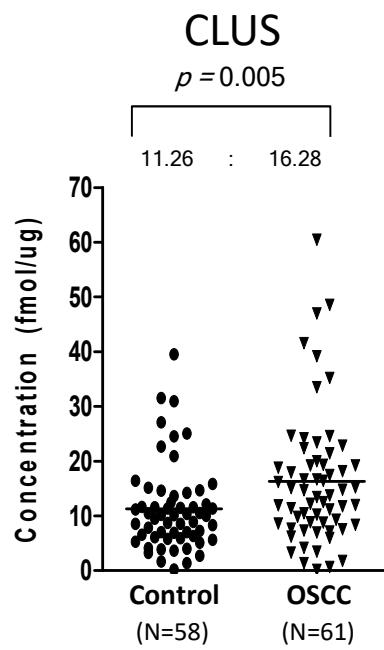
Supplemental Figure S1C

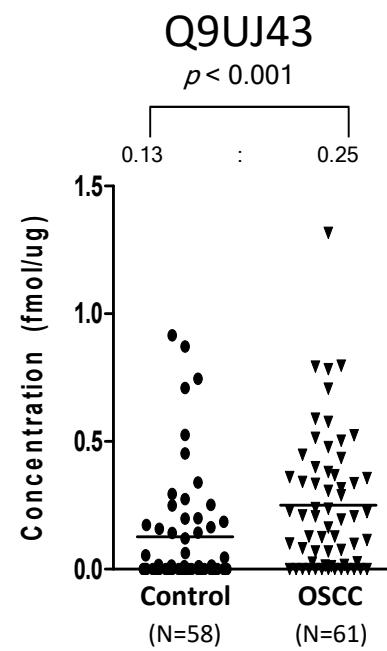
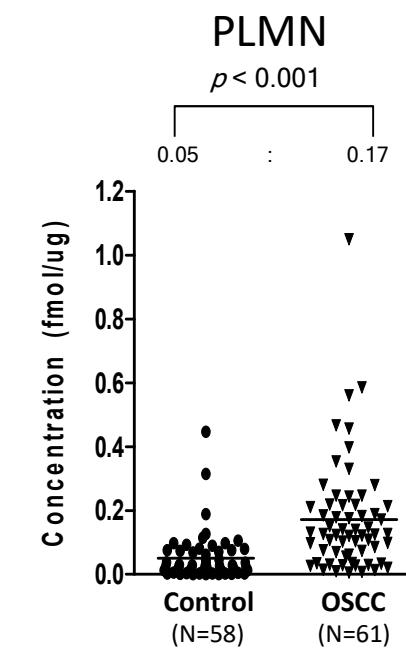
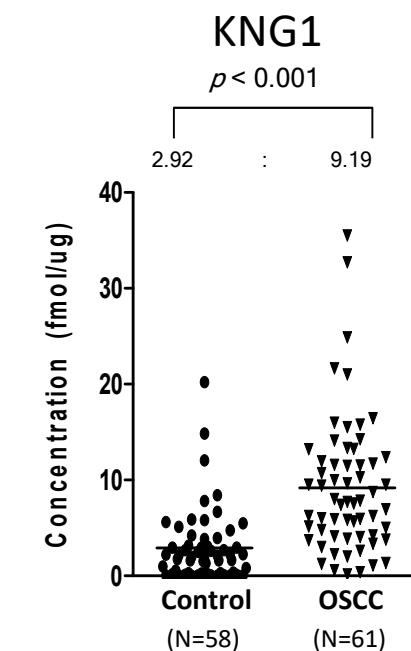
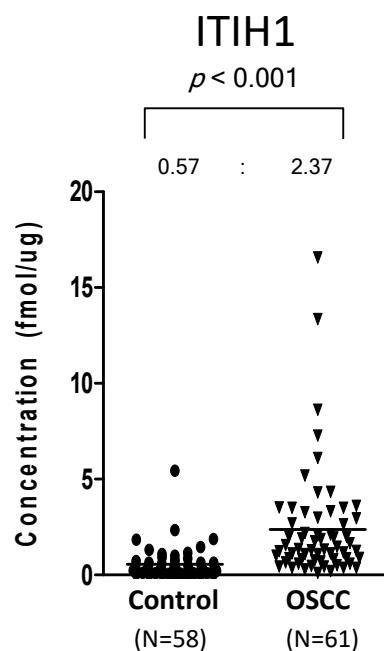
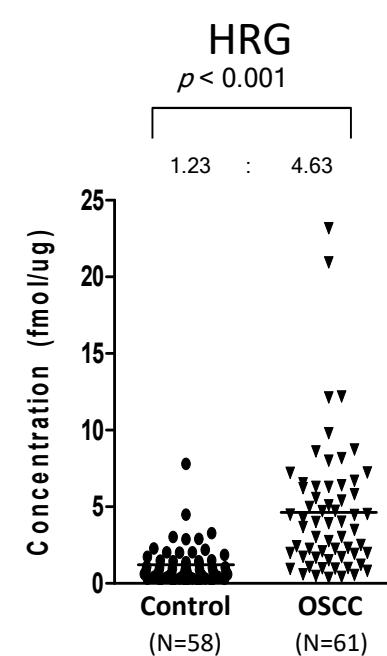
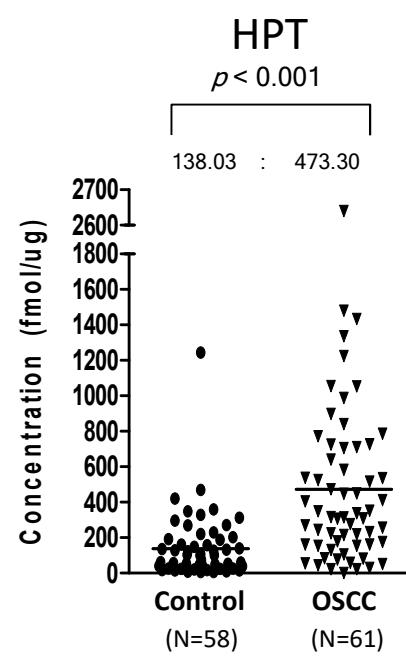
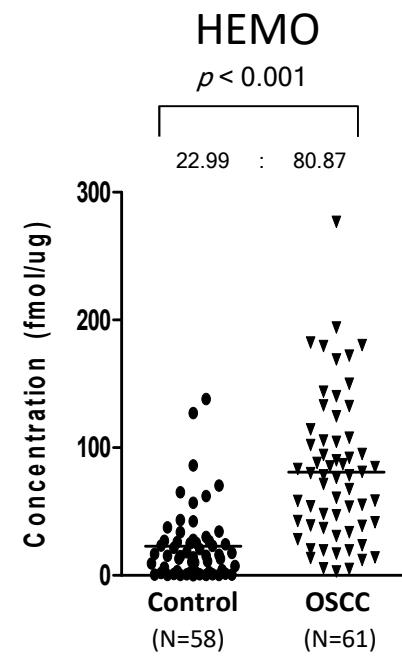
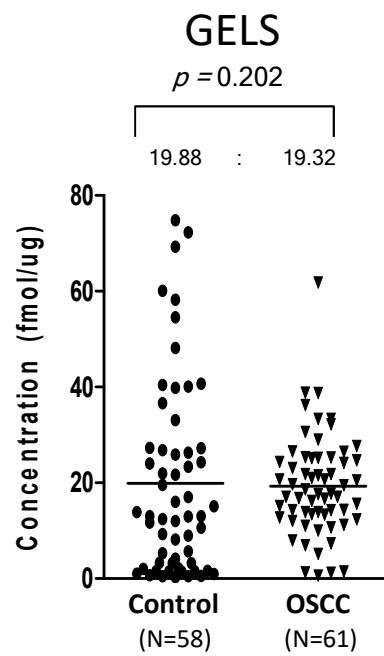


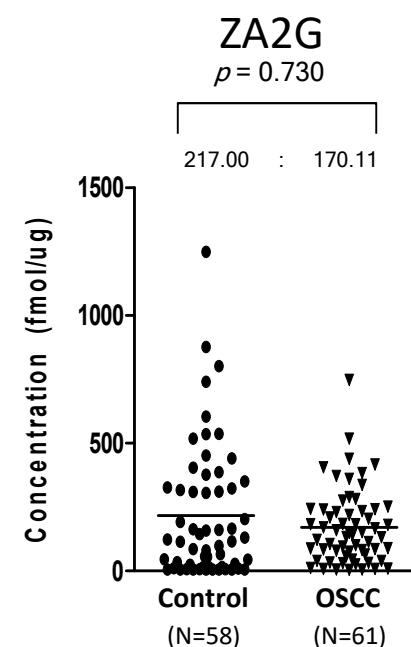
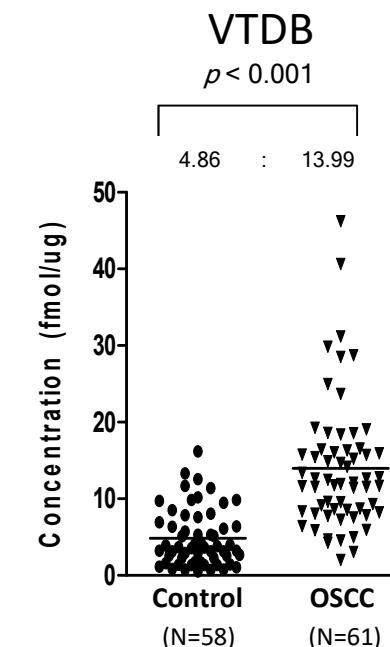
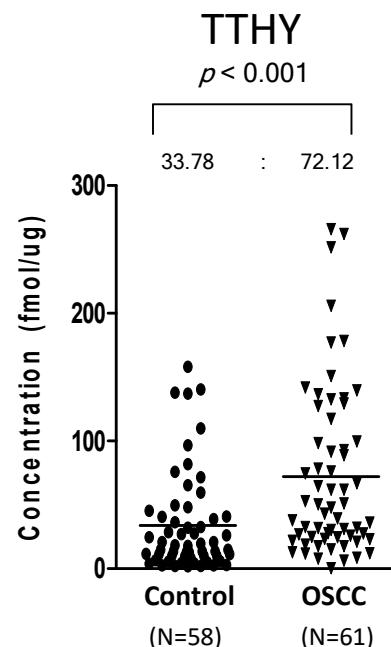
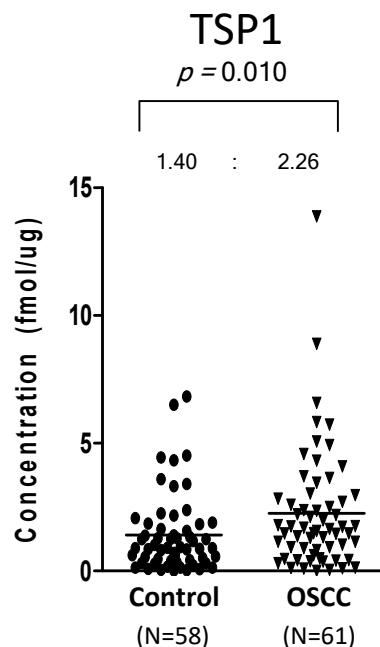
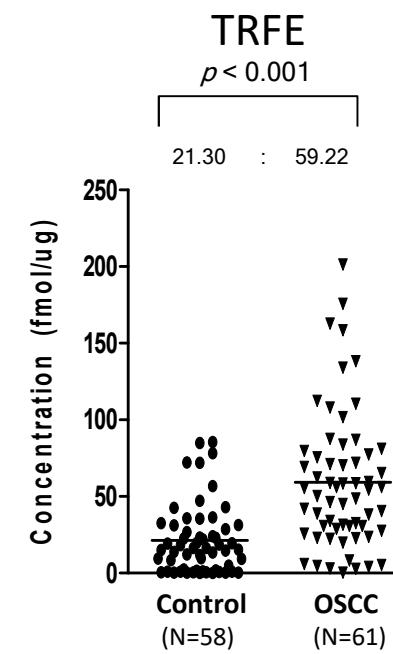
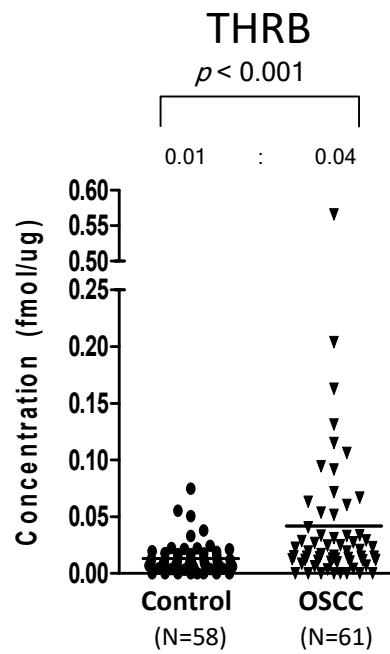
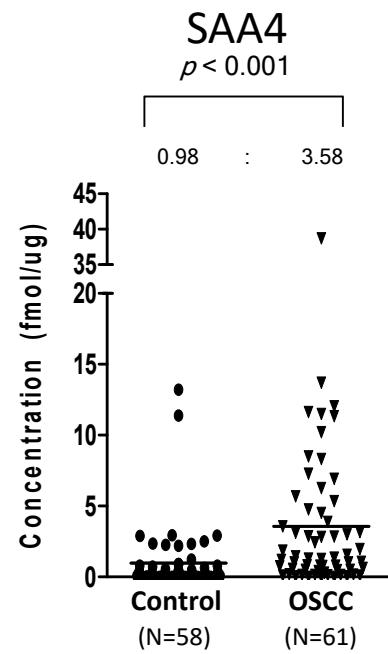
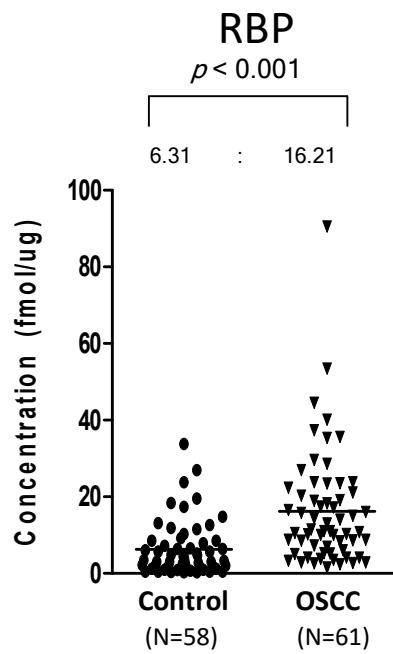




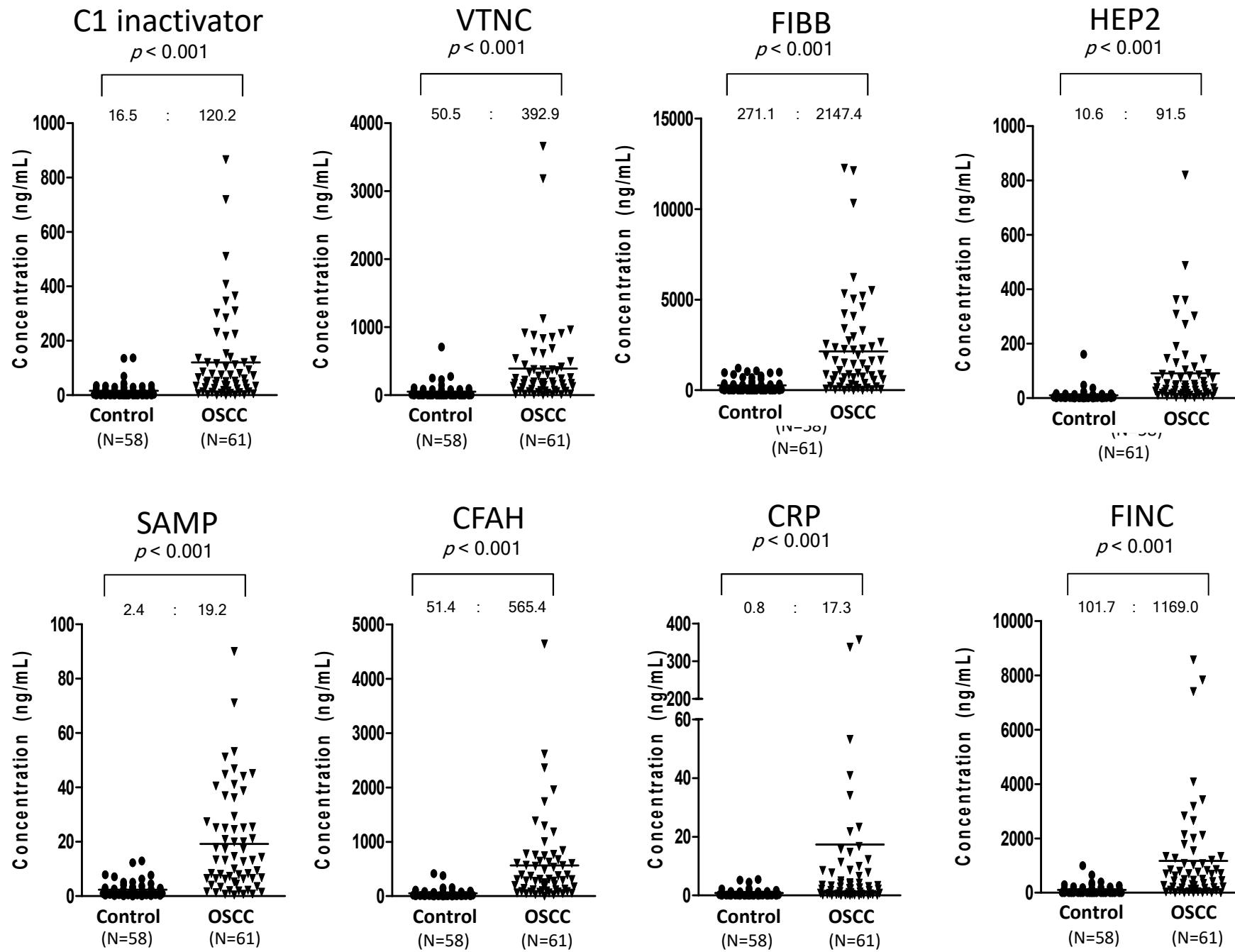


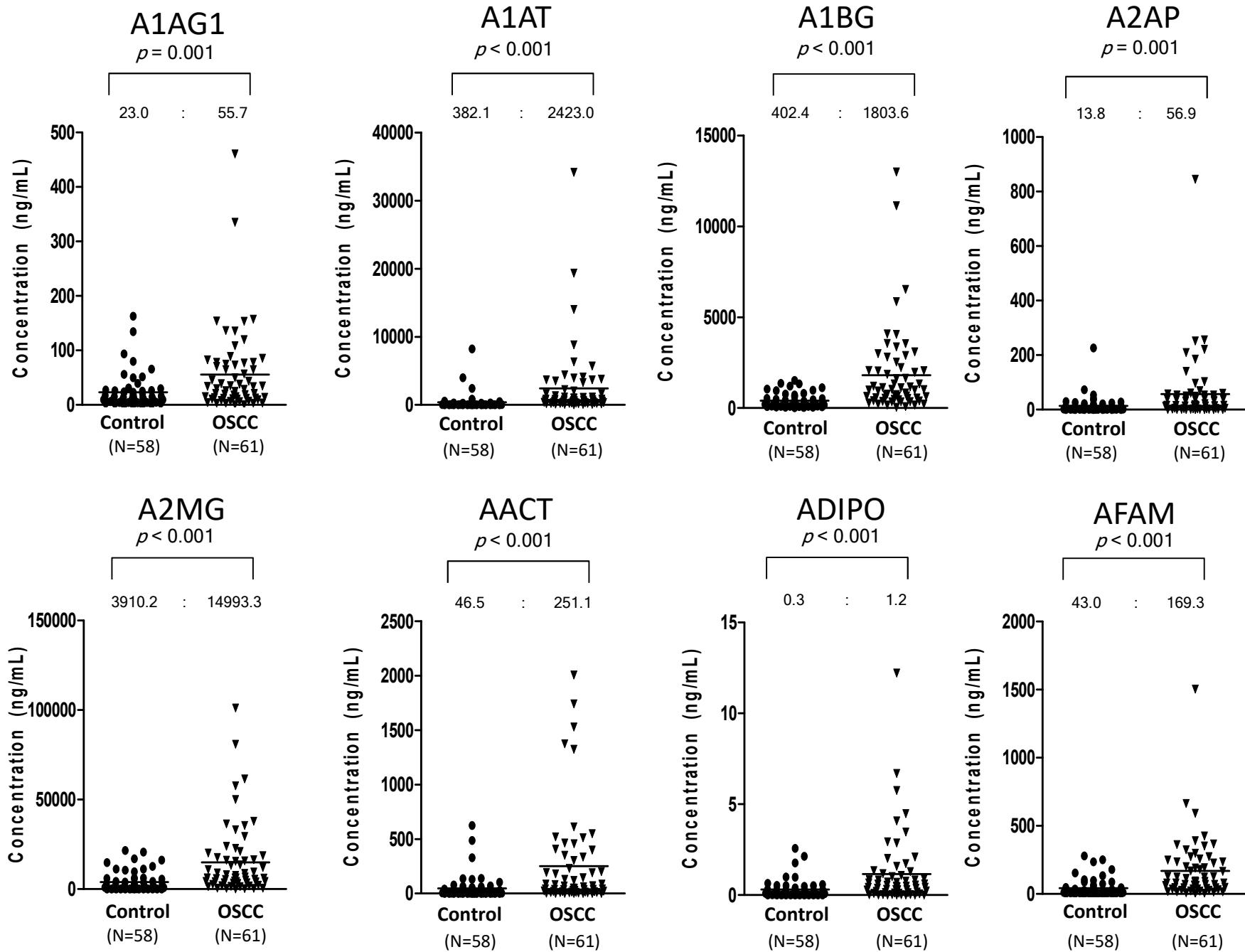


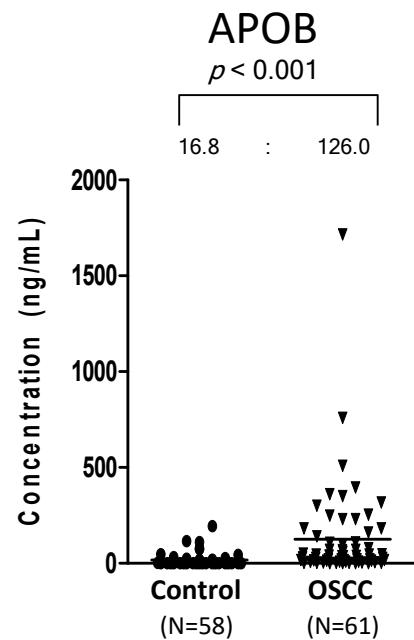
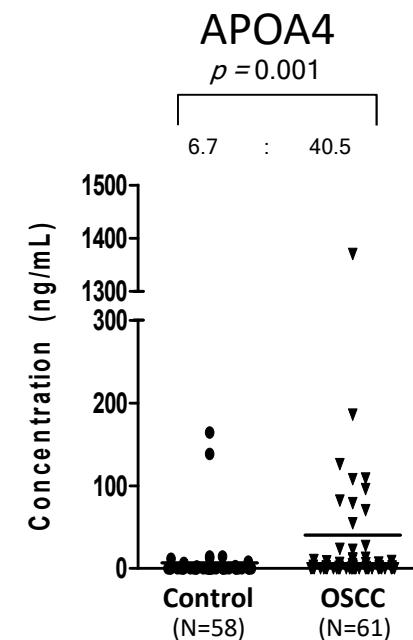
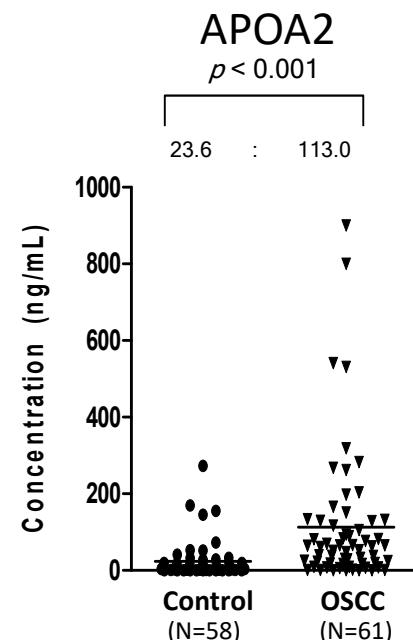
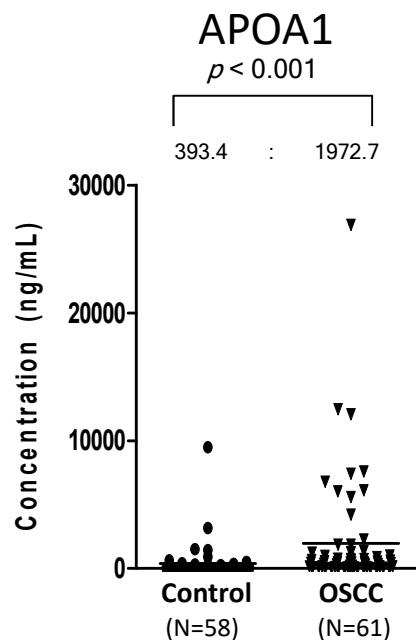
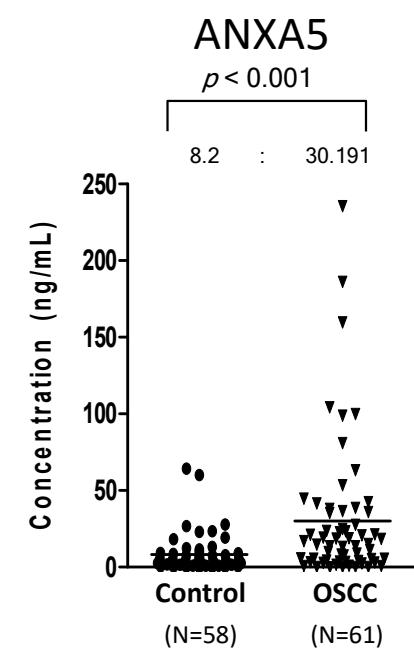
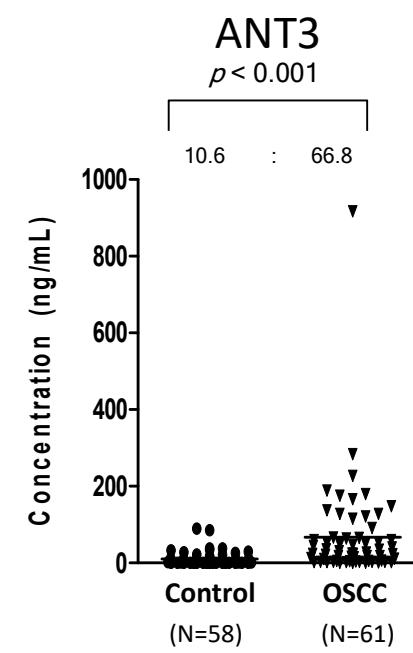
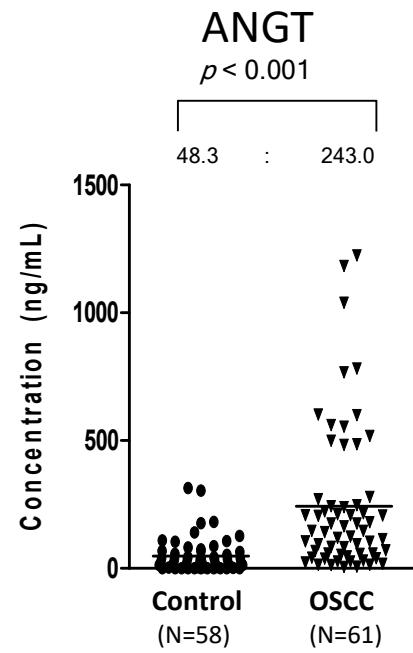
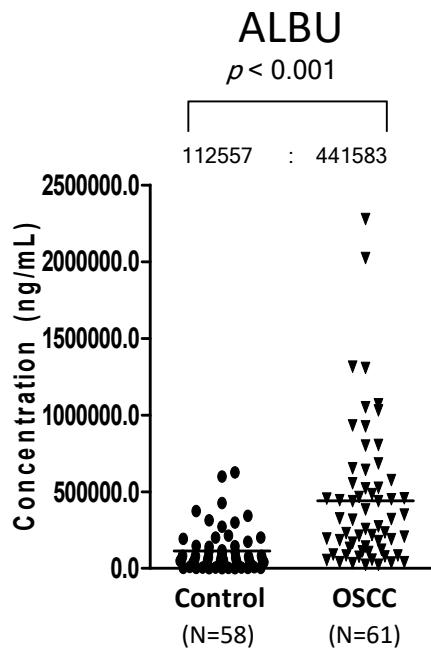


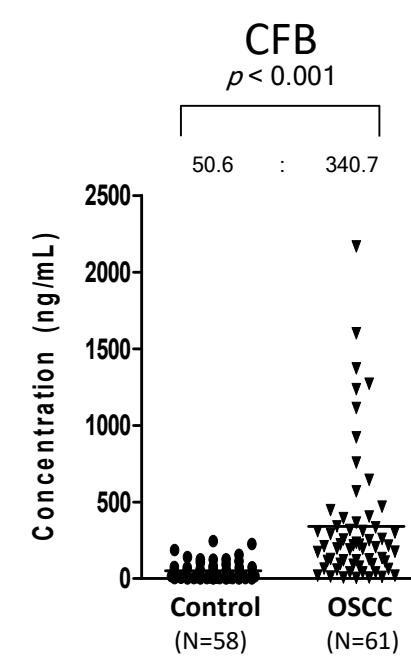
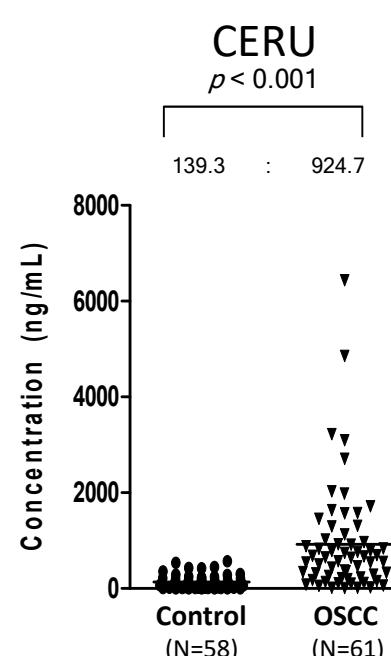
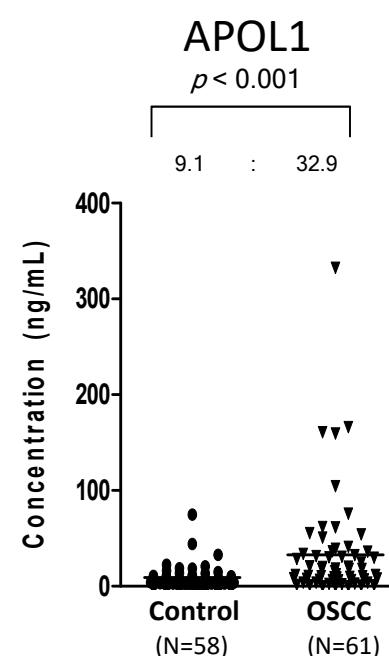
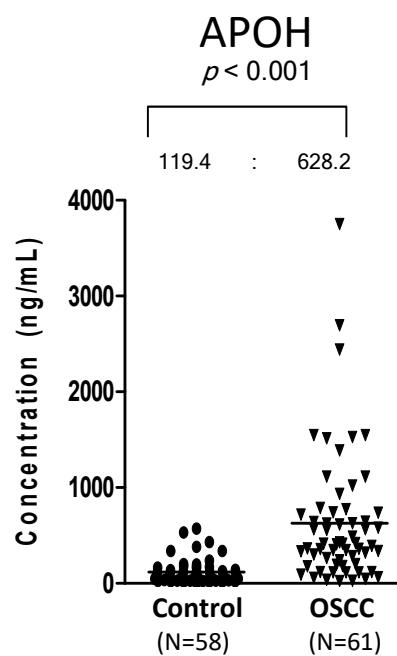
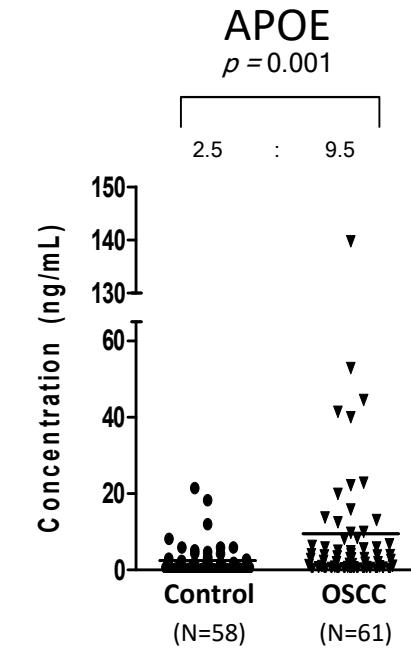
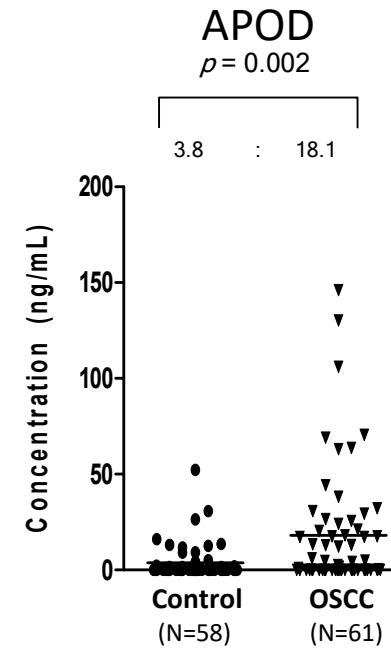
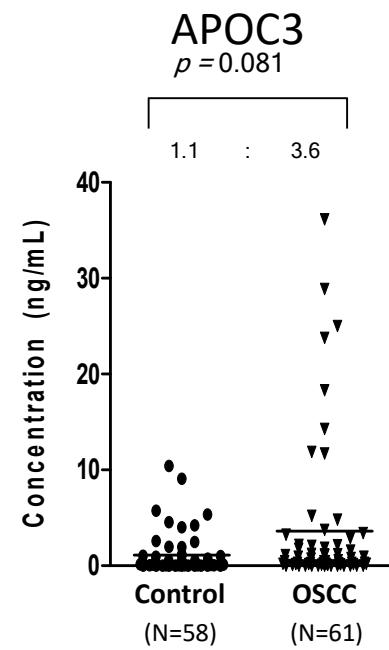
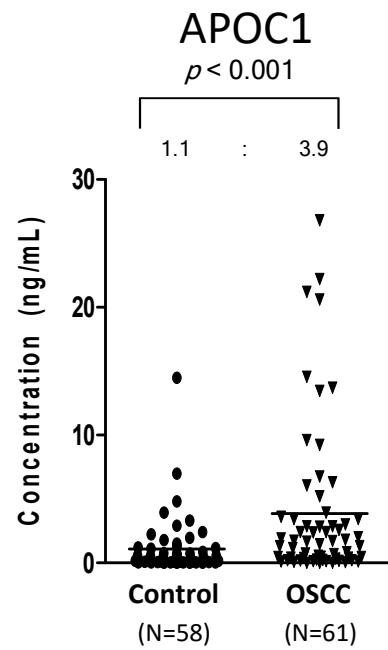


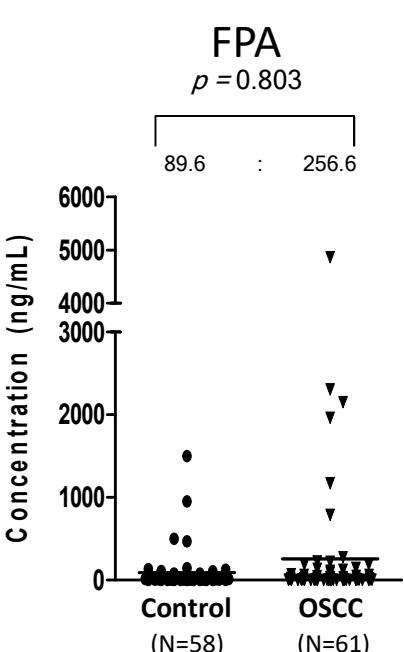
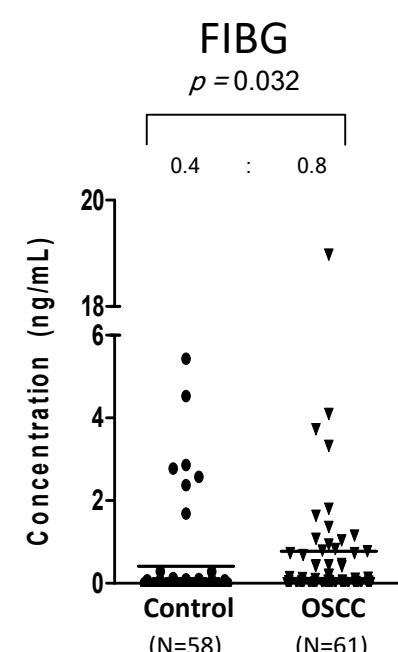
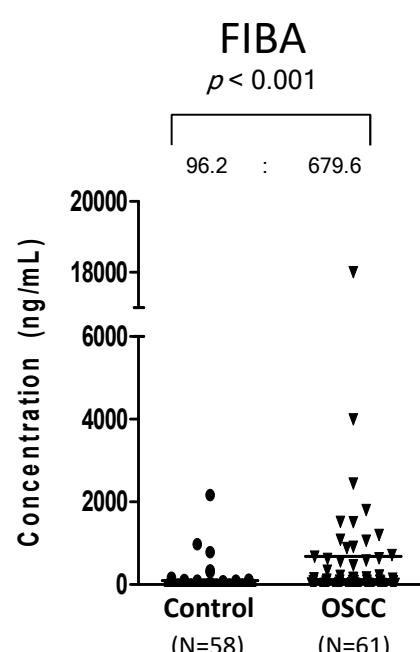
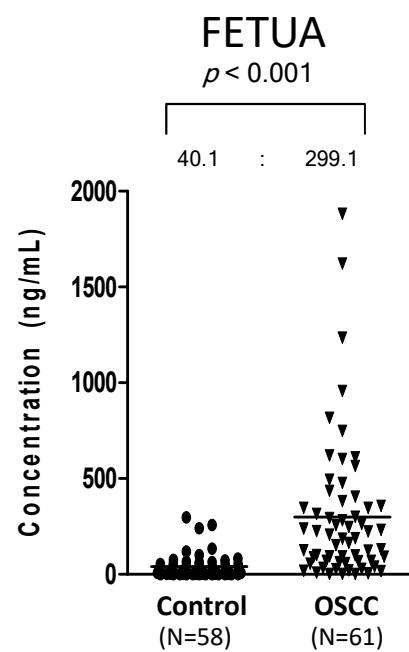
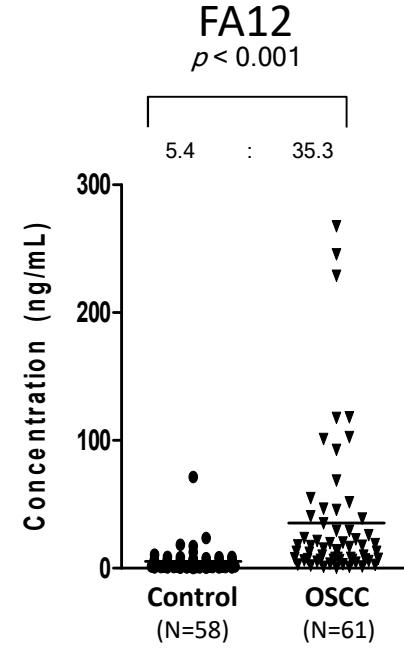
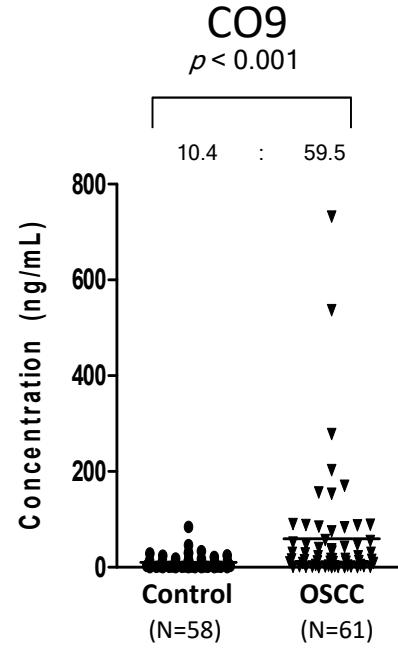
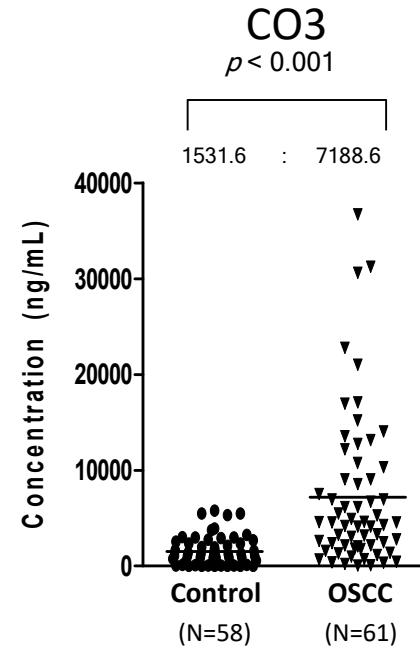
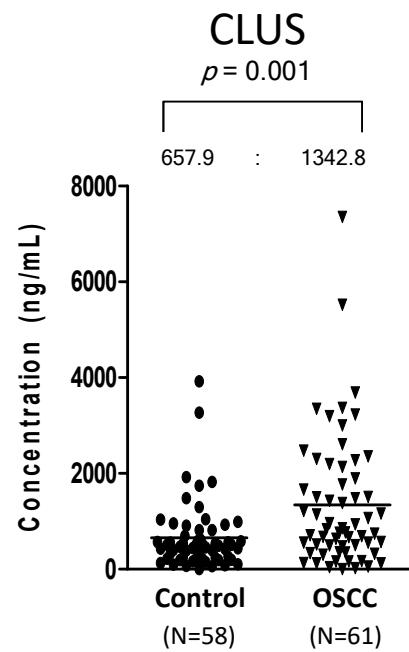
Supplemental Figure S1D

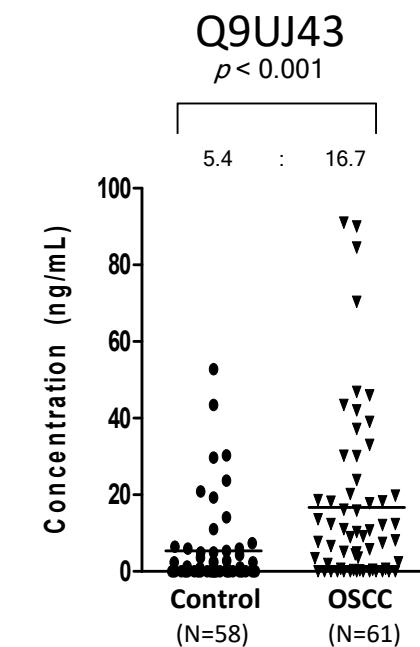
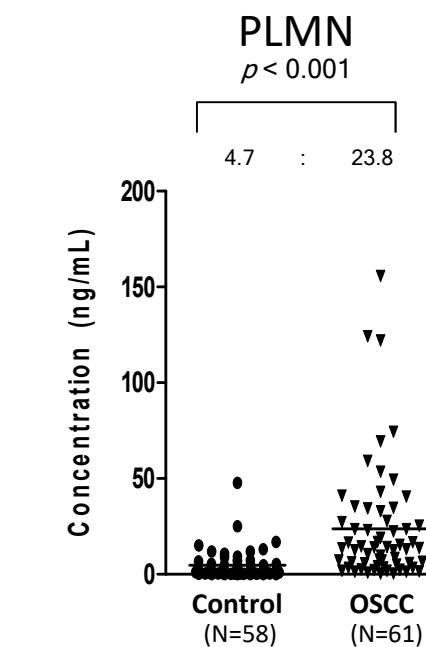
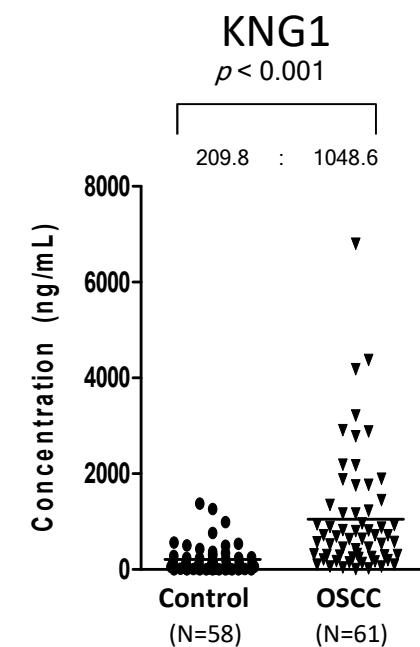
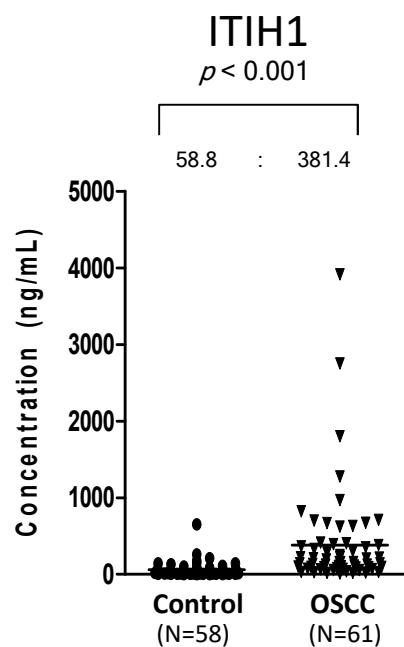
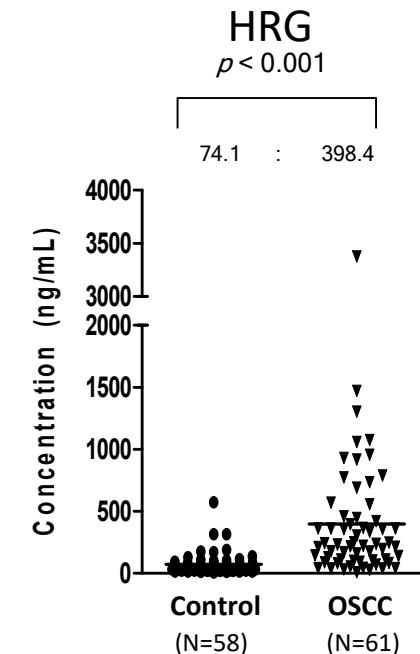
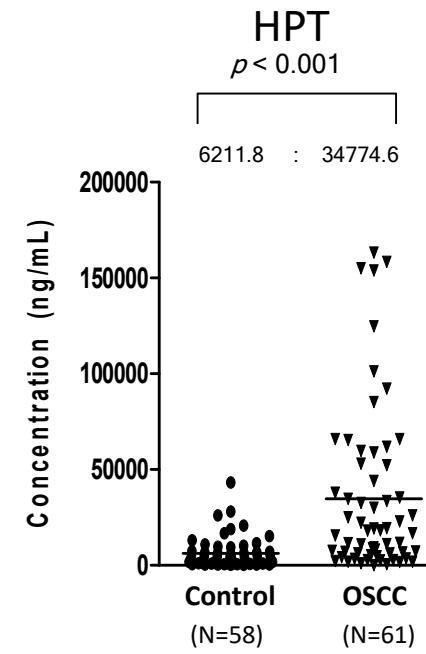
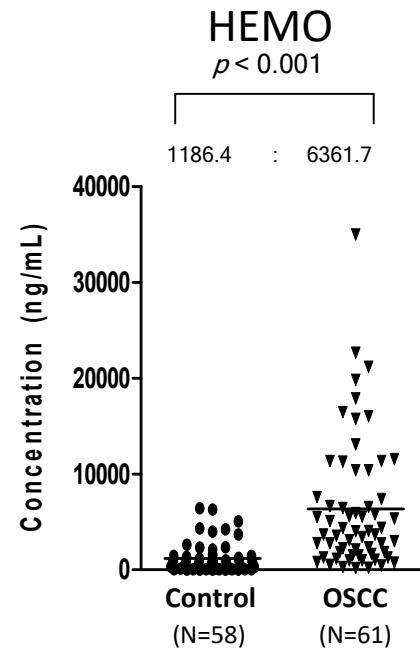
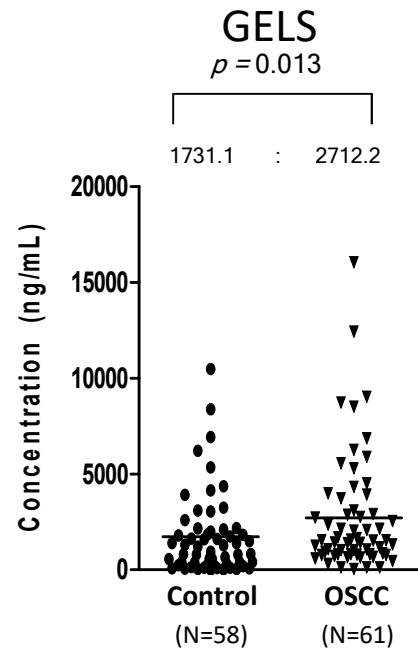


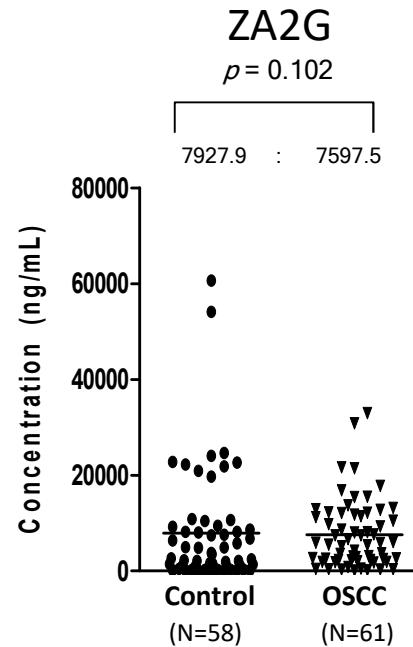
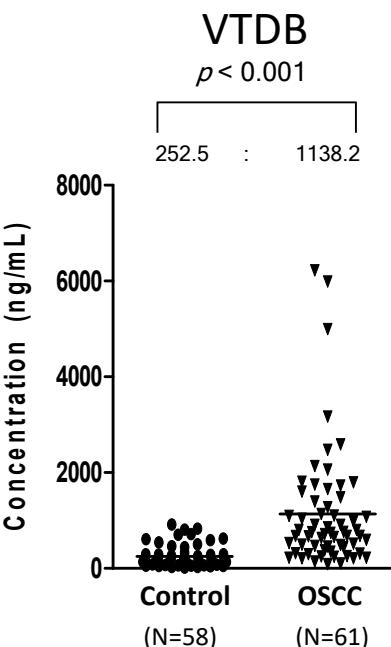
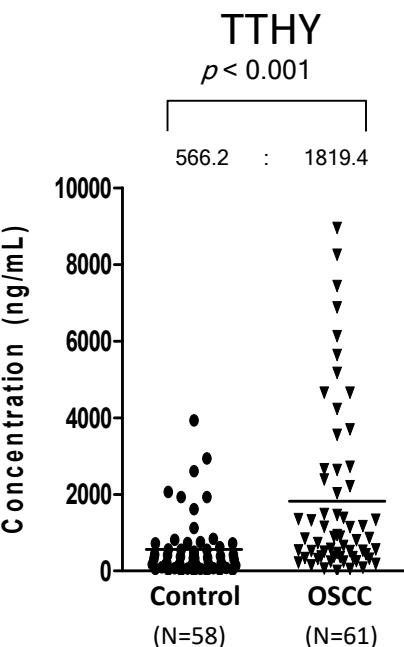
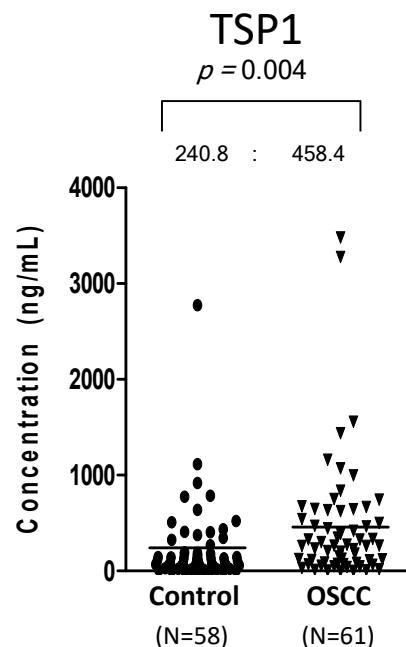
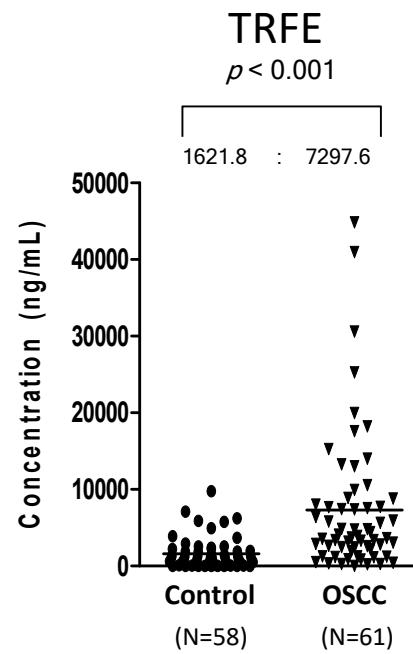
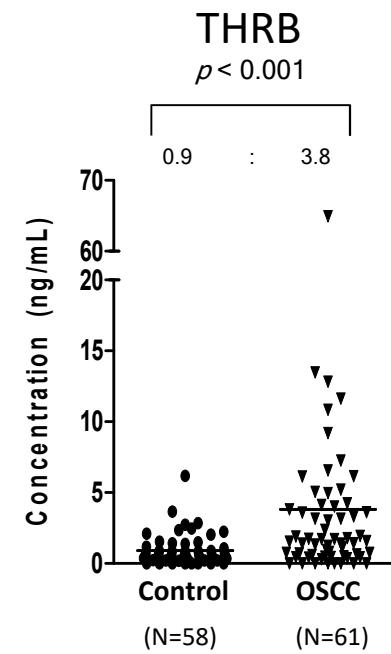
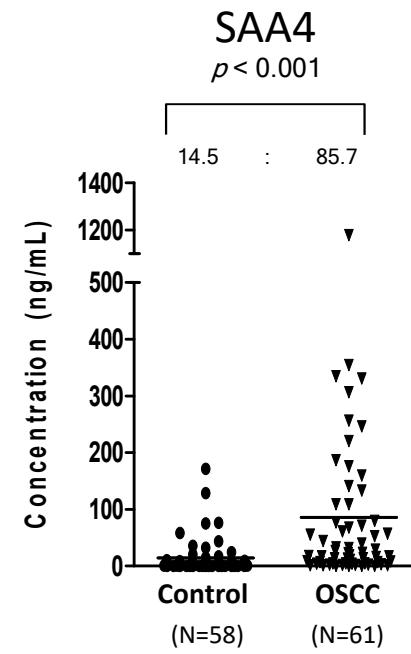
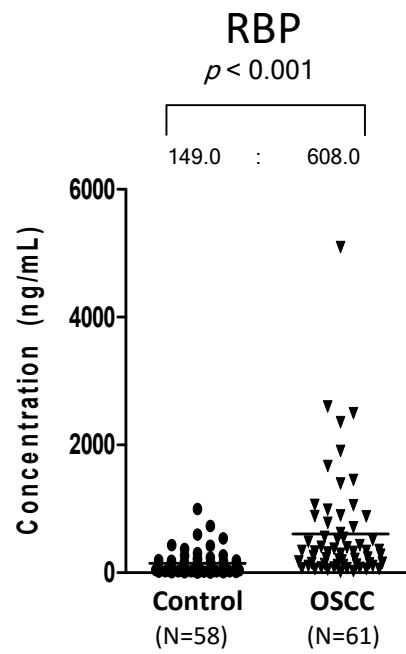




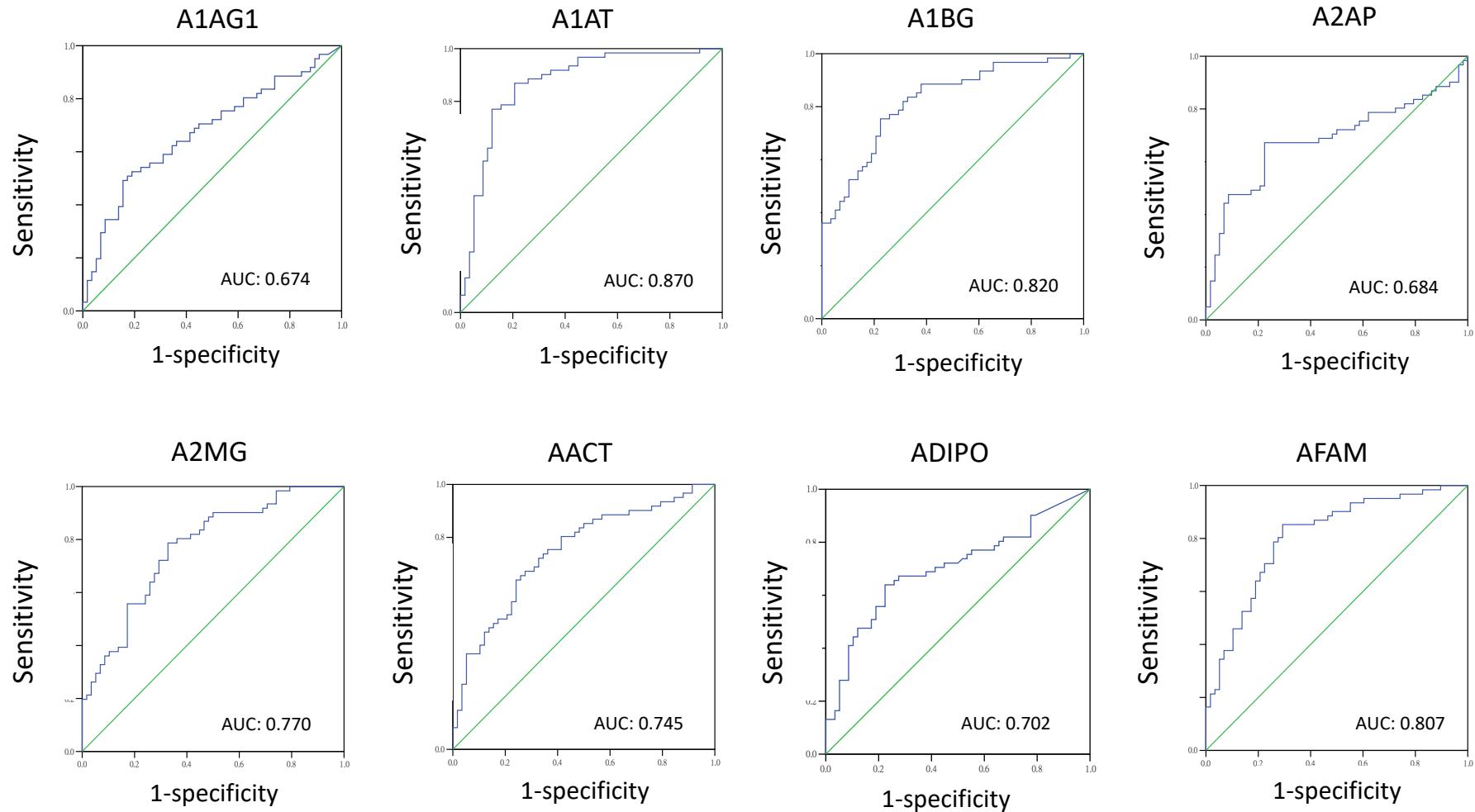


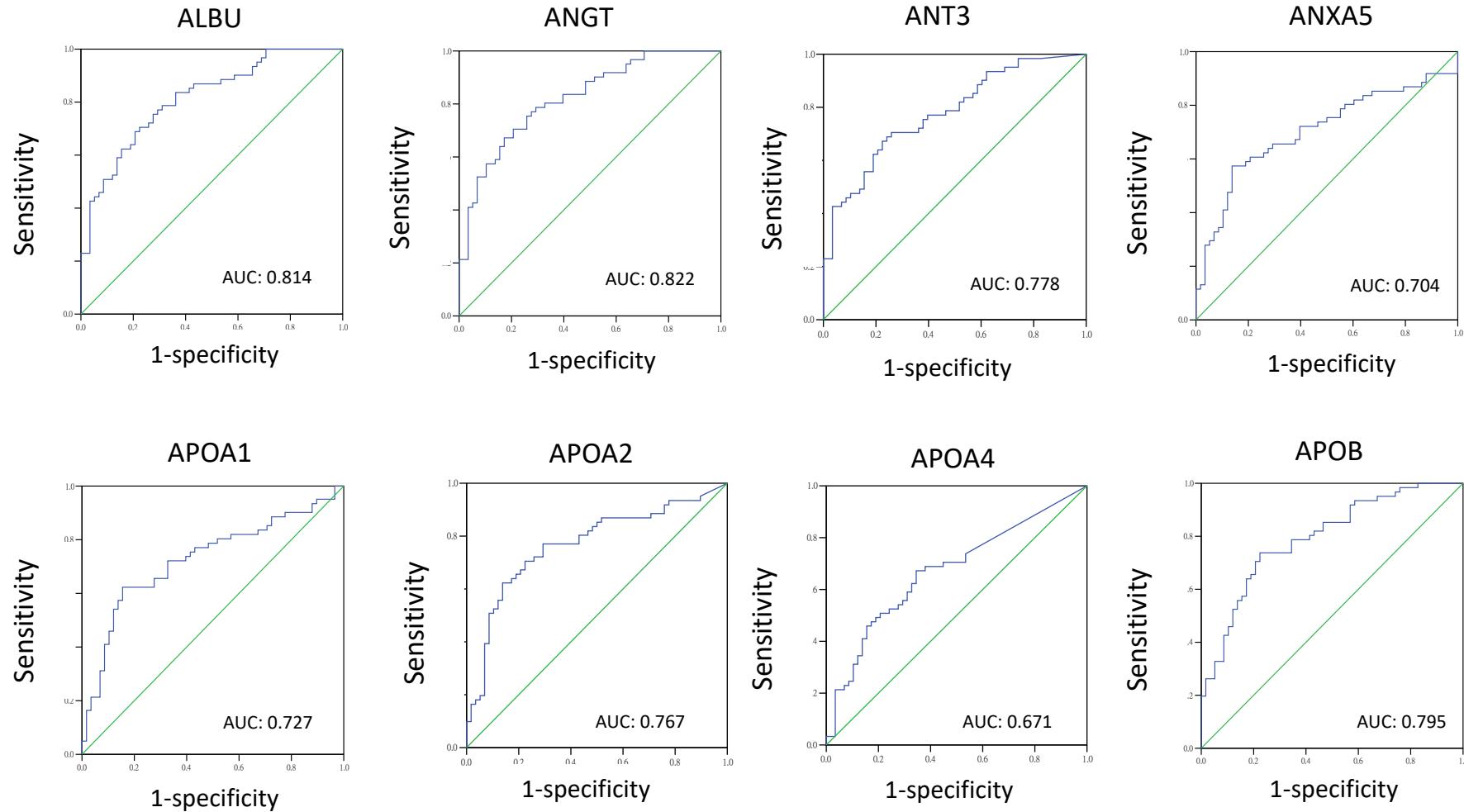


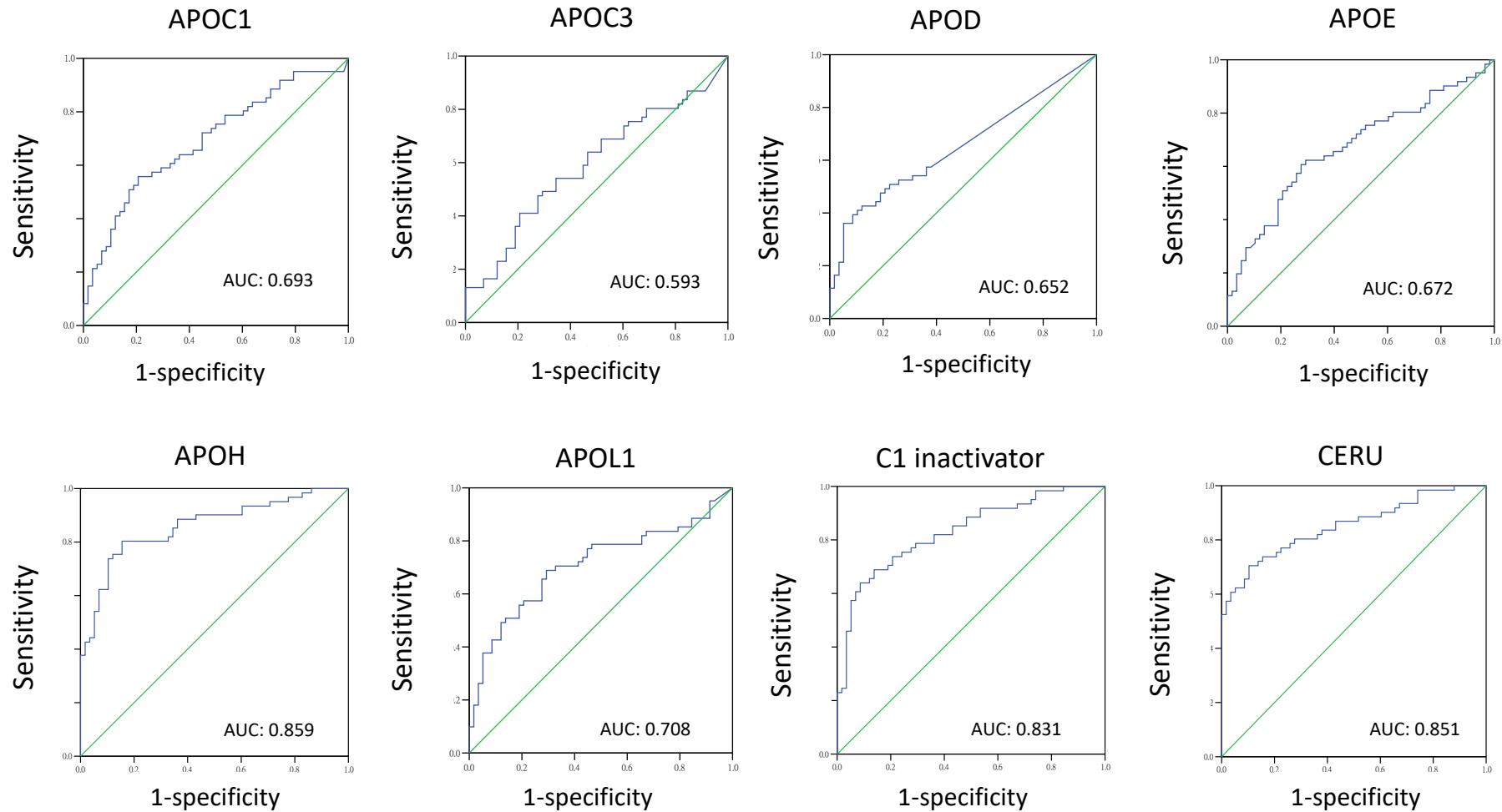


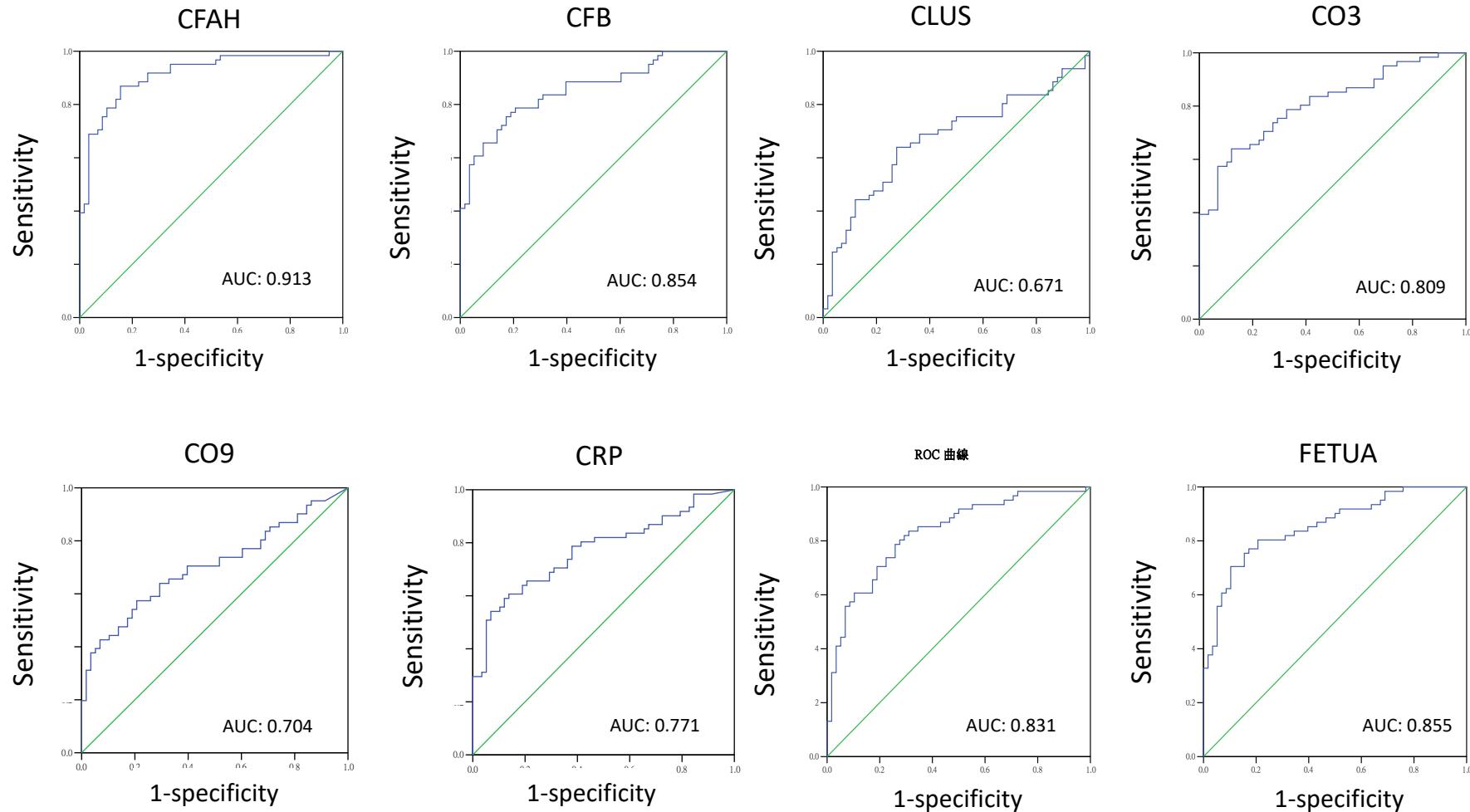


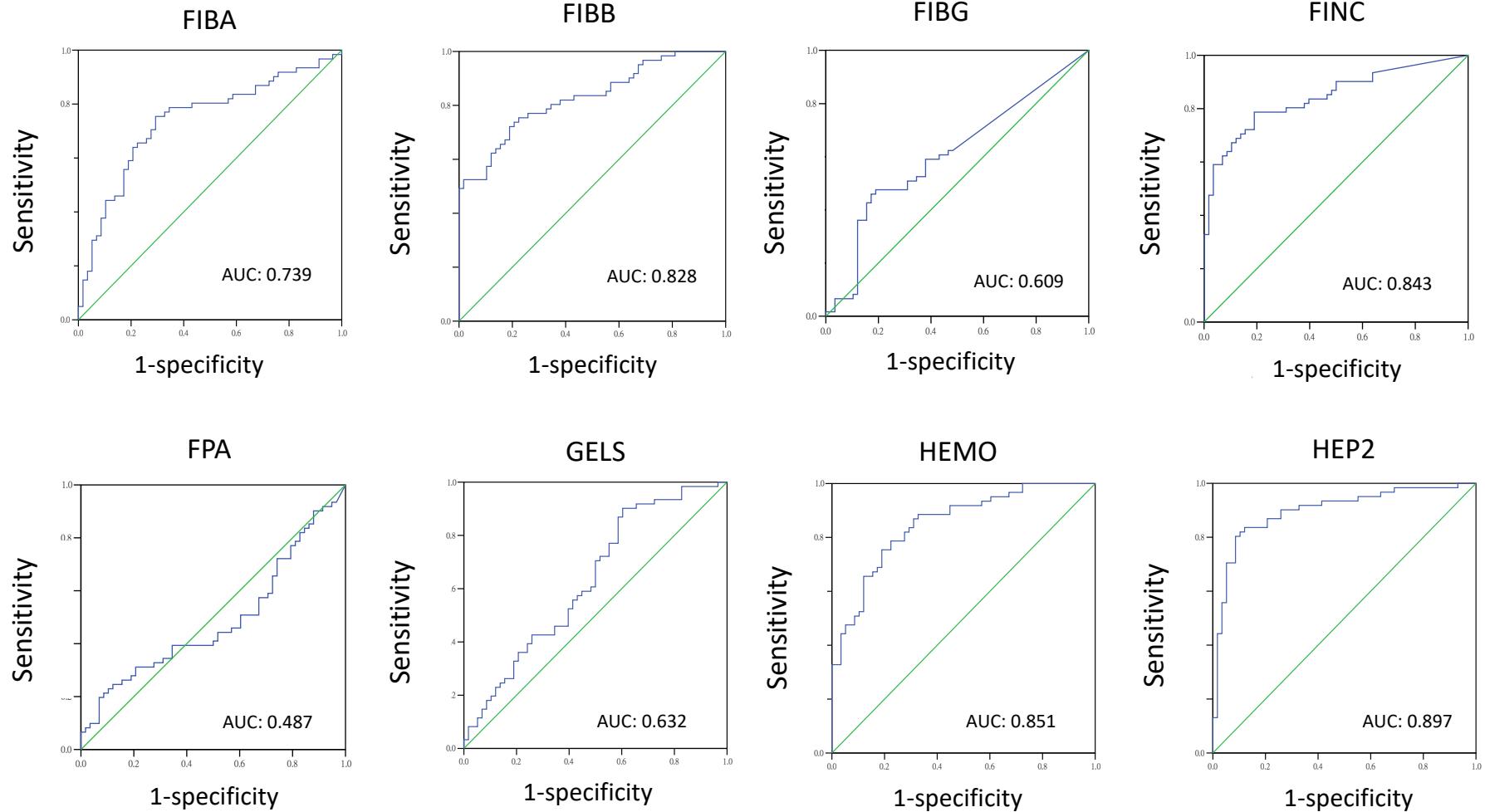
Supplemental Figure S1E

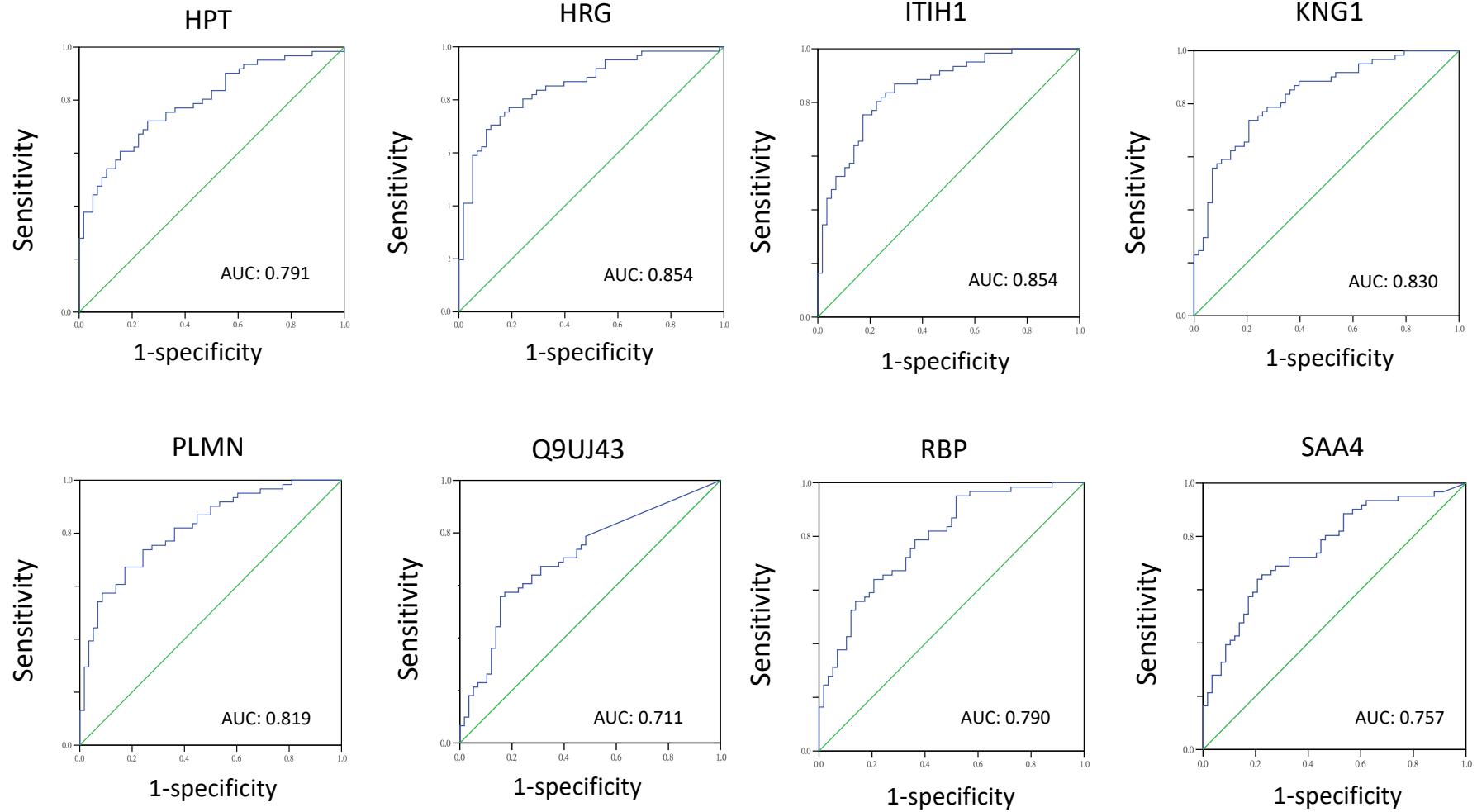


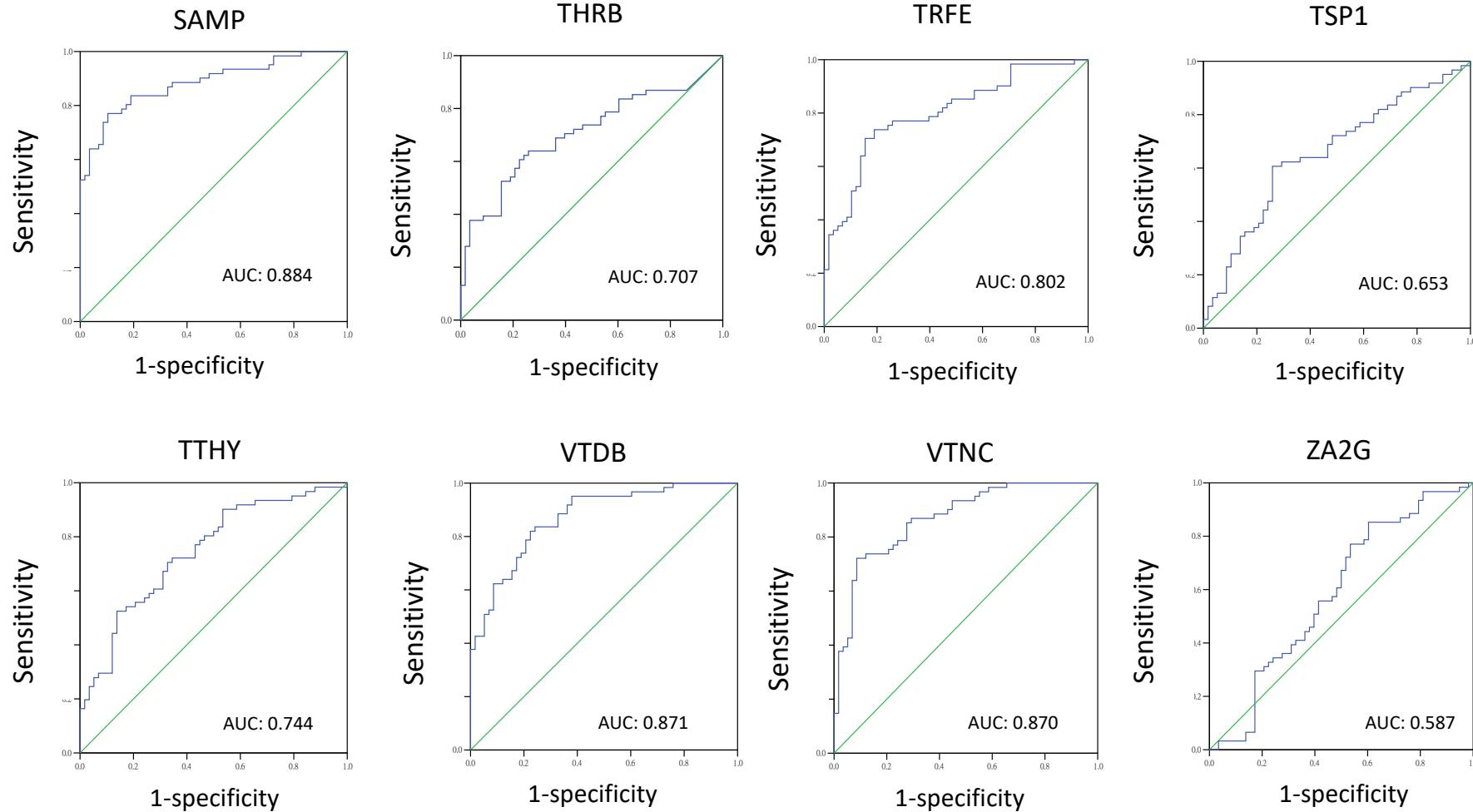




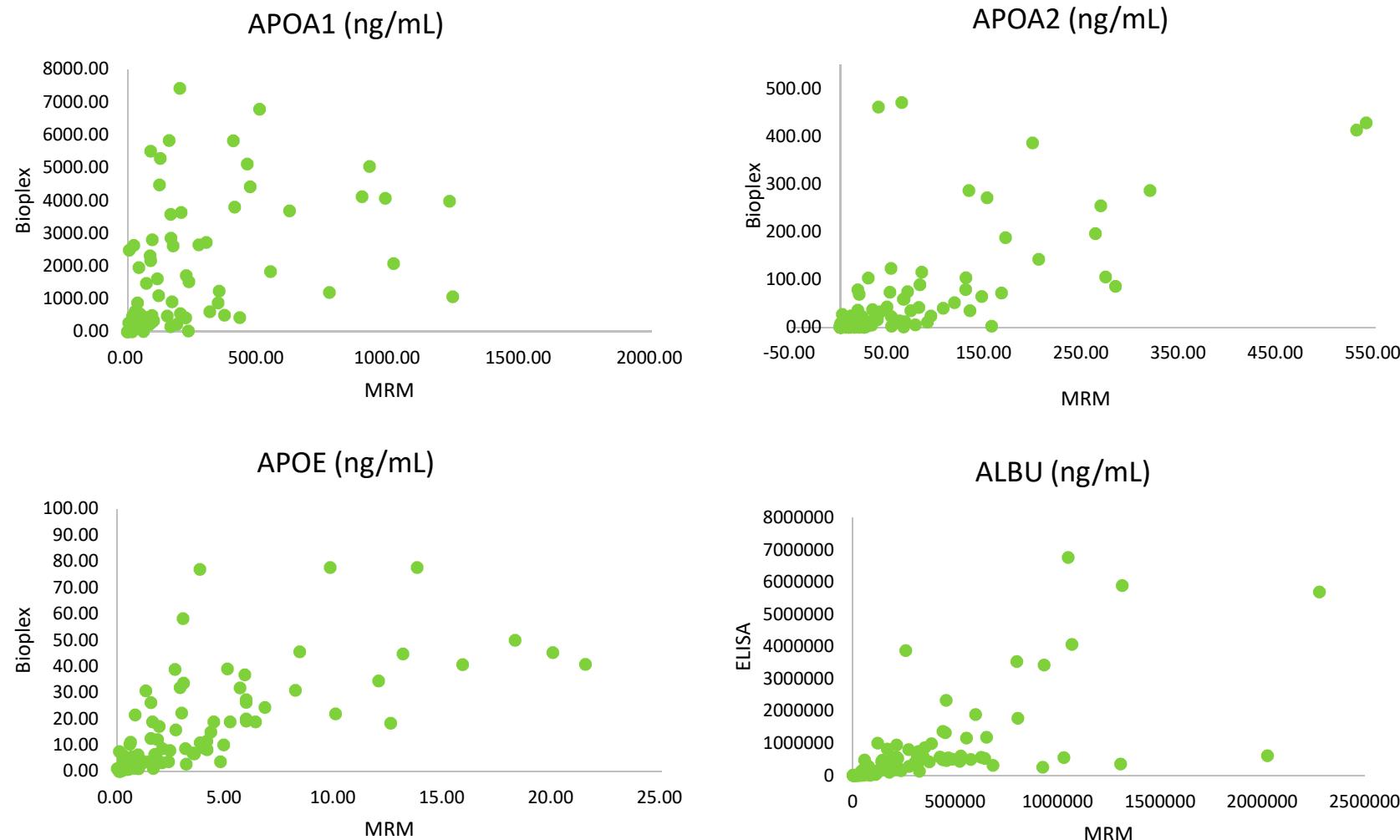








Supplemental Figure S2



Protein name	Assays	Spearman's correlation		n
		Correlation (r)	Significance (p-value)	
APOA1	MRM vs. BioPlex	0.792	< 0.001	117
APOA2	MRM vs. BioPlex	0.779	< 0.001	117
APOE	MRM vs. BioPlex	0.854	< 0.001	117
Albumin	MRM vs. ELISA	0.871	< 0.001	116

Supplemental Figure S3

