

Moving towards endotypes in atopic dermatitis: identification of patient clusters based on serum biomarker analysis

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Table S1. Categorization of serum mediators based on their function.

Serum mediators		
Apoptosis biomarkers	Growth factors	Leukocyte migration biomarkers
Fas	ANG-1	ICAM-1
Fas-L	ANG-2	P-selectin
KIM-1	DKK-1	
	EGF	Neutrophil/granulocyte biomarkers
Atopic biomarker	FGF β	Elastase
Total IgE	HGF	ENA-78 (CXCL5)
	IL-7	GCP-2 (CXCL6)
Chemokines	SCF	GCSF
BLC (CXCL13)	Thrombopoietin	GM-CSF
BRAK (CXCL14)		GRO- α (CXCL1)
CTACK (CCL27)	Immunomodulatory cytokines	IL-6
MPIF (CCL23)	IL-2	NAP-2 (CXCL7)
Eotaxin (CCL11)	IL-10	S100A8
Eotaxin-3 (CCL26)	IL-12	
I-309 (CCL1)	IL-18	Proteases and protease inhibitors
IP-10 (CXCL10)	IL-21	Cathepsin B
ITAC (CXCL11)	IL-22	Cathepsin L
MCP-1 (CCL2)	IL-23	Cathepsin S
MCP-2 (CCL8)	IL-26	Cystatin C
MCP-3 (CCL7)	IL-27	SLPI
MCP-4 (CCL13)	IL-37	MMP-8
MDC (CCL22)	LAIR-1	MMP-9
MPIF (CCL23)	LIF	
MIG (CXCL9)	NGF	Remodelling biomarkers
MIP-1 α (CCL3)	Oncostatin	Endoglin
MIP-1 β (CCL4)	Osteoprotegerin	GDF (MIC-1)
MIP-3 α (CCL20)	PD1	LAP (TGF- β)
PARC (CCL18)	Resistin	LIGHT
PF4 (CXCL4)		MMP-1
RANTES (CCL5)	Inflammatory biomarkers	MMP-3
SDF-1 (CXCL12)	sCD163	TIMP-1
SRPSOX (CXCL16)	M-CSF	
TARC (CCL17)	E-selectin	Th2 cytokines
TECK (CCL25)	IL-1 α	DPP4
XCL-1	IL-1 β	IL-4
	IL-1R1	IL-5
Complement pathway	IL-1R2	IL-9
C5a	IL-1R4	IL-13
	IL-1Ra	Periostin
Epithelial cytokines	sIL-2R α	
IL-25	IL-3	Vascular regulation
IL-33	IL-8 (CXCL8)	PAI-1
S100A12	IL-11	PLGF
TSLP	IL-17	TIE-2
	IL-18BP α	VEGF
Galectins	IL-19	sVEGF-R1
Galectin-1	IL-20	
Galectin-3	IL-31	Viral response
Galectin-9	Osteopontin	IFN- α
	SAA1	IFN- β
Glucose regulation	sSCFR	IFN- γ
Adiponectin	TNF- α	IL-15
Adipsin	TACI	IL-29
Apelin	TNF-R1	TNF- β
Chemerin 4	TNF-R2	
Leptin	TREM1	Other
RBP4	TWEAK	ACE
	VCAM-1	BDNF
	YKL-40 (CHI3L1)	Cathepsin S
		EPO-R
	Innate immunity biomarkers	SOST
	sCD14	
	%Neutralisation of α -toxin-induced lysis	
	Serum Ig targeting alpha toxin	

Table S2. Summary of differentially expressed serum mediators that characterise the 4 AD disease clusters.

Variable	Cluster 1	Cluster 2	Cluster 3	Cluster 4	Healthy controls
Apoptosis biomarkers					
Fas ^{GM}	2229.7 ^{>N,2,3} (1988.6, 2499.7)	1845.1 ^{>N, <1,4} (1701.7, 2000.6)	1717 ^{>N, <1,4} (1551, 1900.8)	2255.4 ^{>N,2,3} (2013.4, 2526.5)	840.9 ^{<1,4, >3} (721.8, 979.7)
Fas-L ^{Med}	40.3 (29.5, 58.4)	54.0 ^{>3} (41.3, 67.4)	30.1 ^{<2,4} (19.2, 56.1)	57.5 ^{>3} (36.5, 93.6)	32.8 (31.0, 37.5)
KIM-1 ^{GM}	7.3x10 ¹ (4.0x10 ¹ , 3.0x10 ²)	8.1x10 ^{1>3} (5.5x10 ¹ , 1.2x10 ²)	3.3x10 ^{1<2,4} (2.2x10 ¹ , 4.9x10 ¹)	1.0x10 ^{2>3} (5.8x10 ¹ , 1.9x10 ²)	4.4x10 ¹ (2.6x10 ¹ , 7.4x10 ¹)
Chemokines					
BLC ^M	68.7 ^{>N,2,3, <4} (55.6, 58.2)	42.4 ^{>3, <1,4} (36.1, 49.4)	28.2 ^{<N,1,2,4} (24.0, 33.0)	116.4 ^{>N,1,2,3} (83.9, 161.5)	43.8 ^{<1,4, >3} (32.9, 58.2)
BRAK ^{Med}	541.8 ^{<4} (318.5, 872.7)	574.5 ^{>N} (389.3, 814.0)	473.7 ^{<4} (243.5, 747.3)	774.7 ^{>N,1,3} (678.3, 1131.9)	310.3 ^{<2,4} (195.5, 487.2)
CTACK ^{Med}	432.4 ^{>3N} (237.6, 720.6)	451.5 ^{>3,N} (287.0, 774.2)	292.5 ^{<2,3,4} (153.9, 555.4)	446.6 ^{>3,N} (283, 821)	2.5 ^{<1,2,3,4} (2.5, 4.3)
Eotaxin ^{Med}	915.3 ^{>N,2,3, <4} (770.5, 1087.4)	543.2 ^{>3,N, <1,4} (458.0, 644.2)	258.3 ^{>N, <1,2,4} (220.8, 302.2)	1747.1 ^{>N,1,2,3} (1407.4, 2168.7)	73.6 ^{<1,2,3,4} (61.4, 88.3)
Eotaxin-3 ^{GM}	1.7 (1.7, 1.7)	1.7 (1.7, 1.7)	1.7 (1.7, 1.7)	1.7 ^{>N,3} (1.7, 6.6)	1.7 ^{<4} (1.7, 1.7)
I309 ^{GM}	75.4 ^{>N,2,3} (61.0, 93.2)	29.4 ^{>N, <1,4} (22.8, 37.9)	25.7 ^{>N, <1,4} (19.2, 34.4)	67 ^{>N,2,3} (47.9, 93.8)	2.1 ^{<1,2,3,4} (2.1, 2.1)
IP-10 ^{GM}	503.7 ^{>N} (442.5, 573.4)	435.7 ^{>N, <4} (397.3, 477.8)	426.4 ^{>N, <4} (383.3, 474.4)	561 ^{>N,2,3} (496.6, 670.1)	280.6 ^{<1,2,3,4} (239.9, 328.1)
ITAC ^{GM}	127.0 ^{>N} (100.5, 160.5)	91.5 ^{>N, <4} (75.6, 110.7)	105.1 ^{>N, <4} (87.2, 126.6)	165.4 ^{>N,2,3} (121.8, 224.6)	38.5 ^{<1,2,3,4} (29.9, 49.4)
MCP-1 ^{GM}	198.4 ^{>N} (177.9, 221.3)	179.2 ^{>N} (164.6, 195.2)	205.0 ^{>N} (186.3, 225.5)	214.2 ^{>N} (190.4, 240.9)	79.1 ^{<1,2,3,4} (67.9, 92.2)
MCP-3 ^{Med}	5.5 ^{<4} (5.5, 39.6)	22.0 ^{<4, >3} (16.8, 49.9)	5.5 ^{<2,4} (5.5, 5.5)	117.8 ^{>N,1,2,3} (27.5, 195)	5.5 ^{<4} (5.5, 29.9)
MCP-4 ^{Med}	55.9 (44.2, 88.8)	61.0 (41.3, 103.5)	57.1 ^{<N} (24.2, 99.1)	67.2 (46.3, 111.1)	103.8 ^{>3} (53.4, 136.7)
MIG ^{GM}	232.4 ^{>N,2,3} (201.6, 268.0)	112.0 ^{<1,4, >3} (98.1, 127.7)	73.1 ^{<N,1,2,4} (60.7, 88.1)	282.7 ^{>N,2,3} (245.5, 325.6)	97.2 ^{<1,4, >3} (90.2, 104.8)
MIP-1α ^{Med}	116.2 ^{>N, <4} (72.7, 154.7)	106.1 ^{>N, <4} (77.6, 152.9)	86.9 ^{>N, <4} (57.4, 143.2)	144.9 ^{>N,1,2,3} (117.1, 191.9)	16.5 ^{<1,2,3,4} (16.5, 16.5)
MIP-3α ^{Med}	10.7 ^{>N,3} (6.5, 15.8)	7.7 ^{>N, <4} (5.4, 10.2)	6.1 ^{>N, <1,4} (2.9, 10.7)	13 ^{>N,2,3} (7.7, 17)	0.6 ^{<1,2,3,4} (0.6, 0.6)
PARC ^{Med}	8.7x10 ^{6>N,2,3,4} (3.9x10 ⁶ , 8.7x10 ⁶)	6.3x10 ^{5<1} (2.7x10 ⁵ , 2.5x10 ⁵)	1.0x10 ^{6<1} (3.7x10 ⁵ , 8.7x10 ⁵)	9.3x10 ^{5<1} (3.8x10 ⁵ , 3.5x10 ⁶)	4.7x10 ^{5<1} (2.8x10 ⁵ , 7.1x10 ⁵)
SDF-1α ^{Med}	7.2x10 ^{2>2,4} (4.3x10 ² , 1.1x10 ³)	2.6x10 ^{1<N,1,3} (2.6x10 ¹ , 2.6x10 ¹)	7.7x10 ^{2>2,4} (4.5x10 ² , 1.3x10 ³)	2.6x10 ^{1<N,1,3} (2.6x10 ¹ , 1.3x10 ²)	3.1x10 ^{2>2,4} (2.4x10 ² , 5.3x10 ²)
TARC ^{Med}	6294.8 ^{>N,2,3} (1765.1, 8943.8)	2880.2 ^{>N, <1} (612.1, 8943.8)	2194.6 ^{>N, <1} (961.4, 8943.8)	2508.1 ^{>N} (1204.2, 5125.6)	96.5 ^{<1,2,3,4} (67.7, 137.1)
TECK ^{Med}	513.5 ^{<N} (323.4, 742.3)	411.3 ^{<N} (286.8, 578.6)	394.9 ^{<N} (292.5, 634.1)	507.5 (412.7, 674.6)	915.9 ^{>1,2,3} (562.1, 1324.1)
XCL-1 ^{Med}	8.5 ^{<N,2,4} (8.5, 8.5)	32.7 ^{>N,1,3, <4} (8.5, 61.3)	8.5 ^{<N,2,4} (8.5, 8.5)	71.9 ^{>N,1,2,3} (50, 118.1)	8.5 ^{>1,3, <4} (8.5, 61.5)
Complement pathway					
C5a ^{GM}	1.7x10 ^{5>2,3,N} (1.4x10 ⁵ , 2.2x10 ⁵)	8.4x10 ^{4<1,4, >N} (7.2x10 ⁴ , 9.8x10 ⁴)	1.1x10 ^{5<1, >N} (8.5x10 ⁴ , 1.3x10 ⁵)	1.4x10 ^{5>N} (1.1x10 ⁵ , 1.8x10 ⁵)	5.2x10 ^{4<1,2,3,4} (4.3x10 ⁴ , 6.3x10 ⁴)
Epithelial cytokines					
IL-25 ^{Med}	1.5x10 ^{4>N,2,3} (1.0x10 ⁴ , 2.2x10 ⁴)	6.7x10 ^{3>N,3, <1,4} (4.9x10 ³ , 9.8x10 ³)	4.5x10 ^{3>N, <1,2,4} (1.6x10 ³ , 6.2x10 ³)	1.8x10 ^{4>N,2,3} (1.5x10 ⁴ , 2.4x10 ⁴)	8.2x10 ^{1<1,2,3,4} (8.2x10 ¹ , 5.8x10 ²)

Variable	Cluster 1	Cluster 2	Cluster 3	Cluster 4	Healthy controls
IL-33 ^{GM}	92.8 ^{>N,2,3} (73.2, 117.8)	38.7 ^{>N,3,<1,4} (31.6, 47.4)	22.5 ^{>N,<1,2,4} (18.8, 27.0)	119.8 ^{>N,2,3} (93.5, 153.5)	9.2 ^{<1,2,3,4} (6.0, 14.1)
S100A12 ^{Med}	4.0x10 ^{6<2,4} (3.4x10 ⁶ , 5.3x10 ⁶)	7.9x10 ^{6>1,3} (6.5x10 ⁶ , 8.7x10 ⁶)	3.7x10 ^{6<2,4} (2.5x10 ⁶ , 5.0x10 ⁶)	8.5x10 ^{6>1,3} (6.6x10 ⁶ , 9.4x10 ⁶)	NA
TSLP ^{GM}	24.4 ^{>N,2,3,<4} (20.9, 28.4)	16.4 ^{>N,3,<1,4} (14.3, 18.7)	6.9 ^{>N,<1,2,4} (5.6, 8.5)	57.0 ^{>N,1,2,3} (45.5, 71.3)	3.3 ^{<1,2,3,4} (1.6, 6.7)
Galectins					
Galectin-1 ^M	2.8x10 ^{4>N,3} (2.6x10 ⁴ , 3.0x10 ⁴)	2.8x10 ^{4>N,3} (2.6x10 ⁴ , 3.0x10 ⁴)	2.4x10 ^{4>N,<1,2} (2.3x10 ⁴ , 2.6x10 ⁴)	2.8x10 ^{4>N} (2.5x10 ⁴ , 3.1x10 ⁴)	1.9x10 ^{4<1,2,3,4} (1.6x10 ⁴ , 2.1x10 ⁴)
Galectin-9 ^{Med}	1x10 ^{1<N} (1x10 ¹ , 1x10 ¹)	1x10 ^{1<N,1,3} (1x10 ¹ , 1x10 ⁵)	1x10 ^{1<N} (1x10 ¹ , 1x10 ⁵)	1x10 ^{1<N,1,3} (1x10 ¹ , 1x10 ⁵)	3.2x10 ^{3>1,2,3,4} (2.3x10 ³ , 4.4x10 ³)
Glucose regulation					
Adipsin ^{Med}	1.7x10 ^{4<4} (1.5x10 ⁴ , 1.7x10 ⁴)	1.7x10 ^{4>N,3} (1.6x10 ⁴ , 1.8x10 ⁴)	1.6x10 ^{4<2,4} (1.5x10 ⁴ , 1.7x10 ⁴)	1.9x10 ^{4>N,1,3} (1.8x10 ⁴ , 2.0x10 ⁴)	1.5x10 ^{4<2,4} (1.4x10 ⁴ , 1.7x10 ⁴)
Chemerin 4 ^{Med}	6.8x10 ^{3<N,4} (6.8x10 ³ , 2.0x10 ⁴)	1.5x10 ^{4<N} (6.8x10 ³ , 2.1x10 ⁴)	6.8x10 ^{3<N,4} (6.8x10 ³ , 1.6x10 ⁴)	2.0x10 ^{4>1,3} (6.8x10 ³ , 2.8x10 ⁴)	1.9x10 ^{4>1,2,3} (1.7x10 ⁴ , 2.5x10 ⁴)
RBP4 ^{Med}	2.7x10 ⁷ (2.5x10 ⁷ , 3.0x10 ⁷)	2.5x10 ^{7<N} (2.2x10 ⁷ , 2.6x10 ⁷)	2.6x10 ^{7<N} (2.3x10 ⁷ , 2.9x10 ⁷)	2.7x10 ⁷ (2.5x10 ⁷ , 2.9x10 ⁷)	2.8x10 ^{7>2,3} (2.7x10 ⁷ , 2.9x10 ⁷)
Growth factors					
EGF ^{GM}	23.0 ^{>2,3} (20.4, 25.8)	12.6 ^{<N,1,4} (11.0, 14.5)	9.5 ^{<N,1,4} (8.3, 10.9)	27.1 ^{>2,3} (23.5, 31.1)	24.3 ^{>2,3} (12.2, 48.6)
FGFb ^{Med}	7.8 ^{<4} (7.8, 7.8)	7.8 ^{<4} (7.8, 7.8)	7.8 ^{<4} (7.8, 7.8)	7.8 ^{>N,1,2,3} (7.8, 136.6)	7.8 ^{<4} (7.8, 7.8)
HGF ^{GM}	1237.5 ^{>N} , >2,3 (1118.1, 1369.6)	824.1 ^{>N,<1,4} (754.3, 901.1)	844.0 ^{>N,<1,4} (770.8, 924.0)	1426.1 ^{>N,2,3} (1249.4, 1627.8)	629.1 ^{<1,2,3,4} (521.8, 758.6)
IL-7 ^M	46.8 ^{>N,2,3} (42.2, 51.4)	30.7 ^{>N,<1,4} (28.3, 33.2)	27.7 ^{>N,<1,4} (25.2, 30.1)	51.6 ^{>N,2,3} (46.0, 57.2)	5.2 ^{<1,2,3,4} (3.8, 6.6)
SCF	29.0 ^{<N,2,4} (16.8, 35.6)	40.4 ^{>1,3} (32.3, 52.1)	22.0 ^{<N,2,4} (14.0, 31.2)	53.6 ^{>1,3} (44.2, 70.0)	50.8 ^{>1,3} (37.7, 59)
Immunomodulatory cytokines					
IL-2 ^{Med}	2.1 ^{<N} (2.1, 2.1)	2.1 (2.1, 2.1)	2.1 ^{<N} (2.1, 2.1)	2.1 ^{>1,2,3} (2.1, 5.4)	2.1 ^{>1,3} (2.1, 2.1)
IL-10 ^{Med}	1.7 ^{<2,4} (1.7, 1.7)	6.4 ^{>N,1,3} (4.4, 8.6)	1.7 ^{<2,4} (1.7, 1.7)	9.0 ^{>N,1,3} (5.5, 12.8)	1.7 ^{<2,4} (1.7, 1.7)
IL-12 ^{Med}	102.7 ^{>N,2,3} (62.3, 154.9)	54.1 ^{>N,3,<1,4} (43.5, 77.3)	29.5 ^{>N,<1,2,4} (11.8, 44.2)	135 ^{>N,2,3} (112.5, 169.2)	3.3 ^{<1,2,3,4} (3.3, 11.7)
IL-18 ^{GM}	370.0 ^{>N,2,3} (318.4, 430.1)	214.2 ^{>N,<1,4} (179.1, 256.2)	284.4 ^{>N,<1} (211.8, 291.3)	314.9 ^{>N,2} (245.3, 404.3)	82.5 ^{<1,2,3,4} (66.4, 102.4)
IL-21 ^{GM}	7.5x10 ^{3>N,3,<4} (6.6x10 ³ , 8.4x10 ³)	6.3x10 ^{3>N,3,<4} (5.7x10 ³ , 3.7x10 ³)	2.4x10 ^{3>N,<1,2,4} (1.7x10 ³ , 3.5x10 ³)	1.7x10 ^{4>N,1,2,3} (1.4x10 ⁴ , 2.2x10 ⁴)	3.5x10 ^{2<1,2,3,4} (1.1x10 ² , 1.2x10 ³)
IL-22 ^{Med}	341.5 ^{>N} (58.4, 554.7)	254.5 ^{>N,3} (106.6, 523.1)	117.0 ^{>N,<2} (31.9, 368.9)	280.8 ^{>N} (104.7, 420.5)	1.8 ^{<1,2,3,4} (1.8, 1.8)
IL-23 ^{Med}	6.8x10 ^{2>N,3} (1.1x10 ² , 1.7x10 ³)	2.3x10 ^{2>N,3} (1.5x10 ² , 9.6x10 ²)	9.8x10 ^{1>1,<4} (1.5x10 ¹ , 7.3x10 ²)	1.3x10 ^{3>N,3} (2.5x10 ² , 4.9x10 ³)	1.5x10 ^{1<1,2,4} (1.5x10 ¹ , 6.9x10 ¹)
IL-26 ^{Med}	5.7x10 ^{2<2,4,>3} (3.8x10 ¹ , 2.2x10 ³)	5.8x10 ^{3>N,1,3} (4.5x10 ³ , 6.9x10 ³)	3.8x10 ^{1<N,1,2,4} (3.8x10 ¹ , 4.2x10 ²)	8.6x10 ^{3>N,1,3} (6.5x10 ³ , 9.9x10 ³)	6.5x10 ^{2<2,4,>3} (1.4x10 ² , 1.5x10 ³)
IL-27 ^{Med}	1.6x10 ^{1<N} (1.6x10 ¹ , 1.6x10 ¹)	1.6x10 ^{1<N,>3} (1.6x10 ¹ , 1.9x10 ¹)	1.6x10 ^{1<N} (1.6x10 ¹ , 1.6x10 ¹)	1.6x10 ^{1<N} (1.6x10 ¹ , 1.9x10 ¹)	1.4x10 ³ (4.7x10 ² , 2.7x10 ³)
IL-37 ^{Med}	1397.3 ^{>N,2,3,<4} (829.1, 2333.5)	950.9 ^{>N,3,<1,4} (554.2, 1276.3)	44.2 ^{>1,2,4} (44.2, 360.4)	2540.7 ^{>N,1,2,3} (2081.5, 3156)	44.2 ^{<1,2,4} (44.2, 44.2)
LAIR-1 ^{Med}	182.1 ^{<4} (182.1, 182.1)	182.1 ^{<4} (182.1, 182.1)	182.1 ^{<4} (182.1, 182.1)	966.4 ^{>N,1,2,3} (182.1, 2330.2)	182.1 ^{<4} (182.1, 182.1)
LIF ^{Med}	4.8 ^{<2,4} (4.8, 4.8)	4.8 ^{>N,1,3} (4.8, 26.2)	4.8 ^{<2,4} (4.8, 4.8)	13.5 ^{>N,1,3} (4.8, 36.9)	4.8 ^{<2,4} (4.8, 4.8)
NGF ^{Med}	4.4 ^{<2,4} (1.8, 10.1)	8.6 ^{<4,>N,1,3} (2.2, 14.5)	1.8 ^{<2,4} (1.8, 5.7)	18.5 ^{>N,1,2,3} (10.4, 24.7)	2.8 ^{<2,4} (1.8, 6.7)

Variable	Cluster 1	Cluster 2	Cluster 3	Cluster 4	Healthy controls
Oncostatin M ^{Med}	11.4 ^{>N} (3.3, 36.7)	20.8 ^{>N} (7.0, 47.7)	3.3 ^{<2,4} (3.3, 11.9)	41.1 ^{>3,N} (14.3, 59.2)	3.3 ^{<1,2,4} (3.3, 3.3)
Osteopetgrin ^{Med}	2091.7 ^{>N} (1885.7, 2388.2)	2738.5 ^{>N,3,<4} (2220.4, 3078.4)	1949.5 ^{>N,<2} (1345.1, 2188.7)	2781.8 ^{>N,2} (2201.4, 3140.6)	7.6 ^{<1,2,3,4} (7.6, 7.6)
PD1 ^{Med}	4.6x10 ⁰ (4.6x10 ⁰ , 5.8x10 ³)	2.3x10 ² (4.6x10 ⁰ , 5.0x10 ³)	4.6x10 ^{0<4} (4.6x10 ⁰ , 3.5x10 ²)	3.1x10 ^{2>3} (9.5x10 ¹ , 2.3x10 ³)	2.1x10 ¹ (4.6x10 ⁰ , 2.0x10 ²)
Resistin ^{Med}	9.8x10 ^{3>2,3} (8.4x10 ³ , 1.2x10 ⁴)	6.5x10 ^{2<N,1,4} (5.6x10 ³ , 7.6x10 ³)	6.2x10 ^{3<N,1} (4.9x10 ³ , 8.1x10 ³)	8.7x10 ^{3>2,3} (7.1x10 ³ , 9.8x10 ³)	9.4x10 ^{3>2,3} (8.9x10 ³ , 1.0x10 ⁴)
Inflammatory biomarkers					
E-selectin ^{Med}	2.3x10 ⁴ (1.8x10 ⁴ , 3.4x10 ⁴)	2.2x10 ^{4<N} (1.5x10 ⁴ , 2.8x10 ⁴)	2.1x10 ^{4<N} (1.6x10 ⁴ , 3.1x10 ⁴)	2.2x10 ^{4<N} (1.7x10 ⁴ , 2.8x10 ⁴)	6.4x10 ^{4>2,3,4} (4.3x10 ⁴ , 8.5x10 ⁴)
IL-1α ^{Med}	22.5 ^{>N,3,<2,4} (8.3, 43.7)	34.6 ^{>N,1,3,<4} (25.4, 50.5)	10.4 ^{<1,2,4} (1.1, 20.1)	77.4 ^{>N,1,2,3} (57.2, 89.5)	4.7 ^{<1,2,4} (3.1, 8.1)
IL-1β ^{GM}	22.7 ^{>N,2,3,<4} (19.7, 26.2)	10.9 ^{>N,3,<1,4} (9.5, 12.4)	7.4 ^{>N,<1,2,4} (6.4, 8.6)	35.5 ^{>N,1,2,3} (28.8, 43.9)	1.1 ^{<1,2,3,4} (0.7, 1.9)
IL-1R1 ^{Med}	88.4 ^{>N,3,<4} (65.2, 112.6)	89.7 ^{>N,3,<4} (68.1, 109.0)	37.2 ^{>N,<1,2,4} (18.8, 55.3)	160.5 ^{>N,1,2,3} (145.6, 201.3)	5.6 ^{<1,2,3,4} (2.7, 7.5)
IL-1R2 ^{Med}	2046.8 (1400.9, 2356.7)	1865.4 ^{>N} (1441.7, 2275.3)	1688.5 (1215.7, 2576.7)	1745 (1366.4, 2120.5)	1473.4 ^{<2} (842.3, 2059.5)
IL-1R4 ^{Med}	3.0 ^{<4} (0.6, 11.6)	9.1 ^{>3} (0.6, 17.2)	0.6 ^{<2,4} (0.6, 9.0)	14.6 ^{>N,3} (4.3, 22.8)	0.6 ^{<4} (0.6, 12.8)
IL-1Ra ^{Med}	37.1 ^{<2,4,>3} (17.3, 69.4)	50.9 ^{>N,1,3,<4} (42.8, 74.0)	4.2 ^{<N,1,2,4} (4.2, 11.2)	118.0 ^{>N,1,2,3} (92.9, 163.0)	4.2 ^{>2,3,4} (4.2, 88.0)
sIL-2Rα ^{Med}	23.5 ^{<N,4} (23.5, 23.5)	23.5 (23.5, 23.5)	23.5 ^{<N,4} (23.5, 23.5)	23.5 ^{>3,3} (23.5, 465.4)	23.5 ^{>3,3} (23.5, 335.7)
IL-6 ^{Med}	13.7 ^{>N,<4} (8.9, 28.5)	23.0 ^{>N,3} (14.9, 51.5)	7.8 ^{>N,<2,4} (3.4, 20.8)	29.7 ^{>N,1,3} (22.6, 53.7)	1.4 ^{<1,2,3,4} (1.4, 10.2)
IL-8 ^{GM}	20.0 ^{>3,<4} (14.2, 28.1)	15.0 ^{<4} (11.4, 17.9)	8.6 ^{<N,1,4} (6.1, 12.0)	68.1 ^{>N,1,2,3} (43.0, 107.9)	20.9 ^{>3,<4} (12.6, 34.4)
IL-11 ^{Med}	4.0 ^{>N,<2,4} (4.0, 26.1)	70.3 ^{>N,1,3} (30.9, 108.8)	4.0 ^{>N,<2,4} (4.0, 9.6)	113.7 ^{>N,1,2,3} (83.5, 227.7)	4 ^{<1,2,3,4} (4, 4)
IL-17 ^{Med}	2.8 ^{<2,4} (2.8, 6.2)	7.7 ^{>N,1,3,<4} (2.8, 17.1)	2.8 ^{<2,4} (2.8, 2.8)	24 ^{>N,1,2,3} (15, 35.4)	2.8 ^{<2,4} (2.8, 2.8)
IL-18BPα ^{Med}	2535.7 ^{>N} (1880.6, 3002.2)	2435.6 ^{<4} (1791.8, 2868.9)	2091.2 ^{<4} (1580.6, 2767.5)	2816.9 ^{>N,2,3} (2604.1, 3281.5)	1416.3 ^{<1,4} (1121.3, 1740.4)
IL-19 ^{Med}	1715.6 ^{>2,3} (1272.7, 2164.4)	1184.9 ^{>3,<1,4} (896.0, 1543.2)	733.7 ^{<1,2,4} (422.2, 911.4)	2316.8 ^{>2,3} (1921.1, 2714.7)	NA
IL-20 ^{Med}	42.4 ^{>N,<4} (42.4, 42.4)	42.4 ^{>N,<4} (42.4, 42.4)	42.4 ^{>N,<4} (42.4, 42.4)	122.2 ^{>N,1,2,3} (102.1, 160.5)	42.4 ^{<1,2,3} (42.4, 42.4)
TNF-α ^{Med}	1.1 ^{<2,4} (1.1, 1.1)	3.6 ^{>N,1,3,<4} (1.1, 6.7)	1.1 ^{<2,4} (1.1, 1.1)	9.3 ^{>N,1,2,3} (2.5, 24.8)	1.1 ^{<2,4} (1.1, 1.1)
TNF-R1 ^M	6.4x10 ^{3>N,2} (5.8x10 ³ , 7.0x10 ³)	5.1x10 ^{3>N,<1} (4.7x10 ³ , 5.5x10 ³)	5.7x10 ^{3>N} (5.2x10 ³ , 6.1x10 ³)	5.5x10 ^{3>N} (4.6x10 ³ , 6.4x10 ³)	5.9x10 ^{2<1,2,3,4} (2.8x10 ² , 9.1x10 ²)
TNF-R2 ^M	2324.9 ^{>N,2,3} (2106.0, 2543.8)	1993.2 ^{>N,<1} (1875.4, 2110.9)	1876.2 ^{>N,<1,4} (1722.1, 2030.4)	2310 ^{>N,3} (2067.3, 2552.6)	1476.4 ^{<1,2,3,4} (1351.6, 1601.1)
TREM1 ^{GM}	8.3x10 ^{1>N,3,<4} (6.3x10 ¹ , 1.1x10 ²)	8.7x10 ^{1>N,3,<4} (6.9x10 ¹ , 1.1x10 ²)	2.4x10 ^{1>N,<1,2,4} (1.8x10 ¹ , 3.3x10 ¹)	1.7x10 ^{2>N,1,2,3} (1.4x10 ² , 1.9x10 ²)	1.4x10 ^{1<1,2,3,4} (9.3x10 ⁰ , 2.1x10 ¹)
YKL40 ^{Med}	5.6x10 ^{4>2} (4.4x10 ⁴ , 8.8x10 ⁴)	3.8x10 ^{4<1} (3.2x10 ⁴ , 4.4x10 ⁴)	4.8x10 ⁴ (3.3x10 ⁴ , 6.7x10 ⁴)	3.9x10 ⁴ (3.2x10 ⁴ , 5.8x10 ⁴)	5.0x10 ⁴ (4.5x10 ⁴ , 6.6x10 ⁴)
Innate immunity biomarkers					
sCD14 ^{Med}	2.0x10 ^{6>2,3,4} (1.5x10 ⁶ , 2.5x10 ⁶)	1.2x10 ^{6<N,1} (1.1x10 ⁶ , 1.3x10 ⁵)	1.6x10 ^{6<N,1} (1.1x10 ⁶ , 2.0x10 ⁶)	1.3x10 ^{6<N,1} (1.1x10 ⁶ , 1.7x10 ⁶)	2.3x10 ^{6>2,3,4} (2.1x10 ⁶ , 2.5x10 ⁶)
%Neutralisation of α-toxin-induced lysis ^{Med}	50.1 ^{>N,3} (28.8, 86.1)	25.1 ^{>N} (15.9, 48.0)	34.0 ^{>N,<1} (9.0, 57.0)	28.6 ^{>N} (14.0, 72.5)	10.8 ^{<1,2,3,4} (0.9, 20.5)
Leukocyte migration biomarkers					
ICAM-1 ^{Med}	6.2x10 ^{5>N,2,3} (5.3x10 ⁵ , 7.7x10 ⁵)	4.1x10 ^{5<1,>N} (3.7x10 ⁵ , 4.9x10 ⁵)	4.9x10 ^{5<1,>N} (3.8x10 ⁵ , 6.7x10 ⁵)	4.8x10 ^{5>N} (3.9x10 ⁵ , 5.6x10 ⁵)	2.9x10 ^{5<1,2,3,4} (2.2x10 ⁵ , 3.4x10 ⁵)

Variable	Cluster 1	Cluster 2	Cluster 3	Cluster 4	Healthy controls
P-selectin	3.6x10 ⁵ (2.0x10 ⁵ , 1.4x10 ⁶)	3.8x10 ^{5<N} (2.1x10 ⁵ , 5.7x10 ⁵)	2.9x10 ^{5<N} (2.0x10 ⁵ , 4.7x10 ⁵)	3.7x10 ⁵ (2.7x10 ⁵ , 5.8x10 ⁵)	5.5x10 ^{5>2,3} (3.5x10 ⁵ , 1.4x10 ⁶)
Neutrophil/Granulocyte biomarkers					
Elastase ^{Med}	2.7x10 ^{4<4,>N} (1.3x10 ⁴ , 8.6x10 ⁴)	2.3x10 ^{4<4,>N} (1.4x10 ⁴ , 8.6x10 ⁴)	2.2x10 ^{4<4,>N} (1.0x10 ⁴ , 8.6x10 ⁴)	8.6x10 ^{4>1,2,3,N} (8.6x10 ⁴ , 8.6x10 ⁴)	1.2x10 ^{4<1,2,3,4} (7.1x10 ³ , 1.6x10 ⁴)
GCP-2 ^{GM}	160.3 ^{>N,<4} (120.9, 212.6)	127.1 ^{>N,<4} (101.5, 159.1)	102.6 ^{<4} (77.1, 136.6)	399.8 ^{>N,1,2,3} (304.8, 524.5)	64.2 ^{<1,2,4} (34.4, 119.8)
GM-CSF ^{Med}	7.6 ^{<4,>N} (7.6, 7.6)	7.6 ^{<4,>N} (7.6, 7.6)	7.6 ^{<4,>N} (7.6, 7.6)	17.3 ^{>N,1,2,3} (7.6, 35.3)	7.6 ^{<1,2,3,4} (7.6, 7.6)
GRO-α ^{GM}	125.5 ^{>3,<4} (110.8, 142.3)	110.3 ^{>3,<4} (102.2, 118.9)	74.7 ^{<1,2,4} (67.3, 82.8)	245.1 ^{>1,2,3} (198.0, 303.4)	NA
S100A8 ^{Med}	1.3x10 ^{5>N} (1.2x10 ⁵ , 1.6x10 ⁵)	1.6x10 ^{5>N,3} (1.4x10 ⁵ , 1.8x10 ⁵)	1.3x10 ^{5>N,<2} (1.1x10 ⁵ , 1.5x10 ⁵)	1.6x10 ^{5>N} (1.5x10 ⁵ , 1.9x10 ⁵)	1.2x10 ^{4<1,2,3,4} (8.4x10 ³ , 1.6x10 ⁴)
Proteases and protease inhibitors					
Cathepsin S ^{Med}	1.6x10 ^{4>2,3} (1.4x10 ⁴ , 2.0x10 ⁴)	1.3x10 ^{4<N,1} (1.1x10 ⁴ , 1.4x10 ⁴)	1.3x10 ^{4<N,1} (1.1x10 ⁴ , 1.6x10 ⁴)	1.4x10 ⁴ (1.2x10 ⁴ , 1.7x10 ⁴)	1.7x10 ^{4>2,3} (1.6x10 ⁴ , 1.9x10 ⁴)
SLPI ^{Med}	9.4x10 ^{3>N,<2,4} (8.2x10 ³ , 1.9x10 ⁴)	1.9x10 ^{4>N,1,3} (1.9x10 ⁴ , 1.9x10 ⁴)	9.2x10 ^{3>N,<2,4} (8.1x10 ³ , 1.9x10 ⁴)	1.9x10 ^{4>N,1,3} (1.9x10 ⁴ , 1.9x10 ⁴)	6.4x10 ^{3<1,2,3,4} (5.9x10 ³ , 7.2x10 ³)
MMP-8 ^{GM}	1.9x10 ^{5<N,>2,3} (1.6x10 ⁵ , 2.2x10 ⁵)	1.0x10 ^{5<N,1,4} (8.9x10 ⁴ , 1.2x10 ⁵)	9.5x10 ^{4<N,1,4} (8.5x10 ⁴ , 1.1x10 ⁵)	1.6x10 ^{5<N,>2,3} (1.3x10 ⁵ , 1.8x10 ⁵)	3.0x10 ^{5>1,2,3,4} (2.7x10 ⁵ , 3.2x10 ⁵)
Remodelling biomarkers					
Endoglin ^{GM}	3.2x10 ^{3>N,2} (2.9x10 ³ , 3.6x10 ³)	2.3x10 ^{3>N,<1,3} (2.0x10 ³ , 2.5x10 ³)	2.8x10 ^{3>N,2} (2.5x10 ³ , 3.1x10 ³)	2.8x10 ^{3>N} (2.4x10 ³ , 3.2x10 ³)	1.2x10 ^{3<1,2,3,4} (1.1x10 ³ , 1.3x10 ³)
LAP-TGFα ^{GM}	3.0x10 ^{5>N,<2,4} (2.8x10 ⁵ , 3.2x10 ⁵)	3.5x10 ^{5>N,1,3} (3.4x10 ⁵ , 3.7x10 ⁵)	2.9x10 ^{5>N,<2,4} (2.7x10 ⁵ , 3.1x10 ⁵)	3.7x10 ^{5>N,1,3} (3.4x10 ⁵ , 4.0x10 ⁵)	1.5x10 ^{5<1,2,3,4} (1.4x10 ⁵ , 1.7x10 ⁵)
LIGHT ^{Med}	4.0 ^{<N,2,4} (4.0, 4.0)	151.7 ^{>1,3,<4} (4.0, 257.3)	4.0 ^{<N,2,4} (4.0, 4.0)	331.3 ^{>N,1,2,3} (209.9, 610.4)	29.8 ^{>1,3,<4} (4, 73.3)
MMP-1 ^{GM}	6.0x10 ^{4>N,2} (5.2x10 ⁴ , 6.9x10 ⁴)	3.40x10 ^{4<1,3} (2.5x10 ⁴ , 3.5x10 ⁴)	4.5x10 ^{4>2} (3.6x10 ⁴ , 5.5x10 ⁴)	4.0x10 ⁴ (3.2x10 ⁴ , 5.0x10 ⁴)	3.5x10 ^{4<1} (2.8x10 ⁴ , 4.3x10 ⁴)
TIMP-1 ^M	6.0x10 ^{5>N,2,3,4} (5.7x10 ⁵ , 6.3x10 ⁵)	4.3x10 ^{5<N,1,3,4} (4.2x10 ⁵ , 4.5x10 ⁵)	5.3x10 ^{5<1,>2} (5.0x10 ⁵ , 5.6x10 ⁵)	4.8x10 ^{5<1} (4.5x10 ⁵ , 5.0x10 ⁵)	5.0x10 ^{5<1,>2} (4.8x10 ⁵ , 5.2x10 ⁵)
Th2 cytokines					
IL-4 ^{Med}	6.7 ^{>N,2,3,<4} (4.1, 9.5)	3.6 ^{>N,3,<1,4} (2.5, 5.5)	1.4 ^{>N,<1,2,4} (0.3, 2.2)	10.2 ^{>N,1,2,3} (7.8, 14.1)	0.3 ^{<1,2,3,4} (0.3, 0.3)
IL-5 ^{GM}	323.7 ^{>N,2,3} (278.9, 375.6)	171.9 ^{>N,<1,4} (151.2, 195.6)	124.9 ^{>N,<1,4} (101.6, 153.6)	506.8 ^{>N,2,3} (415.4, 618.3)	5.0 ^{<1,2,3,4} (2.6, 9.6)
IL-9 ^{GM}	145.9 ^{>N,2,3,<4} (122.0, 174.4)	75.7 ^{>N,3,<1,4} (64.4, 89.0)	34.3 ^{>N,<1,2,4} (25.5, 46.1)	349.2 ^{>N,1,2,3} (265.6, 459)	12.2 ^{<1,2,3,4} (4.8, 31.3)
IL-13 ^{GM}	97.1 ^{>N,2,3,<4} (83.8, 112.6)	51.9 ^{>N,3,<1,4} (45.9, 58.6)	39.9 ^{>N,<1,4} (34.2, 46.7)	170.3 ^{>N,1,2,3} (134.2, 216.2)	23.3 ^{<1,2,3,4} (10.3, 52.8)
Periostin ^{GM}	18.6 ^{>N} (14.5, 24.0)	13.2 ^{>N} (11.4, 15.4)	15.0 ^{>N} (13.1, 17.3)	11.8 (9.4, 14.8)	6.7 ^{<1,2,3} (3.4, 13.4)
Vascular regulation					
PAI-1 ^{Med}	8.7x10 ^{5>4} (7.4x10 ⁵ , 1.2x10 ⁶)	1.0x10 ^{5<N,1,3} (4.4x10 ⁴ , 1.0x10 ⁵)	8.6x10 ^{5>2,3} (6.9x10 ⁵ , 1.1x10 ⁶)	4.4x10 ^{5<N,1,3} (4.4x10 ⁴ , 7.8x10 ⁵)	9.5x10 ^{5>2,4} (6.6x10 ⁵ , 1.3x10 ⁶)
PLGF ^{Med}	40.5 ^{<4} (10.6, 54.2)	45.4 ^{<4} (16.1, 59.7)	31.0 ^{<4} (16.7, 52.3)	64.3 ^{>1,2,3} (40.2, 89.7)	40.8 (26.7, 61.6)
TIE-2 ^M	9.6x10 ^{3>N,2} (8.6x10 ³ , 1.1x10 ⁴)	7.8x10 ^{3>N,<1} (7.3x10 ³ , 8.3x10 ³)	8.6x10 ^{3>N} (7.9x10 ³ , 9.2x10 ³)	8.5x10 ^{3>N} (7.9x10 ³ , 9.1x10 ³)	2.6x10 ^{3<1,2,3,4} (2.3x10 ³ , 2.8x10 ³)
VEGF ^{Med}	873.5 ^{>N,2} (687.4, 1295.9)	493.1 ^{>N,<1,3,4} (317.1, 871.3)	793.0 ^{>N,2} (439.4, 1295.8)	945.5 ^{>N,2} (535.5, 1278.7)	346.4 (170.2, 602.5)
sVEGF-R1 ^{GM}	1.6x10 ^{3>3,<4} (1.3x10 ³ , 1.8x10 ³)	1.2x10 ^{3<4} (1.1x10 ³ , 1.5x10 ³)	1.1x10 ^{3<N,1,4} (9.7x10 ² , 1.2x10 ³)	2.2x10 ^{3>N,1,2,3} (1.9x10 ³ , 2.5x10 ³)	1.5x10 ^{3>3,<4} (1.2x10 ³ , 1.9x10 ³)
Viral response					

Variable	Cluster 1	Cluster 2	Cluster 3	Cluster 4	Healthy controls
IFN- α ^{Med}	95.1 ^{>N,2} (82.0, 100.2)	41.5 ^{>N,<1,3,4} (30.3, 51.2)	93.3 ^{>N,2} (81.8, 101.0)	63.2 ^{>N,2} (55.5, 110.1)	0.7 ^{<1,2,3,4} (0.7, 2.9)
IFN- β ^{Med}	9.4x10 ^{1<N,2,4,>3} (9.7x10 ⁰ , 1.9x10 ²)	1.9x10 ^{2<N,4,>1,3} (1.4x10 ² , 2.9x10 ²)	9.7x10 ^{0<N,1,2,4} (9.7x10 ⁰ , 4.2x10 ¹)	6.5x10 ^{2>1,2,3} (3.4x10 ² , 1.1x10 ³)	3.2x10 ^{2>1,2,3} (1.5x10 ² , 9.1x10 ²)
IFN- γ ^{Med}	4.4 ^{<2,4} (4.4, 4.4)	11.0 ^{<4,>1,3} (4.4, 27.0)	4.4 ^{<2,4} (4.4, 4.4)	95.2 ^{>N,1,2,3} (28.4, 178.2)	4.4 ^{<4} (4.4, 4.4)
IL-15 ^{Med}	174.9 ^{>N,2,3,<4} (101.7, 238.8)	84.6 ^{>N,3,<1,4} (63.0, 132.4)	44.6 ^{>N,<1,2,4} (22.2, 65.4)	242.7 ^{>N,1,2,3} (213, 354.1)	3.7 ^{<1,2,3,4} (1.5, 6.4)
IL-29 ^{Med}	1.0x10 ¹ (1.0x10 ¹ , 1.0x10 ¹)	1.0x10 ^{1>3} (1.0x10 ¹ , 1.1x10 ²)	1.0x10 ^{1<2,4} (1.0x10 ¹ , 1.0x10 ¹)	1.0x10 ^{1>3} (1.0x10 ¹ , 2.0x10 ¹)	1.0x10 ¹ (1.0x10 ¹ , 1.0x10 ¹)
TNF- β ^{Med}	1.5 ^{<4} (1.5, 1.5)	3.5 ^{<4} (1.5, 10.3)	1.5 ^{<N,4} (1.5, 1.5)	44.2 ^{>N,1,2,3} (5.3, 118.6)	1.5 ^{<3,4} (1.5, 12.8)
Other					
ACE ^{GM}	3.6x10 ^{5>2,3} (3.2x10 ⁵ , 4.1x10 ⁵)	2.6x10 ^{5<N,1} (2.4x10 ⁵ , 2.9x10 ⁵)	2.7x10 ^{5<N,1} (2.3x10 ⁵ , 3.2x10 ⁵)	2.9x10 ⁵ (2.5x10 ⁵ , 3.4x10 ⁵)	4.0x10 ^{5>2,3} (3.6x10 ⁵ , 4.3x10 ⁵)
BDNF ^M	2.3x10 ^{5<N} (2.2x10 ⁵ , 2.5x10 ⁵)	2.3x10 ^{5<N} (2.1x10 ⁵ , 2.4x10 ⁵)	2.0x10 ^{5<N,4} (1.8x10 ⁵ , 2.2x10 ⁵)	2.6x10 ^{5<N} (2.5x10 ⁵ , 2.8x10 ⁵)	5.5x10 ^{5>1,2,3,4} (4.7x10 ⁵ , 6.2x10 ⁵)
EPO-R ^{Med}	13.7 ^{<4} (13.7, 225.1)	26.0 ^{<4} (13.7, 473.0)	13.7 ^{<4} (13.7, 45.3)	914.5 ^{>N,1,2,3} (123.9, 1935.1)	13.7 ^{<4} (13.7, 13.7)
MIF ^{GM}	4.4x10 ^{3<N,>2,3} (3.8x10 ³ , 5.0x10 ³)	2.6x10 ^{3<N,1,4} (2.2x10 ³ , 3.1x10 ³)	2.2x10 ^{3<N,1,4} (2.0x10 ³ , 2.5x10 ³)	4.4x10 ^{3<N,>2,3} (3.8x10 ³ , 5.2x10 ³)	7.0x10 ^{3>1,2,3,4} (5.2x10 ³ , 9.3x10 ³)
NAP-2 ^{Med}	4.0x10 ^{7>2,3} (3.3x10 ⁷ , 4.0x10 ⁷)	1.5x10 ^{7<1} (1.3x10 ⁷ , 1.6x10 ⁷)	2.8x10 ^{7<1} (1.5x10 ⁷ , 4.0x10 ⁷)	1.6x10 ⁷ (1.5x10 ⁷ , 1.7x10 ⁷)	1.9x10 ⁷ (1.7x10 ⁷ , 4.0x10 ⁷)
SOST ^{Med}	11.7 ^{<N,2,4} (11.7, 11.7)	11.7 ^{<N,4,>1,3} (11.7, 11.7)	11.7 ^{<N,2,4} (11.7, 11.7)	272.6 ^{>N,1,2,3} (104, 389.5)	11.7 ^{<4,>1,2,3} (11.7, 87.7)
Trappin-2 ^{Med}	37.1 ^{<N,2,4} (14.9, 90.3)	236.4 ^{<N,>1,3} (164.9, 384.5)	36.4 ^{<N,2,4} (13.2, 77.9)	181.5 ^{>N,>1,3} (162.0, 240.0)	2226.0 (1923.7, 2576.3)

Variable types are shown in superscript next to variable name; M (mean) for normally distributed data, Med (median) for log-non normally distributed data, GM (geometric mean) for log-normally distributed data. Numbers shown in brackets after measure of central tendency are 95% confidence intervals (for mean and geomean) or interquartile range (median). Superscript numbers after measure of central tendency are groups that are significantly different.

Table S3. Table showing analytes that do not significantly differ between clusters.

Variable	Cluster 1	Cluster 2	Cluster 3	Cluster 4	Healthy controls
Adiponectin ^{Med}	1.9x10 ⁸ (1.1x10 ⁸ , 4.2x10 ⁸)	1.3x10 ⁸ (8.6x10 ⁷ , 1.7x10 ⁸)	1.8x10 ⁸ (9.6x10 ⁷ , 4.2x10 ⁸)	1.5x10 ⁸ (1.3x10 ⁸ , 1.8x10 ⁸)	1.3x10 ⁸ (9.0x10 ⁷ , 1.6x10 ⁸)
Ang1 ^{Med}	5.4x10 ⁴ (4.2x10 ⁴ , 7.0x10 ⁴)	4.9x10 ⁴ (4.2x10 ⁴ , 5.6x10 ⁴)	5.5x10 ⁴ (4.3x10 ⁴ , 7.1x10 ⁴)	6.2x10 ⁴ (5.0x10 ⁴ , 7.0x10 ⁴)	4.4x10 ⁴ (3.6x10 ⁴ , 6.0x10 ⁴)
Ang2 ^{Med}	1.4x10 ^{4<N} (1.3x10 ⁴ , 1.6x10 ⁴)	1.2x10 ^{4<N} (1.1x10 ⁴ , 1.4x10 ⁴)	1.3x10 ^{4<N} (1.1x10 ⁴ , 1.6x10 ⁴)	1.3x10 ^{4<N} (1.2x10 ⁴ , 1.5x10 ⁴)	1.9x10 ^{4>1,2,3,4} (1.8x10 ⁴ , 2.8x10 ⁴)
Apelin ^{Med}	6.5x10 ^{4>N} (6.1x10 ⁴ , 7.0x10 ⁴)	6.1x10 ^{4>N} (5.7x10 ⁴ , 6.4x10 ⁴)	6.4x10 ^{4>N} (5.8x10 ⁴ , 6.8x10 ⁴)	6.4x10 ^{4>N} (6.3x10 ⁴ , 6.6x10 ⁴)	4.7x10 ^{4<1,2,3,4} (3.8x10 ⁴ , 5.2x10 ⁴)
Cathepsin B ^M	1.7x10 ^{4>N} (1.5x10 ⁴ , 1.9x10 ⁴)	1.6x10 ^{4>N} (1.4x10 ⁴ , 1.7x10 ⁴)	1.7x10 ^{4>N} (1.5x10 ⁴ , 1.9x10 ⁴)	1.6x10 ^{4>N} (1.3x10 ⁴ , 1.8x10 ⁴)	9.9x10 ^{3<1,2,3,4} (8.5x10 ³ , 1.1x10 ⁴)
Cathepsin L ^{Med}	1.3x10 ^{3>2,3,<N} (1.1x10 ³ , 1.7x10 ³)	9.5x10 ^{2<1,N} (6.6x10 ² , 1.2x10 ³)	1.1x10 ^{3<1,N} (7.3x10 ² , 1.4x10 ³)	1.1x10 ^{3<N} (8.2x10 ² , 1.3x10 ³)	6.4x10 ^{3>1,2,3,4} (5.0x10 ³ , 8.0x10 ³)
Cystatin C ^{Med}	2.1x10 ^{5>N} (1.9x10 ⁵ , 2.4x10 ⁵)	1.9x10 ^{5>N} (1.7x10 ⁵ , 2.1x10 ⁵)	2.0x10 ^{5>N} (1.7x10 ⁵ , 2.2x10 ⁵)	1.9x10 ^{5>N} (1.6x10 ⁵ , 2.1x10 ⁵)	1.2x10 ^{5<1,2,3,4} (8.4x10 ⁴ , 1.5x10 ⁵)
DKK-1 ^M	2299.1 (2179.8, 2418.4)	2177.0 (2117.5, 2236.4)	2391.9 (2320.5, 2463.4)	2282.3 (2205, 2359.7)	2403.4 (1913.3, 2893.4)
DPP4 ^{GM}	614.4 (557.0, 678.3)	586.2 (541.2, 634.8)	619.2 (578.8, 662.4)	570.9 (504.2, 646.5)	539.3 (498.5, 583.3)
ENA-78 ^{Med}	2.5x10 ^{3>N} (1.9x10 ³ , 4.0x10 ³)	2.1x10 ^{3>N} (1.2x10 ³ , 2.8x10 ³)	2.7x10 ^{3>N} (2.1x10 ³ , 4.0x10 ³)	3.7x10 ^{3>N} (2.4x10 ³ , 5.3x10 ³)	7.4x10 ^{2<1,2,3,4} (4.8x10 ² , 1.2x10 ³)
Galectin-3 ^M	2.2x10 ^{4>3} (1.9x10 ⁴ , 2.6x10 ⁴)	2.1x10 ^{4<N} (1.8x10 ⁴ , 2.3x10 ⁴)	1.8x10 ^{4<N,1,4} (1.6x10 ⁴ , 1.9x10 ⁴)	2.4x10 ^{4>3} (2.1x10 ⁴ , 2.8x10 ⁴)	2.8x10 ^{4>2,3} (2.6x10 ⁴ , 3.0x10 ⁴)
GCSF ^{GM}	8.4x10 ^{3>N} (6.4x10 ³ , 1.5x10 ⁴)	8.6x10 ^{3>N} (6.3x10 ³ , 1.5x10 ⁴)	7.0x10 ^{3>N} (5.4x10 ³ , 1.1x10 ⁴)	1.1x10 ^{4>N} (8.3x10 ³ , 1.9x10 ⁴)	1.8x10 ^{4<1,2,3,4} (1.8x10 ³ , 1.8x10 ⁴)
GDF ^{Med}	1.1x10 ^{3>2} (8.4x10 ² , 1.6x10 ³)	7.4x10 ^{2<1} (6.7x10 ² , 9.9x10 ²)	1.0x10 ³ (7.3x10 ² , 1.6x10 ³)	8.8x10 ² (7.6x10 ² , 1.1x10 ³)	1.2x10 ³ (1.0x10 ³ , 1.3x10 ³)
IL-3 ^{Med}	2.5 ^{>N} (2.5, 283.2)	2.5 ^{>N} (2.5, 400.9)	2.5 ^{>N} (2.5, 242.4)	167.7 ^{>N} (2.5, 536.1)	2.5 ^{<1,2,3,4} (2.5, 2.5)
IL-16 ^M	2.0x10 ³ (1.7x10 ³ , 2.3x10 ³)	1.9x10 ³ (1.7x10 ³ , 2.1x10 ³)	1.2x10 ³ (1.1x10 ³ , 1.4x10 ³)	2.1x10 ³ (1.9x10 ³ , 2.3x10 ³)	2.0x10 ³ (6.8x10 ² , 3.4x10 ³)
IL-31 ^{Med}	1.2x10 ¹ (1.2x10 ¹ , 3.1x10 ³)	1.2x10 ¹ (1.2x10 ¹ , 2.5x10 ³)	1.2x10 ¹ (1.2x10 ¹ , 1.2x10 ¹)	1.2x10 ¹ (1.2x10 ¹ , 4.6x10 ¹)	1.2x10 ¹ (1.2x10 ¹ , 2.5x10 ¹)
Leptin ^{Med}	8.0x10 ^{2<N} (8.0x10 ² , 2.4x10 ³)	8.0x10 ^{2<N} (8.0x10 ² , 1.4x10 ³)	8.0x10 ^{2<N} (8.0x10 ² , 1.4x10 ³)	8.0x10 ^{2<N} (8.0x10 ² , 2.0x10 ³)	3.6x10 ^{3>1,2,3,4} (2.8x10 ³ , 5.4x10 ³)
MCP-2 ^{Med}	50.4 (42.2, 62.7)	41.8 (34.5, 56.9)	45.4 (33.6, 56.6)	53 ^{>N} (46.1, 65.8)	36.5 ^{<4} (28.7, 44.2)
MCSF ^{GM}	7.6 (7.6, 7.6)	7.6 (7.6, 7.6)	7.6 (7.6, 7.6)	7.6 (7.6, 7.6)	7.6 (7.6, 7.6)
MDC ^{Med}	3352.0 ^{>N} (2210.0, 9001.0)	2493.3 ^{>N} (1764.5, 4146.5)	2330.7 ^{>N} (1784.7, 4040.4)	2975.6 ^{>N} (2125.7, 4217.4)	737.8 ^{<1,2,3,4} (681.2, 873.9)
MIP-1β ^{GM}	138.2 ^{>N} (119.8, 159.4)	108.3 ^{>N} (95.0, 123.4)	127.7 ^{>N} (112.7, 144.7)	132.6 ^{>N} (105.4, 166.9)	41.0 ^{<1,2,3,4} (32.8, 51.4)
MIP-3β ^{Med}	1.6x10 ⁰ (1.6x10 ⁰ , 1.2x10 ¹)	1.6x10 ⁰ (1.6x10 ⁰ , 1.1x10 ⁴)	1.6x10 ⁰ (1.6x10 ⁰ , 1.6x10 ⁰)	1.6x10 ⁰ (1.6x10 ⁰ , 1.5x10 ¹)	6.6x10 ¹ (3.9x10 ¹ , 1.1x10 ²)
MMP-3 ^{GM}	3.5x10 ^{4>2} (3.0x10 ⁴ , 4.2x10 ⁴)	2.4x10 ^{4<1} (2.1x10 ⁴ , 2.8x10 ⁴)	2.6x10 ⁴ (2.2x10 ⁴ , 3.2x10 ⁴)	2.8x10 ⁴ (2.2x10 ⁴ , 3.5x10 ⁴)	2.5x10 ⁴ (2.1x10 ⁴ , 3.0x10 ⁴)
MMP-9 ^{Med}	1.6x10 ⁶ (1.6x10 ⁶ , 1.6x10 ⁶)	1.6x10 ⁶ (1.6x10 ⁶ , 1.6x10 ⁶)	1.6x10 ⁶ (1.6x10 ⁶ , 1.6x10 ⁶)	1.6x10 ⁶ (1.6x10 ⁶ , 1.6x10 ⁶)	1.6x10 ⁶ (1.6x10 ⁶ , 1.6x10 ⁶)
MPIF ^{Med}	480.4 ^{>N} (287.2, 643.2)	341.6 ^{>N} (198.3, 536.6)	398.4 ^{>N} (254.9, 491.8)	362.8 ^{>N} (173.4, 508)	94.4 ^{<1,2,3,4} (40.9, 170.8)

Osteopontin ^{GM}	3.5x10 ^{4<N} (3.1x10 ⁴ , 4.1x10 ⁴)	3.2x10 ^{4<N} (2.7x10 ⁴ , 3.7x10 ⁴)	3.6x10 ^{4<N} (3.0x10 ⁴ , 4.3x10 ⁴)	3.6x10 ^{4<N} (3.1x10 ⁴ , 4.3x10 ⁴)	1.2x10 ^{4>1,2,3,4} (9.2x10 ³ , 1.5x10 ⁴)
PF4 ^{Med}	2.0x10 ^{6<N} (2.0x10 ⁶ , 2.0x10 ⁶)	2.0x10 ^{6<N} (2.0x10 ⁶ , 2.0x10 ⁶)	2.0x10 ^{6<N} (2.0x10 ⁶ , 2.0x10 ⁶)	2.0x10 ^{6<N} (2.0x10 ⁶ , 2.0x10 ⁶)	4.7x10 ^{6>1,2,3,4} (3.2x10 ⁶ , 6.6x10 ⁶)
RANTES ^{Med}	1.6x10 ^{5<N, >2,3} (1.3x10 ⁵ , 2.5x10 ⁵)	8.4x10 ^{4<N,1,3,4} (4.8x10 ⁴ , 1.4x10 ⁵)	1.4x10 ^{5<N,1, >2} (6.3x10 ⁴ , 2.2x10 ⁵)	1.6x10 ^{5<N, >2} (1.1x10 ⁵ , 2.1x10 ⁵)	6.4x10 ^{5>1,2,3,4} (4.0x10 ⁵ , 9.1x10 ⁵)
SAA1 ^{Med}	1.0x10 ⁸ (1.0x10 ⁸ , 1.0x10 ⁸)	1.0x10 ⁸ (1.0x10 ⁸ , 1.0x10 ⁸)	1.0x10 ⁸ (1.0x10 ⁸ , 1.0x10 ⁸)	1.0x10 ⁸ (1.0x10 ⁸ , 1.0x10 ⁸)	1.0x10 ⁸ (5.2x10 ⁷ , 1.0x10 ⁸)
sCD163 ^{Med}	4.3x10 ^{4<N} (4.3x10 ⁴ , 4.3x10 ⁴)	4.3x10 ^{4<N} (4.3x10 ⁴ , 4.3x10 ⁴)	4.3x10 ^{4<N} (4.3x10 ⁴ , 4.3x10 ⁴)	4.3x10 ^{4<N} (4.3x10 ⁴ , 4.3x10 ⁴)	1.4x10 ^{5>1,2,3,4} (1.2x10 ⁵ , 1.7x10 ⁵)
Serum Ig targeting alpha toxin ^{GM}	93.9 ^{>N} (70.6, 125.0)	93.8 ^{>N} (65.5, 134.5)	65.7 ^{>N} (51.7, 83.3)	117.4 ^{>N} (71.6, 192.8)	11.2 ^{<1,2,3,4} (7.0, 18.0)
SRPSOX ^{Med}	8.9x10 ² (6.3x10 ² , 6.0x10 ³)	8.6x10 ² (7.3x10 ² , 2.6x10 ³)	8.0x10 ² (6.1x10 ² , 1.4x10 ³)	8.3x10 ² (6.5x10 ² , 3.5x10 ³)	NA
sSCFR ^{Med}	6.5x10 ^{4>N} (5.8x10 ⁴ , 8.0x10 ⁴)	6.4x10 ^{4>N} (4.9x10 ⁴ , 7.6x10 ⁴)	6.2x10 ^{4>N} (4.9x10 ⁴ , 8.4x10 ⁴)	6.4x10 ^{4>N} (5.4x10 ⁴ , 7.5x10 ⁴)	4.2x10 ^{4<1,2,3,4} (3.8x10 ⁴ , 4.4x10 ⁴)
TACI ^{Med}	8.0x10 ^{2<N} (8.0x10 ² , 8.0x10 ²)	8.0x10 ^{2<N} (8.0x10 ² , 8.0x10 ²)	8.0x10 ^{2<N} (8.0x10 ² , 8.0x10 ²)	8.0x10 ^{2<N} (8.0x10 ² , 8.0x10 ²)	8.9x10 ^{4>1,2,3,4} (6.6x10 ⁴ , 1.1x10 ⁵)
Thrombopoietin ^{Med}	1.2x10 ^{6<N, >2,3,4} (9.8x10 ⁵ , 1.8x10 ⁶)	7.6x10 ^{5<N,1,3,4} (5.6x10 ⁵ , 9.0x10 ⁵)	1.0x10 ^{6<N,1 >2} (8.3x10 ⁵ , 1.2x10 ⁶)	9.2x10 ^{5<N,1, >2} (7.1x10 ⁵ , 1.2x10 ⁶)	8.6x10 ^{6>1,2,3,4} (8x10 ⁶ , 9.1x10 ⁶)
Total IgE ^{Med}	3.4x10 ^{3>N} (1.1x10 ³ , 1.0x10 ⁴)	1.8x10 ^{3>N} (3.1x10 ² , 5.1x10 ³)	1.6x10 ^{3>N} (5.1x10 ² , 5.9x10 ³)	1.8x10 ^{3>N} (2.3x10 ² , 6.3x10 ³)	3.7x10 ^{1<1,2,3,4} (1.5x10 ¹ , 1.8x10 ¹)
TWEAK ^{GM}	1.1x10 ^{4<N, >2} (9.8x10 ³ , 1.2x10 ⁴)	8.5x10 ^{3<N,1,3,4} (8.0x10 ³ , 9.1x10 ³)	1.0x10 ^{4<N, >2} (9.5x10 ³ , 1.1x10 ⁴)	1.1x10 ^{4<N, >2} (9.3x10 ³ , 1.3x10 ⁴)	1.6x10 ^{4>1,2,3,4} (1.4x10 ⁴ , 1.9x10 ⁴)
VCAM-1 ^{Med}	2.4x10 ^{6<N, >2,3} (2.1x10 ⁶ , 3.1x10 ⁶)	1.9x10 ^{6<N,1} (1.6x10 ⁶ , 2.2x10 ⁶)	2.0x10 ^{6<N,1} (1.6x10 ⁶ , 2.5x10 ⁶)	2.3x10 ^{6<N} (2.0x10 ⁶ , 2.6x10 ⁶)	4.3x10 ^{6>1,2,3,4} (3.7x10 ⁵ , 5.3x10 ⁶)

Variable types are shown in superscript next to variable name; M (mean), Med (median), GM (geometric mean). Numbers shown in brackets after measure of central tendency are 95% confidence intervals (for mean and geomean) or interquartile

Table S4. Summary of serum mediators that are differentially regulated in atopic dermatitis versus healthy controls but do not associate with disease severity.

Variables where HC < AD	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
Serum Ig targeting alpha toxin ^{GM}	11.2 ^{<Mod,Sev} (7, 18)	75.7 ^{>HC} (62, 92)	99.4 ^{>HC} (77.5, 127.5)
Apelin ^{Med}	4.7x10 ^{4<Mod,Sev} (3.8x10 ⁴ , 5.2x10 ⁴)	6.3x10 ^{4>HC} (5.8x10 ⁴ , 6.7x10 ⁴)	6.4x10 ^{4>HC} (5.8x10 ⁴ , 6.6x10 ⁴)
C5a ^{GM}	5.2x10 ^{4<Mod,Sev} (4.3x10 ⁴ , 6.3x10 ⁴)	1.3x10 ^{5>HC} (1.1x10 ⁵ , 1.5x10 ⁵)	1.1x10 ^{5>HC} (9.4x10 ⁴ , 1.3x10 ⁵)
Cathepsin B ^M	9.9x10 ^{3<Mod,Sev} (8.5x10 ³ , 1.1x10 ⁴)	1.6x10 ^{4>HC} (1.5x10 ⁴ , 1.8x10 ⁴)	1.7x10 ^{4>HC} (1.5x10 ⁴ , 1.8x10 ⁴)
Cystatin C ^{Med}	1.2x10 ^{5<Mod,Sev} (8.4x10 ⁴ , 1.5x10 ⁵)	1.9x10 ^{5>HC} (1.7x10 ⁵ , 2.2x10 ⁵)	2.0x10 ^{5>HC} (1.7x10 ⁵ , 2.2x10 ⁵)
Elastase ^{Med}	1.2x10 ^{4<Mod,Sev} (7.1x10 ³ , 1.6x10 ⁴)	2.3x10 ^{4>HC} (1.1x10 ⁴ , 8.6x10 ⁴)	3.2x10 ^{4>HC} (1.5x10 ⁴ , 8.6x10 ⁴)
ENA-78 ^{Med}	7.4x10 ^{2<Mod,Sev} (4.8x10 ² , 1.2x10 ³)	2.4x10 ^{3>HC} (1.6x10 ³ , 3.8x10 ³)	2.7x10 ^{3>HC} (1.9x10 ³ , 4.2x10 ³)
Endoglin ^{GM}	1.2x10 ^{3<Mod,Sev} (1.1x10 ³ , 1.3x10 ³)	2.7x10 ^{3>HC} (2.5x10 ³ , 3.0x10 ³)	2.6x10 ^{3>HC} (2.4x10 ³ , 2.9x10 ³)
Eotaxin-1 ^{GM}	73.6 ^{<Mod,Sev} (61, 88)	525.3 ^{>HC} (439, 629)	610.5 ^{>HC} (509, 733)
Fas ^{GM}	840.9 ^{<Mod,Sev} (721, 978)	1889.4 ^{>HC} (1756, 2033)	1987.3 ^{>HC} (1846, 2140)
GCP-2 ^{GM}	64.2 ^{<Mod,Sev} (34, 120)	148.7 ^{>HC} (121, 182)	147.6 ^{>HC} (119, 183)
GCSF ^{Med}	1.8x10 ^{1<Mod,Sev} (1.8x10 ¹ , 1.8x10 ¹)	8.7x10 ^{3>HC} (6.2x10 ³ , 1.7x10 ⁴)	8.6x10 ^{3>HC} (6.2x10 ³ , 1.5x10 ⁴)
HGF ^{GM}	629.1 ^{<Mod,Sev} (522, 759)	935.5 ^{>HC} (868, 1009)	1036.4 ^{>HC} (950, 1130)
ICAM-1 ^{Med}	2.9x10 ^{5<Mod,Sev} (2.2x10 ⁵ , 3.4x10 ⁵)	4.7x10 ^{5>HC} (3.7x10 ⁵ , 6.0x10 ⁵)	5.0x10 ^{5>HC} (4.0x10 ⁵ , 6.7x10 ⁵)
IFN-α ^{Med}	0.7 ^{<Mod,Sev} (1, 3)	82.5 ^{>HC} (53, 99)	81.9 ^{>HC} (48, 98)
IL-1α ^{Med}	4.7 ^{<Mod,Sev} (3, 8)	23.8 ^{>HC} (9, 39)	26.0 ^{>HC} (14, 53)
IL-1β ^{GM}	1.1 ^{<Mod,Sev} (1, 2)	12.5 ^{>HC} (11, 15)	14.4 ^{>HC} (12, 17)
IL-1R1 ^{Med}	5.6 ^{<Mod,Sev} (3, 8)	76.2 ^{>HC} (39, 114)	79.1 ^{>HC} (52, 118)
IL-1R2 ^{Med}	1473.4 ^{<Sev} (842, 2060)	1615.2 (1218, 2126)	2002.6 ^{>HC} (1509, 2624)
IL-3 ^{Med}	2.5 ^{<Mod,Sev} (3, 3)	2.5 ^{>HC} (3, 420)	2.5 ^{>HC} (3, 339)
IL-4 ^{Med}	0.3 ^{<Mod,Sev} (0, 0)	3.6 ^{>HC} (2, 7)	3.7 ^{>HC} (2, 8)
IL-5 ^{Med}	5.0 ^{<Mod,Sev} (3, 10)	196.6 ^{>HC} (164, 236)	220.1 ^{>HC} (191, 254)
IL-6 ^{Med}	1.4 ^{<Mod,Sev} (1, 10)	18.3 ^{>HC} (10, 47)	18.4 ^{>HC} (8, 36)
IL-7 ^M	5.2 ^{<Mod,Sev} (4, 7)	35.4 ^{>HC} (32, 38)	38 ^{>HC} (34, 41)
IL-9 ^{GM}	12.2 ^{<Mod,Sev} (5, 31)	76.4 ^{>HC} (60, 98)	93 ^{>HC} (75, 116)
IL-10 ^{Med}	1.7 ^{<Mod} (2, 2)	1.7 ^{>HC} (2, 8)	1.7 (2, 7)
IL-11 ^{Med}	4.0 ^{<Mod,Sev} (4, 4)	19.7 ^{>HC} (4, 89)	30.6 ^{>HC} (4, 89)
IL-12 ^{Med}	3.3 ^{<Mod,Sev} (3, 12)	58.3 ^{>HC} (29, 109)	58.5 ^{>HC} (38, 104)

IL-13 ^{GM}	23.3 ^{<Mod,Sev} (10, 53)	62.6 ^{>HC} (54, 73)	67.7 ^{>HC} (58, 79)
IL-15 ^{Med}	3.7 ^{<Mod,Sev} (2, 6)	93.4 ^{>HC} (57, 182)	92.6 ^{>HC} (51, 181)
IL-20 ^{Med}	42.4 ^{<Mod,Sev} (42, 42)	42.4 ^{>HC} (42, 42)	42.4 ^{>HC} (42, 42)
IL-21 ^{GM}	3.5x10 ^{2<Mod,Sev} (1.5x10 ² , 1.2x10 ³)	5.5x10 ^{3>HC} (4.4x10 ³ , 6.8x10 ³)	5.6x10 ^{3>HC} (4.4x10 ³ , 7.2x10 ³)
IL-23 ^{Med}	1.5x10 ^{1<Mod,Sev} (1.5x10 ¹ , 6.9x10 ¹)	5.9x10 ^{2>HC} (9.6x10 ¹ , 1.4x10 ³)	2.7x10 ^{2>HC} (1.0x10 ² , 1.5x10 ³)
IL-25 ^{Med}	8.2x10 ^{1<Mod,Sev} (8.2x10 ¹ , 5.8x10 ²)	7.4x10 ^{3>HC} (4.3x10 ³ , 1.5x10 ⁴)	8.3x10 ^{3>HC} (4.7x10 ³ , 1.5x10 ⁴)
IL-33 ^{GM}	9.2 ^{<Mod,Sev} (6, 14)	42.4 ^{>HC} (34, 51)	50.9 ^{>HC} (42, 62)
IL-37 ^{Med}	44.2 ^{<Mod,Sev} (44, 44)	865.9 ^{>HC} (68, 1572)	801.1 ^{>HC} (305, 1645)
ITAC ^{GM}	38.5 ^{<Mod,Sev} (30, 49)	104.1 ^{>HC} (91, 119)	120.4 ^{>HC} (102, 142)
LAP-TGFβ ^M	1.5x10 ^{5<Mod,Sev} (1.4x10 ⁵ , 1.7x10 ⁵)	3.2x10 ^{5>HC} (3.1x10 ⁵ , 3.4x10 ⁵)	3.2x10 ^{5>HC} (3.1x10 ⁵ , 3.4x10 ⁵)
LIF ^{Med}	4.8 ^{<Mod,Sev} (5, 5)	4.8 ^{>HC} (5, 5)	4.8 ^{>HC} (5, 5)
MCP-1 ^{GM}	79.1 ^{<Mod,Sev} (68, 92)	191.1 ^{>HC} (179, 204)	201.7 ^{>HC} (187, 218)

Variables where HC < AD	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
MCP-2 ^{Med}	36.5 ^{<Mod,Sev} (29, 44)	46.6 ^{>HC} (36, 57)	48.4 ^{>HC} (38, 61)
MIG ^{GM}	97.2 ^{<Sev} (90, 105)	119.7 (101, 142)	142.6 ^{>HC} (124, 165)
MIP-1β ^{Med}	41.0 ^{<Mod,Sev} (33, 51)	122.7 ^{>HC} (110, 137)	124.8 ^{>HC} (113, 138)
MPIF ^{Med}	94.4 ^{<Mod,Sev} (41, 171)	371.9 ^{>HC} (199, 530)	404.3 ^{>HC} (246, 562)
Osteopetgrin ^{Med}	7.6 ^{<Mod,Sev} (8, 8)	2109.0 ^{>HC} (1656, 2683)	2278.0 ^{>HC} (1888, 2850)
Oncostatin M ^{Med}	3.3 ^{<Mod,Sev} (3, 3)	7 ^{>HC} (3, 28)	15 ^{>HC} (3, 43)
%Neutralisation of α-toxin-induced lysis ^{Med}	10.8 ^{<Mod,Sev} (1, 21)	26.4 ^{>HC} (15, 56)	35.5 ^{>HC} (23, 80)
S100A8 ^{Med}	1.2x10 ^{4<Mod,Sev} (8.4x10 ³ , 1.6x10 ⁴)	1.5x10 ^{5>HC} (1.3x10 ⁵ , 1.7x10 ⁵)	1.5x10 ^{5>HC} (1.2x10 ⁵ , 1.7x10 ⁵)
SAA1 ^{Med}	1.0x10 ^{8<Mod,Sev} (5.2x10 ⁷ , 1.0x10 ⁸)	1.0x10 ^{8>HC} (1.0x10 ⁸ , 1.0x10 ⁸)	1.0x10 ^{8>HC} (1.0x10 ⁸ , 1.0x10 ⁸)
SLPI ^{Med}	6.4x10 ^{3<Mod,Sev} (5.9x10 ³ , 7.2x10 ³)	1.9x10 ^{4>HC} (8.7x10 ³ , 1.9x10 ⁴)	1.9x10 ^{4>HC} (9.1x10 ³ , 1.9x10 ⁴)
sSCFR ^{Med}	4.2x10 ^{4<Mod,Sev} (3.8x10 ⁴ , 4.4x10 ⁴)	6.3x10 ^{4>HC} (4.9x10 ⁴ , 7.7x10 ⁴)	6.3x10 ^{4>HC} (5.3x10 ⁴ , 7.8x10 ⁴)
TIE-2 ^{GM}	2.6x10 ^{3<Mod,Sev} (2.3x10 ³ , 2.8x10 ³)	8.6x10 ^{3>HC} (8.1x10 ³ , 9.1x10 ³)	8.6x10 ^{3>HC} (8.1x10 ³ , 9.0x10 ³)
TNFR1 ^M	5.9x10 ^{2<Mod,Sev} (2.8x10 ² , 9.1x10 ²)	5.6x10 ^{3>HC} (5.2x10 ³ , 5.9x10 ³)	5.7x10 ^{3>HC} (5.3x10 ³ , 6.1x10 ³)
TREM-1 ^{GM}	1.4x10 ^{1<Mod,Sev} (9.3x10 ⁰ , 2.1x10 ¹)	5.8x10 ^{1>HC} (4.5x10 ¹ , 7.6x10 ¹)	6.6x10 ^{1>HC} (5.4x10 ¹ , 8.2x10 ¹)
TSLP ^{GM}	3.3 ^{<Mod,Sev} (2, 7)	15.1 ^{>HC} (12, 18)	17.0 ^{>HC} (14, 21)
VEGF ^{Med}	346.4 ^{<Mod,Sev} (170, 603)	685.1 ^{>HC} (372, 1085)	856.5 ^{>HC} (500, 1324)

Variables where HC > AD	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
ACE ^{GM}	4.0x10 ^{5>Mod,Sev} (3.6 x10 ⁵ , 4.3 x10 ⁵)	2.8x10 ^{5<HC} (2.6x10 ⁵ , 3.1x10 ⁵)	3.0x10 ^{5<HC} (2.7x10 ⁵ , 3.3x10 ⁵)
Ang2 ^{Med}	1.9x10 ^{4>Mod,Sev} (1.8x10 ⁴ , 2.8x10 ⁴)	1.3x10 ^{4<HC} (1.1x10 ⁴ , 1.5x10 ⁴)	1.4x10 ^{4<HC} (1.2x10 ⁴ , 1.5x10 ⁴)
BDNF ^M	5.5x10 ^{5>Mod,Sev} (4.7x10 ⁵ , 6.2x10 ⁵)	2.2x10 ^{5<HC} (2.1x10 ⁵ , 2.4x10 ⁵)	2.3x10 ^{5<HC} (2.2x10 ⁵ , 2.4x10 ⁵)
Cathepsin S ^M	1.7x10 ^{4>Mod,Sev} (1.6x10 ⁴ , 1.9x10 ⁴)	1.3x10 ^{4<HC} (1.1x10 ⁴ , 1.6x10 ⁴)	1.4x10 ^{4<HC} (1.2x10 ⁴ , 1.7x10 ⁴)
Chemerin 4 ^{Med}	1.9x10 ^{4>Mod,Sev} (1.7x10 ⁴ , 2.5x10 ⁴)	6.8x10 ^{3<HC} (6.8x10 ³ , 2.1x10 ⁴)	1.4x10 ^{4<HC} (6.8x10 ³ , 2.2x10 ⁴)
EGF ^{GM}	24.3 ^{>Mod,Sev} (12, 49)	13.8 ^{<HC} (12, 16)	15.5 ^{<HC} (14, 18)
Galectin-3 ^{GM}	2.8x10 ^{4>Mod,Sev} (2.6x10 ⁴ , 3.0x10 ⁴)	2.0x10 ^{4<HC} (1.8x10 ⁴ , 2.1x10 ⁴)	2.1x10 ^{4<HC} (2.0x10 ⁴ , 2.3x10 ⁴)
Galectin-9 ^{Med}	3.2x10 ^{3>Mod,Sev} (2.3x10 ³ , 4.4x10 ³)	1.0x10 ^{3<HC} (1.0x10 ³ , 8.1x10 ³)	1.0x10 ^{3<HC} (1.0x10 ³ , 1.0x10 ³)
IFN-β ^{Med}	3.2x10 ^{2>Mod,Sev} (1.5x10 ² , 9.1x10 ²)	1.4x10 ^{2<HC} (9.7x10 ⁰ , 3.4x10 ²)	1.4x10 ^{2<HC} (9.7x10 ⁰ , 2.8x10 ²)
IL-27 ^{Med}	1.4x10 ^{3>Mod,Sev} (4.7x10 ² , 2.7x10 ³)	1.6x10 ^{0<HC} (1.6x10 ⁰ , 1.9x10 ⁴)	1.6x10 ^{0<HC} (1.6x10 ⁰ , 1.6x10 ⁰)
Leptin-3 ^{Med}	3.6x10 ^{3>Mod,Sev} (2.8x10 ³ , 5.4x10 ³)	8.0x10 ^{2<HC} (8.0x10 ² , 1.9x10 ³)	8.0x10 ^{2<HC} (8.0x10 ² , 2.0x10 ³)
MIF ^{GM}	7.0x10 ^{3>Mod,Sev} (5.2x10 ³ , 9.3x10 ³)	2.8x10 ^{3<HC} (2.5x10 ³ , 3.1x10 ³)	3.2x10 ^{3<HC} (2.8x10 ³ , 3.6x10 ³)
PAI-1 ^{Med}	9.5x10 ^{5>Mod,Sev} (6.6x10 ⁵ , 1.3x10 ⁶)	6.9x10 ^{5<HC} (2.9x10 ⁵ , 9.6x10 ⁵)	7.5x10 ^{5<HC} (1.7x10 ⁵ , 1.0x10 ⁶)
P-selectin ^{Med}	5.5x10 ^{5>Mod,Sev} (3.5x10 ⁵ , 1.4x10 ⁶)	3.4x10 ^{5<HC} (2.0x10 ⁵ , 5.5x10 ⁵)	3.7x10 ^{5<HC} (2.2x10 ⁵ , 6.6x10 ⁵)
RBP-4 ^{Med}	2.8x10 ^{7>Mod,Sev} (2.7x10 ⁷ , 2.9x10 ⁷)	2.6x10 ^{7<HC} (2.4x10 ⁷ , 2.9x10 ⁷)	2.6x10 ^{7<HC} (2.4x10 ⁷ , 2.8x10 ⁷)
Resistin ^{Med}	9.4x10 ^{3>Mod,Sev} (8.9x10 ³ , 1.0x10 ⁴)	6.8x10 ^{3<HC} (5.7x10 ³ , 9.3x10 ³)	7.5x10 ^{3<HC} (6.0x10 ³ , 9.5x10 ³)
sCD14 ^{Med}	2.3x10 ^{6>Mod,Sev} (2.1x10 ⁶ , 2.5x10 ⁶)	1.3x10 ^{6<HC} (1.1x10 ⁶ , 1.8x10 ⁶)	1.5x10 ^{6<HC} (1.2x10 ⁶ , 2.0x10 ⁶)
sCD163 ^{Med}	1.4x10 ^{5>Mod,Sev} (1.2x10 ⁵ , 1.7x10 ⁵)	4.3x10 ^{4<HC} (4.3x10 ⁴ , 4.3x10 ⁴)	4.3x10 ^{4<HC} (4.3x10 ⁴ , 4.3x10 ⁴)
SCF ^{Med}	50.8 ^{>Mod,Sev} (38, 59)	31.5 ^{<HC} (20, 45)	34.5 ^{<HC} (21, 52)
sIL-2Rα ^{Med}	23.5 ^{>Mod,Sev} (24, 336)	23.5 ^{<HC} (24, 24)	23.5 ^{<HC} (24, 24)
TAC1 ^{Med}	8.9x10 ^{4>Mod,Sev} (6.6x10 ⁴ , 1.1x10 ⁵)	8.0x10 ^{2<HC} (8.0x10 ² , 8.0x10 ²)	8.0x10 ^{2<HC} (8.0x10 ² , 8.0x10 ²)
TECK ^{Med}	915.9 ^{>Mod,Sev} (562, 1324)	405.0 ^{<HC} (294, 601)	481.3 ^{<HC} (306, 755)
TPO ^{Med}	8.6x10 ^{6>Mod,Sev} (8.0x10 ⁶ , 9.1x10 ⁶)	9.2x10 ^{5<HC} (7.2x10 ⁵ , 1.2x10 ⁶)	9.7x10 ^{5<HC} (7.6x10 ⁵ , 1.3x10 ⁶)

Variables where HC > AD	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
Trappin-2 ^{Med}	2226.0 ^{>Mod,Sev} (1924, 2576)	123.1 ^{<HC} (27, 243)	99.8 ^{<HC} (29, 240)
TWEAK ^{GM}	1.6x10 ^{4>Mod,Sev} (1.4x10 ⁴ , 1.9x10 ⁴)	9.9x10 ^{3<HC} (9.3x10 ³ , 1.1x10 ⁴)	1.0x10 ^{4<HC} (9.3x10 ³ , 1.1x10 ⁴)
VCAM-1 ^{Med}	4.3x10 ^{6>Mod,Sev} (3.7x10 ⁶ , 5.3x10 ⁶)	2.0x10 ^{6<HC} (1.6x10 ⁶ , 2.5x10 ⁶)	2.2x10 ^{6<HC} (1.8x10 ⁶ , 2.6x10 ⁶)

Variable types are shown in superscript next to variable name; M (mean) for normally distributed data, Med (median) for log-non normally distributed data, GM (geometric mean) for log-normally distributed

data. Numbers shown in brackets after measure of central tendency are 95% confidence intervals (for mean and geomean) or interquartile range (median). Superscript numbers after measure of central tendency are groups that are significantly different.

Table S5. Summary of serum mediators that are differentially regulated in atopic dermatitis versus healthy controls and associate with disease severity.

Variables where HC < AD	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
BRAK1 ^{Med}	310.3 ^{<Mod,Sev} (196, 487)	522.6 ^{>HC, <Sev} (284, 690)	721.2 ^{>HC, Mod} (421, 948)
CTACK ^{Med}	2.5 ^{<Mod,Sev} (3, 4)	297.6 ^{>HC, <Sev} (168, 521)	480.1 ^{>HC, Mod} (283, 863)
Galectin-1 ^M	1.9x10 ⁴ ^{<Mod,Sev} (1.6x10 ⁴ , 2.1x10 ⁴)	2.5x10 ⁴ ^{>HC, <Sev} (2.4x10 ⁴ , 2.7x10 ⁴)	2.8x10 ⁴ ^{>HC, Mod} (2.7x10 ⁴ , 3.0x10 ⁴)
I-309 ^{GM}	2.1 ^{<Mod,Sev} (2, 2)	29.4 ^{>HC, <Sev} (24, 37)	51.2 ^{>HC, Mod} (42, 62)
IL-18 ^{GM}	82.5 ^{<Mod,Sev} (66, 102)	220.9 ^{>HC, <Sev} (191, 255)	314.7 ^{>HC, Mod} (280, 354)
IL-18BP ^{aMed}	1416.3 ^{<Mod,Sev} (1121, 1740)	2225.8 ^{>HC, <Sev} (1613, 2819)	2597.0 ^{>HC, Mod} (2000, 3135)
IL-22 ^{Med}	1.8 ^{<Mod,Sev} (2, 2)	110.2 ^{>HC, <Sev} (46, 300)	367 ^{>HC, Mod} (140, 602)
IP-10 ^{GM}	280.6 ^{<Mod,Sev} (240, 328)	426.4 ^{>HC, <Sev} (395, 460)	500 ^{>HC, Mod} (458, 546)
MDC ^{Med}	737.8 ^{<Mod,Sev} (681, 874)	2218.2 ^{>HC, <Sev} (1529, 3169)	3730.3 ^{>HC, Mod} (2255, 9001)
MIP-1α ^{Med}	16.5 ^{<Mod,Sev} (17, 17)	90.2 ^{>HC, <Sev} (69, 147)	129.5 ^{>HC, Mod} (85, 162)
MIP-3α ^{Med}	0.6 ^{<Mod,Sev} (1, 1)	7.4 ^{>HC, <Sev} (4, 11)	9.5 ^{>HC, Mod} (6, 14)
Osteopontin ^{GM}	1.2x10 ⁴ ^{<Mod,Sev} (9.2x10 ³ , 1.5x10 ⁴)	3.0x10 ⁴ ^{>HC, <Sev} (2.7x10 ⁴ , 3.4x10 ⁴)	3.9x10 ⁴ ^{>HC, Mod} (3.5x10 ⁴ , 4.4x10 ⁴)
PARC ^{Med}	4.7x10 ⁵ ^{<Mod,Sev} (2.8x10 ⁵ , 7.1x10 ⁵)	7.7x10 ⁵ ^{>HC, <Sev} (3.3x10 ⁵ , 3.8x10 ⁶)	3.1x10 ⁶ ^{>HC, Mod} (5.8x10 ⁵ , 8.7x10 ⁶)
Periostin ^{GM}	6.7 ^{<Mod,Sev} (3, 13)	12.0 ^{>HC, <Sev} (11, 13)	17.8 ^{>HC, Mod} (16, 21)
TARC ^{Med}	96.5 ^{<Mod,Sev} (68, 137)	1504.5 ^{>HC, <Sev} (654, 3200)	3950.0 ^{>HC, Mod} (2104, 8944)
TNFR2 ^M	1476.4 ^{<Mod,Sev} (1352, 1601)	1968.9 ^{>HC, <Sev} (1836, 2102)	2167.0 ^{>HC, Mod} (2053, 2281)
Total IgE ^{Med}	3.7x10 ¹ ^{<Mod,Sev} (1.5x10 ¹ , 8.0x10 ¹)	1.0x10 ³ ^{>HC, <Sev} (2.5x10 ² , 3.2x10 ³)	4.2x10 ³ ^{>HC, Mod} (1.1x10 ³ , 9.4x10 ³)

Variables where HC > AD	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
Cathepsin L ^{Med}	6.4x10 ³ ^{>Mod,Sev} (5.0x10 ³ , 8.0x10 ³)	1.1x10 ³ ^{<HC, Sev} (6.9x10 ² , 1.4x10 ³)	1.1x10 ³ ^{<HC, >Mod} (8.5x10 ² , 1.5x10 ³)
E-selectin ^{Med}	6.4x10 ⁴ ^{>Mod,Sev} (4.3x10 ⁴ , 8.5x10 ⁴)	1.9x10 ⁴ ^{<HC, Sev} (1.4x10 ⁴ , 2.5x10 ⁴)	2.7x10 ⁴ ^{<HC, >Mod} (1.9x10 ⁴ , 3.8x10 ⁴)
MCP-4 ^{Med}	103.8 ^{>Mod} (240, 328)	49.7 ^{<HC, Sev} (29, 70)	76.9 ^{>Mod} (47, 134)
MMP8 ^{Med}	3.0x10 ⁵ ^{>Mod,Sev} (2.7x10 ⁵ , 3.2x10 ⁵)	1.1x10 ⁵ ^{<HC, Sev} (1.0x10 ⁵ , 1.2x10 ⁵)	1.3x10 ⁵ ^{<HC, >Mod} (1.2x10 ⁵ , 1.5x10 ⁵)
RANTES ^{Med}	6.4x10 ⁵ ^{>Mod,Sev}	1.2x10 ⁵ ^{<HC, Sev}	1.5x10 ⁵ ^{<HC, >Mod}

(4.0x10⁵, 9.1x10⁵)

(6.2x10⁴, 1.8x10⁵)

(7.7x10⁴, 2.2x10⁵)

Variable types are shown in superscript next to variable name; M (mean) for normally distributed data, Med (median) for log-non normally distributed data, GM (geometric mean) for log-normally distributed data. Numbers shown in brackets after measure of central tendency are 95% confidence intervals (for mean and geomean) or interquartile range (median). Superscript numbers after measure of central tendency are groups that are significantly different.

Table S6. Summary of serum mediators that are not differentially regulated in atopic dermatitis versus healthy controls but associate with disease severity.

Variable	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
GDF ^{Med}	1.2x10 ³ (1.0x10 ³ , 1.3x10 ³)	8.4x10 ^{2<Sup>Sev} (7.1x10 ² , 1.1x10 ³)	9.6x10 ^{2>Mod} (7.4x10 ² , 1.5x10 ³)
MMP-1 ^{GM}	3.5x10 ⁴ (2.8x10 ⁴ , 4.3x10 ⁴)	3.7x10 ^{4<Sup>Sev} (3.1x10 ⁴ , 4.3x10 ⁴)	4.7x10 ^{4>Mod} (4.1x10 ⁴ , 5.3x10 ⁴)

Variable types are shown in superscript next to variable name; M (mean) for normally distributed data, Med (median) for log-non normally distributed data, GM (geometric mean) for log-normally distributed data. Numbers shown in brackets after measure of central tendency are 95% confidence intervals (for mean and geomean) or interquartile range (median). Superscript numbers after measure of central tendency are groups that are significantly different.

Table S7. Summary of serum mediators that are not differentially regulated in atopic dermatitis versus healthy controls.

Variable	Healthy controls (n=30)	Moderate AD (n=95)	Severe AD (n=98)
Adiponectin ^{Med}	1.3x10 ⁸ (9.0x10 ⁷ , 1.6x10 ⁸)	1.6x10 ⁸ (1.2x10 ⁸ , 4.2x10 ⁸)	1.5x10 ⁸ (9.5x10 ⁷ , 4.2x10 ⁸)
Adipsin ^{Med}	1.5x10 ⁴ (1.4x10 ⁴ , 1.7x10 ⁴)	1.7x10 ⁴ (1.5x10 ⁴ , 1.8x10 ⁴)	1.7x10 ⁴ (1.5x10 ⁴ , 1.8x10 ⁴)
Ang1 ^{Med}	4.4x10 ⁴ (3.6x10 ⁴ , 6.0x10 ⁴)	5.2x10 ⁴ (4.3x10 ⁴ , 6.2x10 ⁴)	5.4x10 ⁴ (4.3x10 ⁴ , 6.9x10 ⁴)
BLC ^{GM}	43.8 (33, 58)	42.8 (36, 51)	53.4 (45, 64)
DKK-1 ^M	2403.4 (1913, 2893)	2260.6 (2195, 2326)	2321 (2268, 2373)
DPP-4 ^{GM}	539.3 (499, 583)	620.7 (583, 661)	584.6 (553, 618)
Eotaxin-3 ^{Med}	1.7 (1.7, 1.7)	1.7 (1.7, 1.7)	1.7 (1.7, 1.7)
EPO-R ^{Med}	13.7 (13.7, 13.7)	13.7 (13.7, 623.1)	13.7 (13.7, 304.2)
FasL ^{Med}	32.8 (31, 38)	45 (28, 68)	45 (26, 63)
FGFB ^{Med}	8 (8, 8)	8 (8, 8)	8 (8, 8)
GMCSF ^{Med}	8 (8, 8)	8 (8, 8)	8 (8, 8)
GRO-α ^{GM}	NA	106.7 (96, 119)	118.2 (105, 133)
IFN-γ ^{Med}	4 (4,4)	4 (4,28)	4 (4,23)
IL1R4 ^{Med}	0.6 (1, 13)	4.3 (1, 14)	5.2 (1, 18)
IL1Ra ^{Med}	4.2 (4, 88)	40.2 (4, 77)	43 (5, 77)
IL-2 ^{Med}	2.1 (2, 2)	2.1 (2, 2)	2.1 (2, 2)
IL-8 ^{GM}	20.9 (13, 34)	15.3 (12, 20)	18.0 (14, 24)
IL-16 ^M	2.0x10 ³ (6.8x10 ² , 3.4x10 ³)	1.6x10 ³ (1.5x10 ³ , 1.8x10 ³)	1.9x10 ³ (1.7x10 ³ , 2.0x10 ³)
IL-17 ^{Med}	2.8 (3, 3)	2.8 (3, 11)	2.8 (3, 12)
IL-19 ^{Med}	NA	1155.4 (787, 1815)	1241.7 (838, 1753)
IL-26 ^{Med}	6.5x10 ² (1.4x10 ² , 1.5x10 ³)	2.1x10 ³ (3.8x10 ¹ , 6.6x10 ³)	2.4x10 ³ (3.8x10 ¹ , 6.4x10 ³)
IL-29 ^{Med}	1.0x10 ¹ (1.0x10 ¹ , 1.0x10 ¹)	1.0x10 ¹ (1.0x10 ¹ , 1.0x10 ¹)	1.0x10 ¹ (1.0x10 ¹ , 1.0x10 ¹)
IL-31 ^{Med}	1.2x10 ¹ (1.2x10 ¹ , 2.5x10 ¹)	1.2x10 ¹ (1.2x10 ¹ , 3.1x10 ³)	1.2x10 ¹ (1.2x10 ¹ , 1.2x10 ¹)
KIM-1 ^{GM}	4.4x10 ¹ (2.6x10 ¹ , 7.4x10 ¹)	6.6x10 ¹ (4.6x10 ¹ , 9.4x10 ¹)	5.7x10 ¹ (4.2x10 ¹ , 7.8x10 ¹)
LAIR-1 ^{Med}	182.1 (182, 182)	182.1 (182, 182)	182.1 (182, 182)
LIGHT ^{Med}	29.8 (4, 73)	4 (1, 128)	4 (4, 206)
MCP-3 ^{Med}	5.5 (6, 30)	14.4 (6, 53)	17.2 (6, 49)
MCSF ^{Med}	7.6 (8, 8)	7.6 (8, 8)	7.6 (8, 8)

MIP-3β^{Med}	6.6x10 ¹ (3.9x10 ¹ , 1.1x10 ²)	1.6x10 ⁰ (1.6x10 ⁰ , 7.4x10 ³)	1.6x10 ⁰ (1.6x10 ⁰ , 1.6x10 ⁰)
MMP-3^{Med}	2.5x10 ⁴ (2.1x10 ⁴ , 3.0x10 ⁴)	2.8x10 ⁴ (2.4x10 ⁴ , 3.2x10 ⁴)	2.8x10 ⁴ (2.5x10 ⁴ , 3.2x10 ⁴)
NAP2^{Med}	1.9x10 ⁷ (1.7x10 ⁷ , 4.0x10 ⁷)	1.7x10 ⁷ (1.3x10 ⁷ , 4.0x10 ⁷)	1.7x10 ⁷ (1.5x10 ⁷ , 4.0x10 ⁷)
NGF^{Med}	2.8 (2, 7)	4.9 (2, 14)	6.4 (2, 13)
PD1^{Med}	2.1x10 ¹ (4.6x10 ⁰ , 2.0x10 ²)	4.6x10 ⁰ (4.6x10 ⁰ , 4.8x10 ³)	3.6x10 ¹ (4.6x10 ⁰ , 1.3x10 ³)
PLGF^{Med}	40.8 (27, 62)	41.5 (19, 59)	35.9 (19, 59)
S100A12^{Med}	NA	5.4x10 ⁶ (3.5x10 ⁵ , 7.8x10 ⁶)	5.3x10 ⁶ (3.6x10 ⁵ , 8.1x10 ⁶)
SDF-1α^{Med}	3.1x10 ² (2.4x10 ² , 5.3x10 ²)	3.7x10 ² (2.6x10 ¹ , 8.7x10 ²)	5.0x10 ² (2.6x10 ¹ , 1.1x10 ³)
SOST^{Med}	11.7 (12, 88)	11.7 (12, 12)	11.7 (12, 12)
SRPSOX^{Med}	NA	8.2x10 ² (6.3x10 ² , 2.9x10 ³)	8.6x10 ² (6.8x10 ² , 2.1x10 ³)
TIMP-1^M	5.0x10 ⁵ (4.8x10 ⁵ , 5.2x10 ⁵)	5.0x10 ⁵ (4.8x10 ⁵ , 5.3x10 ⁵)	5.2x10 ⁵ (4.9x10 ⁵ , 5.4x10 ⁵)
TNF-α^{Med}	1.1 (1, 1)	1.1 (1, 5)	1.1 (1, 5)
TNF-β^{Med}	1.5 (2, 13)	1.5 (2, 9)	1.5 (2, 9)
VEGFR^{GM}	1.5x10 ³ (1.2x10 ³ , 1.9x10 ³)	1.3x10 ³ (1.2x10 ³ , 1.5x10 ³)	1.4x10 ³ (1.3x10 ³ , 1.6x10 ³)
XCL-1^{Med}	8.5 (9, 62)	8.5 (9, 39)	8.5 (9, 50)
YKL40^{Med}	5.0x10 ⁴ (4.5x10 ⁴ , 6.6x10 ⁴)	4.3x10 ⁴ (3.3x10 ⁴ , 6.2x10 ⁴)	4.4x10 ⁴ (3.4x10 ⁴ , 6.3x10 ⁴)

Variable types are shown in superscript next to variable name; M (mean) for normally distributed data, Med (median) for log-non normally distributed data, GM (geometric mean) for log-normally distributed data. Numbers shown in brackets after measure of central tendency are 95% confidence intervals (for mean and geomean) or interquartile range (median). Superscript numbers after measure of central tendency are groups that are significantly different.

Table S8. ISAC sensitization pattern in healthy controls, moderate and severe AD patients.

Component	Sensitisation	Healthy controls (n=30)		Moderate AD (n=95)		Severe AD (n=97)	
		no. of patients	% of patients	no. of patients	% of patients	no. of patients	% of patients
actd1	Negative	29	96.7	81	85.3	79	81.4
	Positive	1	3.3	14	14.7	18	18.6
actd2	Negative	30	100.0	81	85.3	78	80.4
	Positive	0	0.0	14	14.7	19	19.6
actd5	Negative	30	100.0	93	97.9	95	97.9
	Positive	0	0.0	2	2.1	2	2.1
actd8	Negative	28	93.3	61	64.2	55	56.7
	Positive	2	6.7	34	35.8	42	43.3
alng1	Negative	27	90.0	36	37.9	38	39.2
	Positive	3	10.0	59	62.1	59	60.8
alta1	Negative	30	100.0	64	67.4	64	66.0
	Positive	0	0.0	31	63.6	33	34.0
alta6	Negative	30	100.0	85	89.5	82	84.5
	Positive	0	0.0	10	10.5	15	15.5
amba1	Negative	30	100.0	94	99.0	93	95.9
	Positive	0	0.0	1	1.0	4	4.1
anao2	Negative	30	100.0	88	92.6	86	88.7
	Positive	0	0.0	7	7.4	11	11.3
anis1	Negative	30	100.0	93	97.9	93	95.9
	Positive	0	0.0	2	2.1	4	4.1
anis3	Negative	29	96.7	91	95.8	84	86.6
	Positive	1	3.3	4	4.2	13	13.4
apig1	Negative	29	96.7	59	62.1	54	55.7
	Positive	1	3.3	36	37.9	43	44.3
apim1	Negative	30	100.0	91	95.8	90	92.8
	Positive	0	0.0	4	4.2	7	7.2
apim4	Negative	30	100.0	95	100.0	97	100.0
	Positive	0	0.0	0	0.0	0	0.0
arah1	Negative	30	100.0	86	90.5	86	88.7
	Positive	0	0.0	9	9.5	11	11.3
arah2	Negative	30	100.0	84	88.4	85	87.6
	Positive	0	0.0	11	11.6	12	12.4
arah3	Negative	30	100.0	85	89.5	87	89.7
	Positive	0	0.0	10	10.5	10	10.3
arah6	Negative	30	100.0	83	87.4	85	87.6
	Positive	0	0.0	12	12.6	12	12.4
arah8	Negative	28	93.3	41	43.2	42	43.3
	Positive	2	6.7	54	56.8	55	56.7
arah9	Negative	30	100.0	88	92.6	85	87.6
	Positive	0	0.0	7	7.4	12	12.4
artv1	Negative	30	100.0	89	93.7	84	86.6
	Positive	0	0.0	6	6.3	13	13.4
artv3	Negative	30	100.0	88	92.6	81	83.5
	Positive	0	0.0	7	7.4	16	16.5
aspf1	Negative	30	100.0	91	95.8	90	92.8
	Positive	0	0.0	4	4.2	7	7.2
aspf6	Negative	30	100.0	63	66.3	67	69.1
	Positive	0	0.0	32	33.7	30	30.9
aspf3	Negative	28	93.3	87	91.6	87	89.7
	Positive	2	6.7	8	8.4	10	10.3
bere1	Negative	30	100.0	93	97.9	95	97.9
	Positive	0	0.0	2	2.1	2	2.1
betv1	Negative	25	83.3	32	33.7	37	38.1
	Positive	5	16.7	63	66.3	60	61.9
betv2	Negative	30	100.0	84	88.4	81	83.5
	Positive	0	0.0	11	11.6	16	16.5
betv4	Negative	30	100.0	91	95.8	90	92.8
	Positive	0	0.0	4	4.2	7	7.2
blag1	Negative	30	100.0	91	95.8	91	93.8
	Positive	0	0.0	4	4.2	6	6.2
blag2	Negative	30	100.0	69	72.6	54	55.7
	Positive	0	0.0	26	27.4	43	44.3
blag5	Negative	29	96.7	95	100.0	96	99.0
	Positive	1	3.3	0	0.0	1	1.0
blag7	Negative	29	96.7	86	90.5	78	80.4
	Positive	1	3.3	9	9.5	19	19.6
blot5	Negative	29	96.7	75	79.0	64	66.0
	Positive	1	3.3	20	21.1	33	34.0

bosd4	Negative	30	100.0	86	90.5	93	95.9
	Positive	0	0.0	9	9.5	4	4.1
bosd5	Negative	30	100.0	83	87.4	86	88.7
	Positive	0	0.0	12	12.6	11	11.3
bosd6	Negative	30	100.0	89	93.7	86	88.7
	Positive	0	0.0	6	6.3	11	11.3
bosd8	Negative	30	100.0	79	83.2	79	81.4
	Positive	0	0.0	16	16.8	18	18.6
bosdlacto	Negative	30	100.0	93	97.9	88	90.7
	Positive	0	0.0	2	2.1	9	9.3
canf1	Negative	30	100.0	56	59.0	40	41.2
	Positive	0	0.0	39	41.0	57	58.8
canf2	Negative	30	100.0	71	74.7	60	61.9
	Positive	0	0.0	24	25.3	37	38.1
canf3	Negative	30	100.0	70	73.7	57	58.8
	Positive	0	0.0	25	26.3	40	41.2
canf5	Negative	30	100.0	63	66.3	59	60.8
	Positive	0	0.0	32	33.7	38	39.2
chea1	Negative	30	100.0	82	86.3	84	86.6
	Positive	0	0.0	13	13.7	13	13.4
clah8	Negative	30	100.0	83	87.4	86	88.7
	Positive	0	0.0	12	12.6	11	11.3
cora1.0101	Negative	27	90.0	36	37.9	41	42.3
	Positive	3	10.0	59	62.1	56	57.7
cora1.0401	Negative	26	86.7	35	36.8	35	36.1
	Positive	4	13.3	60	63.2	62	63.9
cora8	Negative	30	100.0	85	89.5	81	83.5
	Positive	0	0.0	10	10.5	16	16.5
cora9	Negative	30	100.0	89	93.7	86	88.7
	Positive	0	0.0	6	6.3	11	11.3
cryj1	Negative	29	96.7	81	85.3	71	73.2
	Positive	1	3.3	14	14.7	26	26.8
cupa1	Negative	29	96.7	80	84.2	68	70.1
	Positive	1	3.3	15	15.8	29	29.9
cynd1	Negative	27	90.0	38	40.0	31	32.0
	Positive	3	10.0	57	60.0	66	68.0
derf1	Negative	28	93.3	39	41.0	27	27.8
	Positive	2	6.7	56	59.0	70	72.2
derf2	Negative	27	90.0	35	36.8	34	35.0
	Positive	3	10.0	60	63.2	63	65.0
derp1	Negative	28	93.3	34	35.8	28	28.9
	Positive	2	6.7	61	64.2	69	71.1
derp10	Negative	29	96.7	50	52.6	36	37.1
	Positive	1	3.3	45	47.4	61	62.9
derp2	Negative	27	90.0	36	37.9	35	36.1
	Positive	3	10.0	59	62.1	62	63.9
equc1	Negative	30	100.0	66	69.5	59	60.8
	Positive	0	0.0	29	30.5	38	39.2
equc3	Negative	30	100.0	90	94.7	78	80.4
	Positive	0	0.0	5	5.3	19	19.6
fage2	Negative	30	100.0	93	97.9	96	99.0
	Positive	0	0.0	2	2.1	1	1.0
feld1	Negative	29	96.7	31	32.6	29	29.9
	Positive	1	3.3	64	67.4	68	70.1
feld2	Negative	30	100.0	87	91.6	76	78.3
	Positive	0	0.0	8	8.4	21	21.6
feld4	Negative	30	100.0	64	67.4	52	53.6
	Positive	0	0.0	31	32.6	45	46.4
gadc1	Negative	30	100.0	93	97.9	88	90.7
	Positive	0	0.0	2	2.1	9	9.3
gald1	Negative	30	100.0	78	82.1	71	73.2
	Positive	0	0.0	17	17.9	26	26.8
gald2	Negative	30	100.0	87	91.6	82	84.5
	Positive	0	0.0	8	8.4	15	15.5
gald3	Negative	30	100.0	42	44.2	33	34.0
	Positive	0	0.0	53	55.8	64	66.0
gald5	Negative	30	100.0	92	96.8	84	86.6
	Positive	0	0.0	3	3.2	13	13.4
glym4	Negative	28	93.3	52	54.7	47	48.5
	Positive	2	6.7	43	45.3	50	51.5
glym5	Negative	30	100.0	88	92.6	93	95.9
	Positive	0	0.0	7	7.4	4	4.1
glym6	Negative	30	100.0	85	89.5	86	88.7
	Positive	0	0.0	10	10.5	11	11.3
hev1	Negative	30	100.0	92	96.8	91	93.8

hevb3	Positive	0	0.0	3	3.2	6	6.2
	Negative	30	100.0	94	99.0	89	91.8
hevb5	Positive	0	0.0	1	1.0	8	8.2
	Negative	30	100.0	94	99.0	93	95.9
hevb6.01	Positive	0	0.0	1	1.0	4	4.1
	Negative	30	100.0	83	87.4	78	80.4
hevb8	Positive	0	0.0	12	12.6	19	19.6
	Negative	30	100.0	82	86.3	77	79.4
jujr1	Positive	0	0.0	13	13.7	20	20.6
	Negative	30	100.0	89	93.7	86	88.7
jujr2	Positive	0	0.0	6	6.3	11	11.3
	Negative	30	100.0	81	85.3	63	65.0
jujr3	Positive	0	0.0	14	14.7	34	35.0
	Negative	29	96.7	85	89.5	86	88.7
lepd2	Positive	1	3.3	10	10.5	11	11.3
	Negative	29	96.7	67	70.5	61	62.9
mald1	Positive	1	3.3	28	29.5	36	37.1
	Negative	26	86.7	39	41.0	40	41.2
mera1	Positive	4	13.3	56	59.0	57	58.8
	Negative	30	100.0	85	89.5	81	83.5
musm1	Positive	0	0.0	10	10.5	16	16.5
	Negative	30	100.0	70	73.7	65	67.0
muxf3	Positive	0	0.0	25	26.3	32	33.0
	Negative	30	100.0	81	85.3	76	78.3
olee1	Positive	0	0.0	14	14.7	21	21.6
	Negative	29	96.7	76	80.0	78	80.4
olee7	Positive	1	3.3	19	20.0	19	19.6
	Negative	29	96.7	88	92.6	85	87.6
olee9	Positive	1	3.3	7	7.4	12	12.4
	Negative	30	100.0	57	60.0	41	42.3
parj2	Positive	0	0.0	38	40.0	56	57.7
	Negative	30	100.0	66	69.5	59	60.8
penm1	Positive	0	0.0	29	30.5	38	39.2
	Negative	29	96.7	90	94.7	82	84.5
penm2	Positive	1	3.3	5	5.3	15	15.5
	Negative	30	100.0	77	81.0	72	74.2
penm4	Positive	0	0.0	18	18.9	25	25.8
	Negative	30	100.0	92	96.8	92	94.8
phlp1	Positive	0	0.0	3	3.2	5	5.2
	Negative	26	86.7	35	36.8	32	33.0
phlp11	Positive	4	13.3	60	63.2	65	67.0
	Negative	29	96.7	78	82.1	80	82.5
phlp12	Positive	1	3.3	17	17.9	17	17.5
	Negative	29	96.7	85	89.5	80	82.5
phlp2	Positive	1	3.3	10	10.5	17	17.5
	Negative	29	96.7	56	59.0	56	57.7
phlp4	Positive	1	3.3	39	41.0	41	42.3
	Negative	29	96.7	38	40.0	29	29.9
phlp5	Positive	1	3.3	57	60.0	68	70.1
	Negative	28	93.3	53	55.8	45	46.4
phlp6	Positive	2	6.7	42	44.2	52	53.6
	Negative	28	93.3	61	64.2	55	56.7
phlp7	Positive	2	6.7	34	35.8	42	43.3
	Negative	30	100.0	81	85.3	83	85.6
plaa1	Positive	0	0.0	14	14.7	14	14.4
	Negative	30	100.0	93	97.9	91	93.8
plaa2	Positive	0	0.0	2	2.1	6	6.2
	Negative	30	100.0	73	76.8	55	56.7
plaa3	Positive	0	0.0	22	23.2	42	43.3
	Negative	30	100.0	86	90.5	84	86.6
plal1	Positive	0	0.0	9	9.5	13	13.4
	Negative	30	100.0	91	95.8	90	92.8
pold5	Positive	0	0.0	4	4.2	7	7.2
	Negative	29	96.7	90	94.7	85	87.6
phlp6.1	Positive	1	3.3	5	5.3	12	12.4
	Negative	28	93.3	61	64.2	55	56.7
phlp7.1	Positive	2	6.7	34	35.8	42	43.3
	Negative	30	100.0	81	85.3	83	85.6
plaa1.1	Positive	0	0.0	14	14.7	14	14.4
	Negative	30	100.0	93	97.9	91	93.8
plaa2.1	Positive	0	0.0	2	2.1	6	6.2
	Negative	30	100.0	73	76.8	55	56.7
plaa3.1	Positive	0	0.0	22	23.2	42	43.3
	Negative	30	100.0	86	90.5	84	86.6
	Positive	0	0.0	9	9.5	13	13.4

plal1.1	Negative	30	100.0	91	95.8	90	92.8
	Positive	0	0.0	4	4.2	7	7.2
pold5.1	Negative	29	96.7	90	94.7	85	87.6
	Positive	1	3.3	5	5.3	12	12.4
prup1	Negative	28	93.3	40	42.1	42	43.3
	Positive	2	6.7	55	57.9	55	56.7
prup3	Negative	30	100.0	57	60.0	46	47.4
	Positive	0	0.0	38	40.0	51	52.6
salk1	Negative	30	100.0	90	94.7	89	91.8
	Positive	0	0.0	5	5.3	8	8.2
sesi1	Negative	30	100.0	91	95.8	92	94.8
	Positive	0	0.0	4	4.2	5	5.2
tria14	Negative	30	100.0	91	95.8	81	83.5
	Positive	0	0.0	4	4.2	16	16.5
tria19.0101	Negative	30	100.0	95	100.0	92	94.8
	Positive	0	0.0	0	0.0	5	5.2
triaaati	Negative	30	100.0	92	96.8	89	91.8
	Positive	0	0.0	3	3.2	8	8.2
vesv5	Negative	28	93.3	90	94.7	80	82.5
	Positive	2	6.7	5	5.3	17	17.5

Sensitisation is binominally displayed; sensitisation of >0.3 was classified as being positive

Table S9. ISAC sensitization pattern for each of the four cluster and healthy controls

Component	Sensitisation	Healthy controls (n=30)		Cluster 1 (n=44)		Cluster 2 (n=54)		Cluster 3 (n=62)		Cluster 4 (n=28)	
		No. of HC	% of HC	No. of patients	% of patients	No. of patients	% of patients	No. of patients	% of patients	No. of patients	% of patients
actd1	Negative	29	96.7	37	84.1	47	87.0	49	79.0	23	82.1
	Positive	1	3.3	7	15.9	7	13.0	13	21.0	5	17.9
actd2	Negative	30	100.0	33	75.0	45	83.3	55	88.7	55	78.6
	Positive	0	0.0	11	25.0	9	16.7	7	11.3	6	21.4
actd5	Negative	30	100.0	44	100.0	54	100.0	60	96.8	26	92.9
	Positive	0	0.0	0	0.0	0	0.0	2	3.2	2	7.1
actd8	Negative	28	93.3	23	52.3	33	61.1	41	66.1	15	53.6
	Positive	2	6.7	21	47.7	21	38.9	21	33.9	13	46.4
alng1	Negative	27	90.0	18	40.9	18	33.3	26	41.9	9	32.1
	Positive	3	10.0	26	59.1	36	66.7	36	58.1	19	67.9
alta1	Negative	30	100.0	30	68.2	31	57.4	49	79.0	16	57.1
	Positive	0	0.0	14	31.8	23	42.6	13	21.0	12	42.9
alta6	Negative	30	100.0	34	77.3	52	96.3	56	90.3	21	75.0
	Positive	0	0.0	10	22.7	2	3.7	6	9.7	7	25.0
amba1	Negative	30	100.0	41	93.2	54	100.0	62	100.0	26	92.9
	Positive	0	0.0	3	6.8	0	0.0	0	0.0	2	7.1
anao2	Negative	30	100.0	40	90.9	46	85.2	60	96.8	24	85.7
	Positive	0	0.0	4	9.1	8	14.8	2	3.2	4	14.3
anis1	Negative	30	100.0	41	93.2	54	100.0	61	98.4	26	92.9
	Positive	0	0.0	3	6.8	0	0.0	1	1.6	2	7.1
anis3	Negative	29	96.7	40	90.9	50	92.6	55	88.7	26	92.9
	Positive	1	3.3	4	9.1	4	7.4	7	11.3	2	7.1
apig1	Negative	29	96.7	19	43.2	31	57.4	41	66.1	18	64.3
	Positive	1	3.3	25	56.8	23	42.6	21	33.9	10	35.7
apim1	Negative	30	100.0	39	88.6	53	98.2	58	93.5	27	96.4
	Positive	0	0.0	5	11.4	1	1.8	4	6.4	1	3.6
apim4	Negative	30	100.0	44	100.0	54	100.0	62	100.0	28	100.0
	Positive	0	0.0	0	0.0	0	0.0	0	0.0	0	0.0
arah1	Negative	30	100.0	40	90.9	46	85.2	57	91.9	25	89.3
	Positive	0	0.0	4	9.1	8	14.8	5	8.1	3	10.7
arah2	Negative	30	100.0	37	84.1	45	83.3	59	95.2	25	89.3
	Positive	0	0.0	7	15.9	9	16.7	3	4.8	3	10.7
arah3	Negative	30	100.0	37	84.1	47	87.0	59	95.2	25	89.3
	Positive	0	0.0	7	15.9	7	13.0	3	4.8	3	10.7
arah6	Negative	30	100.0	35	79.5	46	85.2	58	93.5	25	89.3
	Positive	0	0.0	9	20.4	8	14.8	4	6.4	3	10.7
arah8	Negative	28	93.3	18	40.9	20	37.0	31	50.0	11	39.3
	Positive	2	6.7	26	59.1	34	63.0	31	50.0	17	60.7
arah9	Negative	30	100.0	37	84.1	51	94.4	57	91.9	24	85.7
	Positive	0	0.0	7	15.9	3	5.6	5	8.1	4	14.3
artv1	Negative	30	100.0	39	88.6	49	90.7	55	88.7	26	92.9
	Positive	0	0.0	5	11.4	5	9.3	7	11.3	2	7.1
artv3	Negative	30	100.0	36	81.8	50	92.6	56	90.3	23	82.1
	Positive	0	0.0	8	18.2	4	7.4	6	9.7	5	17.9
aspf1	Negative	30	100.0	41	93.2	53	98.2	57	91.9	26	92.9
	Positive	0	0.0	3	6.8	1	1.8	5	8.1	2	7.1
aspf6	Negative	30	100.0	27	61.4	39	72.2	42	67.7	19	67.9
	Positive	0	0.0	17	38.6	15	27.8	20	32.3	9	32.1
aspf3	Negative	28	93.3	39	88.6	47	87.0	59	95.2	26	92.9
	Positive	2	6.7	5	11.4	7	13.0	3	4.8	2	7.1
bere1	Negative	30	100.0	44	100.0	53	98.2	61	98.4	26	92.9
	Positive	0	0.0	0	0.0	1	1.8	1	1.6	2	7.1
betv1	Negative	25	83.3	16	36.4	16	29.6	27	43.5	7	25.0
	Positive	5	16.7	28	63.6	38	70.4	35	56.5	21	75.0
betv2	Negative	30	100.0	37	84.1	48	88.9	55	88.7	21	75.0
	Positive	0	0.0	7	15.9	6	11.1	7	11.3	7	25.0
betv4	Negative	30	100.0	40	90.9	52	96.3	59	95.2	27	96.4
	Positive	0	0.0	4	9.1	2	3.7	3	4.8	1	3.6
blag1	Negative	30	100.0	40	90.9	52	96.3	59	95.2	27	96.4
	Positive	0	0.0	4	9.1	2	3.7	3	4.8	1	3.6
blag2	Negative	30	100.0	22	50.0	33	61.1	45	72.6	19	67.9
	Positive	0	0.0	22	50.0	21	38.9	17	27.4	9	32.1
blag5	Negative	29	96.7	44	100.0	54	100.0	61	98.4	28	100.0
	Positive	1	3.3	0	0.0	0	0.0	1	1.6	0	0.0
blag7	Negative	29	96.7	36	81.8	49	90.7	51	82.3	24	85.7

blot5	Positive	1	3.3	8	18.2	5	9.3	11	17.7	4	14.3
	Negative	29	96.7	28	63.6	43	79.6	42	67.7	23	82.1
bosd4	Positive	1	3.3	16	36.4	11	20.4	20	32.3	5	17.9
	Negative	30	100.0	42	95.5	50	92.6	57	91.9	26	92.9
bosd5	Positive	0	0.0	2	4.5	4	7.4	5	8.1	2	7.1
	Negative	30	100.0	38	86.4	50	92.6	53	85.5	24	85.7
bosd6	Positive	0	0.0	6	13.6	4	7.4	9	14.5	4	14.3
	Negative	30	100.0	40	90.9	50	92.6	55	88.7	27	96.4
bosd8	Positive	0	0.0	4	9.1	4	7.4	7	11.3	1	3.6
	Negative	30	100.0	34	77.3	42	77.8	52	83.9	26	92.9
bosdlacto	Positive	0	0.0	10	22.7	12	22.2	10	16.1	2	7.1
	Negative	30	100.0	40	90.9	51	94.4	58	93.5	28	100.0
canf1	Positive	0	0.0	4	9.1	3	5.6	4	6.4	0	0.0
	Negative	30	100.0	16	36.4	29	53.7	33	53.2	14	50.0
canf2	Positive	0	0.0	28	63.6	25	46.3	29	46.8	14	50.0
	Negative	30	100.0	25	56.8	38	70.4	46	74.2	19	67.9
canf3	Positive	0	0.0	19	43.2	16	29.6	16	25.8	9	32.1
	Negative	30	100.0	25	56.8	38	70.4	42	67.7	18	64.3
canf5	Positive	0	0.0	19	43.2	16	29.6	20	32.3	10	35.7
	Negative	30	100.0	25	56.8	33	61.1	39	62.9	22	78.6
chea1	Positive	0	0.0	19	43.2	21	38.9	23	37.1	6	21.4
	Negative	30	100.0	36	81.8	45	83.3	55	88.7	26	92.9
clah8	Positive	0	0.0	8	18.2	9	16.7	7	11.3	2	7.1
	Negative	30	100.0	36	81.8	50	92.6	56	90.3	23	82.1
cora1	Positive	0	0.0	8	18.2	4	7.4	6	9.7	5	17.9
	Negative	27	90.0	18	40.9	19	35.2	27	43.5	10	35.7
cora1.0401	Positive	3	10.0	26	59.1	35	64.8	35	56.5	18	64.3
	Negative	26	86.7	15	34.1	19	35.2	23	37.1	10	35.7
cora8	Positive	4	13.3	29	65.9	35	64.8	39	62.9	18	64.3
	Negative	30	100.0	34	77.3	50	92.6	55	88.7	23	82.1
cora9	Positive	0	0.0	10	22.7	4	7.4	7	11.3	5	17.9
	Negative	30	100.0	39	88.6	50	92.6	59	95.2	23	82.1
cryj1	Positive	0	0.0	5	11.4	4	7.4	3	4.8	5	17.9
	Negative	29	96.7	34	77.3	42	77.8	55	88.7	18	64.3
cupa1	Positive	1	3.3	10	22.7	12	22.2	7	11.3	10	35.7
	Negative	29	96.7	32	72.7	39	72.2	55	88.7	18	64.3
cynd1	Positive	1	3.3	12	27.3	15	27.8	7	11.3	10	35.7
	Negative	27	90.0	11	25.0	12	22.2	31	50.0	11	39.3
derf1	Positive	3	10.0	33	75.0	42	77.8	31	50.0	17	60.7
	Negative	28	93.3	10	22.7	21	38.9	20	32.3	12	42.9
derf2	Positive	2	6.7	34	77.3	33	61.1	42	67.7	16	57.1
	Negative	27	90.0	13	29.6	22	40.7	18	29.0	13	46.4
derp1	Positive	3	10.0	31	70.5	32	59.3	44	71.0	15	53.6
	Negative	28	93.3	9	20.4	19	35.2	20	32.3	11	39.3
derp10	Positive	2	6.7	35	79.5	35	64.8	42	67.7	17	60.7
	Negative	29	96.7	15	34.1	26	48.1	28	45.2	13	46.4
derp2	Positive	1	3.3	29	65.9	28	51.9	34	54.8	15	53.6
	Negative	27	90.0	13	29.6	24	44.4	17	27.4	14	50.0
equc1	Positive	3	10.0	31	70.5	30	55.6	45	72.6	14	50.0
	Negative	30	100.0	23	52.3	36	66.7	45	72.6	17	60.7
equc3	Positive	0	0.0	21	47.7	18	33.3	17	27.4	11	39.3
	Negative	30	100.0	37	84.1	48	88.9	52	83.9	27	96.4
fage2	Positive	0	0.0	7	15.9	6	11.1	10	16.1	1	3.6
	Negative	30	100.0	42	95.5	53	98.2	62	100.0	28	100.0
feld1	Positive	0	0.0	2	4.5	1	1.8	0	0.0	0	0.0
	Negative	29	96.7	12	27.3	19	35.2	21	33.9	6	21.4
feld2	Positive	1	3.3	32	72.7	35	64.8	41	66.1	22	78.6
	Negative	30	100.0	36	81.8	45	83.3	53	85.5	25	89.3
feld4	Positive	0	0.0	8	18.2	9	16.7	9	14.5	3	10.7
	Negative	30	100.0	23	52.3	33	61.1	43	69.3	14	50.0
gadc1	Positive	0	0.0	21	47.7	21	38.9	19	30.6	14	50.0
	Negative	30	100.0	40	90.9	51	94.4	59	95.2	27	96.4
gald1	Positive	0	0.0	4	9.1	3	5.6	3	4.8	1	3.6
	Negative	30	100.0	31	70.5	42	77.8	47	75.8	25	89.3
gald2	Positive	0	0.0	13	29.6	12	22.2	15	24.2	3	10.7
	Negative	30	100.0	38	86.4	46	85.2	55	88.7	26	92.9
gald3	Positive	0	0.0	6	13.6	8	14.8	7	11.3	2	7.1
	Negative	30	100.0	14	31.8	25	46.3	19	30.6	14	50.0
gald5	Positive	0	0.0	30	68.2	29	53.7	43	69.3	14	50.0
	Negative	30	100.0	40	90.9	49	90.7	57	91.9	26	92.9
glym4	Positive	0	0.0	4	9.1	5	9.3	5	8.1	2	7.1
	Negative	28	93.3	21	47.7	27	50.0	34	54.8	14	50.0

glym5	Positive	2	6.7	23	52.3	27	50.0	28	45.2	14	50.0
	Negative	30	100.0	41	93.2	51	94.4	61	98.4	24	85.7
glym6	Positive	0	0.0	3	6.8	3	5.6	1	1.6	4	14.3
	Negative	30	100.0	39	88.6	47	87.0	59	95.2	22	78.6
hevb1	Positive	0	0.0	5	11.4	7	13.0	3	4.8	6	21.4
	Negative	30	100.0	41	93.2	49	90.7	62	100.0	27	96.4
hevb3	Positive	0	0.0	3	6.8	5	9.3	0	0.0	1	3.6
	Negative	30	100.0	42	95.5	49	90.7	61	98.4	27	96.4
hevb5	Positive	0	0.0	2	4.5	5	9.3	1	1.6	1	3.6
	Negative	30	100.0	43	97.7	52	96.3	62	100.0	26	92.9
hevb6.01	Positive	0	0.0	1	2.3	2	3.7	0	0.0	2	7.1
	Negative	30	100.0	39	88.6	44	81.5	51	82.3	23	82.1
hevb8	Positive	0	0.0	5	11.4	10	18.5	11	17.7	5	17.9
	Negative	30	100.0	36	81.8	42	77.8	56	90.3	21	75.0
jugr1	Positive	0	0.0	8	18.2	12	22.2	6	9.7	7	25.0
	Negative	30	100.0	40	90.9	49	90.7	58	93.5	24	85.7
jugr2	Positive	0	0.0	4	9.1	5	9.3	4	6.4	4	14.3
	Negative	30	100.0	32	72.7	41	75.9	47	75.8	20	71.4
jugr3	Positive	0	0.0	12	27.3	13	24.1	15	24.2	8	28.6
	Negative	29	96.7	39	88.6	48	88.9	56	90.3	24	85.7
lepd2	Positive	1	3.3	5	11.4	6	11.1	6	9.7	4	14.3
	Negative	29	96.7	27	61.4	39	72.2	34	54.8	25	89.3
mald1	Positive	1	3.3	17	38.6	15	27.8	28	45.2	3	10.7
	Negative	26	86.7	18	40.9	19	35.2	29	46.8	10	35.7
mera1	Positive	4	13.3	26	59.1	35	64.8	33	53.2	18	64.3
	Negative	30	100.0	37	84.1	47	87.0	57	91.9	21	75.0
musm1	Positive	0	0.0	7	15.9	7	13.0	5	8.1	7	25.0
	Negative	30	100.0	28	63.6	36	66.7	49	79.0	18	64.3
muxf3	Positive	0	0.0	16	36.4	18	33.3	13	21.0	10	35.7
	Negative	30	100.0	31	70.5	43	79.6	55	88.7	24	85.7
olee1	Positive	0	0.0	13	29.6	11	20.4	7	11.3	4	14.3
	Negative	29	96.7	33	75.0	41	75.9	54	87.1	22	78.6
olee7	Positive	1	3.3	11	25.0	13	24.1	8	12.9	6	21.4
	Negative	29	96.7	41	93.2	45	83.3	57	91.9	26	92.9
olee9	Positive	1	3.3	3	6.8	9	16.7	5	8.1	2	7.1
	Negative	30	100.0	17	38.6	29	53.7	35	56.5	14	50.0
parj2	Positive	0	0.0	27	61.4	25	46.3	27	43.5	14	50.0
	Negative	30	100.0	24	54.5	34	63.0	44	71.0	19	67.9
penm1	Positive	0	0.0	20	45.5	20	37.0	18	29.0	9	32.1
	Negative	29	96.7	39	88.6	48	88.9	56	90.3	25	89.3
penm2	Positive	1	3.3	5	11.4	6	11.1	6	9.7	3	10.7
	Negative	30	100.0	31	70.5	44	81.5	48	77.4	22	78.6
penm4	Positive	0	0.0	13	29.6	10	18.5	14	22.6	6	21.4
	Negative	30	100.0	44	100.0	51	94.4	58	93.5	27	96.4
phlp1	Positive	0	0.0	0	0.0	3	5.6	4	6.4	1	3.6
	Negative	26	86.7	12	27.3	10	18.5	32	51.6	9	32.1
phlp11	Positive	4	13.3	32	72.7	44	81.5	30	48.4	19	67.9
	Negative	29	96.7	35	79.5	40	74.1	55	88.7	24	85.7
phlp12	Positive	1	3.3	9	20.4	14	25.9	7	11.3	4	14.3
	Negative	29	96.7	38	86.4	46	85.2	57	91.9	20	71.4
phlp2	Positive	1	3.3	6	13.6	8	14.8	5	8.1	8	28.6
	Negative	29	96.7	23	52.3	26	48.1	45	72.6	14	50.0
phlp4	Positive	1	3.3	21	47.7	28	51.9	17	27.4	14	50.0
	Negative	29	96.7	11	25.0	10	18.5	30	48.4	13	46.4
phlp5	Positive	1	3.3	33	75.0	44	81.5	32	51.6	15	53.6
	Negative	28	93.3	19	43.2	23	42.6	37	59.7	16	57.1
phlp6	Positive	2	6.7	25	56.8	31	57.4	25	40.3	12	42.9
	Negative	28	93.3	23	52.3	26	48.1	46	74.2	17	60.7
phlp7	Positive	2	6.7	21	47.7	28	51.9	16	25.8	11	39.3
	Negative	30	100.0	36	81.8	45	83.3	57	91.9	22	78.6
plaa1	Positive	0	0.0	8	18.2	9	16.7	5	8.1	6	21.4
	Negative	30	100.0	42	95.5	50	92.6	61	98.4	27	96.4
plaa2	Positive	0	0.0	2	4.5	4	7.4	1	1.6	1	3.6
	Negative	30	100.0	22	50.0	37	68.5	45	72.6	20	71.4
plaa3	Positive	0	0.0	22	50.0	17	31.5	17	27.4	8	28.6
	Negative	30	100.0	36	81.8	49	90.7	57	91.9	24	85.7
plal1	Positive	0	0.0	8	18.2	5	9.3	5	8.1	4	14.3
	Negative	30	100.0	39	88.6	53	98.2	60	96.8	25	89.3
pold5	Positive	0	0.0	5	11.4	1	1.8	2	3.2	3	10.7
	Negative	29	96.7	40	90.9	48	88.9	56	90.3	28	100.0
phlp6.1	Positive	1	3.3	4	9.1	6	11.1	6	9.7	0	0.0
	Negative	28	93.3	23	52.3	26	48.1	46	74.2	17	60.7

phlp7.1	Positive	2	6.7	21	47.7	28	51.9	16	25.8	11	39.3
	Negative	30	100.0	36	81.8	45	83.3	57	91.9	22	78.6
plaa1.1	Positive	0	0.0	8	18.2	9	16.7	5	8.1	6	21.4
	Negative	30	100.0	42	95.5	50	92.6	61	98.4	27	96.4
plaa2.1	Positive	0	0.0	2	4.5	4	7.4	1	1.6	1	3.6
	Negative	30	100.0	22	50.0	37	68.5	45	72.6	20	71.4
plaa3.1	Positive	0	0.0	22	50.0	17	31.5	17	27.4	8	28.6
	Negative	30	100.0	36	81.8	49	90.7	57	91.9	24	85.7
plal1.1	Positive	0	0.0	8	18.2	5	9.3	5	8.1	4	14.3
	Negative	30	100.0	39	88.6	53	98.2	60	96.8	25	89.3
pold5.1	Positive	0	0.0	5	11.4	1	1.8	2	3.2	3	10.7
	Negative	29	96.7	40	90.9	48	88.9	56	90.3	28	100.0
prup1	Positive	1	3.3	4	9.1	6	11.1	6	9.7	0	0.0
	Negative	28	93.3	18	40.9	21	38.9	30	48.4	10	35.7
prup3	Positive	2	6.7	26	59.1	33	61.1	32	51.6	18	64.3
	Negative	30	100.0	17	38.6	29	53.7	39	62.9	14	50.0
salk1	Positive	0	0.0	27	61.4	25	46.3	23	37.1	14	50.0
	Negative	30	100.0	39	88.6	50	92.6	61	98.4	25	89.3
sesi1	Positive	0	0.0	5	11.4	4	7.4	1	1.6	3	10.7
	Negative	30	100.0	42	95.5	54	100.0	58	93.5	25	89.3
tria14	Positive	0	0.0	2	4.5	0	0.0	4	6.4	3	10.7
	Negative	30	100.0	41	93.2	49	90.7	55	88.7	23	82.1
tria19	Positive	0	0.0	3	6.8	5	9.3	7	11.3	5	17.9
	Negative	30	100.0	42	95.5	53	98.2	60	96.8	28	100.0
triaaati	Positive	0	0.0	2	4.5	1	1.8	2	3.2	0	0.0
	Negative	30	100.0	41	93.2	52	96.3	60	96.8	24	85.7
vesv5	Positive	0	0.0	3	6.8	2	3.7	2	3.2	4	14.3
	Negative	28	93.3	39	88.6	47	87.0	55	88.7	26	92.9
	Positive	2	6.7	5	11.4	7	13.0	7	11.3	2	7.1

Sensitisation is binominally displayed; sensitisation of >0.3 was classified as being positive

Supplemental Movie E10. Principal component analysis of AD patients.

3D animation of the first three PCs showing the separation of the four patient clusters generated by unsupervised k-means cluster analysis (see Figure 1C). Patients in cluster 1 are represented by black dots, patients in cluster 2 by red dots, patients in cluster 3 by green dots and patients in cluster 4 by blue dots.