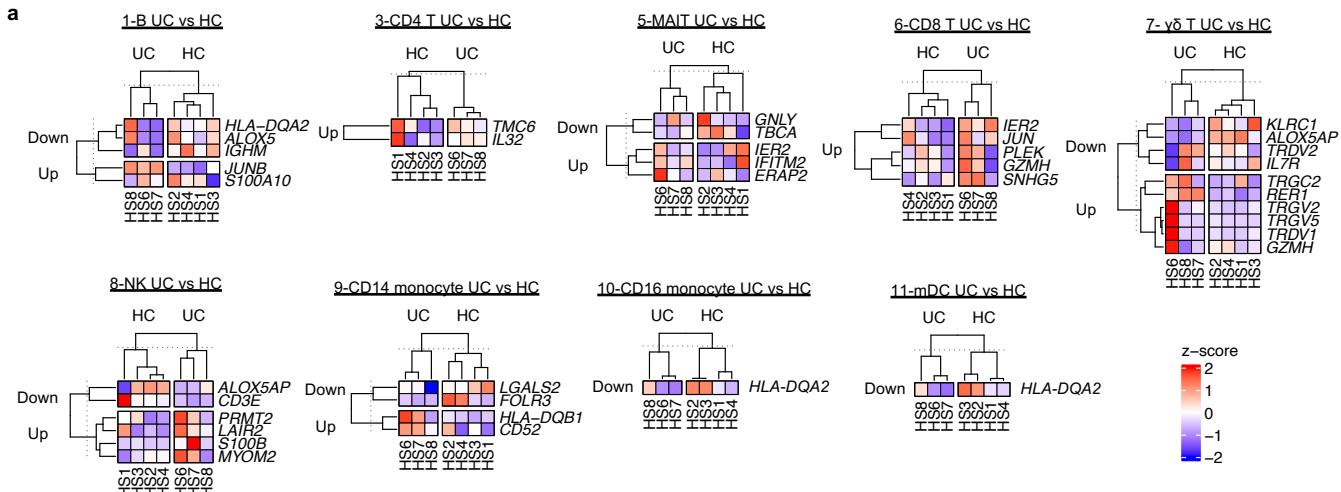
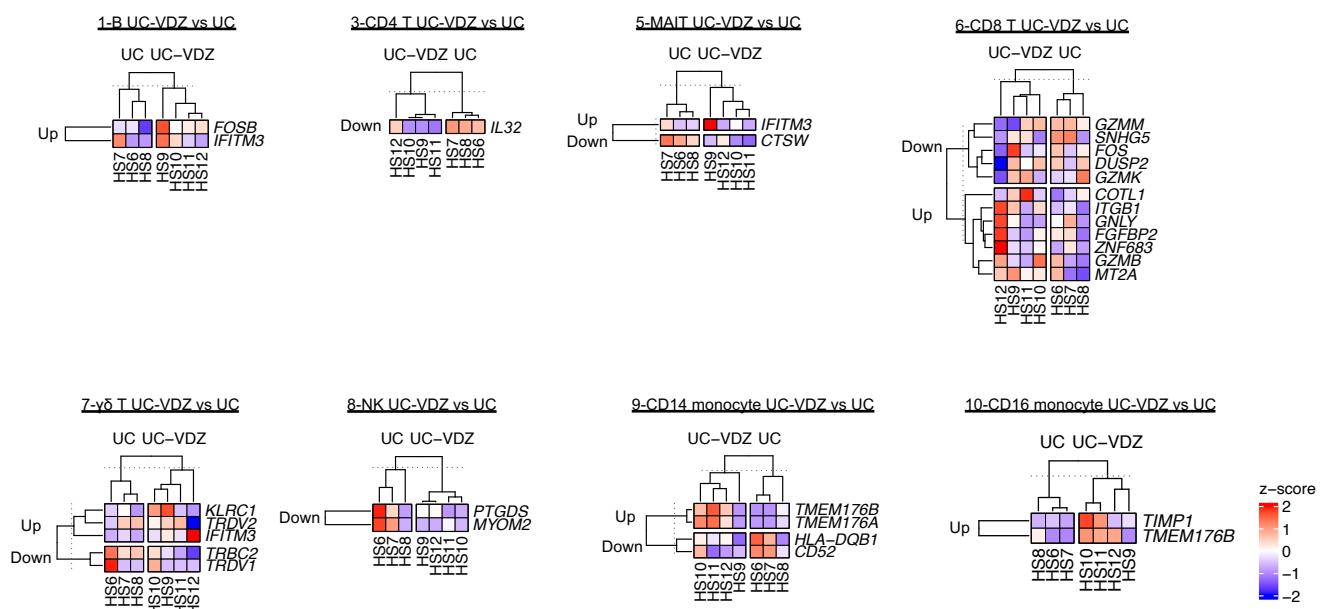
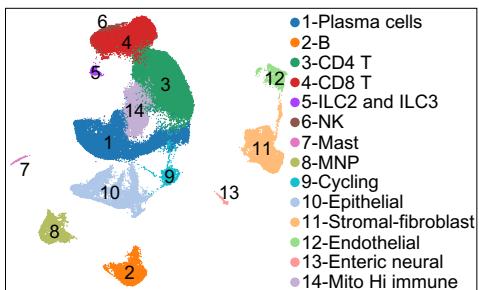


Supplementary Fig.1| Peripheral blood leukocyte (PBL) scRNA-seq coarse cell subset annotations and both coarse and fine cell subset landmark genes. **a**, UMAP visualization of pooled multiplex PBL scRNA-seq for 20,130 cells from HC (n=4), UC (n=3), and UC-VDZ (n=4) patients highlighting coarse cell subset annotations. **b-e**, Dot plot representation of landmark genes expressed by **(b)** coarse cell subsets and **(c-e)** fine PBL cell subsets. Genes ranked by Wilcoxon rank-sum. Cell subsets without landmark genes above selected minimum log fold change are not displayed.

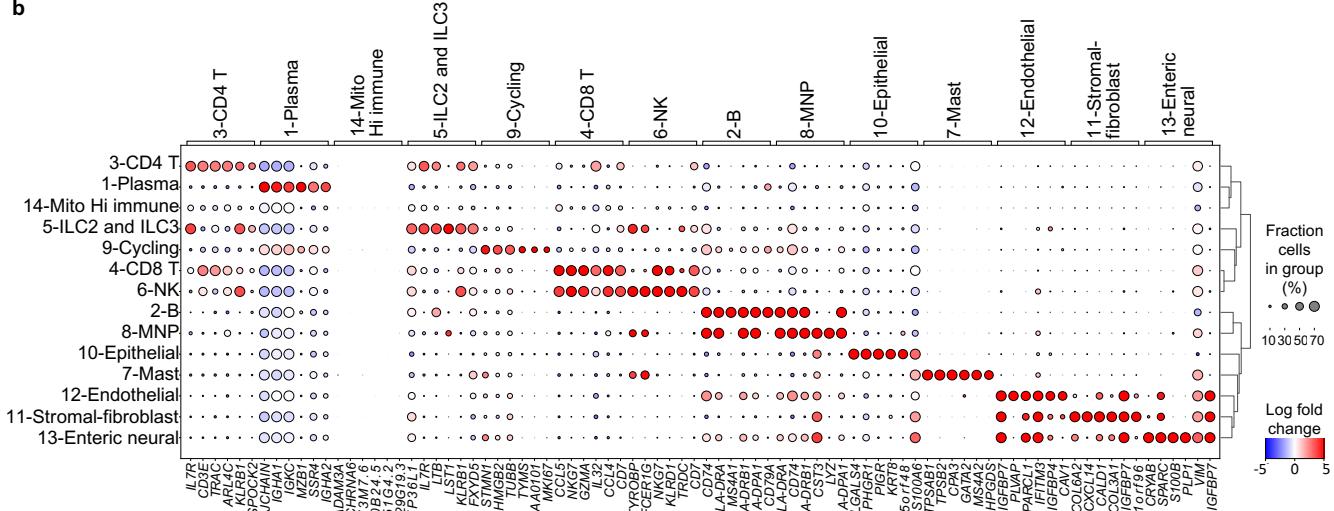
a**b**

Supplementary Fig.2| PBL scRNA-seq DE gene analysis comparing UC versus HC and UC-VDZ versus UC. **a,b,** Heatmap of expression z-scores for scRNA-seq DE genes in the indicated PBL cell subsets with $\log_2 \text{fc} > 2$ or < -2 and Bonferroni-corrected $p\text{-value} < 0.1$ in (a) UC (Up/Down) relative to HC and (b) UC-VDZ (Up/Down) relative to UC identified by MAST analysis. Ribosomal and mitochondrial genes are not displayed; cell subsets without DE genes are not displayed.

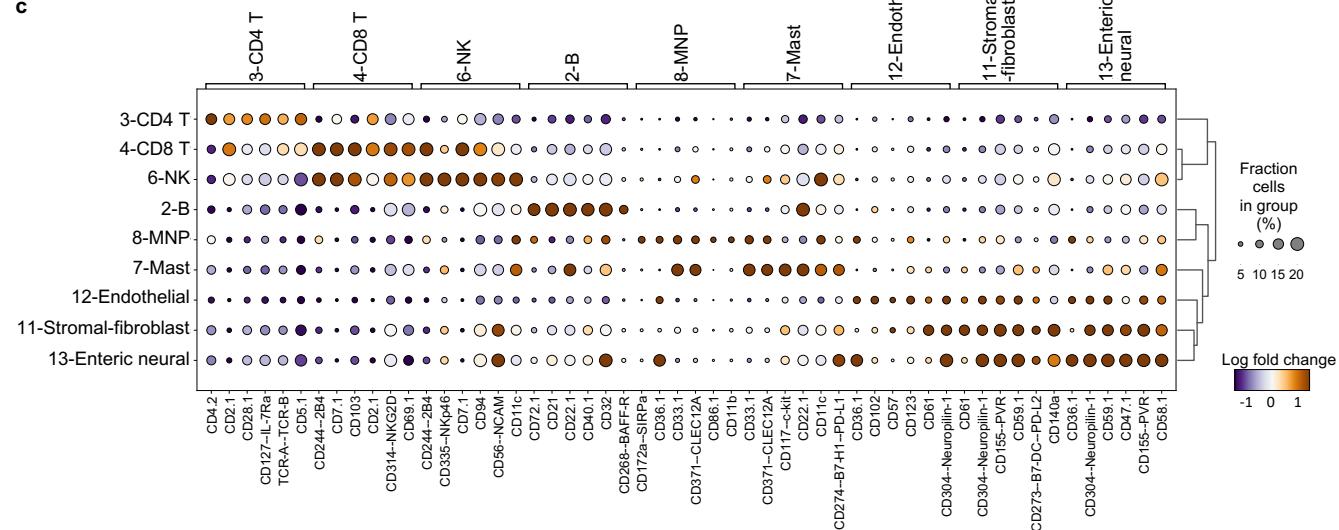
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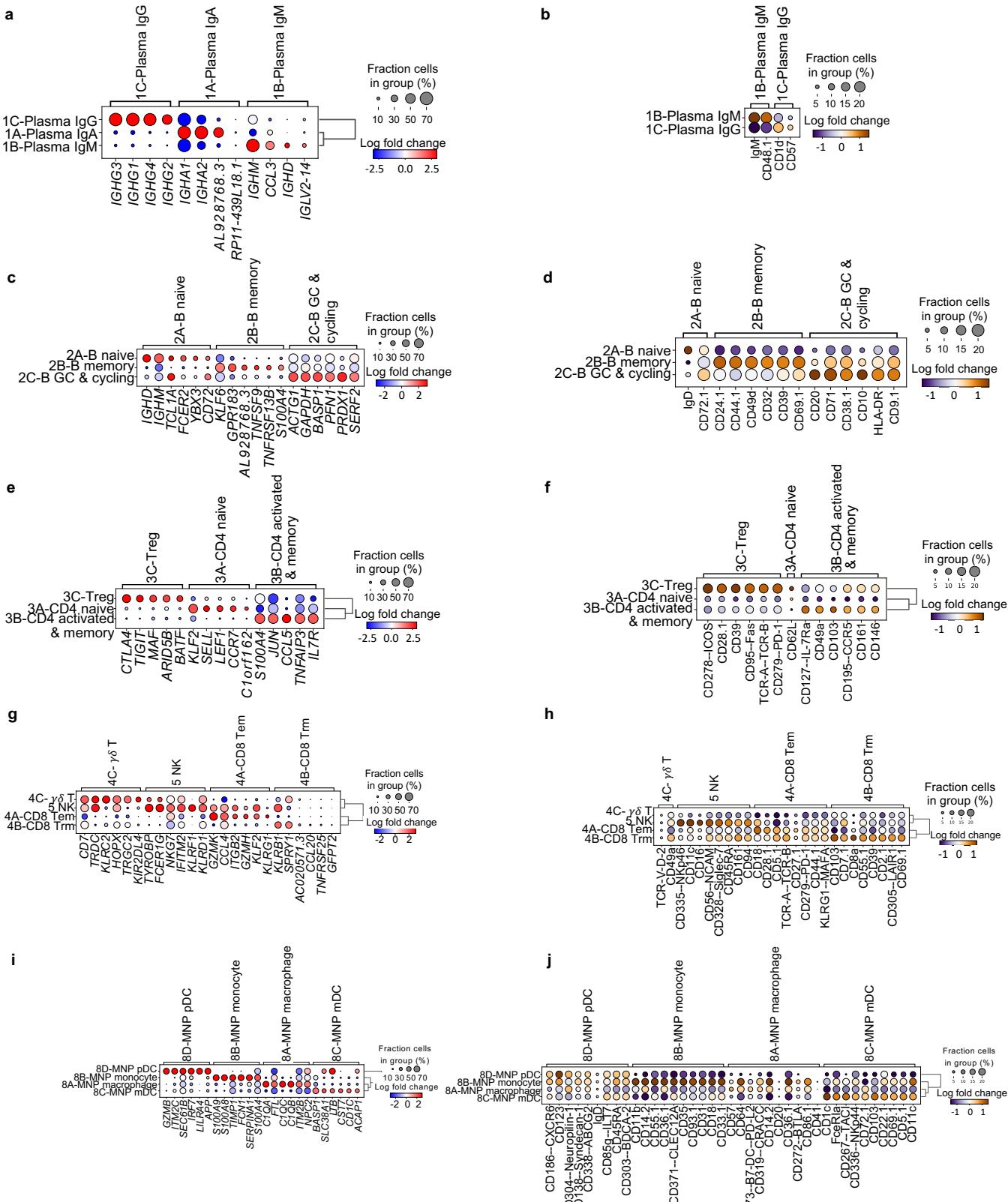
h



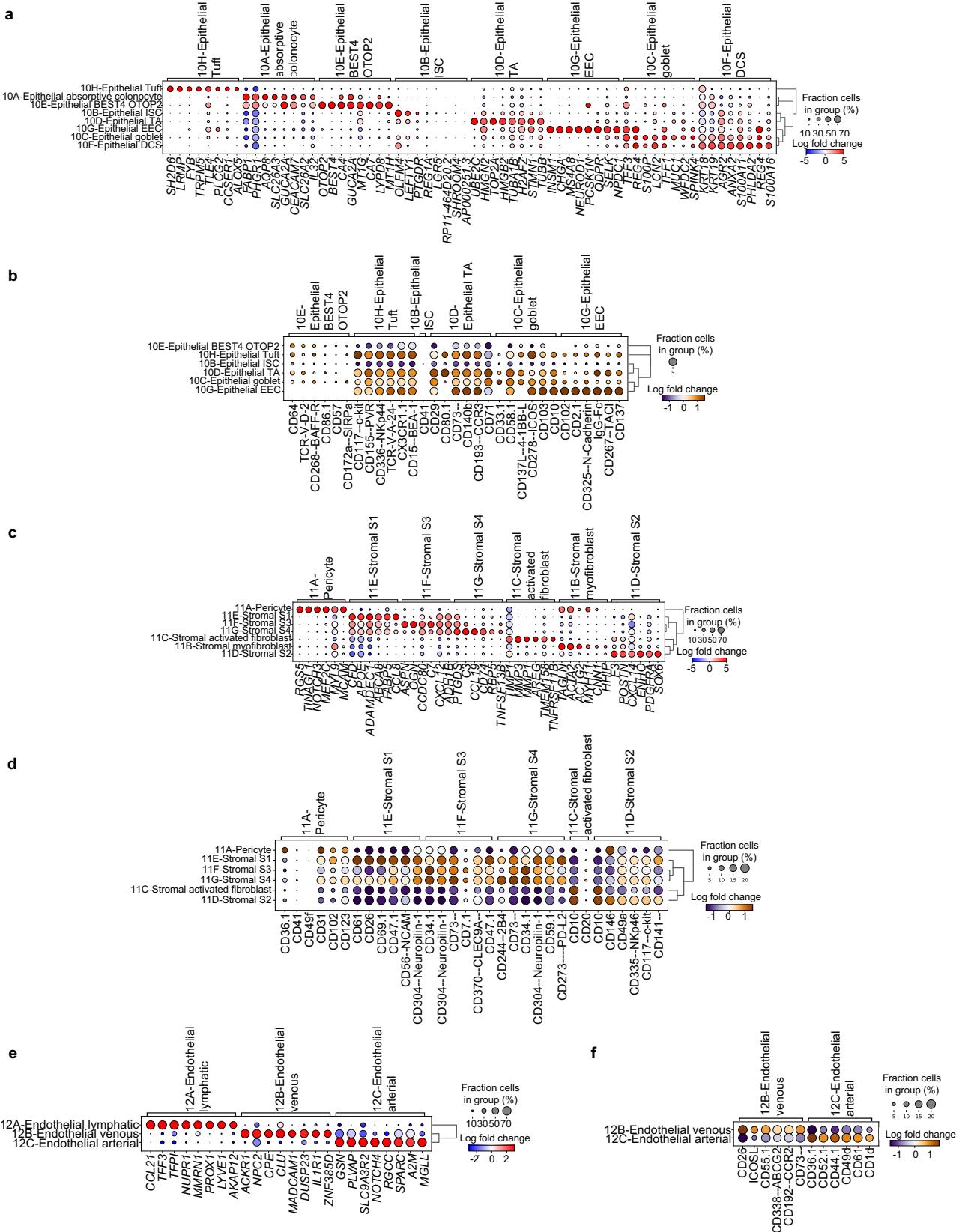
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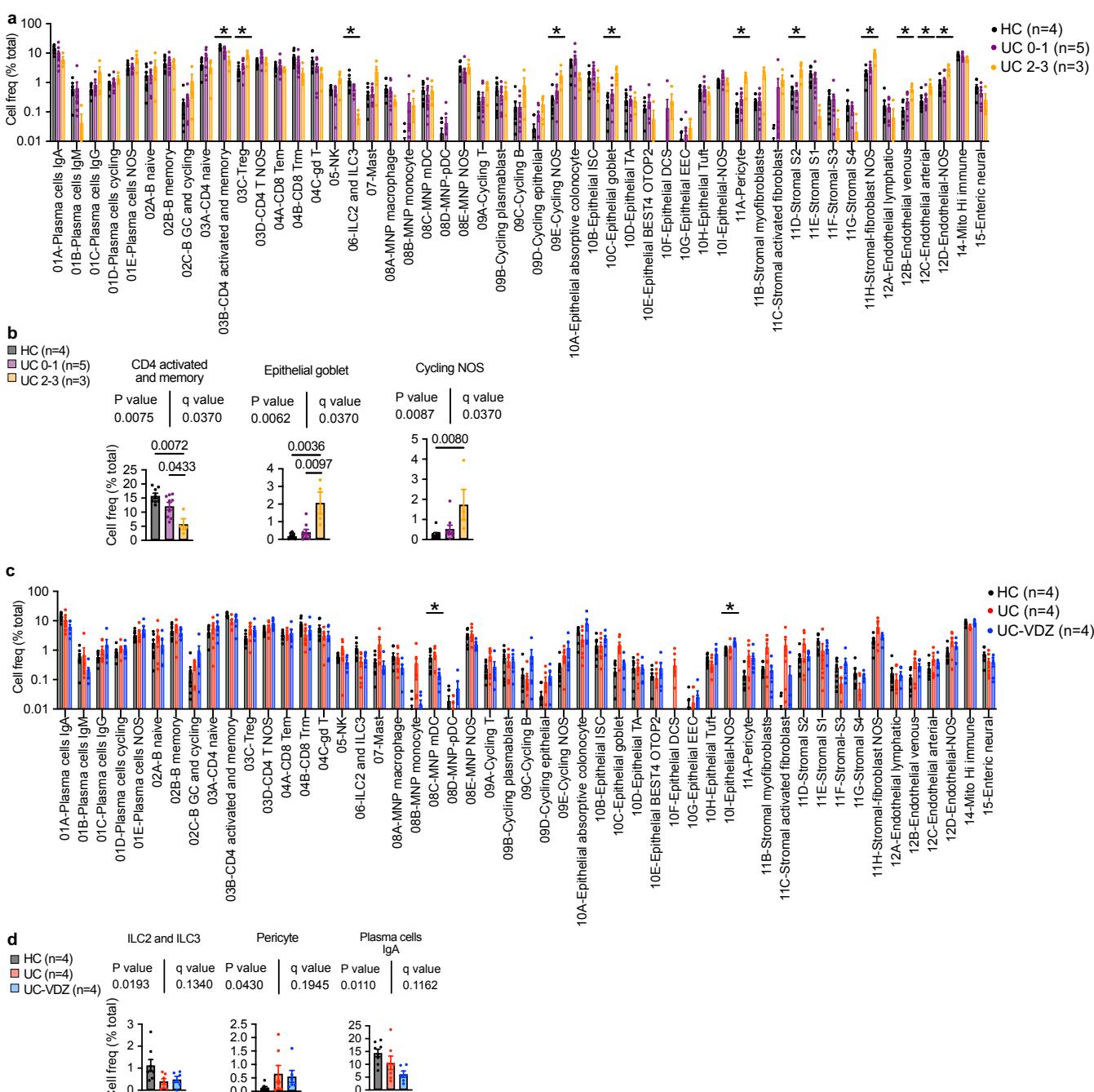
Supplementary Fig.3| Biopsy scRNA-seq and CITE-seq coarse cell subset annotations and landmark genes and proteins. **a**, UMAP visualization of scRNA-seq and CITE-seq for 93,900 cells from HC (n=4), UC (n=4), and UC-VDZ (n=4) patients highlighting coarse cell subset annotations. **b-c**, Dot plot representation of landmark (**b**) genes and (**c**) proteins expressed by indicated coarse cell subsets. Genes and proteins were ranked by Wilcoxon rank-sum. Cell subsets without landmark genes or proteins above selected minimum log fold change are not displayed.



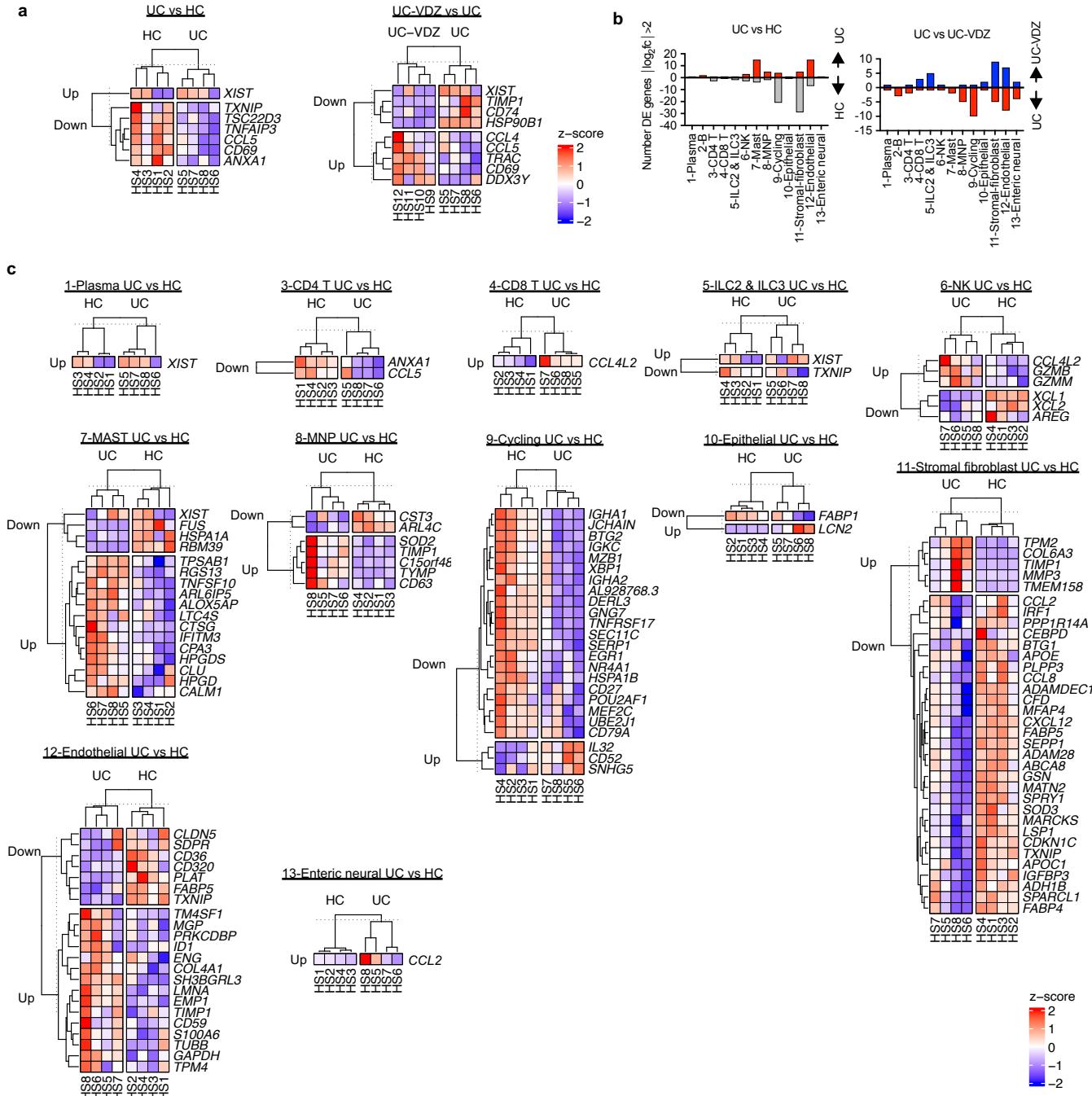
Supplementary Fig.4| Landmark genes and proteins associated with fine immune cell subsets in biopsy scRNA-seq and CITE-seq data. **a-j**, Dot plot representation of landmark (**a,c,e,g,i**) genes and (**b,d,f,h,j**) proteins expressed by indicated fine cell subsets. Genes and proteins were ranked by Wilcoxon rank-sum. Cell subsets without landmark genes or proteins above selected minimum log fold change are not displayed.



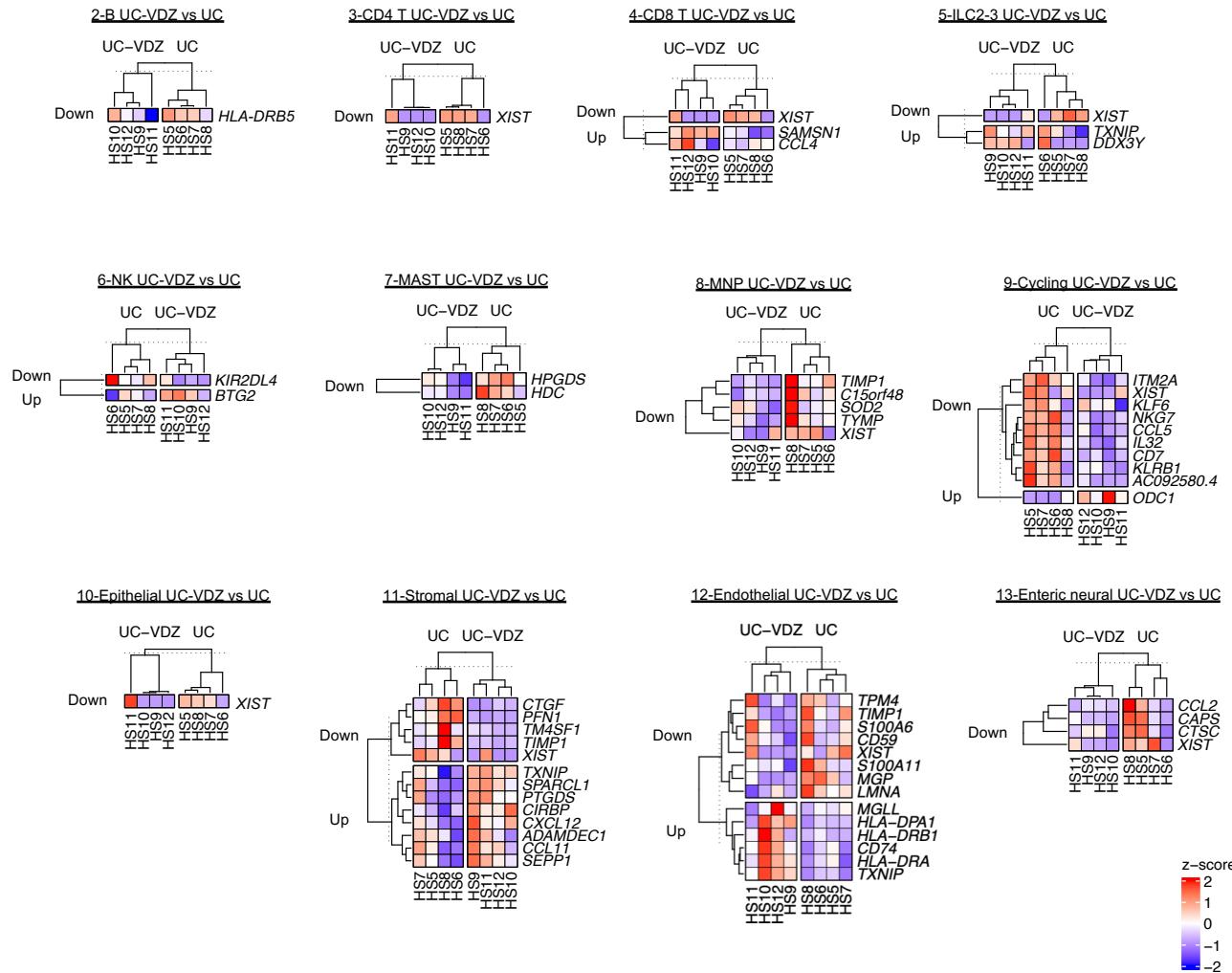
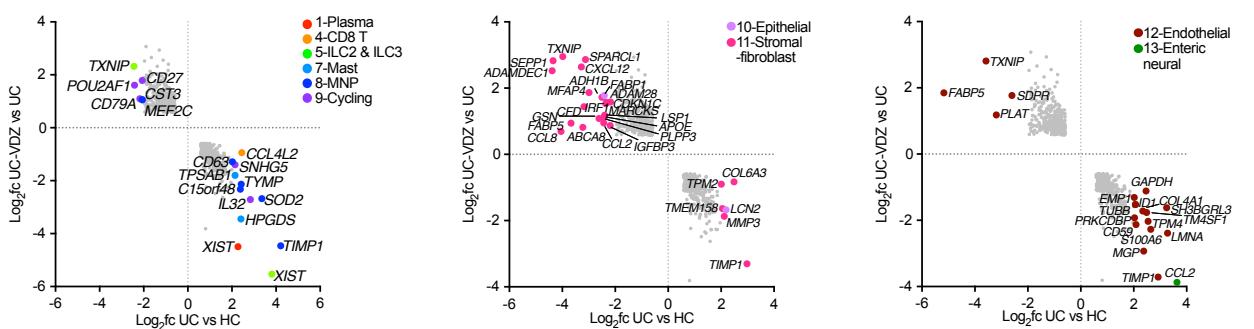
Supplementary Fig. 5 | Landmark genes and proteins associated fine non-immune cell subsets in biopsy biopsy scRNA-seq and CITE-seq data. **a-f**, Dot plot representation of landmark (**a,c,e**) genes and (**b,d,f**) proteins expressed by indicated biopsy fine cell subsets. Genes and proteins were ranked by Wilcoxon rank-sum. Cell subsets without landmark genes or proteins above selected minimum log fold change are not displayed.



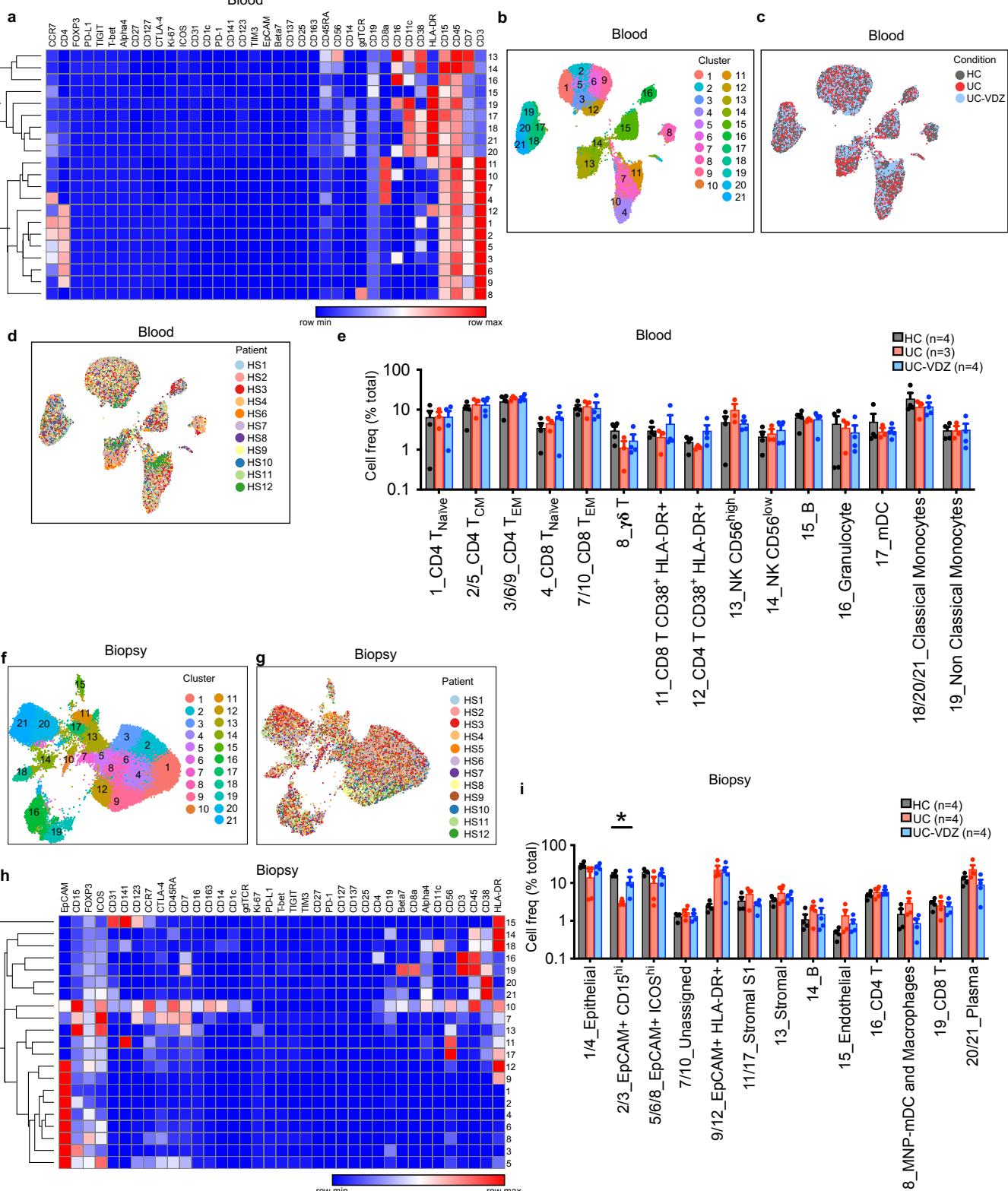
Supplementary Fig. 6 | Biopsy CITE-seq cell subsets that correlated with inflammatory severity, disease status, and VDZ treatment. **a-d**, Cell frequency as a percent of total cells per study subject and fine cell annotation stratified by (a,b) endoscopic severity and (c,d) disease and treatment status (mean \pm SEM; n=number of patients; each dot represents one patient sample). One-way ANOVA Kruskal-Wallis test was performed for each cell subset individually in (a, c), followed by global FDR correction of all p-values for all cell subsets, with q<0.1 threshold for discovery (*=q<0.1, only significant differences are indicated). A nested one-way ANOVA test was also performed on log transformed values treating biopsies as replicates with an unadjusted p<0.05 as an additional threshold for discovery (*). Select subsets are shown in (b, d) with exact p-value and q-value; NOS="not otherwise specified".



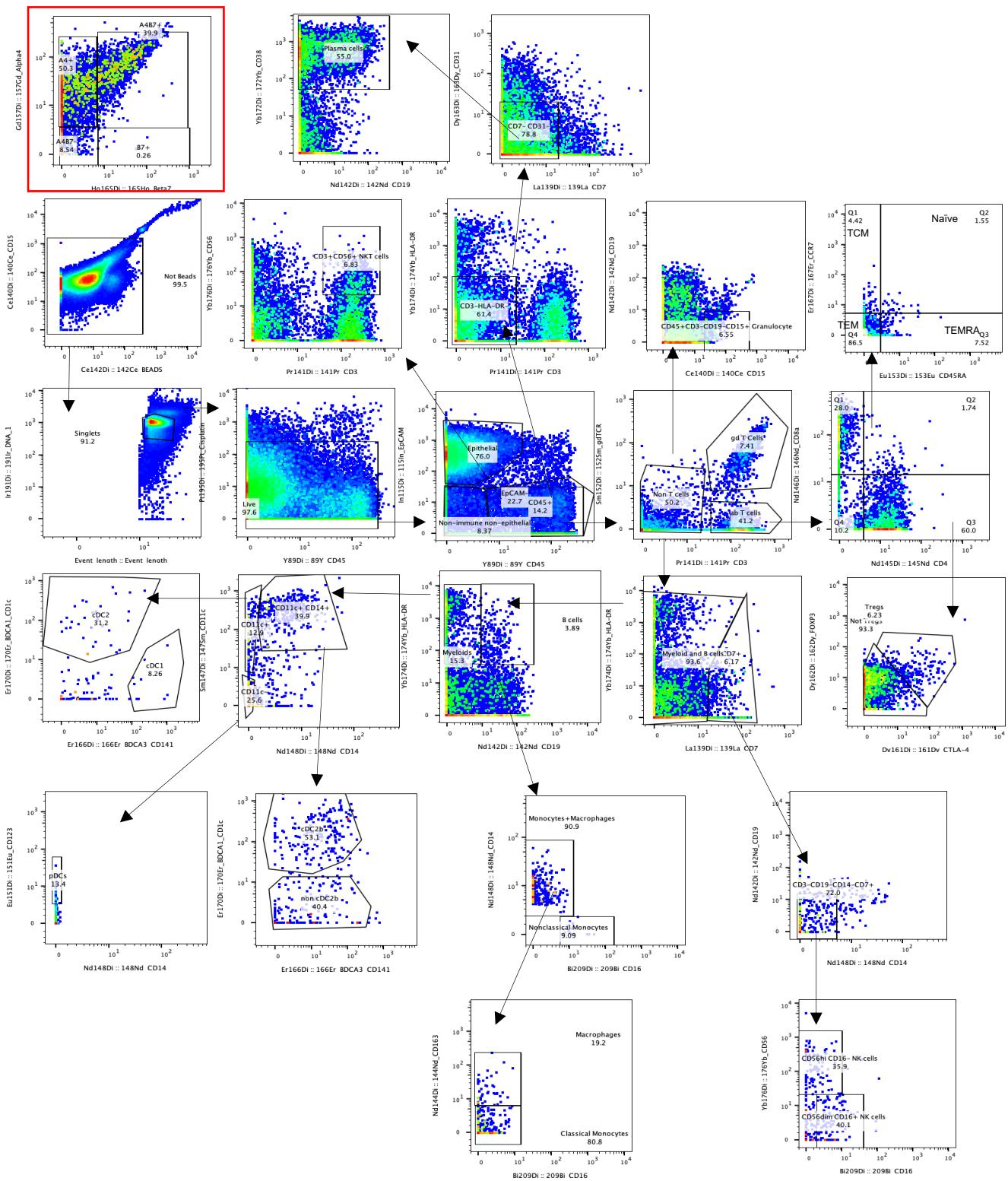
Supplementary Fig.7| Biopsy scRNA-seq DE genes in cell subsets comparing UC vs HC and UC-VDZ vs UC. **a**, Heatmap of expression z-scores for scRNA-seq DE genes for all biopsy cells, with $\log_2\text{fc} > 1$ or < -1 and Bonferroni-corrected p-value < 0.1 , comparing UC (Up/Down) relative to HC and UC-VDZ (Up/Down) relative to UC identified by MAST analysis. **b**, Number of scRNA-seq DE genes in biopsy cell subsets with $\log_2\text{fc} > 2$ or < -2 in UC versus HC and UC-VDZ versus UC identified by MAST analysis. **c**, Heatmap of expression z-scores for scRNA-seq DE genes for the indicated cell subsets with $\log_2\text{fc} > 2$ or < -2 and Bonferroni-corrected p-value < 0.1 in UC (Up/Down) relative to HC identified by MAST analysis. Ribosomal and mitochondrial genes are not displayed; cell subsets without DE genes are not displayed.

a**b**

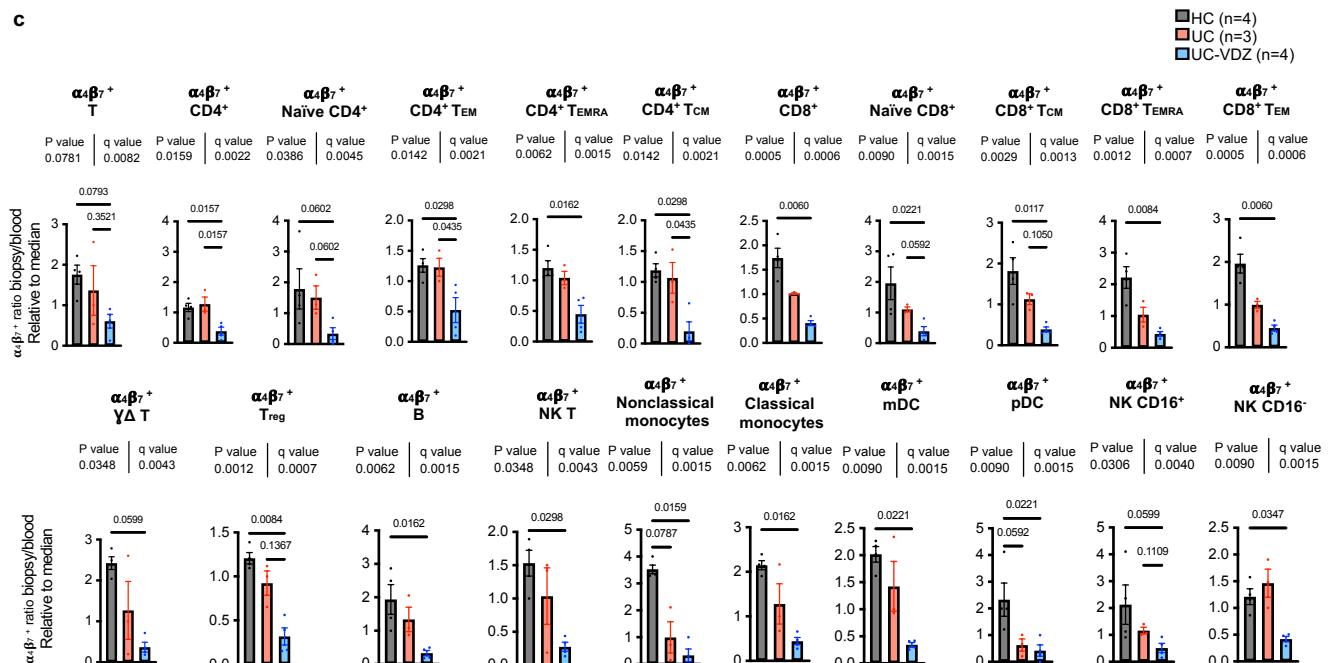
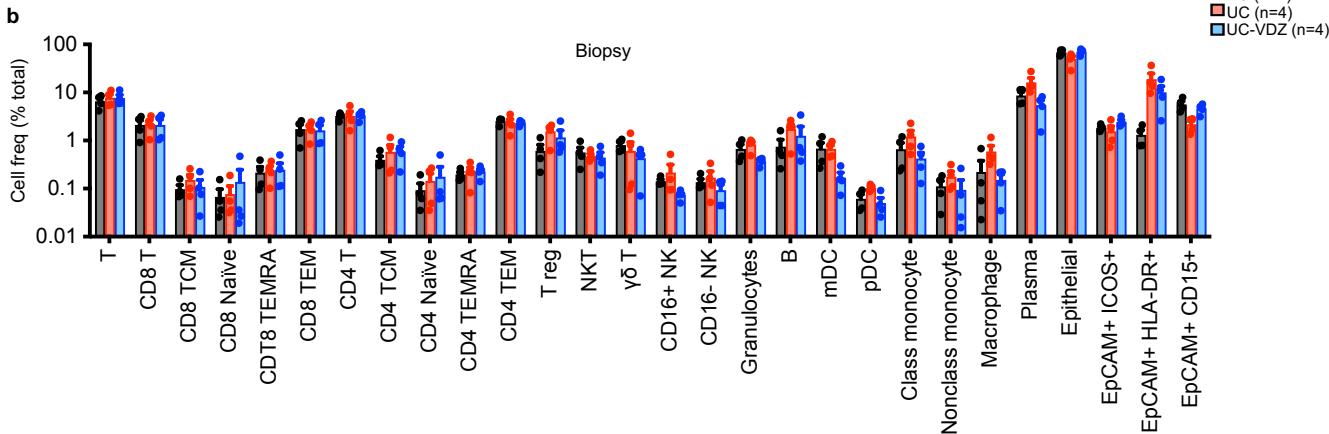
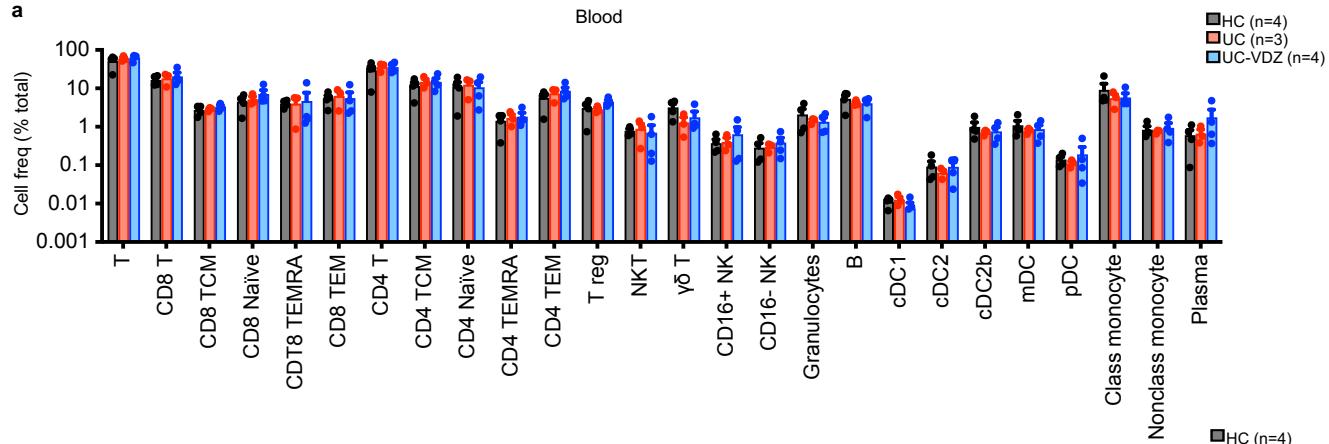
Supplementary Fig.8| Biopsy scRNA-seq DE genes in cell subsets comparing UC-VDZ vs UC and reciprocal expression analysis. **a**, Heatmap of expression z-scores for scRNA-seq DE genes for the indicated cell subsets with log₂fc >2 or <-2 and Bonferroni-corrected p-value <0.1 in UC-VDZ (Up/Down) relative to UC identified by MAST analysis. **b**, scRNA-seq DE genes in biopsy coarse cell annotation subsets with log₂fc >2 or <-2 in UC versus HC and an inverse log₂fc for UC-VDZ versus UC with log₂fc <-0.5 or >0.5 identified by MAST analysis, respectively. Ribosomal and mitochondrial genes are not displayed; cell subsets without DE genes are not displayed.



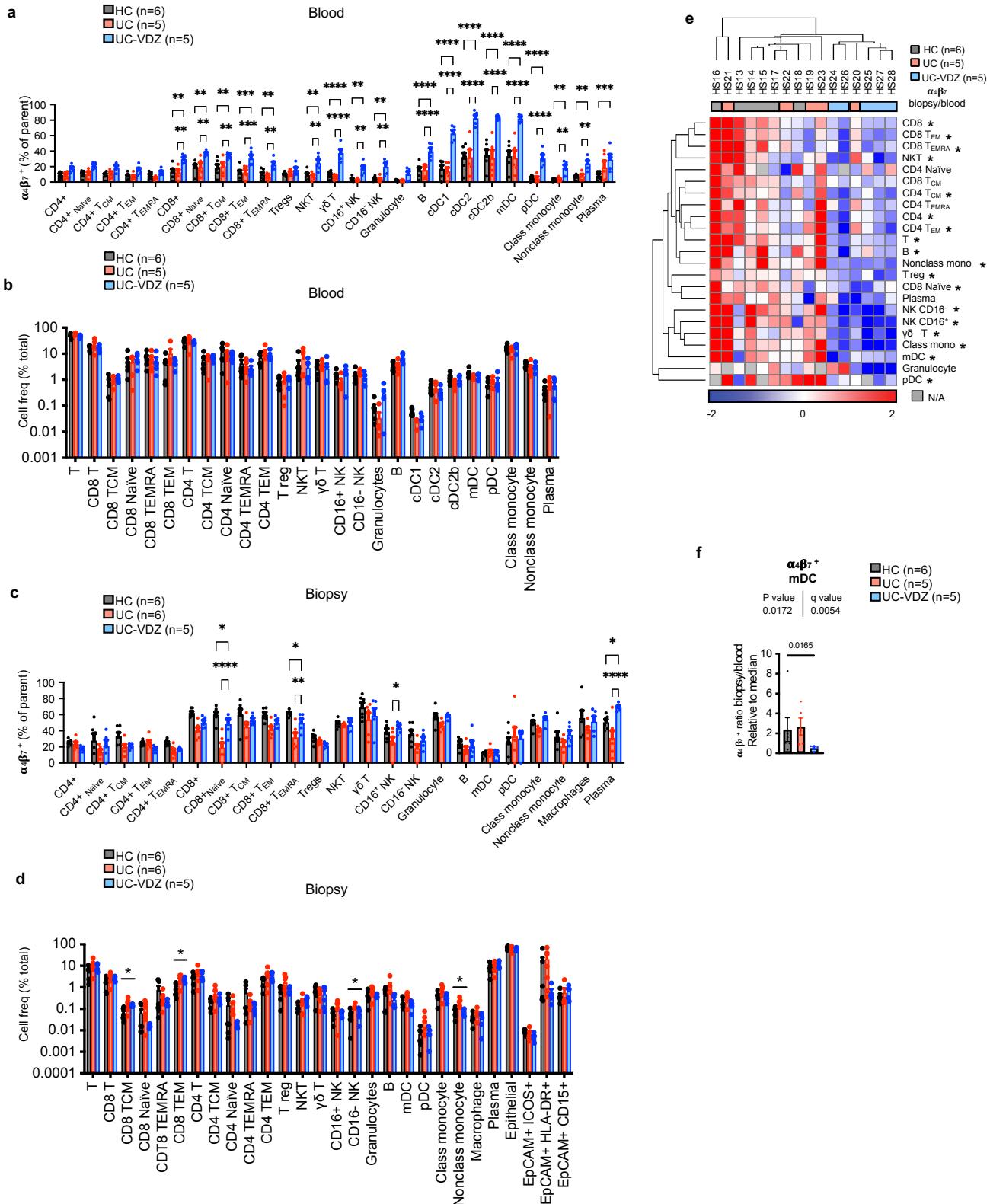
Supplementary Fig. 9| Heatmaps and total frequencies obtained from unsupervised analysis of CyTOF data from blood and biopsies. **a,h**, Heatmap visualization of mean intensity of each marker for the 21 identified clusters in blood and biopsies, respectively. **b-d**, UMAP visualization of the indicated samples (100,000 out of 1,390,156 live cell events displayed for blood) highlighting **(b)** FlowSOM identified clusters, **(c)** disease and treatment status, and **(d)** patient sample. **e,i**, Cell frequencies for each annotated cell subset per condition for **(e)** blood and **(i)** biopsy samples (mean \pm SEM; n=number of patients; each dot represents one patient sample). **f,g**, UMAP visualization of the indicated samples (60,000 out of 684,249 live cell events displayed for biopsies) highlighting **(f)** FlowSOM identified clusters, and **(g)** patient sample. One-way ANOVA Kruskal-Wallis test was performed for each cell subset individually in panels **(e)** and **(i)**, followed by global FDR correction of all p-values for all cell subsets, with $q < 0.1$ threshold for discovery (* $q < 0.1$, only significant differences are indicated).



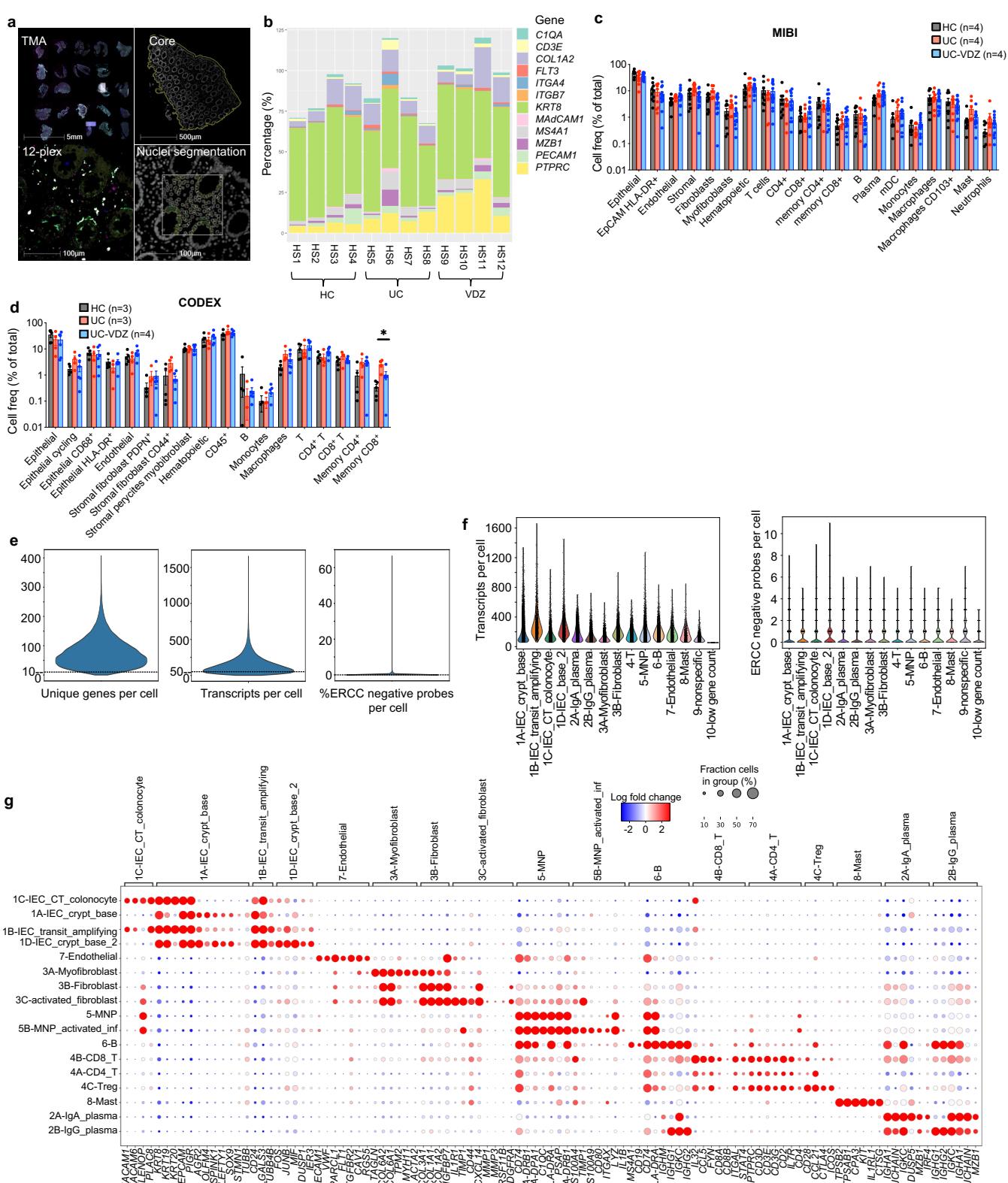
Supplementary Fig. 10 | Representative supervised CyTOF gating scheme.



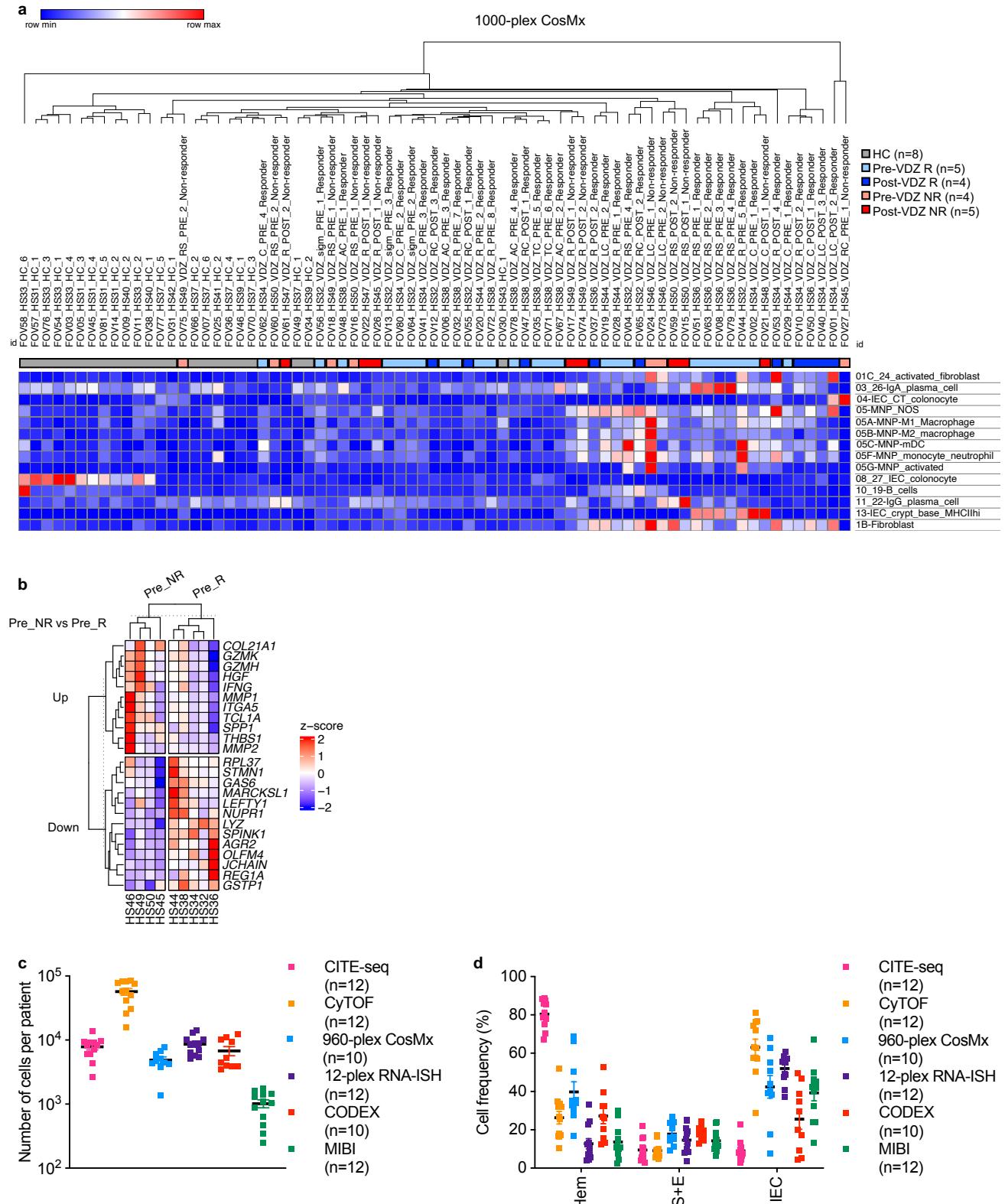
Supplementary Fig. 11| Total frequencies obtained from supervised analysis of CyTOF data from blood and biopsies and significant $\alpha_4\beta_7^+$ cell subsets. **a,b**, Cell frequencies for each annotated cell subset per condition in blood and biopsy, respectively. **c**, $\alpha_4\beta_7^+$ cell frequency in the biopsy versus blood for the indicated cell subsets relative to the median for each cell subset. For panels **a-b**, mean \pm SEM; n=number of patients; each dot represents one patient sample; multiple one-way ANOVA Kruskal-Wallis test with FDR correction; q<0.1 threshold for discovery; individual subsets are shown in **(c)** with exact p-value and q-value; individual inter-column q-values are displayed only for cell subsets with overall FDR corrected q<0.01. Class monocyte-classical monocyte; Nonclass monocyte-nonclassical monocyte; mDC-cDC1,cDC2,cDC2b.



Supplementary Fig. 12| $\alpha_4\beta_7^+$ cell subsets and total frequencies obtained from supervised analysis of CyTOF data from blood and biopsies in a second case control study of patients in near or complete endoscopic remission (HS13-HS30). a,c, Percentage of $\alpha_4\beta_7^+$ cells in each defined cell subset per condition for blood and biopsy samples, respectively (mean \pm SEM; n=number of patients; each dot represents one patient sample; two-way ANOVA comparing HC vs UC-VDZ and UC vs UC-VDZ with FDR correction; q<0.1 threshold for discovery; *q<0.05; **q<0.01; ***q<0.001; and ****q<0.0001; only significant differences are indicated. b,d, Cell frequencies for each annotated cell subset per condition in blood and biopsy, respectively (mean \pm SEM; n=number of patients, multiple one-way ANOVA Kruskal-Wallis test with FDR correction; q<0.1 threshold for discovery). e, Heatmap of biopsy/blood ratio of $\alpha_4\beta_7^+$ cells for each cell subset by patient (hierarchically clustered by Euclidian distance, average linkage). f, $\alpha_4\beta_7^+$ cell frequency in the biopsy versus blood for mDC relative to the median (mean \pm SEM; n=number of patients; each dot represents one patient sample).



Supplementary Fig.13| Spatial transcriptomics and proteomics on FFPE tissue microarray (TMA). **a**, Image analysis workflow for 12plex RNA-ISH assay using HALO, scan of the entire TMA (0.5X), with representative images (4X), representative overlay of the 12 markers (20X), and representative nuclei segmentation and localization of the 12 different markers (20X), which are representative of 19 cores. **b**, Stacked bar plots for cells expressing the indicated gene; values are greater than 100% due to cells co-expressing genes of interest. **c,d**, Cell frequencies as a percent of total for the indicated cell subsets identified by (c) MIBI and (d) CODEX; multiple one-way ANOVA Kruskal-Wallis test with FDR correction; $q < 0.1$ threshold for discovery; * $=q < 0.1$. **e**, Violin plots showing unique genes, transcripts, and %ERCC negative probes per cell. Cells with <10 unique genes or <50 transcripts per cell were excluded (dashed line). **f**, Transcripts and ERCC negative probes per cell, stratified by coarse annotation. **g**, Dot plot representation of curated landmark genes for annotated subsets. Genes ranked by Wilcoxon rank-sum. ERCC, External RNA Controls Consortium.



Supplementary Fig.14| CosMx spatial transcriptomic on TMA from retrospective longitudinal study and comparison of total cells analyzed per patient and cell frequency per tissue compartment among study subjects HS1-HS12. **a**, Heatmap of selected cell subsets by patient fov (hierarchically clustered by Euclidian distance, average linkage, columns color-coded by group). **b**, Heatmap of expression z-scores for the indicated genes in VDZ non-responders (Pre_NR) (Up/Down) relative to responders (Pre_R) before treatment identified by DESeq2. **c**, Total number of cells analyzed per patient using different multi-omic methods. **d**, Cell frequency per patient expressed as percent of total for the indicated cellular compartment among different multi-omic methods. For (**c**, **d**), mean \pm SEM, n=number of patients, each dot represents one patient sample. Hem-hematopoietic; S+E-stromal + endothelial; IEC-intestinal epithelial cell.

Patient ID	Age (y)	Sex	Disease status	UC Medication	Disease duration (y)	Duration VDZ (mo)	Prior anti-TNF exposure	Montreal classification	Mayo Endoscopic subscore	Responder to VDZ
HS1	50-59	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS2	40-49	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS3	20-29	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS4	50-59	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS5	20-29	F	UC	5-ASA	15	n/a	No	E2	2	n/a
HS6	40-49	M	UC	5-ASA	25	n/a	Yes	E3	2	n/a
HS7	60-69	F	UC	5-ASA	7	n/a	No	E2	1	n/a
HS8	50-59	F	UC	5-ASA	5	n/a	No	E1	1	n/a
				Median	11 (7-18)					
HS9	50-59	M	UC	VDZ	25	8	Yes	E2	1	Yes
HS10	30-39	M	UC	VDZ	8	7	No	E2	1	Yes
HS11	20-29	F	UC	VDZ	8	64	Yes	E2	3	No
HS12	50-59	M	UC	VDZ	10	2	Yes	E3	1	Yes
				9 (8-14)						
p-value	ns	ns	p=0.0025	p<0.0001	ns	n/a	ns	ns	p=0.01	
HS13	20-29	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS14	30-39	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS15	20-29	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS16	60-69	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS17	50-59	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS18	40-49	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS19	40-49	M	UC	ADA	27	n/a	Yes	E2	0	n/a
HS20*	30-39	F	UC	ADA	20	n/a	Yes	E2	1-2	n/a
HS21	50-59	M	UC	IFX/MTX	6	n/a	Yes	E3	0	n/a
HS22	30-39	F	UC	5-ASA	6	n/a	No	E3	0	n/a
HS23	40-49	M	UC	IFX/AZA	10	n/a	Yes	E3	1	n/a
HS30	40-49	F	UC	ADA	2	n/a	Yes	E3	0	n/a
				8 (5-22)						
HS24	50-59	M	UC	VDZ	5.5	45	Yes	E2	0	Yes
HS25	60-69	M	UC	VDZ	37	5	No	E3	0	Yes
HS26	30-39	F	UC	VDZ	21	55	Yes	E3	0	Yes
HS27	30-39	F	UC	VDZ	1	6	No	E1	0	Yes
HS28	40-49	M	UC	VDZ	8	45	Yes	E1	0	Yes
				8 (3-29)						
p-value	ns	ns	p=0.0002	p<0.0001	ns	n/a	p=0.0329	ns	ns	
HS31 (HS16)	60-69	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS33 (HS14)	30-39	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS35 (HS15)	20-29	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS37 (HS13)	20-29	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS39	20-29	F	HC	None	n/a	n/a	No	n/a	0	n/a
HS40	60-69	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS41	50-59	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS42	40-49	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS43	50-59	M	HC	None	n/a	n/a	No	n/a	0	n/a
HS32	20-29	F	UC	VDZ	2	4	No	E2	pre 2 ; post 0	Yes
HS34 (HS27)	30-39	F	UC	VDZ	1	6	No	E1	pre 2; post 0	Yes
HS36	40-49	M	UC	VDZ	3	2	Yes	E2	pre 3; post 0	Yes
HS38 (HS9)	50-59	M	UC	VDZ	25	7	Yes	E3	pre 2; post 0	Yes
HS44	30-39	F	UC	VDZ	2	2	No	E1	pre 1	Yes
HS45	40-49	M	UC	VDZ	14	19	Yes	E2	pre 1; post 3	No
HS46	40-49	n/a	UC	VDZ	21	2	Yes	E2	pre 3; post 3	No
HS47	30-39	M	UC	VDZ	11	5	Yes	E2	post 3	No
HS48	70-79	M	UC	VDZ	8	6	Yes	E3	pre 2; post 2	No
HS49	30-39	F	UC	VDZ	2	3	Yes	E2	pre 2; post 3	No
HS50	20-29	M	UC	VDZ	0	5	Yes	E2	pre 2; post 2	No
				3 (2-14)						
p-value	ns	ns	p<0.0001	p<0.0001	n/a	n/a	p=0.0010	ns	pre, ns; post, p=0.0016	

Supplementary Table 1. Baseline demographic and clinical data for study participants. Categorical variables were analyzed by Chi-square test and continuous variables were compared using one-way ANOVA with FDR correction or Mann-Whitney test where appropriate. ns, not significant; n/a, not applicable; pre, pre-VDZ treatment; post, post-VDZ treatment. *HS20 had a short segment of moderate proctitis, otherwise in endoscopic remission with no change in therapy. HS31, HS33, HS35, HS37, HS34, HS38 were additional samples from HS16, HS14, HS15, HS13, HS27, and HS9, respectively. ADA-adalimumab; IFX-infliximab; VDZ-vedolizumab; AZA-azathioprine; 5-ASA-5-aminosalicylic acid.

Patient ID	Disease status	Medication	Biopsy location-severity	Mayo Endoscopic subscore	Geboes grade	Biopsy sample ID	Blood scRNA-seq	Biopsy scRNA-seq	CITE-seq	Blood CyTOF	Biopsy CyTOF	12-plex RNA-ISH	CODEX	MIBI	CosMx SMI 960-plex
HS1	HC	N/A	R-0	0	0.0	HS1-CLR1	Y	Y		Y	Y	Y (1-core)	N/A	N/A	Y (1-FOV)
			L-0		1.1	HS1-CLL1		Y				Y (1-core)	N/A	Y (2-FOV)	Y (2-FOV)
HS2	HC	N/A	R-0	0	1.1	HS2-CLR1	Y	Y		Y	Y	Y (1-core)	Y (1-core)	Y (3-FOV)	Y (1-FOV)
			L-0		1.1	HS2-CLL1		Y				Y (1-core)	Y (1-core)	Y (3-FOV)	Y (1-FOV)
HS3	HC	N/A	R-0	0	0.0	HS3-CLR1	Y	Y		Y	Y	N/A	Y (1-core)	N/A	Y (1-FOV)
			L-0		0.0	HS3-CLL1		Y				Y (1-core)	Y (1-core)	Y (1-FOV)	N/A
HS4	HC	N/A	R-0	0	0.0	HS4-CLR1	Y	Y		Y	Y	Y (1-core)	N/A	N/A	Y (1-FOV)
			L-0		0.0	HS4-CLL1		Y				Y (1-core)	Y (1-core)	Y (1-FOV)	Y (1-FOV)
HS5	UC	5-ASA	R-0	2	0.0	HS5-CLR1	N/A	Y		N/A	Y	Y (1-core)	Y (1-core)	Y (2-FOV)	Y (1-FOV)
			L-2		0.2	HS5-CLL1		Y				Y (1-core)	N/A	Y (1-FOV)	N/A
HS6	UC	5-ASA	R-2	2	2A.2	HS6-CLR1	Y	Y		Y	Y	Y (1-core)	N/A	Y (2-FOV)	N/A
			L-2		3.1	HS6-CLL1		Y				N/A	N/A	N/A	N/A
HS7	UC	5-ASA	R-1	1	1.1	HS7-CLR1	Y	Y		Y	Y	Y (1-core)	Y (1-core)	N/A	Y (1-FOV)
			L-0		0.0	HS7-CLL1		Y				Y (1-core)	Y (1-core)	Y (2-FOV)	Y (1-FOV)
HS8	UC	5-ASA	L-1	1	3.2	HS8-CLL1	Y	Y		Y	Y	Y (1-core)	Y (1-core)	Y (1-FOV)	Y (1-FOV)
HS9	UC	VDZ	R-0	1	0.0	HS9-CLR1	Y	Y		Y	Y	Y (1-core)	Y (1-core)	Y (3-FOV)	N/A
			L-1		0.0	HS9-CLL1		Y				Y (1-core)	Y (1-core)	Y (3-FOV)	N/A
HS10	UC	VDZ	R-0	1	1.1	HS10-CLR1	Y	Y		Y	Y	Y (1-core)	Y (1-core)	Y (1-FOV)	Y (1-FOV)
			L-1		0.1	HS10-CLL1		N/A				N/A	N/A	N/A	Y (1-FOV)
HS11	UC	VDZ	R-0	3	0.1	HS11-CLR1	Y	Y		Y	Y	N/A	N/A	N/A	Y (1-FOV)
			L-3		3.2	HS11-CLL1		Y				Y (1-core)	Y (1-core)	Y (1-FOV)	Y (1-FOV)
HS12	UC	VDZ	R-1	1	1.2	HS12-CLR1	Y	Y		Y	Y	Y (1-core)	Y (1-core)	Y (3-FOV)	N/A
			L-1		1.3	HS12-CLL1		N/A				Y (1-core)	Y (1-core)	Y (3-FOV)	Y (1-FOV)

Supplementary Table 2. Biospecimens analyzed by each multi-omics method. R, right colon; L, left colon; 5-ASA, 5-aminosalicylate; CLR1, right colon biopsy 1; CLL1, left colon biopsy; Y, yes; N/A not applicable. FOV-field of view

Assay type	Target	Ion channel / DNA barcode	Metal / DNA barcode ID	Clone/Probe	Concentration (µg/ml) /Dilution (ratio)	h5ad file symbol	Catalog #	Vendor
CITE-seq (TotalSeq)	CD103 (Integrin αE)	GACCTCATTTGTAAT	A0145	Ber-ACT8	proprietary	CD103	399907	BioLegend
	CD35	ACTTCCCGTCGATCTT	A0167	E11	proprietary	CD35	399907	BioLegend
	CD25	TTTGCTCTGCACTGCC	A0085	BC96	proprietary	CD25	399907	BioLegend
	CD62L	GTCCTTGCAACTTGA	A0147	DREG-56	proprietary	CD62L	399907	BioLegend
	CD3	CTCATTGTAACCTCT	A0034	UCHT1	proprietary	CD3	399907	BioLegend
	CD33	TAACTCAGGGCTAT	A0052	P67.6	proprietary	CD33.1	399907	BioLegend
	CD4	TGTTCCCGCTCAACT	A0072	RPA-T4	proprietary	CD4.2	399907	BioLegend
	CD58 (LFA-3)	GTTCCATATGGACGAC	A0174	TS2/9	proprietary	CD58.1	399907	BioLegend
	CD8a	GCTGCCCTTTCATT	A0080	RPA-T8	proprietary	CD8a	399907	BioLegend
	CD56	TCTTCTTGTGATAGG	A0047	5.1H11	proprietary	CD56-NCAM	399907	BioLegend
	CD11b	GACAAGTGATCTGCA	A0161	ICRF44	proprietary	CD11b	399907	BioLegend
CD183 (CXCR3)		GCGATGGTAGATTAT	A0140	G025H7	proprietary	CD183-CXCR3	399907	BioLegend
	CD30	TCAGGGTGTGCTGTA	A0028	BY88	proprietary	CD30	399907	BioLegend
CD336 (NKp44)		GGGCAATTAGGAGGT	A0802	P44-8	proprietary	CD336-NKp44	399907	BioLegend
	CD69	GTCCTTGGCTTAA	A0146	FN50	proprietary	CD69.1	399907	BioLegend
CD335 (NKp46)		ACAATTGAAACAGCG	A0101	9E2	proprietary	CD335-NKp46	399907	BioLegend
	CD49f	TTCGAGGATGATCT	A0070	G0H3	proprietary	CD49f	399907	BioLegend
	CD161	GTACGCAGTCCTCT	A0149	HP-3G10	proprietary	CD161	399907	BioLegend
CD45RO		CTCCGAATCATGTTG	A0087	UCHL1	proprietary	CD45RO	399907	BioLegend
	CD20	TTCTGGTGTGCTAGA	A0100	2H7	proprietary	CD20	399907	BioLegend
KLRG1 (MAFA)		CTTATTCTCTGGCCT	A0153	SA231A2	proprietary	KLRG1-MAFA	399907	BioLegend
	CD19	CTGGGCAATTACTCG	A0050	HIB19	proprietary	CD19.1	399907	BioLegend
	CD32	GCTTCGGAATTACCG	A0142	FUN-2	proprietary	CD32	399907	BioLegend
CD276 (B7-H3)		GACTGGAGGGATT	A0010	DCN.70	proprietary	CD276-B7-H3	399907	BioLegend
CD279 (PD-1)		ACAGCGCCGTATT	A0088	EH12.2H7	proprietary	CD279-PD-1	399907	BioLegend
CD197 (CCR7)		AGTTAGCTGACCGA	A0148	G043H7	proprietary	CD197-CCR7	399907	BioLegend
CD252 (OX40L)		TTTAGTGATCCGACT	A0021	11C3.1	proprietary	N/A	399907	BioLegend
CD223 (LAG-3)		CATTGTCGCCGGT	A0152	11C3C65	proprietary	CD223-LAG-3	399907	BioLegend
HLA-DR		AAATCGGAGCAAGTA	A0159	L243	proprietary	HLA-DR	399907	BioLegend
CD45RA		TCAATCCTCTGGCTT	A0063	HI100	proprietary	CD45RA	399907	BioLegend
CD196 (CCR6)		GATCCCTTGTCACT	A0143	G034E3	proprietary	CD196-CCR6	399907	BioLegend
	CD2	TACGATTTGTCAGG	A0367	TS1/8	proprietary	CD2.1	399907	BioLegend
CD326 (Ep-CAM)		TTCCGAGCAAGTAC	A0123	9C4	proprietary	CD326-EpCAM	399907	BioLegend
CD169 (Sialoadhesin, Siglec-1)		TACTCAGCGTGTG	A0206	7-239	proprietary	CD169-Sialoadhesin-Siglec-1	399907	BioLegend
	CD41	ACGTTGTGCGCTGT	A0353	HIP8	proprietary		399907	BioLegend
	CD49b	GCTTTCTTCACTATG	A0371	P1E6-C5	proprietary	CD49b	399907	BioLegend
CD303 (BDCA-2)		GAGATGTCCGAATT	A0370	201A	proprietary	CD303-BDCA-2	399907	BioLegend
	CD29	GTATTCCCTCAGTC	A0369	TS2/16	proprietary	CD29	399907	BioLegend
CD184 (CXCR4)		TCAGGTCTTTAAC	A0366	12G5	proprietary	CD184-CXCR4	399907	BioLegend
	CD98	GCACCAACAGCCATT	A0374	MEM-108	proprietary	CD98	399907	BioLegend
CD124 (IL-4R α)		CCGTCTCTGATAGT	A0363	G077F6	proprietary	N/A	399907	BioLegend
CD135 (Flt-3/Fk-2)		CAGTAGATGGAGCAT	A0351	BV10A4H2	proprietary	CD135	399907	BioLegend
CD370 (CLEC9A/DNGR1)		CTGCATTTCAGTAAG	A0207	8F9	proprietary	CD370-CLEC9A-DNGR1	399907	BioLegend
	CD21	ACACTAGTGTGTCGG	A0181	Bu32	proprietary	CD21	399907	BioLegend
	CD39	TTACCTGGTATCCGT	A0176	A1	proprietary	CD39	399907	BioLegend
CD141 (Thrombomodulin)		GGATAACCGCGCTT	A0163	M80	proprietary	CD141-Thrombomodulin	399907	BioLegend
	CD64	AAAGTATGCCCTAGGA	A0162	10.1	proprietary	CD64	399907	BioLegend
CD152 (CTLA-4)		ATGGTTCACGTAATC	A0151	BN13	proprietary	CD152	399907	BioLegend
	CD61	AGGTTGGAGTAGACT	A0372	VI-PL2	proprietary	CD61	399907	BioLegend
	CD163	GCTTTCTTCCCTCTTA	A0358	GH1/61	proprietary	CD163.1	399907	BioLegend
CD357 (GITR)		ACCTTTGACACTCG	A0360	10B-17	proprietary	CD357-GITR	399907	BioLegend
CD137 (4-1BB)		CAGTAAGTTCGGGAC	A0355	4B4-1	proprietary	CD137	399907	BioLegend
CD81 (TAPA-1)		GTATCCTTCTTCGTC	A0373	5A6	proprietary	CD81-TAPA-1	399907	BioLegend
CD366 (Tim-3)		TGTCCTACCCAATT	A0169	F38-2E2	proprietary	CD366-Tim-3	399907	BioLegend
CD57 Recombinant		AACTCCCTATGGAGG	A0168	QA17A04	proprietary	CD57	399907	BioLegend
CD278 (ICOS)		CGCGCACCCATTAAA	A0171	C398.4A	proprietary	CD278-ICOS	399907	BioLegend
CD95 (Fas)		CCAGCTTCAAGAGC	A0156	DX2	proprietary	CD95-Fas	399907	BioLegend
CD80		ACGAATCAATCTGTG	A0005	2D10	proprietary	CD80.1	399907	BioLegend
CD138 (Syndecan-1)		ACTCTTCGTTTACG	A0055	MI15	proprietary	CD138-Syndecan-1	399907	BioLegend
	CD70	CGCGAACATAAGAAC	A0027	113-16	proprietary	CD70.1	399907	BioLegend
CD269 (BCMA)		CAGATGATCCCCATT	A0056	19F2	proprietary	CD269-BCMA	399907	BioLegend
IgM		TAGCGAGCCCTATA	A0136	MHM-88	proprietary	IgM	399907	BioLegend
CD59		AATTAGCCTGTCGAGA	A0361	p282 (H19)	proprietary	CD59.1	399907	BioLegend
CD194 (CCR4)		AGCTTACTTCGACGA	A0071	L291H4	proprietary	CD194-CCR4	399907	BioLegend
CD275 (B7-H2, ICOSL)		GTGCAATTCAACAGTA	A0009	2D3	proprietary	CD275-B7-H2-ICOSL	399907	BioLegend
CD314 (NKG2D)		CGTGTTCCTCTCTCA	A0165	1D11	proprietary	CD314-NKG2D	399907	BioLegend
	CD86	GCTTTGTCAGTGC	A0006	IT2.2	proprietary	CD86.1	399907	BioLegend
Galectin-9		ACTCACTGGAGCTC	A0016	9M1-3	proprietary	N/A	399907	BioLegend
CD195 (CCR5)		CCAAAGTAAGGCCA	A0141	J418F1	proprietary	CD195-CCR5	399907	BioLegend
CD1c		GAGCTACTTCACCTG	A0160	L161	proprietary	CD1c	399907	BioLegend
CD28		TGAGAACGACCCCTA	A0386	CD28.2	proprietary	CD28.1	399907	BioLegend
TSL PR (TSLPR-R)		CAGTCCTCTCTGTC	A0387	1D3	proprietary	TSL PR	399907	BioLegend
	CD38	TGTACCCGCTTGTGA	A0389	HIT2	proprietary	CD38.1	399907	BioLegend
CD155 (PVR)		ATCACATGTTGTCGA	A0023	SKII.4	proprietary	CD155-PVR	399907	BioLegend
CD270 (HVEM, TR2)		TTAGAGAAACAGCC	A0020	122	proprietary	CD270-HVEM-TR2	399907	BioLegend
CD178 (Fas-L)		CCGGTCTCTGTATT	A0177	NOK-1	proprietary	CD178-FasL	399907	BioLegend
CD127 (IL-7Ra)		TGTTGTTGTCCTATG	A0390	A019D5	proprietary	CD127-IL-7Ra	399907	BioLegend
	CD7	TGGATTCCCGGAGCT	A0066	CD7-6B7	proprietary	CD7.1	399907	BioLegend
CD117 (c-kit)		AGACTAATAGCTGAC	A0061	104D2	proprietary	CD117-c-kit	399907	BioLegend
	CD10	CAGCCATTCTATTAG	A0062	HI10a	proprietary	CD10	399907	BioLegend
CD40		CTCAGATGAGATGATG	A0031	5C3	proprietary	CD40.1	399907	BioLegend
CD48		CTACGACGCTAGAGA	A0029	BJ40	proprietary	CD48.1	399907	BioLegend
CD154		GCTAGATAGATGCAA	A0032	24-31	proprietary	CD154	399907	BioLegend
	CD47	GCATTCCTGTCACCTA	A0026	CC2C6	proprietary	CD47.1	399907	BioLegend
CD112 (Nectin-2)		AACTCCGCTCAAG	A0024	TX31	proprietary	CD112-Nectin-2	399907	BioLegend
	CD52	CTTGTACGAGCAGAA	A0033	HI186	proprietary	CD52.1	399907	BioLegend
CD71		CCGTGTTCTCTATTA	A0394	CY1G4	proprietary	CD71	399907	BioLegend
CD22		GGGTTGTTGTCCTTG	A0393	S-HCL-1	proprietary	CD22.1	399907	BioLegend
CD15 (SSEA-1)		TCACCAAGTACCTAGT	A0392	W6D3	proprietary	CD15-BEA-1	399907	BioLegend

XCR1	AAGACGCTATGTCAAC	A0208	S15046E	proprietary	XCR1.1	399907	BioLegend
CD26	GGTGGCTAGATAATG	A0396	BA5b	proprietary	CD26	399907	BioLegend
B7-H4	TGTATGCTGCCTTG	A0395	MIH43	proprietary	B7-H4	399907	BioLegend
CD31	ACCTTTATGCCACCG	A0124	WM59	proprietary	CD31	399907	BioLegend
CD14	TCTCAGACCTCCGTA	A0081	M5E2	proprietary	CD14.2	399907	BioLegend
TIGIT (VSTM3)	TTGCTTACCGCCAGA	A0089	A15153G	proprietary	TIGIT-VSTM3	399907	BioLegend
CD294 (CRTH2)	TGTTACCGAGGCC	A0102	BM16	proprietary	CD294-CRTH2	399907	BioLegend
CD45	TCCCTTGGATTTAC	A0048	2D1	proprietary	CD45	399907	BioLegend
CD34	GCAGAAATCTCCCT	A0054	581	proprietary	CD34.1	399907	BioLegend
CD244 (2B4)	TGCGCTTGGATGGTAG	A0189	C1.7	proprietary	CD244-2B4	399907	BioLegend
CD185 (CXCR5)	AATTCAACCCTGCC	A0144	J252D4	proprietary	CD185-CXCR5	399907	BioLegend
CD18	TATGGGGACACTCT	A0385	TS1/18	proprietary	CD18	399907	BioLegend
CD119 (IFN-γ R ? chain)	TGTTATTCCTCTGT	A0219	GIR-208	proprietary	CD119-IFN-G-R-A-chain	399907	BioLegend
CD134 (OX40)	AACCCACCGTTGTTA	A0158	Ber-ACT35 (ACT35)	proprietary	CD134-OX40	399907	BioLegend
CD24	AGATTCCCTCGTGT	A0180	ML5	proprietary	CD24.1	399907	BioLegend
TCR γ/δ	CTTCCGATTCTTCA	A0139	B1	proprietary	TCR-G-TCR-D	399907	BioLegend
CD115 (CSF-1R)	AAATCACGGCTCTGT	A0398	9-4D2-E4	proprietary	CD115-CSF-1R	399907	BioLegend
CD1d	TCGAGTCGCTTATCA	A0164	51.1	proprietary	CD1d	399907	BioLegend
CD192 (CCR2)	GAGTTCCCTTACCTG	A0242	K036C2	proprietary	CD192-CCR2	399907	BioLegend
CD105	ATCGTCGAGAGCTAG	A0068	43A3	proprietary	CD105	399907	BioLegend
CD144 (VE-Cadherin)	TCCACTCATCTGTA	A0400	BV9	proprietary	CD144-VE-cadherin	399907	BioLegend
CD193 (CCR3)	ACCAATCCCTTCGTC	A0397	5E8	proprietary	CD193-CCR3	399907	BioLegend
CD55	GCTCATTAACCCATTA	A0383	JS11	proprietary	CD55.1	399907	BioLegend
CD79b (Igβ)	ATTCTTCAACCGAAG	A0187	CB3-1	proprietary	CD79b-Ig-B	399907	BioLegend
CD123	CTTCACTCTGTCAGG	A0064	6H6	proprietary	CD123	399907	BioLegend
CD11c	TACGCCATAAACCTG	A0053	S-HCL-3	proprietary	CD11c	399907	BioLegend
CD54	CTGATAGACCTTGGAT	A0217	HA58	proprietary	CD54	399907	BioLegend
CD268 (BAFF-R)	CGAAGTCGATCCGTA	A0215	11C1	proprietary	CD268-BAFF-R	399907	BioLegend
TIM4	CGTCTATAGATGTTG	A0428	9F4	proprietary	Tim4	399907	BioLegend
CD1a	GATCGTGTGTTGTTA	A0402	HI149	proprietary	CD1a	399907	BioLegend
CD301 (CLEC10A)	ACCTAGAAATCAGCA	A0401	H037G3	proprietary	CD301-CLEC10A	399907	BioLegend
CD63	GAGATGTCGAACT	A0404	H5C6	proprietary	CD63.1	399907	BioLegend
CD44	AATCCCTCGAATGTT	A0125	BJ18	proprietary	CD44.1	399907	BioLegend
IgD	CAGTCCTCGTAGAGT	A0384	IA6-2	proprietary	IgD	399907	BioLegend
CD72	CAGTCGTGTTGAGATA	A0419	3F3	proprietary	CD72.1	399907	BioLegend
CD93	GCGCTACTCCCTGA	A0446	VIMD2	proprietary	CD93.1	399907	BioLegend
CD85g (ILT7)	TGTCAGTTCCATGTA	A0409	17G10.2	proprietary	CD85g-ILT7	399907	BioLegend
CD172a (SIRPa)	CGTGTITTAACITGAG	A0408	15-414	proprietary	CD172a-SIRPa	399907	BioLegend
CD284 (TLR4)	GCTTAGCTGTATCCG	A0405	HTA125	proprietary	CD284-TLR4	399907	BioLegend
CD304 (Neuropilin-1)	GGACTAAGTTCTGTT	A0406	12C2	proprietary	CD304-Neuropilin-1	399907	BioLegend
CD36	TTCTTGCCTTGC	A0407	5-271	proprietary	CD36.1	399907	BioLegend
CD102 (ICAM-2)	TGACCTTCTCTCTC	A0244	CBR-IC2/2	proprietary	CD102	399907	BioLegend
TCR α/β	CGTAACTGAGGGCA	A0224	IP26	proprietary	TCRA-TCR-B	399907	BioLegend
CD122 (IL-2R?)	TCATTTCCTCCGATT	A0246	TU27	proprietary	N/A	399907	BioLegend
CD83	CCACTCATTCGGT	A0359	HB15e	proprietary	CD83.1	399907	BioLegend
CD267 (TACI)	AGTGTATGGAGGAA	A0247	1A1	proprietary	CD267-TACI	399907	BioLegend
anti-human IgA	AAGATGTCGAGCAA	A0186	#N/A	proprietary	N/A	399907	BioLegend
CD107a (LAMP-1)	CAGCCCCACTGCAATA	A0155	H4A3	proprietary	CD107a-LAMP-1	399907	BioLegend
CD146	CCTTGATAAACATCA	A0134	P1H12	proprietary	CD146	399907	BioLegend
CD140b (PDGFR?)	CAATGGTTCACTGCC	A0129	18A2	proprietary	CD140b	399907	BioLegend
CD49a	ACTGTATGGACTCAGA	A0575	TS2/7	proprietary	CD49a	399907	BioLegend
α	AACTCTGTGTTGAG	A0584	6B11	proprietary	TCR-V-A-24-JA-18-InKT cell	399907	BioLegend
CD9	GAGTCACCAACTCTG	A0579	H19a	proprietary	CD9.1	399907	BioLegend
TCR Vα7.2	TACGAGCAGTATTCA	A0581	3C10	proprietary	TCR-V-A-7.2	399907	BioLegend
TCR Vγ9	AAGTATGTGTTCTG	A0583	B3	proprietary	TCR-V-G-9	399907	BioLegend
CD49d	CCATTCAACTCCGG	A0576	9F10	proprietary	CD49d	399907	BioLegend
C5L2	ACAATTGGTCTGCGA	A0572	1D9-M12	proprietary	C5L2	399907	BioLegend
CD338 (ABCg2)	TAAGACTTGGCCGTC	A0569	5D3	proprietary	CD338-ABCg2	399907	BioLegend
CD73 (Ecto-5'-nucleotidase)	CAGTTCTCAGTCTG	A0577	AD2	proprietary	CD73-Ecto-5'-nucleotidase	399907	BioLegend
anti-human CD79a (Ig?)	CTTATCACCGCTT	A0578	#N/A	proprietary	N/A	399907	BioLegend
CD158b (KIR2DL2/L3, NKAT2)	GACCCGTAGTTGAT	A0592	DX27	proprietary	CD158b-KIR2DL2-KIR2DL3-		
CD226 (DNAM-1)	AGACCAACTCTATCA	A0805	TX25	proprietary	NKAT2	399907	BioLegend
CD186 (CXCR6)	GACAGTCGATGCCA	A0804	K041E5	proprietary	CD226-DNAM-1.1	399907	BioLegend
CD158f (KIR2DL5)	AAAGTGATGCCACTG	A0600	UP-R1	proprietary	CD186-CXCR6	399907	BioLegend
CD158e1 (KIR3DL1, NKB1)	GGACGCTTCTCTGA	A0599	DX9	proprietary	CD158f-KIR2DL5	399907	BioLegend
CD140a (PDGFR?)	ATGCGCCGAGAATT	A0128	16A1	proprietary	CD158e1-KIR3DL1-NKB1	399907	BioLegend
CD354 (TREM-1)	TAGCCGTTCTCTTG	A0586	TREM-26	proprietary	CD140a	399907	BioLegend
TCR Vδ2	TCAGTCAGATGTTGAT	A0582	B6	proprietary	CD140b	399907	BioLegend
CD202b (Tie2/Tek)	CGATGCCCTAACCTAT	A0588	33.1 (Ab33)	proprietary	TCR-V-D-2	399907	BioLegend
CD96 (TACTILE)	TGGCCCTATAAACGGT	A0175	NK92.39	proprietary	CD202b-Tie2-Tek	399907	BioLegend
CD158 (KIR2DL1-KIR2DS1-)				proprietary	CD96-TACTILE	399907	BioLegend
CD158b (KIR2DL1-KIR2DS1-)				proprietary	CD96-KIR2DL1-KIR2DS1-		
CD158 (KIR2DL1/S1/S3/S5)	TATCAACCAACGCTT	A0420	HP-MA4	proprietary	KIR2DS3-KIR2DS5	399907	BioLegend
CD337 (NKP30)	AAAAGTCCTCTGCG	A0801	P30-15	proprietary	CD337-NKP30	399907	BioLegend
CD253 (Trail)	GCCATTCTCTGCTAA	A0803	RIK-2	proprietary	CD253-TRAIL	399907	BioLegend
CD319 (CRACK)	AGTATGCCATGCTT	A0830	162.1	proprietary	CD319-CRACK	399907	BioLegend
CD305 (LAIR1)	ATTTCCTATCCCTGT	A0590	NKTA255	proprietary	CD305-LAIR1	399907	BioLegend
CD325 (N-Cadherin)	CCTCTCTCTCTCT	A0433	8C11	proprietary	CD325-N-Cadherin	399907	BioLegend
mast cell tryptase	ACTGATAGACCCGCT	A0580	AA1	proprietary	N/A	399907	BioLegend
CLEC12A	CATTAGAGCTGCGCA	A0853	50C1	proprietary	CD325-N-Cadherin	399907	BioLegend
CD90 (Thy1)	GCATTGATACGATTCA	A0060	5E10	proprietary	CD325-N-Cadherin	399907	BioLegend
CD273 (B7-DC, PD-L2)	TCAACGCTTGGCTAG	A0008	24F.10C12	proprietary	CD327-B7-DC-PD-L2	399907	BioLegend
CD272 (BTLA)	TTTATGGACTAAGG	A0170	MIH26	proprietary	CD272-BTLA	399907	BioLegend
CD5	CATTAACGGGATGCC	A0138	UCHT2	proprietary	CD5.1	399907	BioLegend
CD23	TCTGTATAACCGCT	A0897	EBVCS-5	proprietary	CD23	399907	BioLegend
CD85j (ILT2)	CCTTGTAGGCTATG	A0896	GH1/75	proprietary	CD85j-ILT2	399907	BioLegend
CD94	CTTTCGGCTCTTACA	A0867	DX22	proprietary	CD94	399907	BioLegend
CD27	GCACCTCTGCATGTA	A0154	O323	proprietary	CD27.1	399907	BioLegend
CD328 (Siglec-7)	CTTAGCATTCTCTG	A0902	6-434	proprietary	CD328-Siglec-7	399907	BioLegend
CD82	TCCCACCTCCGCTT	A0920	ASL-24	proprietary	N/A	399907	BioLegend
CD16	AAGTTCACTCTTGC	A0083	3G8	proprietary	CD16	399907	BioLegend
MERTK	TCCCTGATGACCCCA	A0423	590H1G1E3	proprietary	MERTK.1	399907	BioLegend

IgG Fc	CTGGAGCGATTAGAA	A0375	M1310G05	proprietary	IgG-Fc	399907	BioLegend
FcεRⅠα	CTCGTTCCGTATCG	A0352	AER-37 (CRA-1)	proprietary	FcεRⅠα	399907	BioLegend
CD274 (B7-H1, PD-L1)	GTTGTCCGACAATAC	A0007	29E.2A3	proprietary	CD274-B7-H1-PD-L1	399907	BioLegend
CD254 (TRANCE, RANKL)	TCCGTGTTAGTTGT	A0356	MH24	proprietary	CD254-TRANCE-RANKL	399907	BioLegend
CX3CR1	AGTATCGTCTCTGG	A0179	K0124E1	proprietary	CX3CR1.1	399907	BioLegend
CD137L (4-1BB Ligand)	ATTCGCCTTACGCCA	A0022	5F4	proprietary	CD137L-4-1BB-Ligand	399907	BioLegend
Siglec-8	CTTCTCCCTCACGCAA	A0199	7C9	proprietary	N/A	399907	BioLegend
CD45	89	Y	H130	1.5		304002	BioLegend
EpCAM	115	In	9C4	1.5		324202	BioLegend
CD7	139	La	M-T701	1.5		555359	BD Bioscience
CD15	140	Ce	W6D3	1.5		323002	BioLegend
CD3	141	Pr	UCHT1	0.4		300402	BioLegend
CD19	142	Nd	HIB19	0.4		302202	BioLegend
CD163	144	Nd	GH161	3		333602	BioLegend
CD4	145	Nd	RPA-T4	0.4		14-0049-82	Thermo
CD8a	146	Nd	RPA-T8	0.4		301002	BioLegend
CD11c	147	Sm	BU15	0.4		MA1-82142	Thermo
CD14	148	Nd	M5E2	1.5		301802	BioLegend
CD127	149	Sm	A019D5	1.5		351302	BioLegend
CD123	151	Eu	6H6	0.8		14-1239-82	Thermo
gdTCR	152	Sm	5A6.E9	0.4		HB-9772	ATCC
CD45RA	153	Eu	HI100	1.5		304102	BioLegend
TIM3	154	Sm	F38-2E2	3		345002	BioLegend
TIGIT	155	Gd	A15153G	1.5		372720	BioLegend
PD-L1	156	Gd	29E.2A3	1.5		329702	BioLegend
CyTOF	Integrin a4	157	Gd	9F10	1.5	304302	BioLegend
	CD27	158	Gd	O323	0.4	302802	BioLegend
CD137L-4-1BB	CD137L-4-1BB	159	Tb	4B4-1	1.5	14-9056-82	Thermo
	Tbet	160	Gd	4B10	0.4	14-5825-82	Thermo
	CTLA-4	161	Dy	14D3	0.4	14-1529-82	Thermo
	FOXP3	162	Dy	PCH101	0.8	14-4776-82	Thermo
	CD31	163	Dy	WM59	0.8	303102	BioLegend
	Integrin b7	165	Ho	FIB504	1.5	321202	BioLegend
	CD141_BDCA3	166	Er	M80	0.8	344102	BioLegend
	CCR7	167	Er	GO43H7	1.5	353202	BioLegend
	KI-67	168	Er	KI-67	0.8	350502	BioLegend
	CD25	169	Tm	MA251	0.4	356102	BioLegend
CD1c_BDCA1	CD1c_BDCA1	170	Er	L161	0.8	331502	BioLegend
	CD38	172	Yb	HIT2	0.4	303502	BioLegend
	ICOS	173	Yb	C398.4A	0.8	313502	BioLegend
	HLA-DR	174	Yb	L243	1.5	307665	BioLegend
	PD-1	175	Lu	EH12.2H7	1.5	329902	BioLegend
	CD56	176	Yb	NCAM16.2	0.8	559043	BD Bioscience
	CD16	209	Bi	3G8	0.4	302049	BioLegend
	COL1A2	488		432721-T1			ACD bio-tecne
	MS4A1	550		426771-T2			ACD bio-tecne
	FLT3	650		419491-T3			ACD bio-tecne
RNA-ISH	ITG α 4	488		408591-T4			ACD bio-tecne
	MZB1	550		579241-T5			ACD bio-tecne
	C1Q4	650		485451-T6			ACD bio-tecne
	ITG β 7	488		605941-T7			ACD bio-tecne
	MAjCAM-1	550		404711-T8			ACD bio-tecne
	KRT8	650		310261-T9			ACD bio-tecne
	PECAM-1	488		487381-T10			ACD bio-tecne
	CD3E	550		553971-T11			ACD bio-tecne
	PTPRC	650		601991-T12			ACD bio-tecne
	HLA-A	AF750	BX004	EP1395Y	1:200	ab216653	Abcam
CODEX	CD34	ATT0550	BX020	QBEND/10	1:150	MA1-10205	Thermofisher
	CD4	Cy5	BX003	EPR6855	1:200	ab133616	Abcam
	CD20	AF750	BX007	L26	1:200	14-0202-82	Thermofisher
	CD14	ATT0550	BX037	EPR3653	1:500	ab133335	Abcam
	CD68	Cy5	BX015	KP1	1:200	MA5-13324	Thermofisher
	Vimentin	AF750	BX022	091D3	1:200	677802	BioLegend
	CD8	ATT0550	BX026	CB1/144B	1:200	372908	BioLegend
	CD11c	Cy5	BX024	118A5	1:200	14-9761-82	Thermofisher
	CD31	AF750	BX001	EP3095	1:100	ab134168	Abcam
	E-Cadherin	ATT0550	BX014	4A2C7	1:300	33-4000	Thermofisher
CODEX	CD45	Cy5	BX021	D9M8I	1:300	#13917	CST
	SMA	AF750	BX013	1A4	1:200	ab7817	Abcam
	CD45RO	ATT0550	BX017	UCHL1	1:200	304202	BioLegend
	CD3e	Cy5	BX045	EPR449E	1:200	ab52959	Abcam
	Pan-cytokeratin	AF750	BX019	AE1/AE3	1:200	914204	BioLegend
	CD44	ATT0550	BX005	156-3C11	1:400	#3570	CST
	HLA-DR	Cy5	BX033	EPR3692	1:200	ab92511	Abcam
	Granzyme B	ATT0550	BX041	D6E9W	1:50	#46890	CST
	Collagen IV	Cy5	BX042	EPR209660	1:200	ab214417	Abcam
	Podoplanin	ATT0550	BX023	NC-08	1:200	337002	BioLegend
E-Cadherin	FOXP3	AF647	BX031	259D/C7	1:100	560044	BD Pharmigen
	KI67	ATT0550	BX047	B56	1:300	556003	BD Pharmigen
	CD163	AF647	BX016	D6U1J	1:50	#93498	CST
	CD19	ATT0550	BX028	RM32	1:50	31-1219-00	RevMab
	CD11b	Cy5	BX030	EP1345Y	1:50	ab52478	Abcam
	CD21	ATT0550	BX032	EP3093	1:400	ab75985	Abcam
	CD57	AF647	BX049	HNK-1	1:100	359610	BioLegend
E-Cadherin	dsDNA	89	Y	3519 DNA	0.2	ab27156	Abcam
	Histone H3	113	In	36	0.5	ab287971	Abcam
	Foxp3-ΔF488	115	In	D1H2	1:100	711501-100	Ionpath
	anti-488	141	Pr	236A/E7	2	561181	BD Pharmigen
	CD163	142	Nd	EPR14643-36	0.6	#A-11094	Thermofisher
	CD4	143	Nd	EPR6855	1:100	ab215976	Abcam
						714301-100	Ionpath

CD11c	144	Nd	EP1347Y	1:100	714401-100	Ionpath	
CD14	145	Nd	D7A2T	0.6	#43878	CST	
CD16	146	Nd	D1N9L	0.6	#72204	CST	
LAG3	147	Sm	17B4	5	NBP1-97657	Novus	
PD-1	148	Nd	D4W2J	1:100	714801-100	Ionpath	
PD-L1	149	Sm	E1L3N	1:100	714902-100	Ionpath	
Granzyme B	150	Nd	D6E9W	1:100	715002-100	Ionpath	
CD56	151	Eu	MRQ-42	1:100	715101-100	Ionpath	
CD31	152	Sm	EP3095	1:100	715201-100	Ionpath	
Ki67	153	Eu	D2H10	0.15	#44092	CST	
CD138	154	Sm	EPR6454	5	ab2261108	Abcam	
CD117	155	Gd	YR145	1:100	715501-100	Ionpath	
MIBI	156	Gd	D4B9C	1:100	715601-100	Ionpath	
CD68	157	Gd	EPR4166(2)	15	ab271889	Abcam	
CD103	158	Gd	C8/144B	1:100	715801-100	Ionpath	
CD3	159	Tb	D7A6E	1:100	715903-100	Ionpath	
Tbet	160	Gd	D6N8B	5	#27112	CST	
CD45RO	161	Dy	UCHL1	1:100	716101-100	ionpath	
TIM3	162	Dy	EPR22241	1:100	716201-100	Ionpath	
Vimentin	163	Dy	D21H3	1:100	716301-100	Ionpath	
CD27	164	Dy	EPR8569	5	ab256583	Abcam	
Pan-Keratin	165	Ho	AE1/AE3	1:100	716501-100	Ionpath	
aSMA	166	Er	SP171	2.5	ab242395	Abcam	
CD20	167	Er	L26	1:100	716701-100	Abcam	
CD11b	168	Er	D6X1N	0.5	ab187537	Abcam	
BDC43 (CD141)	169	Tm	E7Y9P	10	#34149	CST	
CD21	170	Er	EP3093	1:100	717001-100	Ionpath	
IDO1	171	Yb	EPR20374	1:100	717101-100	Ionpath	
HLA-DR	172	Yb	EPR3692	1:100	717201-100	Ionpath	
EpCAM	173	Yb	D9S3P	0.5	#55725	CST	
CD45	175	Yb	2B11/PD7/26	1:100	717501-100	Ionpath	
HLA Class 1, ABC	176	Yb	EMR8-5	1:100	717602-100	Ionpath	
CosMx	AATK					Nanostring	
SMI 960-	ABL1					Nanostring	
plex panel	ABL2					Nanostring	
	ACE					Nanostring	
	ACE2					Nanostring	
	ACKR1					Nanostring	
	ACKR3					Nanostring	
	ACKR4					Nanostring	
	ACTA2					Nanostring	
	ACTG2					Nanostring	
	ACVR1					Nanostring	
	ACVR1B					Nanostring	
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	ACVRL1					Nanostring	
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	ADGRE5					Nanostring	
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	ADIPOQ					Nanostring	
	ADIRF					Nanostring	
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	AH1					Nanostring	
	AHR					Nanostring	
	AKT1					Nanostring	
	ALCAM					Nanostring	
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	APOB					Nanostring	
	APOD					Nanostring	
	APP					Nanostring	
	AQP3					Nanostring	
	AR					Nanostring	
	AREG					Nanostring	
	ARF1					Nanostring	

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<i>FABP5</i>	Nanostring
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<i>FASN</i>	Nanostring
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<i>FCGBP</i>	Nanostring
<i>FCGR3A</i>	Nanostring
<i>FCRLA</i>	Nanostring
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<i>FGF7</i>	Nanostring
<i>FGF9</i>	Nanostring
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<i>FGFR2</i>	Nanostring
<i>FGFR3</i>	Nanostring
<i>FGG</i>	Nanostring
<i>FGR</i>	Nanostring
<i>FKBP11</i>	Nanostring
<i>FLT1</i>	Nanostring
<i>FLT3LG</i>	Nanostring
<i>FN1</i>	Nanostring
<i>FOS</i>	Nanostring
<i>FOXF1</i>	Nanostring
<i>FOXP3</i>	Nanostring
<i>FPR1</i>	Nanostring
<i>FYB1</i>	Nanostring
<i>FYN</i>	Nanostring
<i>FZD1</i>	Nanostring
<i>FZD3</i>	Nanostring
<i>FZD4</i>	Nanostring
<i>FZD5</i>	Nanostring
<i>FZD6</i>	Nanostring
<i>FZD7</i>	Nanostring
<i>FZD8</i>	Nanostring
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<i>G6PD</i>	Nanostring
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<i>GNLY</i>	Nanostring
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<i>GPER1</i>	Nanostring
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<i>HBA1</i>	Nanostring
<i>HBB</i>	Nanostring
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<i>HLA-DQA1</i>	Nanostring
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<i>HSP90B1</i>	Nanostring
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<i>KRT6C</i>	Nanostring
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<i>LPARS</i>	Nanostring
<i>LTB</i>	Nanostring
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<i>LUM</i>	Nanostring
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<i>MPO</i>	Nanostring
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<i>MX1</i>	Nanostring
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<i>PDGFB</i>	Nanostring
<i>PDGFC</i>	Nanostring
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<i>PDGFRB</i>	Nanostring
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<i>PPBP</i>	Nanostring
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<i>RGCC</i>	Nanostring
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	<i>TNF</i>	Nanostring
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	<i>TNFRSF10A</i>	Nanostring
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	<i>TNFRSF11B</i>	Nanostring
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	<i>TNFSF10</i>	Nanostring
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	<i>TNFSF4</i>	Nanostring
	<i>TNFSF8</i>	Nanostring
	<i>TNFSF9</i>	Nanostring
	<i>TOP2A</i>	Nanostring
	<i>TOX</i>	Nanostring
	<i>TP53</i>	Nanostring
	<i>TPM1</i>	Nanostring
	<i>TPM2</i>	Nanostring
	<i>TPSAB1</i>	Nanostring
	<i>TPSB2</i>	Nanostring
	<i>TSC2D1</i>	Nanostring
	<i>TSHZ2</i>	Nanostring
	<i>TSLP</i>	Nanostring
	<i>TTR</i>	Nanostring
	<i>TUBB</i>	Nanostring
	<i>TUBB4B</i>	Nanostring
	<i>TWIST1</i>	Nanostring
	<i>TWIST2</i>	Nanostring
	<i>TXK</i>	Nanostring
	<i>TYK2</i>	Nanostring
	<i>TYMS</i>	Nanostring
	<i>TYROBP</i>	Nanostring
	<i>UBE2C</i>	Nanostring
	<i>UCP1</i>	Nanostring
	<i>UPK3A</i>	Nanostring
	<i>VCAM1</i>	Nanostring
	<i>VCAN</i>	Nanostring
	<i>VEGFA</i>	Nanostring
	<i>VEGFB</i>	Nanostring
	<i>VEGFC</i>	Nanostring
	<i>VEGFD</i>	Nanostring
	<i>VHL</i>	Nanostring
	<i>VIM</i>	Nanostring
	<i>VPREB3</i>	Nanostring
	<i>VSIR</i>	Nanostring
	<i>VTN</i>	Nanostring
	<i>VWF</i>	Nanostring
	<i>WIF1</i>	Nanostring
	<i>WNT10B</i>	Nanostring
	<i>WNT11</i>	Nanostring
	<i>WNT2</i>	Nanostring
	<i>WNT2B</i>	Nanostring
	<i>WNT3</i>	Nanostring
	<i>WNT5A</i>	Nanostring
	<i>WNT5B</i>	Nanostring
	<i>WNT7A</i>	Nanostring
	<i>WNT7B</i>	Nanostring
	<i>WNT9A</i>	Nanostring
	<i>XBP1</i>	Nanostring
	<i>XCL1</i>	Nanostring
	<i>XCL2</i>	Nanostring
	<i>YBX3</i>	Nanostring
	<i>YES1</i>	Nanostring
	<i>ZFP36</i>	Nanostring
CosMx	<i>AATK</i>	Nanostring
SMI 1000-	<i>ABL1</i>	Nanostring
plex panel	<i>ABL2</i>	Nanostring
	<i>ACACB</i>	Nanostring
	<i>ACE</i>	Nanostring
	<i>ACKR1</i>	Nanostring
	<i>ACKR3</i>	Nanostring
	<i>ACKR4</i>	Nanostring
	<i>ACP5</i>	Nanostring

<i>ACTA2</i>	Nanostring
<i>ACTG2</i>	Nanostring
<i>ACVR1</i>	Nanostring
<i>ACVR1B</i>	Nanostring
<i>ACVR2A</i>	Nanostring
<i>ACVRL1</i>	Nanostring
<i>ADGRA2</i>	Nanostring
<i>ADGRA3</i>	Nanostring
<i>ADGRE2</i>	Nanostring
<i>ADGRES</i>	Nanostring
<i>ADGRF1</i>	Nanostring
<i>ADGRF3</i>	Nanostring
<i>ADGRF5</i>	Nanostring
<i>ADGRG1</i>	Nanostring
<i>ADGRG3</i>	Nanostring
<i>ADGRG5</i>	Nanostring
<i>ADGRG6</i>	Nanostring
<i>ADGRL1</i>	Nanostring
<i>ADGRL2</i>	Nanostring
<i>ADGRL4</i>	Nanostring
<i>ADGRV1</i>	Nanostring
<i>ADIPOQ</i>	Nanostring
<i>ADIRF</i>	Nanostring
<i>ADM2</i>	Nanostring
<i>AGR2</i>	Nanostring
<i>AH1</i>	Nanostring
<i>AHR</i>	Nanostring
<i>AIF1</i>	Nanostring
<i>AKT1</i>	Nanostring
<i>ALCAM</i>	Nanostring
<i>ALOX5AP</i>	Nanostring
<i>ANGPT1</i>	Nanostring
<i>ANGPT2</i>	Nanostring
<i>ANGPTL1</i>	Nanostring
<i>ANKRD1</i>	Nanostring
<i>ANXA1</i>	Nanostring
<i>ANXA2</i>	Nanostring
<i>ANXA4</i>	Nanostring
<i>APOA1</i>	Nanostring
<i>APOC1</i>	Nanostring
<i>APOD</i>	Nanostring
<i>APOE</i>	Nanostring
<i>APP</i>	Nanostring
<i>AQP3</i>	Nanostring
<i>AR</i>	Nanostring
<i>AREG</i>	Nanostring
<i>ARF1</i>	Nanostring
<i>ARG1</i>	Nanostring
<i>ARHGDI1B</i>	Nanostring
<i>ARID5B</i>	Nanostring
<i>ATF3</i>	Nanostring
<i>ATG10</i>	Nanostring
<i>ATG12</i>	Nanostring
<i>ATG5</i>	Nanostring
<i>ATM</i>	Nanostring
<i>ATP5F1B</i>	Nanostring
<i>ATP5F1E</i>	Nanostring
<i>ATR</i>	Nanostring
<i>AXL</i>	Nanostring
<i>AZGP1</i>	Nanostring
<i>AZU1</i>	Nanostring
<i>B2M</i>	Nanostring
<i>B3GNT7</i>	Nanostring
<i>BAG3</i>	Nanostring
<i>BASP1</i>	Nanostring
<i>BAX</i>	Nanostring
<i>BBLN</i>	Nanostring
<i>BCL2</i>	Nanostring
<i>BCL2L1</i>	Nanostring
<i>BECN1</i>	Nanostring
<i>BEST1</i>	Nanostring
<i>BGN</i>	Nanostring
<i>BID</i>	Nanostring
<i>BIRC3</i>	Nanostring
<i>BIRC5</i>	Nanostring
<i>BMP1</i>	Nanostring
<i>BMP2</i>	Nanostring
<i>BMP3</i>	Nanostring
<i>BMP4</i>	Nanostring
<i>BMP5</i>	Nanostring
<i>BMP7</i>	Nanostring
<i>BMPR1A</i>	Nanostring
<i>BMPR2</i>	Nanostring
<i>BRAF</i>	Nanostring
<i>BRCA1</i>	Nanostring
<i>BST1</i>	Nanostring
<i>BST2</i>	Nanostring
<i>BTF3</i>	Nanostring
<i>BTG1</i>	Nanostring
<i>BTK</i>	Nanostring
<i>C11orf96</i>	Nanostring

<i>C1QA</i>	Nanostring
<i>C1QB</i>	Nanostring
<i>C1QC</i>	Nanostring
<i>CSAR2</i>	Nanostring
<i>CACNA1C</i>	Nanostring
<i>CALB1</i>	Nanostring
<i>CALD1</i>	Nanostring
<i>CALM1</i>	Nanostring
<i>CALM2</i>	Nanostring
<i>CALM3</i>	Nanostring
<i>CAMP</i>	Nanostring
<i>CARMN</i>	Nanostring
<i>CASP3</i>	Nanostring
<i>CASP8</i>	Nanostring
<i>CASR</i>	Nanostring
<i>CAV1</i>	Nanostring
<i>CCDC80</i>	Nanostring
<i>CCL11</i>	Nanostring
<i>CCL13</i>	Nanostring
<i>CCL15</i>	Nanostring
<i>CCL17</i>	Nanostring
<i>CCL18</i>	Nanostring
<i>CCL19</i>	Nanostring
<i>CCL2</i>	Nanostring
<i>CCL20</i>	Nanostring
<i>CCL21</i>	Nanostring
<i>CCL22</i>	Nanostring
<i>CCL26</i>	Nanostring
<i>CCL28</i>	Nanostring
<i>CCL3/L1/L3</i>	Nanostring
<i>CCL4/L1/L2</i>	Nanostring
<i>CCL5</i>	Nanostring
<i>CCL8</i>	Nanostring
<i>CCND1</i>	Nanostring
<i>CCR1</i>	Nanostring
<i>CCR10</i>	Nanostring
<i>CCR2</i>	Nanostring
<i>CCR5</i>	Nanostring
<i>CCR7</i>	Nanostring
<i>CCL2</i>	Nanostring
<i>CD14</i>	Nanostring
<i>CD163</i>	Nanostring
<i>CD164</i>	Nanostring
<i>CD19</i>	Nanostring
<i>CD1C</i>	Nanostring
<i>CD2</i>	Nanostring
<i>CD209</i>	Nanostring
<i>CD22</i>	Nanostring
<i>CD24</i>	Nanostring
<i>CD27</i>	Nanostring
<i>CD274</i>	Nanostring
<i>CD276</i>	Nanostring
<i>CD28</i>	Nanostring
<i>CD300A</i>	Nanostring
<i>CD33</i>	Nanostring
<i>CD34</i>	Nanostring
<i>CD36</i>	Nanostring
<i>CD37</i>	Nanostring
<i>CD38</i>	Nanostring
<i>CD3D</i>	Nanostring
<i>CD3E</i>	Nanostring
<i>CD3G</i>	Nanostring
<i>CD4</i>	Nanostring
<i>CD40</i>	Nanostring
<i>CD40LG</i>	Nanostring
<i>CD44</i>	Nanostring
<i>CD47</i>	Nanostring
<i>CD48</i>	Nanostring
<i>CD52</i>	Nanostring
<i>CD53</i>	Nanostring
<i>CD55</i>	Nanostring
<i>CD58</i>	Nanostring
<i>CD59</i>	Nanostring
<i>CD5L</i>	Nanostring
<i>CD63</i>	Nanostring
<i>CD68</i>	Nanostring
<i>CD69</i>	Nanostring
<i>CD70</i>	Nanostring
<i>CD74</i>	Nanostring
<i>CD79A</i>	Nanostring
<i>CD80</i>	Nanostring
<i>CD81</i>	Nanostring
<i>CD83</i>	Nanostring
<i>CD84</i>	Nanostring
<i>CD86</i>	Nanostring
<i>CD8A</i>	Nanostring
<i>CD8B</i>	Nanostring
<i>CD9</i>	Nanostring
<i>CD93</i>	Nanostring
<i>CDH1</i>	Nanostring
<i>CDH11</i>	Nanostring

<i>CDH19</i>	Nanostring
<i>CDH5</i>	Nanostring
<i>CDKN1A</i>	Nanostring
<i>CDKN3</i>	Nanostring
<i>CEACAM1</i>	Nanostring
<i>CEACAM6</i>	Nanostring
<i>CELSR1</i>	Nanostring
<i>CELSR2</i>	Nanostring
<i>CENPF</i>	Nanostring
<i>CFD</i>	Nanostring
<i>CFLAR</i>	Nanostring
<i>CHEK1</i>	Nanostring
<i>CHEK2</i>	Nanostring
<i>CHI3L1</i>	Nanostring
<i>CIDEA</i>	Nanostring
<i>CIITA</i>	Nanostring
<i>CLCF1</i>	Nanostring
<i>CLDN4</i>	Nanostring
<i>CLEC10A</i>	Nanostring
<i>CLEC12A</i>	Nanostring
<i>CLEC14A</i>	Nanostring
<i>CLEC1A</i>	Nanostring
<i>CLEC2B</i>	Nanostring
<i>CLEC2D</i>	Nanostring
<i>CLEC4A</i>	Nanostring
<i>CLEC4D</i>	Nanostring
<i>CLEC4E</i>	Nanostring
<i>CLEC5A</i>	Nanostring
<i>CLEC7A</i>	Nanostring
<i>CLOCK</i>	Nanostring
<i>CLU</i>	Nanostring
<i>CMKLR1</i>	Nanostring
<i>CNTFR</i>	Nanostring
<i>COL11A1</i>	Nanostring
<i>COL12A1</i>	Nanostring
<i>COL14A1</i>	Nanostring
<i>COL15A1</i>	Nanostring
<i>COL16A1</i>	Nanostring
<i>COL17A1</i>	Nanostring
<i>COL18A1</i>	Nanostring
<i>COL1A1</i>	Nanostring
<i>COL1A2</i>	Nanostring
<i>COL21A1</i>	Nanostring
<i>COL27A1</i>	Nanostring
<i>COL3A1</i>	Nanostring
<i>COL4A1</i>	Nanostring
<i>COL4A2</i>	Nanostring
<i>COL4A5</i>	Nanostring
<i>COL5A1</i>	Nanostring
<i>COL5A2</i>	Nanostring
<i>COL5A3</i>	Nanostring
<i>COL6A1</i>	Nanostring
<i>COL6A2</i>	Nanostring
<i>COL6A3</i>	Nanostring
<i>COL8A1</i>	Nanostring
<i>COL9A2</i>	Nanostring
<i>COL9A3</i>	Nanostring
<i>COTL1</i>	Nanostring
<i>COX4I2</i>	Nanostring
<i>CPA3</i>	Nanostring
<i>CPB1</i>	Nanostring
<i>CRIP1</i>	Nanostring
<i>CRP</i>	Nanostring
<i>CRYAB</i>	Nanostring
<i>CSF1</i>	Nanostring
<i>CSF1R</i>	Nanostring
<i>CSF2</i>	Nanostring
<i>CSF2RA</i>	Nanostring
<i>CSF2RB</i>	Nanostring
<i>CSF3</i>	Nanostring
<i>CSF3R</i>	Nanostring
<i>CSHL1</i>	Nanostring
<i>CSK</i>	Nanostring
<i>CSPG4</i>	Nanostring
<i>CT7</i>	Nanostring
<i>CSTB</i>	Nanostring
<i>CTL44</i>	Nanostring
<i>CTNNB1</i>	Nanostring
<i>CTSD</i>	Nanostring
<i>CTSG</i>	Nanostring
<i>CTSW</i>	Nanostring
<i>CUDZ1</i>	Nanostring
<i>CX3CL1</i>	Nanostring
<i>CX3CR1</i>	Nanostring
<i>CXCL1/2/3</i>	Nanostring
<i>CXCL10</i>	Nanostring
<i>CXCL12</i>	Nanostring
<i>CXCL13</i>	Nanostring
<i>CXCL14</i>	Nanostring
<i>CXCL16</i>	Nanostring
<i>CXCL17</i>	Nanostring

<i>CXCL5</i>	Nanostring
<i>CXCL8</i>	Nanostring
<i>CXCL9</i>	Nanostring
<i>CXCR1</i>	Nanostring
<i>CXCR2</i>	Nanostring
<i>CXCR3</i>	Nanostring
<i>CXCR4</i>	Nanostring
<i>CXCR5</i>	Nanostring
<i>CXCR6</i>	Nanostring
<i>CYP1B1</i>	Nanostring
<i>CYP2U1</i>	Nanostring
<i>CYSTM1</i>	Nanostring
<i>CYTOR</i>	Nanostring
<i>DCN</i>	Nanostring
<i>DDC</i>	Nanostring
<i>DDIT3</i>	Nanostring
<i>DDR1</i>	Nanostring
<i>DDR2</i>	Nanostring
<i>DDX58</i>	Nanostring
<i>DHRS2</i>	Nanostring
<i>DLL1</i>	Nanostring
<i>DLL4</i>	Nanostring
<i>DMBT1</i>	Nanostring
<i>DNMT1</i>	Nanostring
<i>DNMT3A</i>	Nanostring
<i>DPP4</i>	Nanostring
<i>DPT</i>	Nanostring
<i>DST</i>	Nanostring
<i>DUSP1</i>	Nanostring
<i>DUSP2</i>	Nanostring
<i>DUSP4</i>	Nanostring
<i>DUSP5</i>	Nanostring
<i>DUSP6</i>	Nanostring
<i>EFNA1</i>	Nanostring
<i>EFNA4</i>	Nanostring
<i>EFNA5</i>	Nanostring
<i>EFNB1</i>	Nanostring
<i>EFNB2</i>	Nanostring
<i>EGF</i>	Nanostring
<i>EGFR</i>	Nanostring
<i>EIF5A1/1</i>	Nanostring
<i>ELANE</i>	Nanostring
<i>EMP3</i>	Nanostring
<i>ENG</i>	Nanostring
<i>ENO1</i>	Nanostring
<i>ENTPD1</i>	Nanostring
<i>EOMES</i>	Nanostring
<i>EPCAM</i>	Nanostring
<i>EPHA2</i>	Nanostring
<i>EPHA3</i>	Nanostring
<i>EPHA4</i>	Nanostring
<i>EPHA7</i>	Nanostring
<i>EPHB2</i>	Nanostring
<i>EPHB3</i>	Nanostring
<i>EPHB4</i>	Nanostring
<i>EPHB6</i>	Nanostring
<i>EPOR</i>	Nanostring
<i>ERBB2</i>	Nanostring
<i>ERBB3</i>	Nanostring
<i>ESAM</i>	Nanostring
<i>ESR1</i>	Nanostring
<i>ETS1</i>	Nanostring
<i>ETV4</i>	Nanostring
<i>ETV5</i>	Nanostring
<i>EZH2</i>	Nanostring
<i>EZR</i>	Nanostring
<i>FABP4</i>	Nanostring
<i>FABP5</i>	Nanostring
<i>FAM30A</i>	Nanostring
<i>FAP</i>	Nanostring
<i>FAS</i>	Nanostring
<i>FASLG</i>	Nanostring
<i>FASN</i>	Nanostring
<i>FAU</i>	Nanostring
<i>FCER1G</i>	Nanostring
<i>FCGBP</i>	Nanostring
<i>FCGR3A/B</i>	Nanostring
<i>FCRLA</i>	Nanostring
<i>FES</i>	Nanostring
<i>FFAR2</i>	Nanostring
<i>FFAR3</i>	Nanostring
<i>FFAR4</i>	Nanostring
<i>FGF1</i>	Nanostring
<i>FGF12</i>	Nanostring
<i>FGF13</i>	Nanostring
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<i>FGF2</i>	Nanostring
<i>FGF7</i>	Nanostring
<i>FGF9</i>	Nanostring
<i>FGFR1</i>	Nanostring
<i>FGFR1</i>	Nanostring

<i>FGFR3</i>	Nanostring
<i>FGG</i>	Nanostring
<i>FGR</i>	Nanostring
<i>FHIT</i>	Nanostring
<i>FKBP11</i>	Nanostring
<i>FKBP5</i>	Nanostring
<i>FLT1</i>	Nanostring
<i>FLT3LG</i>	Nanostring
<i>FN1</i>	Nanostring
<i>FOS</i>	Nanostring
<i>FOXF1</i>	Nanostring
<i>FOXP3</i>	Nanostring
<i>FPR1</i>	Nanostring
<i>FYB1</i>	Nanostring
<i>FYN</i>	Nanostring
<i>FZD1</i>	Nanostring
<i>FZD3</i>	Nanostring
<i>FZD4</i>	Nanostring
<i>FZD5</i>	Nanostring
<i>FZD6</i>	Nanostring
<i>FZD7</i>	Nanostring
<i>FZD8</i>	Nanostring
<i>G0S2</i>	Nanostring
<i>G6PD</i>	Nanostring
<i>GADD45B</i>	Nanostring
<i>GAS6</i>	Nanostring
<i>GATA3</i>	Nanostring
<i>GC</i>	Nanostring
<i>GCG</i>	Nanostring
<i>GDF15</i>	Nanostring
<i>GLUD1</i>	Nanostring
<i>GLUL</i>	Nanostring
<i>GNLY</i>	Nanostring
<i>GPBAR1</i>	Nanostring
<i>GPER1</i>	Nanostring
<i>GPNMB</i>	Nanostring
<i>GPR183</i>	Nanostring
<i>GPX1</i>	Nanostring
<i>GPX3</i>	Nanostring
<i>GSK3B</i>	Nanostring
<i>GSN</i>	Nanostring
<i>GSTP1</i>	Nanostring
<i>GZMA</i>	Nanostring
<i>GZMB</i>	Nanostring
<i>GZMH</i>	Nanostring
<i>GZMK</i>	Nanostring
<i>H2A.Z1</i>	Nanostring
<i>H4C3</i>	Nanostring
<i>HAVCR2</i>	Nanostring
<i>HBA1/2</i>	Nanostring
<i>HBB</i>	Nanostring
<i>HCAR2/3</i>	Nanostring
<i>HCK</i>	Nanostring
<i>HCST</i>	Nanostring
<i>HDAC1</i>	Nanostring
<i>HDAC11</i>	Nanostring
<i>HDAC3</i>	Nanostring
<i>HDAC4</i>	Nanostring
<i>HDAC5</i>	Nanostring
<i>HEXB</i>	Nanostring
<i>HEY1</i>	Nanostring
<i>HGF</i>	Nanostring
<i>HIF1A</i>	Nanostring
<i>HILDA</i>	Nanostring
<i>HLA-DPA1</i>	Nanostring
<i>HLA-DPB1</i>	Nanostring
<i>HLA-DQA1</i>	Nanostring
<i>HLA-DQB1/2</i>	Nanostring
<i>HLA-DRA</i>	Nanostring
<i>HLA-DRB</i>	Nanostring
<i>HMGFB2</i>	Nanostring
<i>HMGCS1</i>	Nanostring
<i>HMGN2</i>	Nanostring
<i>HPGDS</i>	Nanostring
<i>HSD17B2</i>	Nanostring
<i>HSP90AA1</i>	Nanostring
<i>HSP90AB1</i>	Nanostring
<i>HSP90B1</i>	Nanostring
<i>HSPA1A/B</i>	Nanostring
<i>HSPB1</i>	Nanostring
<i>HSPG2</i>	Nanostring
<i>HTT</i>	Nanostring
<i>IAPP</i>	Nanostring
<i>ICA1</i>	Nanostring
<i>ICAM1</i>	Nanostring
<i>ICAM2</i>	Nanostring
<i>ICAM3</i>	Nanostring
<i>ICOS</i>	Nanostring
<i>ICOSLG</i>	Nanostring
<i>IDO1</i>	Nanostring
<i>IER3</i>	Nanostring

<i>IFI27</i>	Nanostring
<i>IFI44L</i>	Nanostring
<i>IFI6</i>	Nanostring
<i>IFIH1</i>	Nanostring
<i>IFIT1</i>	Nanostring
<i>IFIT3</i>	Nanostring
<i>IFITM1</i>	Nanostring
<i>IFITM3</i>	Nanostring
<i>IFNA1/13</i>	Nanostring
<i>IFNAR1</i>	Nanostring
<i>IFNAR2</i>	Nanostring
<i>IFNG</i>	Nanostring
<i>IFNGR1</i>	Nanostring
<i>IFNGR2</i>	Nanostring
<i>IFNL2/3</i>	Nanostring
<i>IGF1</i>	Nanostring
<i>IGF1R</i>	Nanostring
<i>IGF2</i>	Nanostring
<i>IGF2R</i>	Nanostring
<i>IGFBP3</i>	Nanostring
<i>IGFBP5</i>	Nanostring
<i>IGFBP6</i>	Nanostring
<i>IGFBP7</i>	Nanostring
<i>IGHA1</i>	Nanostring
<i>IGHD</i>	Nanostring
<i>IGHG1</i>	Nanostring
<i>IGHG2</i>	Nanostring
<i>IGHM</i>	Nanostring
<i>IGKC</i>	Nanostring
<i>IKZF3</i>	Nanostring
<i>IL10</i>	Nanostring
<i>IL10RA</i>	Nanostring
<i>IL10RB</i>	Nanostring
<i>IL11</i>	Nanostring
<i>IL11RA</i>	Nanostring
<i>IL12A</i>	Nanostring
<i>IL12B</i>	Nanostring
<i>IL12RB1</i>	Nanostring
<i>IL12RB2</i>	Nanostring
<i>IL13RA1</i>	Nanostring
<i>IL15</i>	Nanostring
<i>IL15RA</i>	Nanostring
<i>IL16</i>	Nanostring
<i>IL17A</i>	Nanostring
<i>IL17B</i>	Nanostring
<i>IL17D</i>	Nanostring
<i>IL17RA</i>	Nanostring
<i>IL17RB</i>	Nanostring
<i>IL17RE</i>	Nanostring
<i>IL18</i>	Nanostring
<i>IL18R1</i>	Nanostring
<i>IL1A</i>	Nanostring
<i>IL1B</i>	Nanostring
<i>IL1R1</i>	Nanostring
<i>IL1R2</i>	Nanostring
<i>IL1RAP</i>	Nanostring
<i>IL1RL1</i>	Nanostring
<i>IL1RN</i>	Nanostring
<i>IL2</i>	Nanostring
<i>IL20</i>	Nanostring
<i>IL20RA</i>	Nanostring
<i>IL22RA1</i>	Nanostring
<i>IL23A</i>	Nanostring
<i>IL24</i>	Nanostring
<i>IL27RA</i>	Nanostring
<i>IL2RA</i>	Nanostring
<i>IL2RB</i>	Nanostring
<i>IL2RG</i>	Nanostring
<i>IL32</i>	Nanostring
<i>IL33</i>	Nanostring
<i>IL34</i>	Nanostring
<i>IL36G</i>	Nanostring
<i>IL3RA</i>	Nanostring
<i>IL4R</i>	Nanostring
<i>IL6</i>	Nanostring
<i>IL6R</i>	Nanostring
<i>IL6ST</i>	Nanostring
<i>IL7</i>	Nanostring
<i>IL7R</i>	Nanostring
<i>INHA</i>	Nanostring
<i>INHBA</i>	Nanostring
<i>INHBB</i>	Nanostring
<i>INS</i>	Nanostring
<i>INSIG1</i>	Nanostring
<i>INSR</i>	Nanostring
<i>IRF3</i>	Nanostring
<i>IRF4</i>	Nanostring
<i>ISG15</i>	Nanostring
<i>ITGA1</i>	Nanostring
<i>ITGA2</i>	Nanostring
<i>ITGA3</i>	Nanostring

<i>ITGA5</i>	Nanostring
<i>ITGA6</i>	Nanostring
<i>ITGA8</i>	Nanostring
<i>ITGA9</i>	Nanostring
<i>ITGAE</i>	Nanostring
<i>ITGAL</i>	Nanostring
<i>ITGAM</i>	Nanostring
<i>ITGAV</i>	Nanostring
<i>ITGAX</i>	Nanostring
<i>ITGB1</i>	Nanostring
<i>ITGB2</i>	Nanostring
<i>ITGB4</i>	Nanostring
<i>ITGB5</i>	Nanostring
<i>ITGB6</i>	Nanostring
<i>ITGB8</i>	Nanostring
<i>ITK</i>	Nanostring
<i>ITM2A</i>	Nanostring
<i>ITM2B</i>	Nanostring
<i>JAG1</i>	Nanostring
<i>JAK1</i>	Nanostring
<i>JAK2</i>	Nanostring
<i>JCHAIN</i>	Nanostring
<i>JUN</i>	Nanostring
<i>JUNB</i>	Nanostring
<i>KDR</i>	Nanostring
<i>KIT</i>	Nanostring
<i>KITLG</i>	Nanostring
<i>KLF2</i>	Nanostring
<i>KLK3</i>	Nanostring
<i>KLRB1</i>	Nanostring
<i>KLRF1</i>	Nanostring
<i>KLRK1</i>	Nanostring
<i>KRAS</i>	Nanostring
<i>KRT1</i>	Nanostring
<i>KRT10</i>	Nanostring
<i>KRT13</i>	Nanostring
<i>KRT14</i>	Nanostring
<i>KRT15</i>	Nanostring
<i>KRT16</i>	Nanostring
<i>KRT17</i>	Nanostring
<i>KRT18</i>	Nanostring
<i>KRT19</i>	Nanostring
<i>KRT20</i>	Nanostring
<i>KRT23</i>	Nanostring
<i>KRT4</i>	Nanostring
<i>KRT5</i>	Nanostring
<i>KRT6A/B/C</i>	Nanostring
<i>KRT7</i>	Nanostring
<i>KRT8</i>	Nanostring
<i>KRT80</i>	Nanostring
<i>KRT86</i>	Nanostring
<i>LAG3</i>	Nanostring
<i>LAIR1</i>	Nanostring
<i>LAMA4</i>	Nanostring
<i>LAMP2</i>	Nanostring
<i>LAMP3</i>	Nanostring
<i>LCN2</i>	Nanostring
<i>LDB2</i>	Nanostring
<i>LDHA</i>	Nanostring
<i>LDLR</i>	Nanostring
<i>LEFTY1</i>	Nanostring
<i>LEP</i>	Nanostring
<i>LGALS1</i>	Nanostring
<i>LGALS3</i>	Nanostring
<i>LGALS3BP</i>	Nanostring
<i>LGALS9</i>	Nanostring
<i>LGR5</i>	Nanostring
<i>LIF</i>	Nanostring
<i>LIFR</i>	Nanostring
<i>LINC01781</i>	Nanostring
<i>LINC01857</i>	Nanostring
<i>LINC02446</i>	Nanostring
<i>LMNA</i>	Nanostring
<i>LPAR5</i>	Nanostring
<i>LTB</i>	Nanostring
<i>LTBR</i>	Nanostring
<i>LTF</i>	Nanostring
<i>LUM</i>	Nanostring
<i>LY6D</i>	Nanostring
<i>LY75</i>	Nanostring
<i>LYN</i>	Nanostring
<i>LYVE1</i>	Nanostring
<i>LYZ</i>	Nanostring
<i>MAF</i>	Nanostring
<i>MALAT1</i>	Nanostring
<i>MAML2</i>	Nanostring
<i>MAP1LC3B/2</i>	Nanostring
<i>MAP2K1</i>	Nanostring
<i>MAPK13</i>	Nanostring
<i>MAPK14</i>	Nanostring
<i>MARCKSL1</i>	Nanostring

<i>MARCO</i>	
<i>MB</i>	Nanostring
<i>MECOM</i>	Nanostring
<i>MEG3</i>	Nanostring
<i>MERTK</i>	Nanostring
<i>MET</i>	Nanostring
<i>MFAP5</i>	Nanostring
<i>MGP</i>	Nanostring
<i>MHC I</i>	Nanostring
<i>MFIF</i>	Nanostring
<i>MIR4435-2HG</i>	Nanostring
<i>MKI67</i>	Nanostring
<i>MMP1</i>	Nanostring
<i>MMP12</i>	Nanostring
<i>MMP14</i>	Nanostring
<i>MMP19</i>	Nanostring
<i>MMP2</i>	Nanostring
<i>MMP7</i>	Nanostring
<i>MMP9</i>	Nanostring
<i>MPO</i>	Nanostring
<i>MRC1</i>	Nanostring
<i>MRC2</i>	Nanostring
<i>MS4A1</i>	Nanostring
<i>MS4A4A</i>	Nanostring
<i>MS4A6A</i>	Nanostring
<i>MSMB</i>	Nanostring
<i>MSR1</i>	Nanostring
<i>MT1R</i>	Nanostring
<i>MT1X</i>	Nanostring
<i>MT2A</i>	Nanostring
<i>MTOR</i>	Nanostring
<i>MX1</i>	Nanostring
<i>MXRA8</i>	Nanostring
<i>MYC</i>	Nanostring
<i>MYH11</i>	Nanostring
<i>MYH6</i>	Nanostring
<i>MYL12A</i>	Nanostring
<i>MYL4</i>	Nanostring
<i>MYL7</i>	Nanostring
<i>MYL9</i>	Nanostring
<i>MZB1</i>	Nanostring
<i>MZT2A/B</i>	Nanostring
<i>NACA</i>	Nanostring
<i>NANOG</i>	Nanostring
<i>NCAM1</i>	Nanostring
<i>NCR1</i>	Nanostring
<i>NDRG1</i>	Nanostring
<i>NDUFA4L2</i>	Nanostring
<i>NEAT1</i>	Nanostring
<i>NEIL2</i>	Nanostring
<i>NFKB1</i>	Nanostring
<i>NFKBIA</i>	Nanostring
<i>NGFR</i>	Nanostring
<i>NKG7</i>	Nanostring
<i>NLRC4</i>	Nanostring
<i>NLRCS</i>	Nanostring
<i>NLRP1</i>	Nanostring
<i>NLRP2</i>	Nanostring
<i>NLRP3</i>	Nanostring
<i>NOD2</i>	Nanostring
<i>NOSIP</i>	Nanostring
<i>NOTCH1</i>	Nanostring
<i>NOTCH2</i>	Nanostring
<i>NOTCH3</i>	Nanostring
<i>NPPC</i>	Nanostring
<i>NPR1</i>	Nanostring
<i>NPR2</i>	Nanostring
<i>NPR3</i>	Nanostring
<i>NR1H2</i>	Nanostring
<i>NR1H3</i>	Nanostring
<i>NR2F2</i>	Nanostring
<i>NR3C1</i>	Nanostring
<i>NRG1</i>	Nanostring
<i>NRXN1</i>	Nanostring
<i>NRXN3</i>	Nanostring
<i>NTRK2</i>	Nanostring
<i>NUPR1</i>	Nanostring
<i>NUSAP1</i>	Nanostring
<i>OAS1</i>	Nanostring
<i>OAS2</i>	Nanostring
<i>OAS3</i>	Nanostring
<i>OASL</i>	Nanostring
<i>OLFM4</i>	Nanostring
<i>OLR1</i>	Nanostring
<i>OSM</i>	Nanostring
<i>OSMR</i>	Nanostring
<i>P2RX5</i>	Nanostring
<i>PARP1</i>	Nanostring
<i>PCNA</i>	Nanostring
<i>PDCD1</i>	Nanostring
<i>PDCD1LG2</i>	Nanostring

<i>PDGFA</i>	Nanostring
<i>PDGFB</i>	Nanostring
<i>PDGFC</i>	Nanostring
<i>PDGFD</i>	Nanostring
<i>PDGFRA</i>	Nanostring
<i>PDGFRB</i>	Nanostring
<i>POSSA</i>	Nanostring
<i>PECAM1</i>	Nanostring
<i>PF4/V1</i>	Nanostring
<i>PFN1</i>	Nanostring
<i>PGF</i>	Nanostring
<i>PGK1</i>	Nanostring
<i>PGR</i>	Nanostring
<i>PHLDA2</i>	Nanostring
<i>PIGR</i>	Nanostring
<i>PLAC8</i>	Nanostring
<i>PLAC9</i>	Nanostring
<i>PLCG1</i>	Nanostring
<i>PLD3</i>	Nanostring
<i>PNOC</i>	Nanostring
<i>POU5F1</i>	Nanostring
<i>PPARA</i>	Nanostring
<i>PPARD</i>	Nanostring
<i>PPARG</i>	Nanostring
<i>PPIA</i>	Nanostring
<i>PREF1</i>	Nanostring
<i>PROX1</i>	Nanostring
<i>PRSS2</i>	Nanostring
<i>PRTN3</i>	Nanostring
<i>PSAP</i>	Nanostring
<i>PSCA</i>	Nanostring
<i>PSD3</i>	Nanostring
<i>PTEN</i>	Nanostring
<i>PTGDR2</i>	Nanostring
<i>PTGDS</i>	Nanostring
<i>PTGES</i>	Nanostring
<i>PTGES2</i>	Nanostring
<i>PTGES3</i>	Nanostring
<i>PTGIS</i>	Nanostring
<i>PTGS1</i>	Nanostring
<i>PTGS2</i>	Nanostring
<i>PTK2</i>	Nanostring
<i>PTK6</i>	Nanostring
<i>PTPRC</i>	Nanostring
<i>PTPRCAP</i>	Nanostring
<i>PTTG1</i>	Nanostring
<i>PXDN</i>	Nanostring
<i>QRFPR</i>	Nanostring
<i>RAC1</i>	Nanostring
<i>RAC2</i>	Nanostring
<i>RACK1</i>	Nanostring
<i>RAG1</i>	Nanostring
<i>RAMP1</i>	Nanostring
<i>RAMP2</i>	Nanostring
<i>RAMP3</i>	Nanostring
<i>RARA</i>	Nanostring
<i>RARB</i>	Nanostring
<i>RARG</i>	Nanostring
<i>RARRES1</i>	Nanostring
<i>RARRES2</i>	Nanostring
<i>RB1</i>	Nanostring
<i>RBM47</i>	Nanostring
<i>RPBP1</i>	Nanostring
<i>REG1A</i>	Nanostring
<i>RELA</i>	Nanostring
<i>RELT</i>	Nanostring
<i>RGCC</i>	Nanostring
<i>RGS1</i>	Nanostring
<i>RGS13</i>	Nanostring
<i>RGS2</i>	Nanostring
<i>RGS5</i>	Nanostring
<i>RNF43</i>	Nanostring
<i>ROR1</i>	Nanostring
<i>RORA</i>	Nanostring
<i>RPL21</i>	Nanostring
<i>RPL22</i>	Nanostring
<i>RPL32</i>	Nanostring
<i>RPL34</i>	Nanostring
<i>RPL37</i>	Nanostring
<i>RPS4Y1</i>	Nanostring
<i>RSP03</i>	Nanostring
<i>RUNX3</i>	Nanostring
<i>RXRA</i>	Nanostring
<i>RXRΒ</i>	Nanostring
<i>RYK</i>	Nanostring
<i>RYR2</i>	Nanostring
<i>S100A10</i>	Nanostring
<i>S100A2</i>	Nanostring
<i>S100A4</i>	Nanostring
<i>S100A6</i>	Nanostring
<i>S100A8</i>	Nanostring

<i>S100A9</i>	Nanostring
<i>S100B</i>	Nanostring
<i>S100P</i>	Nanostring
<i>SAA1/2</i>	Nanostring
<i>SAT1</i>	Nanostring
<i>SCG5</i>	Nanostring
<i>SCGB3A1</i>	Nanostring
<i>SEC23A</i>	Nanostring
<i>SEC61G</i>	Nanostring
<i>SELENOP</i>	Nanostring
<i>SELL</i>	Nanostring
<i>SELPLG</i>	Nanostring
<i>SERPINA1</i>	Nanostring
<i>SERPINA3</i>	Nanostring
<i>SERPINB5</i>	Nanostring
<i>SERPINH1</i>	Nanostring
<i>SFN</i>	Nanostring
<i>SH3BGRL3</i>	Nanostring
<i>SIGIRR</i>	Nanostring
<i>SLA</i>	Nanostring
<i>SLC2A1</i>	Nanostring
<i>SLC40A1</i>	Nanostring
<i>SLCO2B1</i>	Nanostring
<i>SLPI</i>	Nanostring
<i>SMAD2</i>	Nanostring
<i>SMAD3</i>	Nanostring
<i>SMAD4</i>	Nanostring
<i>SMARC81</i>	Nanostring
<i>SMO</i>	Nanostring
<i>SNAI1</i>	Nanostring
<i>SNAI2</i>	Nanostring
<i>SOD1</i>	Nanostring
<i>SOD2</i>	Nanostring
<i>SORBS1</i>	Nanostring
<i>SOSTDC1</i>	Nanostring
<i>SOX2</i>	Nanostring
<i>SOX4</i>	Nanostring
<i>SOX9</i>	Nanostring
<i>SPARCL1</i>	Nanostring
<i>SPINK1</i>	Nanostring
<i>SPOCK2</i>	Nanostring
<i>SPP1</i>	Nanostring
<i>SPRY2</i>	Nanostring
<i>SPRY4</i>	Nanostring
<i>SQLE</i>	Nanostring
<i>SQSTM1</i>	Nanostring
<i>SRC</i>	Nanostring
<i>SREBF1</i>	Nanostring
<i>SRGN</i>	Nanostring
<i>SRSF2</i>	Nanostring
<i>SST</i>	Nanostring
<i>ST6GAL1</i>	Nanostring
<i>ST6GALNAC3</i>	Nanostring
<i>STAT1</i>	Nanostring
<i>STAT3</i>	Nanostring
<i>STAT4</i>	Nanostring
<i>STAT5A</i>	Nanostring
<i>STAT5B</i>	Nanostring
<i>STAT6</i>	Nanostring
<i>STMN1</i>	Nanostring
<i>SYK</i>	Nanostring
<i>TACSTD2</i>	Nanostring
<i>TAGLN</i>	Nanostring
<i>TAP1</i>	Nanostring
<i>TAP2</i>	Nanostring
<i>TBX21</i>	Nanostring
<i>TCAP</i>	Nanostring
<i>TCF7</i>	Nanostring
<i>TCL1A</i>	Nanostring
<i>TEK</i>	Nanostring
<i>TFEB</i>	Nanostring
<i>TGFB1</i>	Nanostring
<i>TGFB2</i>	Nanostring
<i>TGFB3</i>	Nanostring
<i>TGFB1I</i>	Nanostring
<i>TGFB1R1</i>	Nanostring
<i>TGFB1R2</i>	Nanostring
<i>THBS1</i>	Nanostring
<i>THBS2</i>	Nanostring
<i>THSD4</i>	Nanostring
<i>TIE1</i>	Nanostring
<i>TIGIT</i>	Nanostring
<i>TIMP1</i>	Nanostring
<i>TLR1</i>	Nanostring
<i>TLR2</i>	Nanostring
<i>TLR3</i>	Nanostring
<i>TLR4</i>	Nanostring
<i>TLR5</i>	Nanostring
<i>TLR7</i>	Nanostring
<i>TLR8</i>	Nanostring
<i>TM4SF1</i>	Nanostring

<i>TNF</i>	Nanostring
<i>TNFAIP6</i>	Nanostring
<i>TNFRSF10A</i>	Nanostring
<i>TNFRSF10B</i>	Nanostring
<i>TNFRSF10D</i>	Nanostring
<i>TNFRSF11A</i>	Nanostring
<i>TNFRSF11B</i>	Nanostring
<i>TNFRSF12A</i>	Nanostring
<i>TNFRSF13B</i>	Nanostring
<i>TNFRSF14</i>	Nanostring
<i>TNFRSF17</i>	Nanostring
<i>TNFRSF18</i>	Nanostring
<i>TNFRSF19</i>	Nanostring
<i>TNFRSF1A</i>	Nanostring
<i>TNFRSF1B</i>	Nanostring
<i>TNFRSF21</i>	Nanostring
<i>TNFRSF4</i>	Nanostring
<i>TNFRSF9</i>	Nanostring
<i>TNFSF10</i>	Nanostring
<i>TNFSF12</i>	Nanostring
<i>TNFSF13B</i>	Nanostring
<i>TNFSF14</i>	Nanostring
<i>TNFSF15</i>	Nanostring
<i>TNFSF4</i>	Nanostring
<i>TNFSF8</i>	Nanostring
<i>TNFSF9</i>	Nanostring
<i>TNNC1</i>	Nanostring
<i>TNT2</i>	Nanostring
<i>TNXA/B</i>	Nanostring
<i>TOP2A</i>	Nanostring
<i>TOX</i>	Nanostring
<i>TP53</i>	Nanostring
<i>TP1I</i>	Nanostring
<i>TPM1</i>	Nanostring
<i>TPM2</i>	Nanostring
<i>TPSAB1/B2</i>	Nanostring
<i>TPT1</i>	Nanostring
<i>TSC2D1</i>	Nanostring
<i>TSHZ2</i>	Nanostring
<i>TTN</i>	Nanostring
<i>TTR</i>	Nanostring
<i>TUBB</i>	Nanostring
<i>TUBB4B</i>	Nanostring
<i>TWIST1</i>	Nanostring
<i>TWIST2</i>	Nanostring
<i>TXK</i>	Nanostring
<i>TYK2</i>	Nanostring
<i>TYMS</i>	Nanostring
<i>TYROBP</i>	Nanostring
<i>UBA52</i>	Nanostring
<i>UBE2C</i>	Nanostring
<i>UPK3A</i>	Nanostring
<i>VCAM1</i>	Nanostring
<i>VCAN</i>	Nanostring
<i>VEGFA</i>	Nanostring
<i>VEGFB</i>	Nanostring
<i>VEGFC</i>	Nanostring
<i>VEGFD</i>	Nanostring
<i>VHL</i>	Nanostring
<i>VIM</i>	Nanostring
<i>VPREB3</i>	Nanostring
<i>VSIR</i>	Nanostring
<i>VTN</i>	Nanostring
<i>VWA1</i>	Nanostring
<i>VWF</i>	Nanostring
<i>WIF1</i>	Nanostring
<i>WNT10B</i>	Nanostring
<i>WNT11</i>	Nanostring
<i>WNT3</i>	Nanostring
<i>WNT5A</i>	Nanostring
<i>WNT5B</i>	Nanostring
<i>WNT7A</i>	Nanostring
<i>WNT7B</i>	Nanostring
<i>WNT9A</i>	Nanostring
<i>XBP1</i>	Nanostring
<i>XCL1/2</i>	Nanostring
<i>XKR4</i>	Nanostring
<i>YBX3</i>	Nanostring
<i>YES1</i>	Nanostring
<i>ZBTB16</i>	Nanostring
<i>ZFP36</i>	Nanostring

Supplementary Table 3. Antibodies and reagents.

Assay type	Cell subsets	Gating Strategy
CyTOF supervised	Epithelial	EpCAM ⁺ CD45 ⁻
	EpCAM ⁺ CD15 ⁺	EpCAM ⁺ CD15 ⁺
	EpCAM ⁺ ICOS ⁺	EpCAM ⁺ ICOS ⁺
	T	CD45 ⁺ EpCAM ⁻ CD3 ⁺ gdTCR ⁻
	gd T	CD45 ⁺ EpCAM ⁻ CD3 ⁺ gdTCR ⁺
	CD8 ⁺ T	CD45 ⁺ EpCAM ⁻ CD3 ⁺ gdTCR ⁻ CD4 ⁻ CD8a ⁺
	Naïve CD8 ⁺	CD45 ⁺ CD3 ⁺ CD4 ⁻ CD8a ⁺ CD45RA ⁺ CCR7 ⁺
	CD8 ⁺ TCM	CD45 ⁺ CD3 ⁺ CD4 ⁻ CD8a ⁺ CD45RA ⁻ CCR7 ⁺
	CD8 ⁺ TEM	CD45 ⁺ CD3 ⁺ CD4 ⁻ CD8a ⁺ CD45RA ⁻ CCR7 ⁻
	CD8 ⁺ TEMRA	CD45 ⁺ CD3 ⁺ CD4 ⁻ CD8a ⁺ CD45RA ⁻ CCR7 ⁻
	CD4 ⁺ T	CD45 ⁺ EpCAM ⁻ CD3 ⁺ gdTCR ⁻ CD4 ⁺ CD8a ⁻
	Naïve CD4 ⁺	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ CD45RA ⁺ CCR7 ⁺
	CD4 ⁺ TCM	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ CD45RA ⁻ CCR7 ⁺
	CD4 ⁺ TEM	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ CD45RA ⁻ CCR7 ⁻
	CD4 ⁺ TEMRA	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ CD45RA ⁺ CCR7 ⁻
	T _{reg}	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ FOXP3 ⁺ CTLA-4 ⁺ CD127 ⁻
	NKT	CD45 ⁺ EpCAM ⁻ CD3 ⁺ CD16 ⁺ CD56 ⁺
	CD16 ⁻ NK	CD45 ⁺ EpCAM ⁻ CD3 ⁻ gdTCR ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁻ CD7 ⁺ CD56 ^{hi} CD16 ⁻
	CD16 ⁺ NK	CD45 ⁺ EpCAM ⁻ CD3 ⁻ gdTCR ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁻ CD7 ⁺ CD56 ^{dim} CD16 ⁺
CyTOF unsupervised blood	Granulocytes	CD45 ⁺ CD3 ⁻ CD19 ⁻ CD15 ⁺
	B	CD45 ⁺ EpCAM ⁻ CD3 ⁻ gdTCR ⁻ HLA-DR ⁺ CD7 ⁻ CD19 ⁺
	Plasma	CD45 ⁺ EpCAM ⁻ CD3 ⁻ HLA-DR ⁻ CD7 ⁻ CD31 ⁻ CD19 ⁻ CD38 ⁺
	cDC1	CD45 ⁺ CD3 ⁻ CD7 ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁻ CD11c ⁺ CD1c ⁻ CD141 ⁺
	cDC2	CD45 ⁺ CD3 ⁻ CD7 ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁻ CD11c ⁺ CD1c ⁺ CD141 ⁻
	cDC2b	CD45 ⁺ CD3 ⁻ CD7 ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁺ CD11c ⁺ CD1c ⁺
	pDC	CD45 ⁺ CD3 ⁻ CD7 ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁻ CD123 ⁺ CD11c ⁻ CD1c ⁺
	Classical Monocytes	CD45 ⁺ CD3 ⁻ CD7 ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁻ CD16 ⁻ CD163 ⁻
	Non-Classical Monocytes	CD45 ⁺ CD3 ⁻ CD7 ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁻ CD16 ⁺
	Macrophages	CD45 ⁺ CD3 ⁻ CD7 ⁻ HLA-DR ⁺ CD19 ⁻ CD14 ⁺ CD16 ⁻ CD163 ⁺
CyTOF unsupervised biopsy	1_CD4 ^{naive}	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ CD45RA ⁺ CCR7 ⁺
	2,5_CD4 ^{TCM}	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ CD45RA ⁻ CCR7 ⁺
	3,6,9_CD4 ^{TEM}	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻ CD45RA ⁻ CCR7 ⁻
	4_CD8 ^{naive}	CD45 ⁺ CD3 ⁺ CD8a ⁺ CD4 ⁻ CD45RA ⁺ CCR7 ⁺
	7,10_CD8 ^{TEM}	CD45 ⁺ CD3 ⁺ CD8a ⁺ CD4 ⁻ CD45RA ⁻ CCR7 ⁻
	8_gd T	CD45 ⁺ CD3 ⁺ gdTCR ⁺
	11_CD8 T CD38 ⁺ HLA-DR ⁺	CD45 ⁺ CD8a ⁺ HLA-DR ⁺ CD38 ⁺
	12_CD4 T CD38 ⁺ HLA-DR ⁺	CD45 ⁺ CD4 ⁺ HLA-DR ⁺ CD38 ⁺
	13_NK CD56 ^{high}	CD45 ⁺ CD7 ⁺ CD56 ^{hi} CD16 ^{hi}
	14_NK CD56 ^{low}	CD45 ⁺ CD7 ⁺ CD56 ^{low} CD16 ^{low}
	15_B	CD45 ⁺ HLA-DR ⁺ CD7 ⁻ CD19 ⁺
	16_Granulocyte	CD45 ⁺ CD16 ⁺ CD15 ⁺
17_mDC	17_mDC	CD45 ⁺ CD7 ⁻ HLA-DR ⁺ CD14 ^{low} CD123 ⁻ CD11c ⁺
	18,20,21_Classical Monocytes	CD45 ⁺ CD7 ⁻ HLA-DR ⁺ CD14 ⁺ CD16 ⁻ CD163 ⁻
	19_Non classical Monocytes	CD45 ⁺ CD7 ⁻ HLA-DR ⁺ CD14 ⁻ CD16 ⁺
	1,4_Epithelial	EpCAM ⁺
	2,3_EpCAM ⁺ CD15 ^{high}	EpCAM ⁺ CD15 ^{hi}
Epithelial	5,6,8_EpCAM ⁺ ICOS ^{high}	EpCAM ⁺ ICOS ^{hi}
	9,12_EpCAM ⁺ HLA-DR ⁺	EpCAM ⁺ HLA-DR ⁺
	11,17_Stromal S1	CD45 ⁻ CD31 ⁻ EpCAM ⁻ CD56 ⁺
	13_Stromal	CD45 ⁻ CD31 ⁻ EpCAM ⁻
	14_B	CD45 ⁺ HLA-DR ⁺ CD19 ⁺
	15_Endothelial	CD45 ⁻ HLA-DR ⁺ EpCAM ⁻ CD31 ⁺
	16_CD4 T	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻
	18_MNP-mDC and Macrophages	CD45 ⁺ CD7 ⁻ HLA-DR ⁺ CD14 ^{low} CD16 ⁻ CD11c ⁺ CD19 ⁻ CD163 ⁺
	19_CD8 T	CD45 ⁺ CD3 ⁺ CD8a ⁺ CD4 ⁻
	20,21_Plasma	CD45 ⁺ EpCAM ⁻ CD3 ⁻ HLA-DR ⁻ CD7 ⁻ CD19 ⁻ CD38 ⁺
Epithelial	Epithelial	E-cadherin ⁺ CD45 ⁻ CD31 ⁻
	Epithelial CD68 ⁺	E-cadherin ⁺ CD68 ⁺ CD3 ⁻ CD163 ⁻
	Epithelial HLA-DR ⁺	E-cadherin ⁺ HLA-DR ⁺ CD3 ⁻ CD163 ⁻

	Epithelial cycling	E-cadherin ⁺ Ki67 ⁺
	Endothelial	CD31 ⁺ E-cadherin ⁻ CD45 ⁻
	Stromal fibroblast CD44 ⁺	Vimentin ⁺ CD44 ⁺ CD45 ⁻ CD31 ⁻ CD20 ⁻ CD14 ⁻ CD3 ⁻ E-cadherin ⁻ PDPN ⁻
	Stromal pericytes myofibroblasts	Vimentin ⁺ SMA ⁺ Collagen IV ⁺
	Stromal fibroblast PDPN ⁺	Vimentin ⁺ PDPN ⁺ CD45 ⁻ CD31 ⁻ CD20 ⁻ CD14 ⁻ CD3 ⁻ E-cadherin ⁻ CD44 ⁻
CODEX supervised	Hematopoietic	CD45 ⁺ E-cadherin ⁻ CD31 ⁻
	CD45 ⁺	CD45 ⁺
	Monocyte	CD45 ⁺ HLA-DR ⁺ CD3 ⁻ CD163 ⁺
	Macrophages	CD45 ⁺ HLA-DR ⁺ CD20 ⁺ CD3 ⁻ E-cadherin ⁻ CD163 ⁻
	B	CD45 ⁺ CD3 ⁺ E-cadherin ⁻
	T	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8 ⁻
	CD4 ⁺ T	CD45 ⁺ CD3 ⁺ CD4 ⁻ CD8 ⁺
	CD8 ⁺ T	
	Memory CD4 ⁺ T	CD45 ⁺ CD8 ⁻ CD3 ⁺ CD4 ⁺ CD45RO ⁺
	Memory CD8 ⁺ T	CD45 ⁺ CD4 ⁻ CD3 ⁺ CD8 ⁺ CD45RO ⁺
	1_B	CD45 ⁺ HLA-DR ⁺ CD19 ⁺ CD20 ⁺
CODEX unsupervised	2_Endothelial	E-cadherin ⁻ CD45 ⁻ CD31 ⁺
	3_Epithelial	E-cadherin ⁺ PCK ⁺ CD45 ⁺ CD31 ⁻
	4_Epithelial CD68 ⁺	E-cadherin ⁺ PCK ⁺ CD68 ⁺
	5_Epithelial cycling	E-cadherin ⁺ PCK ⁺ Ki67 ⁺
	6_Macrophages	CD45 ⁺ HLA-DR ⁺ CD163 ⁺
	7_Memory T	CD45 ⁺ CD3 ⁺ CD45RO ⁺
	8_Stromal fibroblast CD44 ⁺	Vimentin ⁺ CD44 ⁺ CD45 ⁻ CD3 ⁻ E-cadherin ⁻
	9_Stromal pericytes myofibroblast	Vimentin ⁺ SMA ⁺ Collagen IV ⁺
	10_Stromal fibroblast PDPN ⁺	Vimentin ⁺ PDPN ⁺ CD45 ⁻ CD3 ⁻ E-cadherin ⁻
	11_T CD45RO ⁻	CD45 ⁺ CD3 ⁺ CD45RO ⁻
MIBI supervised	Hematopoietic	CD45 ⁺ EpCAM ⁻ CD31 ⁻
	Epithelial	EpCAM ⁺ CD45 ⁻ CD31 ⁻
	Epithelial HLA-DR ⁺	EpCAM ⁺ HLA-DR ⁺
	Endothelial	CD31 ⁺ EpCAM ⁻ CD45 ⁻
	Stromal	Vimentin ⁺ EpCAM ⁻ CD45 ⁻ CD31 ⁻
	T	CD45 ⁺ CD3 ⁺
	CD4 ⁺ T	CD45 ⁺ CD3 ⁺ CD4 ⁺ CD8a ⁻
	CD8 ⁺ T	CD45 ⁺ CD3 ⁺ CD4 ⁻ CD8a ⁺
	Memory CD4 ⁺ T	CD45 ⁺ CD8 ⁻ CD3 ⁺ CD4 ⁺ CD45RO ⁺
	Memory CD8 ⁺ T	CD45 ⁺ CD4 ⁻ CD3 ⁺ CD8 ⁺ CD45RO ⁺
	Neutrophils	CD45 ⁺ EpCAM ⁻ CD3 ⁻ CD20 ⁻ CD16 ⁺ CD56 ⁻ HLA-DR ⁻
	B	CD45 ⁺ CD3 ⁻ HLA-DR ⁺ CD20 ⁺
	Plasma	CD45 ⁺ CD3 ⁻ CD20 ⁻ CD138 ⁺
	mDC	EpCAM ⁻ CD45 ⁺ CD3 ⁻ CD20 ⁻ CD11c ⁺
	Mast	EpCAM ⁻ CD45 ⁺ CD3 ⁻ CD20 ⁻ CD117 ⁺
	Monocytes	HLA-DR ⁺ CD163 ⁻ CD3 ⁻ CD20 ⁻ CD14 ⁺ CD68 ⁺
	Macrophages	CD163 ⁺ CD68 ⁺ CD3 ⁻
	Macrophages CD103 ⁺	CD163 ⁺ CD3 ⁻ CD68 ⁺ CD103 ⁺
	Fibroblasts	CD45 ⁻ EpCAM ⁻ CD31 ⁻ Vimentin ⁺ CD56 ⁻
	Myofibroblasts	CD45 ⁻ EpCAM ⁻ CD31 ⁻ Vimentin ⁺ SMA ⁺ CD56 ⁻

Supplementary Table 4. Cell subset phenotypes for supervised and unsupervised analyses for each multi-omics method.

Cell type	Gene list	Numeric ID
MNP-pDC	<i>HLA-DRA</i>	8118548
	<i>CD74</i>	8115147
	<i>HLA-DRB1</i>	8125445
	<i>CST3</i>	8065403
	<i>LYZ</i>	7957023
	<i>HLA-DPA1</i>	8125556
	<i>GZMB</i>	7978366
	<i>ITM2C</i>	8048995
	<i>SEC61B</i>	8156838
	<i>IRF1</i>	8114010
	<i>LILRA4</i>	8039246
	<i>APP</i>	8069644
	<i>C12orf75</i>	7958253
	<i>PPP1R14B</i>	7949060
	<i>JCHAIN</i>	8100827
	<i>TCF4</i>	8023415
	<i>UGC G</i>	8157216
	<i>ALOX5AP</i>	7968344
	<i>SERPINF1</i>	8003667
	<i>CCDC50</i>	8084818
MNP-Monocytes	<i>HLA-DRA</i>	8118548
	<i>CD74</i>	8115147
	<i>HLA-DRB1</i>	8125445
	<i>CST3</i>	8065403
	<i>LYZ</i>	7957023
	<i>HLA-DPA1</i>	8125556
	<i>S100A9</i>	7905571
	<i>S100A8</i>	7920244
	<i>TIMP1</i>	8167185
	<i>FCN1</i>	8165011
	<i>SERPINA1</i>	7981068
	<i>S100A4</i>	7920271
	<i>LILRA5</i>	8039236
	<i>S100A6</i>	7920258
	<i>TYMP</i>	8077103
	<i>VCAN</i>	8106743
	<i>SOD2</i>	8130556
	<i>CD44</i>	7939341
	<i>CD14</i>	8114612
	<i>EREG</i>	8095728
	<i>TSPO</i>	8073612
MNP-Macrophage	<i>HLA-DRA</i>	8118548
	<i>CD74</i>	8115147
	<i>HLA-DRB1</i>	8125445
	<i>CST3</i>	8065403
	<i>LYZ</i>	7957023
	<i>HLA-DPA1</i>	8125556
	<i>C1Q A</i>	7898793
	<i>FTL</i>	8030171
	<i>C1QC</i>	7898799
	<i>C1QB</i>	7898805
	<i>ITM2B</i>	7969003
	<i>NPC2</i>	7980146
	<i>SLC40A1</i>	8057677
	<i>CCL3</i>	8014369
	<i>MS4A7</i>	7940259
	<i>CTSS</i>	7919800
	<i>FGFR7</i>	8030374
	<i>CCL4</i>	8006602
MNP-mDC	<i>HLA-DRA</i>	8118548
	<i>CD74</i>	8115147
	<i>HLA-DRB1</i>	8125445
	<i>CST3</i>	8065403
	<i>LYZ</i>	7957023
	<i>HLA-DPA1</i>	8125556
	<i>BASP1</i>	8104601
	<i>SLC38A1</i>	7962516
	<i>LTB</i>	8124950
	<i>CST7</i>	8061416
	<i>CD1C</i>	7906348
	<i>ACAP1</i>	8004331
	<i>DAPP1</i>	8096602
	<i>STK17A</i>	8132503
	<i>ADAM8</i>	7937150
	<i>FCER1A</i>	7906443
	<i>RHOF</i>	7967202
	<i>FSCN1</i>	8131339
	<i>ICAM3</i>	8033987
	<i>CD86</i>	7896700
	<i>CXCR4</i>	8055465
	<i>COL6A2</i>	8069301
	<i>CXCL14</i>	8114249
	<i>CALD1</i>	8136347
	<i>COL3A1</i>	8046922

	<i>IGFBP7</i>	8100541
	<i>C11orf96</i>	7939492
	<i>TIMP1</i>	8167185
	<i>MMP3</i>	7951284
	<i>MMP1</i>	7951271
Stromal- Activated Fibroblast	<i>AREG</i>	8095736
	<i>TMEM158</i>	8086530
	<i>TNFRSF11B</i>	8152512
	<i>ACSL4</i>	8174474
	<i>PLAU</i>	7928429
	<i>HGF</i>	8140556
	<i>STC1</i>	8149825
	<i>GJA1</i>	8121749
	<i>CD82</i>	7939546
	<i>ISG20</i>	7985777
	<i>NRG1</i>	8145766
	<i>COL6A2</i>	8069301
	<i>CXCL14</i>	8114249
	<i>CALD1</i>	8136347
	<i>COL3A1</i>	8046922
	<i>IGFBP7</i>	8100541
	<i>C11orf96</i>	7939492
	<i>F3</i>	7917875
	<i>POSTN</i>	7971077
Stromal S2	<i>ENHO</i>	8160816
	<i>PDGFRA</i>	8095080
	<i>SOX6</i>	7946757
	<i>PLAT</i>	8150509
	<i>NSG1</i>	8093839
	<i>HSD17B2</i>	7997491
	<i>DDHD1</i>	7979223
	<i>PDGFD</i>	7951351
	<i>AGT</i>	7924987
	<i>WNT5B</i>	7953012
	<i>IGFBP7</i>	8100541
	<i>PLVAP</i>	8035297
	<i>SPARCL1</i>	8101659
	<i>IFTM3</i>	7945371
	<i>IGFBP4</i>	8007100
	<i>CAV1</i>	8135594
	<i>CCL21</i>	8160889
	<i>TFF3</i>	8070567
	<i>TFP1</i>	8057599
	<i>NUPR1</i>	8000574
	<i>MMRN1</i>	8096415
	<i>PROX1</i>	7909681
	<i>LYVE1</i>	7946579
	<i>AKAP12</i>	8122807
	<i>TBX1</i>	8071276
	<i>FABP4</i>	8151532
	<i>PPFIBP1</i>	7954559
	<i>RBP1</i>	8091078
	<i>EFEMP1</i>	8052355
	<i>NR2F1</i>	8106923
	<i>ACKR1</i>	7906435
	<i>NPC2</i>	7980146
	<i>CPE</i>	8098204
	<i>CLU</i>	8149927
Endothelial	<i>MADCAM1</i>	8023941
	<i>DUSP23</i>	7906469
	<i>IL1R1</i>	8043995
	<i>ZNF385D</i>	8085774
	<i>KCTD12</i>	7972055
	<i>IL33</i>	8154295
	<i>ADGRG6</i>	8122365
	<i>LPCAT4</i>	7987230
	<i>SELP</i>	7922200
	<i>CTNNAL1</i>	8163063
	<i>TLL1</i>	8098214
	<i>GSN</i>	8157582
	<i>SLC9A3R2</i>	7992463
	<i>NOTCH4</i>	8125383
	<i>RGCC</i>	7968789
	<i>A2M</i>	7960947
	<i>MGLL</i>	8090433
	<i>VAMP5</i>	8043203
	<i>FLT1</i>	7970763
	<i>TSC22D1</i>	7971350
	<i>RBP7</i>	7897522
	<i>ADGRF5</i>	8126798
	<i>ADGRL4</i>	7917182
	<i>SRGN</i>	7927964
	<i>IL7R</i>	8104901
	<i>CD3E</i>	7944179
	<i>ARL4C</i>	8059854
	<i>KLRB1</i>	7961059

	<i>SPOCK2</i>	7934215
	<i>KLF2</i>	8026564
	<i>SELL</i>	7896687
	<i>LEF1</i>	8102232
	<i>CCR7</i>	8015031
	<i>C1orf162</i>	7903980
	<i>LYPD3</i>	8037301
	<i>ACTN1</i>	7979824
	<i>TRABD2A</i>	8053379
CD4 T	<i>GCNT4</i>	8112668
	<i>SLC40A1</i>	8057677
	<i>FHT</i>	8088458
	<i>FAM117B</i>	8047565
	<i>S100A4</i>	7920271
	<i>JUN</i>	7916609
	<i>CCL5</i>	8014316
	<i>TNFAIP3</i>	8122265
	<i>ANXA1</i>	8155849
	<i>ANKRD28</i>	8085628
	<i>ID2</i>	8040103
	<i>EGR1</i>	8108370
	<i>GPR65</i>	7976080
	<i>LGALS3</i>	7974461
T reg	<i>HOPX</i>	8100507
	<i>IL7R</i>	8104901
	<i>CD3E</i>	7944179
	<i>ARL4C</i>	8059854
	<i>KLRB1</i>	7961059
	<i>SPOCK2</i>	7934215
	<i>CTLA4</i>	8047692
	<i>TIGIT</i>	8081799
	<i>MAF</i>	8002969
	<i>ARID5B</i>	7927732
	<i>BATF</i>	7975793
	<i>TBC1D4</i>	7972021
	<i>PHACTR2</i>	8122426
	<i>RNF19A</i>	8152041
	<i>TNFRSF4</i>	7911413
	<i>UCP2</i>	7950307
Cytotoxic lymphocyte	<i>ICA1</i>	8138202
	<i>TNFRSF1B</i>	7896721
	<i>TNFRSF18</i>	7911403
	<i>DUSP4</i>	8150076
	<i>TOX2</i>	8062782
	<i>CCL5</i>	8014316
	<i>NKG7</i>	8038809
	<i>GZMA</i>	8105340
	<i>IL32</i>	7992828
	<i>CCL4</i>	8006602
	<i>CD7</i>	8019478
	<i>GZMK</i>	8105331
	<i>ITGB2</i>	8070826
B	<i>GZMH</i>	7978360
	<i>KLF2</i>	8026564
	<i>KLRG1</i>	7953835
	<i>KLRB1</i>	7961059
	<i>SPRY1</i>	8097282
	<i>CMC1</i>	8078312
	<i>SH2D1A</i>	8169792
	<i>HCS7</i>	8028104
	<i>SH3BGRL3</i>	7899153
	<i>KLRD1</i>	7953949
	<i>CD3E</i>	7944179
	<i>CD3D</i>	7952056
	<i>CD69</i>	7896693
B	<i>CD3G</i>	7944185
	<i>CD8A</i>	8053584
	<i>CD74</i>	8115147
	<i>HLA-DRA</i>	8118548
	<i>MS4A1</i>	7940287
	<i>HLA-DRB1</i>	8125445
	<i>HLA-DPA1</i>	8125556
	<i>CD79A</i>	8029136
	<i>IGHM</i>	7896697
	<i>TCL1A</i>	7981183
	<i>FCER2</i>	8033420
	<i>YBX3</i>	7961230
	<i>KLF4</i>	8163002
	<i>KLF6</i>	7931810
	<i>GPR183</i>	7972557
	<i>TNFSF9</i>	8025053
	<i>CD27</i>	7953333
	<i>LDLRAD4</i>	8020308
	<i>ACTG1</i>	8019183
	<i>GAPDH</i>	7953385
	<i>BASP1</i>	8104601

	<i>PFN1</i>	8011759
	<i>HMGN1</i>	8070389
	<i>YBX1</i>	7900585
	<i>LCP1</i>	7971461
	<i>SLC25A5</i>	8169640
	<i>ARPC1B</i>	8134552
	<i>ANP32B</i>	8156750
	<i>ACTR3</i>	8044686
	<i>MARCKSL1</i>	7914525
Plasma	<i>JCHAIN</i>	8100827
	<i>MZB1</i>	8114511
	<i>SSR4</i>	8170775
	<i>CD79A</i>	8029136
	<i>IGHM</i>	7896697
	<i>CD44</i>	7939341
	<i>CD9</i>	7953291
	<i>CD48</i>	7921667
	<i>THY1</i>	7952268
	<i>LAG3</i>	7953418
Epithelial	<i>LGALS4</i>	8036591
	<i>PIGR</i>	7923929
	<i>KRT8</i>	7963567
	<i>C15orf48</i>	7983478
	<i>SH2D6</i>	8043157
	<i>TRPM5</i>	7945742
	<i>TLE4</i>	8156060
	<i>FABP1</i>	8053654
	<i>AQP8</i>	7994252
	<i>SLC26A3</i>	8142171
	<i>GUCA2A</i>	7915404
	<i>CEACAM7</i>	8037053
	<i>SLC26A2</i>	8109194
	<i>OTOP2</i>	8009705
	<i>BEST4</i>	7915598
	<i>CA4</i>	8008900
	<i>OLFM4</i>	7969288
	<i>LEFTY1</i>	7924663
	<i>PTGDR</i>	7974363
	<i>UBE2C</i>	8063043
Neutrophil	<i>HMGN2</i>	7899187
	<i>TOP2A</i>	8014974
	<i>HMBG2</i>	8103728
	<i>TUBA1B</i>	7963046
	<i>S100A8</i>	7920244
	<i>S100A9</i>	7905571
	<i>FCGR3A</i>	7921868
	<i>FCGR3B</i>	7921873
	<i>IL1R2</i>	8043981
	<i>GCA</i>	8046003
Pre-VDZ response signature	<i>CSF3R</i>	7914950
	<i>S100P</i>	8093950
	<i>BASP1</i>	8104601
	<i>FPR1</i>	8038899
	<i>IFITM2</i>	7937330
	<i>ELANE</i>	8024956
	<i>GAS6</i>	7970329
	<i>LEFTY1</i>	7924663
	<i>STMN1</i>	7913864
	<i>RPL37</i>	81111814
Pre-VDZ non- response signature	<i>MARCKSL1</i>	7914525
	<i>NUPR1</i>	8000574
	<i>GSTP1</i>	7941936
	<i>LYZ</i>	7957023
	<i>SPINK1</i>	8114964
	<i>AGR2</i>	8138381
	<i>OLFM4</i>	7969288
	<i>JCHAIN</i>	8100827
	<i>REG1A</i>	8042986
	<i>MMP2</i>	7995681
	<i>THBS1</i>	7982597
	<i>MMP1</i>	7951271
	<i>TCL1A</i>	7981183
	<i>COL21A1</i>	8127201
	<i>SPP1</i>	8096301
	<i>ITGA5</i>	7963786
	<i>HGF</i>	8140556
	<i>GZMK</i>	8105331
	<i>GZMH</i>	7978360
	<i>IFNG</i>	7964787

Supplementary Table 5. Cell subset-specific gene signatures used for Gene Set Enrichment Analysis (GSEA). Numeric ID, Affymetrix numeric probe identifier corresponding to each gene.

Assay type	Patient ID	Specimen/imaged	Number of cells	% of hematopoietic	% of Stromal and Endothelial	% IEC	Multi-omic method
CITE-seq	HS1	Blood	899				
	HS2	Blood	3120				
	HS3	Blood	1699				
	HS4	Blood	1761				
	HS5	N/A	N/A				
	HS6	Blood	3474				
	HS7	Blood	2356				
	HS8	Blood	1111				
	HS9	Blood	1473				
	HS10	Blood	1669				
	HS11	Blood	1003				
	HS12	Blood	1565				
CyTOF	HS1	Biopsy	9572	87.4	3.9	7.7	single cell transcriptomic and proteomic
	HS2	Biopsy	2648	85.7	5.4	8.3	
	HS3	Biopsy	8256	88.6	7.4	2.8	
	HS4	Biopsy	13662	77.8	10.5	10.7	
	HS5	Biopsy	6103	75.2	15.9	7.8	
	HS6	Biopsy	9036	70.2	22.2	6.3	
	HS7	Biopsy	9548	88.2	3.0	7.9	
	HS8	Biopsy	6076	79.1	15.8	4.3	
	HS9	Biopsy	4393	67.4	8.7	23.0	
	HS10	Biopsy	9192	80.6	5.7	13.3	
	HS11	Biopsy	7072	86.1	5.2	6.1	
	HS12	Biopsy	8342	78.9	10.8	9.6	
Multiplexed RNA-ISH	HS1	Blood	57374				
	HS2	Blood	136673				
	HS3	Blood	167045				
	HS4	Blood	151381				
	HS5	N/A	N/A				
	HS6	Blood	75079				
	HS7	Blood	162640				
	HS8	Blood	121482				
	HS9	Blood	139270				
	HS10	Blood	85768				
	HS11	Blood	176299				
	HS12	Blood	117145				
CosMx SMI 960-plex	HS1	Biopsy	82382	16.6	6.1	76.4	single cell proteomic
	HS2	Biopsy	83878	17.1	4.7	76.7	
	HS3	Biopsy	82480	32.0	8.9	57.7	
	HS4	Biopsy	57637	25.7	9.7	63.2	
	HS5	Biopsy	50659	26.7	9.4	62.9	
	HS6	Biopsy	30428	52.1	17.3	28.7	
	HS7	Biopsy	59387	34.6	5.6	57.2	
	HS8	Biopsy	41843	31.9	16.4	50.3	
	HS9	Biopsy	78302	18.0	7.0	74.1	
	HS10	Biopsy	25923	10.6	6.6	80.9	
	HS11	Biopsy	75650	33.8	7.4	57.7	
	HS12	Biopsy	15680	16.8	10.0	71.8	
CODEX	HS1	Core	8816	4.2	3.8	57.0	
	HS2	Core	10198	4.3	8.1	58.2	
	HS3	Core	6314	6.5	19.0	61.0	
	HS4	Core	5091	5.5	24.9	47.5	
	HS5	Core	8490	8.8	15.4	49.4	
	HS6	Core	8868	12.4	18.0	48.9	FFPE spatial transcriptomic
	HS7	Core	5779	7.2	10.3	55.4	
	HS8	Core	5365	13.1	13.0	37.1	
	HS9	Core	10212	22.9	8.7	57.9	
	HS10	Core	6975	24.5	8.4	55.3	
	HS11	Core	13223	33.2	25.1	41.0	
	HS12	Core	14161	10.6	20.8	55.0	
MBI	HS1	FOV	5131	30.6	14.3	55.2	
	HS2	FOV	6021	27.7	9.4	62.9	
	HS3	FOV	1353	50.2	11.2	38.6	
	HS4	FOV	4262	16.9	15.1	68.0	
	HS5	FOV	3774	32.8	16.4	50.8	
	HS6	FOV	N/A				FFPE spatial transcriptomic
	HS7	FOV	4451	35.1	20.8	44.1	
	HS8	FOV	4888	68.7	23.6	7.7	
	HS9	FOV	N/A				
	HS10	FOV	7765	35.7	24.0	40.3	
	HS11	FOV	6819	66.2	16.1	17.8	
	HS12	FOV	4319	33.9	27.0	39.1	
CODEX	HS1	Core	N/A				
	HS2	Core	10302	12.6	15.5	49.6	
	HS3	Core	3987	32.5	17.4	29.6	
	HS4	Core	3868	25.9	17.6	17.4	
	HS5	Core	3451	20.5	13.0	47.0	
	HS6	Core	N/A				FFPE spatial proteomic
	HS7	Core	4487	13.2	19.8	20.1	
	HS8	Core	5214	39.5	20.7	5.2	
	HS9	Core	3937	15.6	14.3	38.9	
	HS10	Core	11201	26.8	24.6	34.7	
	HS11	Core	12286	52.9	16.4	4.5	
	HS12	Core	9351	32.5	21.7	9.2	
MBI	HS1	FOV	893	10.7	23.6	41.9	
	HS2	FOV	1455	14.4	10.3	45.4	
	HS3	FOV	1295	7.6	6.4	50.1	
	HS4	FOV	250	2.4	8.8	67.2	
	HS5	FOV	636	10.2	18.4	43.4	
	HS6	FOV	1630	29.9	16.0	13.4	FFPE spatial proteomic
	HS7	FOV	349	11.0	24.2	42.5	
	HS8	FOV	473	18.8	14.8	23.9	
	HS9	FOV	1090	11.4	12.1	40.2	
	HS10	FOV	1041	4.3	10.2	46.5	
	HS11	FOV	1732	27.7	11.8	24.2	
	HS12	FOV	1401	15.3	14.6	32.4	

Supplementary Table 6. Cell counts and percentages of hematopoietic, stromal and endothelial, and intestinal epithelial cell (IEC) subsets for each patient biospecimen and multi-omic method.

Gene	per FOV				FOVs as replicates				Biopsies as replicates				Include in gene set
	baseMean	log2FoldChange	p-value	padj	baseMean	log2FoldChange	p-value	padj	baseMean	log2FoldChange	p-value	padj	
<i>KRT20</i>	526.97688	1.04072071	0.06350846	0.41395902	1053.37471	1.829395189	0.01106976	0.29072685	1259.97142	1.759530615	0.03333355	0.58904936	
<i>MMP2</i>	214.781871	0.779233372	8.81E-05	0.08818248	408.395547	0.650608071	0.02130012	0.40684052	489.294934	0.704799783	0.04645766	0.64526877	Yes
<i>CLDN4</i>	759.82305	0.745929417	0.02768875	0.31826169	1483.20505	1.290087005	0.00365172	0.2046753	1776.28492	1.241179209	0.02295442	0.54436599	
<i>TIMP1</i>	276.204316	0.737634615	0.01112788	0.20232501	508.769011	0.667081209	0.12191453	0.76158335	602.95769	0.541749624	0.25696256	0.82642476	
<i>PLAC8</i>	935.934625	0.720560727	0.06581948	0.41395902	1848.43313	1.347484715	0.00690973	0.27583642	2314.7905	1.208635846	0.03799885	0.62106315	
<i>THBS1</i>	344.101653	0.672537259	0.00119803	0.04991792	647.142023	0.611143584	0.02401064	0.40684052	750.480415	0.745730473	0.00962025	0.40743756	
<i>JUN</i>	453.634985	0.660664794	0.00672182	0.16894968	870.901615	0.956538051	0.00759029	0.28460002	1009.87437	0.1096086482	NA	NA	
<i>CEACAM1</i>	294.859297	0.64336382	0.03858965	0.35695204	557.851558	1.107380773	0.00344384	0.20217367	673.708459	1.040242363	0.0190342	0.48659234	
<i>MALAT1</i>	5162.4939	0.636166653	0.19155692	0.58401499	10254.0368	0.404230675	0.0184969	0.69225416	12095.2239	0.1062920598	0.11062494	0.73528713	
<i>COL3A1</i>	392.521296	0.633436878	0.0360134	0.41395902	739.851456	0.541377924	0.02273887	0.8427126	860.298456	0.433934687	0.42279269	0.95203628	
<i>MMP1</i>	207.589933	0.618646275	1.89E-07	0.00014484	385.571488	0.550571237	0.00041056	0.14987715	466.18652	0.56304112	0.0033652	0.23712057	
<i>FOS</i>	352.469296	0.606341877	0.02583546	0.4033241	643.747795	0.86933204	0.00585602	0.69787787	714.941229	0.965660035	0.06011965	0.7039651	
<i>SOD2</i>	296.546792	0.598764496	0.01659691	0.23644298	564.82725	0.538386908	0.11886787	0.76044926	676.89997	0.47775366	0.20524567	0.79313927	
<i>NEAT1</i>	866.390363	0.572790109	0.122019	0.48996338	1726.885005	1.12080306	0.01984024	0.40684052	2078.89566	1.00077444	0.06888335	0.71379068	
<i>MT2A</i>	934.428365	0.571389758	0.00954193	0.19083854	1775.58372	0.495068787	0.00284265	0.70715244	2111.81567	0.565782914	0.11290352	0.74055794	
<i>SQSTM1</i>	474.161774	0.512084338	0.00661297	0.16894968	930.493327	0.48882786	0.00154193	0.15418283	1115.58897	0.813628662	0.01383623	0.47568006	
<i>MMP9</i>	255.898007	0.511030856	0.00172926	0.06404677	487.427814	0.380037984	0.006231327	0.61346199	538.529708	0.426210316	0.06352648	0.7039651	
<i>ATF3</i>	246.568056	0.505668525	0.00875501	0.18140874	459.530162	0.74200607	0.00798477	0.28460002	544.420923	0.785730082	0.01587284	0.48605613	
<i>DUSP1</i>	372.731731	0.486584761	0.00782812	0.18140874	678.460406	0.675381703	0.00409363	0.20427463	798.9052	0.704769132	0.00981204	0.40743756	
<i>EZR</i>	464.523069	0.436439785	0.13530634	0.50487476	906.187312	0.834827562	0.16867739	0.374303	1086.545009	0.747947535	0.07461127	0.71379068	
<i>KRT16</i>	609.755295	0.425423051	0.05657241	0.40890046	975.556748	0.445061083	0.145596945	0.78606569	1220.02616	0.428064554	0.22402087	0.80631339	
<i>CDKN1A</i>	378.607985	0.411588752	0.04890184	0.39601595	709.030108	0.777604211	0.00430737	0.20470263	865.725998	0.67779355	0.03367684	0.58904936	
<i>CCL18</i>	181.931178	0.399177236	0.0096405	0.04191509	333.952038	0.355215006	0.02214067	0.40684052	402.185354	0.365597111	0.05635691	0.70234803	
<i>CRIP1</i>	579.174589	0.375098957	0.06923619	0.41774829	1180.98787	0.665194598	0.043153953	0.59532003	1378.27245	0.741358386	0.06495967	0.7039651	
<i>OAS1</i>	262.689164	0.372964912	0.07135545	0.41825833	483.452543	0.72808809	0.00948100	0.29072685	588.271886	0.683797817	0.04216207	0.63067923	
<i>CCL21</i>	283.216261	0.34774063	0.11548864	0.48534695	543.974464	0.124593639	0.675556403	0.97295875	614.389899	0.311076366	0.18767891	0.79313927	
<i>TCL1A</i>	161.962139	0.324429339	4.83E-05	0.0604232	297.745557	0.286217762	0.01287283	0.15418283	358.914476	0.31327535	0.00066112	0.13182804	
<i>FKBP5</i>	156.655595	0.319563737	0.01916562	0.20630378	288.313877	0.24083704	0.16907511	0.30350933	338.101572	0.241524611	0.21504822	0.80164351	
<i>CXCL13</i>	261.757772	0.304432077	0.02551588	0.31291833	486.537263	0.20020202	0.2058663	0.81386279	586.178145	0.259310993	0.15162784	0.77280279	
<i>MYL12A</i>	504.843393	0.303777309	0.22682194	0.61367037	993.525404	0.720716688	0.03159557	0.47063252	1160.67198	0.666088033	0.09287529	0.7263558	
<i>OASL</i>	258.014594	0.298766106	0.00554993	0.15416448	474.548174	0.456788851	0.00318844	0.2017367	577.31661	0.465282666	0.01485569	0.4777821	
<i>ZFP36</i>	466.041918	0.29221624	0.13335282	0.50487439	831.650665	0.475025375	0.00596948	0.60978778	988.611548	0.487474463	0.1091259	0.73528713	
<i>RSP01</i>	65.2856523	0.290966063	0.04109392	0.35695204	122.094016	0.257934678	0.11869042	0.76044922	143.914325	0.308995191	0.09661188	0.7263558	
<i>COL21A1</i>	96.5411045	0.290774928	0.00083991	0.03999575	178.277478	0.319372181	0.00147965	0.15418283	213.722185	0.33860771	0.0022411	0.22083429	
<i>ARHGDI</i>	205.966227	0.287861641	0.04096562	0.35695204	382.444048	0.257510969	0.10132545	0.70175124	455.340504	0.218177936	0.16459059	0.78052343	
<i>COL1A1</i>	623.954591	0.284562889	0.33760458	0.71526394	1151.763134	0.251325271	0.49427984	0.95529411	1379.0178	0.06702726	0.88072016	0.97848848	
<i>IL4R</i>	220.007153	0.280776618	0.01092497	0.20231429	416.543453	0.343877681	0.01407256	0.32991373	495.25051	0.385286083	0.01478337	0.4777821	
<i>SPP1</i>	193.795024	0.274902042	0.00265226	0.08037161	352.005521	0.3195950212	0.0020901	0.17382665	424.562732	0.317695632	0.00524	0.26121389	
<i>ITG45</i>	182.035166	0.273837899	0.00211068	0.07271833	340.852052	0.226084023	0.0456812	0.596394157	402.26067	0.25635497	0.03038688	0.5716175	
<i>IERS3</i>	401.465412	0.267372809	0.08111708	0.43161934	733.389792	0.512339802	0.01421472	0.32991373	878.336888	0.438805459	0.08685337	0.72408968	
<i>KLF2</i>	409.778611	0.259101857	0.03253334	0.338889	753.184873	0.297886963	0.02445933	0.40684052	882.492339	0.283721192	0.04871918	0.65261317	
<i>CD163</i>	117.519516	0.257532928	0.026352748	0.35695204	215.741815	0.235189381	0.010572635	0.72768895	256.5577475	0.22120639	0.20230369	0.79313927	
<i>CD52</i>	178.342569	0.249309495	0.03080577	0.33484528	329.444325	0.171281424	0.02209890	0.84271262	397.674623	0.260074683	0.00454881	0.25195327	Yes
<i>VEGFC</i>	166.304572	0.241700243	0.034815416	0.39610595	238.033362	0.19786047	0.019683197	0.38836279	263.045931	0.244462294	0.15182682	0.77280279	
<i>MARCO</i>	125.215062	0.23998011	0.00838667	0.18140874	231.83625	0.232269458	0.023826927	0.40684052	281.024054	0.265672579	0.02692013	0.55278634	
<i>KRT8</i>	330.983952	0.230305325	0.56872815	0.89607096	643.013228	0.685827652	0.16679373	0.80350933	752.251250	0.30805751	0.87752381		
<i>COL6A3</i>	260.388401	0.228403251	0.553764748	0.172202858	320.563744	0.41757954	0.335751454	0.92653553	565.933646	0.133127424	0.53617693	0.95668154	
<i>IGHG1</i>	11861.3389	0.227212704	0.72987109	0.97488453	2151.257799	0.226551772	0.181740402	0.98267871	2486.8984	0.222486389	0.83419358	0.97848848	
<i>P2RX5</i>	212.30091	0.225002807	0.03911167	0.35695204	387.684989	0.186952907	0.16030120	0.80350933	470.149445	0.158137544	0.30897581	0.87763215	
<i>MSA41</i>	141.663715	0.224287822	0.11323991	0.48178198	266.691471	0.145828475	0.030703973	0.93483037	317.340523	0.200614537	0.28703487	0.84792809	
<i>ADGRL2</i>	156.091385	0.222645818	0.06134205	0.41301069	289.506168	0.260626292	0.01292833	0.32449596	347.203971	0.27990094	0.022295265	0.54436599	
<i>FGR</i>	135.270051	0.218239284	0.0179528	0.20630378	248.924379	0.197403655	0.03606925	0.52169727	297.93001	0.218763278	0.04238266	0.63067923	
<i>MMP14</i>	223.439612	0.215868778	0.0230019	0.31251846	412.802454	0.170848854	0.014367474	0.78606569	486.266457	0.166358881	0.22042274	0.80164351	
<i>LINC01781</i>	161.499891	0.213090821	0.05752892</td										