

Supplementary information for the manuscript:**“Sjögren’s and non-Sjögren’s sicca share a similar symptom burden but with a distinct symptom-associated proteomic signature”**

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Supplementary Table 1 – Associations between proteins and symptoms (ESSPRI). Values shown are the unstandardized coefficients and 95% confidence intervals.

Proteins vs ESSPRI	SjS (n = 53)		Sicca (n = 60)	
	B (95% CI)	P value	B (95% CI)	P value
ADM	0.15 (-1.22 to 1.53)	0.83	2.2 (1.44 to 3)	<0.0001
TNFRSF13B	-0.3 (-1.5 to 0.9)	0.60	1.9 (0.64 to 3.17)	0.004
FGF_23	0.7 (-0.42 to 1.82)	0.21	1.9 (0.70 to 3.1)	0.005
CD40	-0.1 (-2.1 to 1.9)	0.92	2.9 (1.5 to 4.4)	0.0001
IL_10RB	0.35 (-1.77 to 2.48)	0.74	3.5 (1.6 to 5.4)	0.001
CD5	-0.5 (-2.65 to 1.64)	0.63	1.9 (0.53 to 3.34)	0.008
CD4	-1.3 (-3.44 to 0.82)	0.22	2.6 (1.2 to 4.04)	0.001
ACE2	0.75 (-0.48 to 1.99)	0.23	1.7 (0.88 to 2.47)	<0.0001
LEP	0.2 (-0.64 to 1.04)	0.80	0.8 (0.21 to 1.35)	0.009
TRAIL_R2	-0.04 (-1.60 to 1.52)	0.75	2.1 (0.90 to 3.33)	0.001

SPON2	-4.3 (-8.28 to -0.33)	0.03	4.3 (1.1 to 7.8)	0.01
PD_L1	-0.24 (-0.42 to 0.19)	0.42	1.35 (0.26 to 2.45)	0.02
SLAMF1	0.4 (-1.15 to 2.01)	0.73	1.4 (0.32 to 2.5)	0.01
CSF_1	-0.03 (-3 to 2.94)	0.98	4.4 (1.82 to 7.05)	0.001
IL_1ra	0.35 (-0.73 to 1.43)	0.52	0.96 (0.1 to 1.8)	0.03
Gal_9	-0.7 (-2.67 to 1.13)	0.42	2.3 (0.82 to 3.86)	0.003
TNFRSF10A	-0.3 (-1.97 to 1.35)	0.71	1.7 (0.37 to 3)	0.01
IL_12B	-0.02 (-0.98 to 0.93)	0.06	0.4 (-0.04 to 0.90)	0.07
TNFRSF9	0.01 (-1.28 to 1.3)	0.98	1.4 (0.39 to 2.43)	0.008
TNFRSF11A	0.01 (-1.51 to 1.69)	0.91	1.4 (0.28 to 2.91)	0.02

Supplementary Table 2 – Results of multiple regression analysis model (forward) for proteins and symptoms in sicca patients. Values shown are the unstandardized coefficients and 95% confidence intervals.

Variables vs ESSPRI	Model (Forward)	
	B (95% CI)	P value
ADM	2.5 (1.6-3.6)	< 0.0001
CD40	2.5 (1.1-3.8)	< 0.0001
SPON2	-4.6 (-8.6-1.2)	0.009

Supplementary Table 3 - Correlations between ADM and clinical/immunological variables. Values shown are the unstandardized coefficients and 95% confidence intervals.

ADM versus clinical parameters	SjS (n = 53)		Sicca (n = 60)	
	B (95% CI)	P value	B (95% CI)	P value
Age	0.01 (0.003 to 0.02)	0.01	0.02 (0.01 to 0.03)	0.001
BMI	0.03 (0.01 to 0.05)	0.01	0.04 (0.02 to 0.07)	0.001
Weight	0.01 (0.0001 to 0.01)	0.04	0.01 (0.003 to 0.20)	0.01
Fat mass	-0.005 (-0.003 to 0.01)	0.22	0.004 (-0.003 to 0.01)	0.25
Free fat mass	0.003 (-0.01 to 0.02)	0.62	0.02 (-0.003 to 0.36)	0.09
Leptin	0.2 (0.1 to 0.36)	0.004	0.3 (0.15 to 0.41)	<0.0001
VAS global (patient)	-0.003 (-0.01 to 0.005)	0.43	-0.01 (-0.02 to -0.002)	0.008
EQ5D	0.10 (-0.35 to 0.55)	0.65	-0.7 (-1.16 to -0.28)	0.002
IgG	-0.004 (-0.21 to 0.13)	0.66	0.06 (-0.01 to 0.13)	0.07
C3	0.7 (0.19 to 1.23)	0.008	0.6 (0.06 to 1.13)	0.03

C4	0.7 (-0.64 to 1.41)	0.45	0.6 (-0.25 to 2.96)	0.1
K/L free light chain quotient	0.01 (-0.12 to 0.14)	0.86	0.33 (0.10 to 0.71)	0.01
Focus score	0.01 (-0.14 to 0.17)	0.86	0.01 (-0.47 to 0.48)	0.98
Schirmer's test	-0.01 (-0.02 to 0.005)	0.09	-0.01 (-0.02 to 0.003)	0.13
Unstimulated salivary flow	-0.1 (-0.35 to 0.09)	0.24	-0.15 (-0.27 to -0.02)	0.02

Supplementary Table 4 – Multiple regression analysis models for proteins and symptoms in sicca patients. Values shown are the unstandardized coefficients and 95% confidence intervals.

Variables vs ESSPRI	Individual predictors		Model 1 (Multiple regression)		Model 2 (Forward)	
	β (95% CI)	P value	β (95% CI)	P value	β (95% CI)	P value
Age	0.06 (0.01-0.1)	0.02	-0.01 (-0.1-0.1)	0.44	-	-
BMI	0.16 (0.06-0.26)	0.002	0.06 (-0.1-0.2)	0.16	-	-
HADS D	0.16 (0.03-0.3)	0.01	0.1 (-0.1-0.3)	0.07	0.2 (0.04-0.3)	0.008
C3	1.6 (-0.9-4.2)	0.20	-0.7 (-2.9-5.6)	0.42	-	-
Unstimulated salivary flow	-0.6 (-1.1—0.1)	0.02	-0.12 (-1.7-1.8)	0.50	-	-
ADM	2.2 (1.4-3)	< 0.0001	2.3 (1.3-3.5)	<0.0001	2.5 (1.5-3.4)	< 0.0001
CD40	2.9 (1.5-4.4)	< 0.0001	2.5 (1.2-3.8)	0.001	2.5 (1.2-3.8)	< 0.0001
SPON2	4.3 (1.1-7.7)	0.01	-4.5 (-8--1.3)	0.01	-4.6 (-8.6-1.2)	0.009