nature portfolio

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Reporting Summary

Nature Portfolio wishes to improve the reproducibility of the work that we publish. This form provides structure for consistency and transparency in reporting. For further information on Nature Portfolio policies, see our <u>Editorial Policies</u> and the <u>Editorial Policy Checklist</u>.

Statistics

For	all st	atistical analyses, confirm that the following items are present in the figure legend, table legend, main text, or Methods section.			
n/a	Cor	Confirmed			
	×	The exact sample size (n) for each experimental group/condition, given as a discrete number and unit of measurement			
	x	A statement on whether measurements were taken from distinct samples or whether the same sample was measured repeatedly			
	×	The statistical test(s) used AND whether they are one- or two-sided Only common tests should be described solely by name; describe more complex techniques in the Methods section.			
×		A description of all covariates tested			
	x	A description of any assumptions or corrections, such as tests of normality and adjustment for multiple comparisons			
	×	A full description of the statistical parameters including central tendency (e.g. means) or other basic estimates (e.g. regression coefficient) AND variation (e.g. standard deviation) or associated estimates of uncertainty (e.g. confidence intervals)			
	×	For null hypothesis testing, the test statistic (e.g. <i>F</i> , <i>t</i> , <i>r</i>) with confidence intervals, effect sizes, degrees of freedom and <i>P</i> value noted Give <i>P</i> values as exact values whenever suitable.			
X		For Bayesian analysis, information on the choice of priors and Markov chain Monte Carlo settings			
	×	For hierarchical and complex designs, identification of the appropriate level for tests and full reporting of outcomes			
	x	Estimates of effect sizes (e.g. Cohen's d, Pearson's r), indicating how they were calculated			
	Our web collection on statistics for biologists contains articles on many of the points above.				

Software and code

Policy information about availability of computer code						
Data collection	Commercial Fluidigm CYTOF software was used for CyTOF data acquisition.					
Data analysis	All code used in this study including R markdowns and Jupyter notebooks are available on the Ulcerative Colitis project GitHub repository (https://github.com/mkattah/UC_VDZ). Seurat v3, Cellranger 3.0.2, scanpy==1.9.1, anndata==0.8.0, squidpy==1.2.3, excellxgene==2.9.2, and MAST==1.20, were used for scRNA-seq and CosMx 960-plex RNA-ISH analysis. R-based Cytometry Clustering Optimization aNd Evaluation (Cyclone) pipeline was developed by the UCSF Data Science CoLab (https://github.com/UCSF-DSCOLAB/cyclone). Cyclone, FlowSOM 2.6.0 and FlowJo v10 were used for CyTOF and CODEX analysis. Cell segmentation and quantitation for 12-plex RNA-ISH, MIBI, and CODEX were performed using HALO version 3.4. Additional statistical analyses were performed in GraphPad Prism v9. Conda enivronments are specified in .yaml files provided at our GitHub link to facilitate reproducibility.					

For manuscripts utilizing custom algorithms or software that are central to the research but not yet described in published literature, software must be made available to editors and reviewers. We strongly encourage code deposition in a community repository (e.g. GitHub). See the Nature Portfolio guidelines for submitting code & software for further information.

Policy information about availability of data

All manuscripts must include a data availability statement. This statement should provide the following information, where applicable:

- Accession codes, unique identifiers, or web links for publicly available datasets
- A description of any restrictions on data availability
- For clinical datasets or third party data, please ensure that the statement adheres to our policy

The raw sequencing data generated in this study have been deposited on the dbGaP database under accession code phs003502.v1.p1. The raw sequencing data are available under controlled access for privacy concerns; access can be requested through dbGaP. Processed data are deposited as a GEO Super Series under accession code GSE250498 and are publicly available. All additional data generated in this study are provided in the Supplementary Information/Source Data file and on Figshare. Processed and annotated objects are saved in AnnData (h5ad) format78. These AnnData objects are accessible in Figshare (fresh versus cryopreserved scRNA-seq 10.6084/m9.figshare.21936240; blood scRNA-seq 10.6084/m9.figshare.21900948; biopsy scRNA-seq 10.6084/m9.figshare.21919425; biopsy CITE-seq 10.6084/m9.figshare.21919356; PBL CyTOF 10.6084/m9.figshare.21977834; biopsy CyTOF 10.6084/m9.figshare.21977798; secondary CyTOF PBL and biopsy analysis 10.6084/m9.figshare.23902065; CosMx 960-plex RNA-ISH 1st run 10.6084/m9.figshare.21919338; CosMx 1000-plex RNA-ISH 2nd longitudinal analysis 10.6084/m9.figshare.23896959). Publicly available microarray data (GSE73661) were downloaded from the NCBI gene expression omnibus. GRCh38 was used as the reference genome.

Research involving human participants, their data, or biological material

Policy information about studies with human participants or human data. See also policy information about sex, gender (identity/presentation), and sexual orientation and race, ethnicity and racism.

Reporting on sex and gender	Baseline demographic and clinical data for the study participants are provided in Supplementary Table 1. Demographic options were defined by the investigators and participants chose their classifications. We attempted to balance sex across disease and treatment groups. The study design is insufficient to determine the association of sex with disease or treatment response.
Reporting on race, ethnicity, or other socially relevant groupings	Baseline demographic and clinical data for the study participants are available through dbGaP. Demographic options were defined by the investigators and participants chose their classifications. We recruited consecutively eligible patients regardless of race or ethnicity across disease and treatment groups.
Population characteristics	Baseline demographic and clinical data for the study participants are provided in Supplementary Table 1. Demographic options were defined by the investigators and participants chose their classifications.
Recruitment	Patients undergoing colonoscopy or sigmoidoscopy for standard of care indications were screened for study eligibility. Eligible patients were recruited consecutively to minimize self-selection bias. Patients were compensated \$50 for each sample collection event. For this study, HC patients were patients without known or suspected IBD undergoing elective colonoscopy or sigmoidoscopy for various indications (e.g. colorectal cancer screening). UC patients were eligible if they had an established diagnosis of UC by UCSF Gastroenterology faculty, had at least mild endoscopic disease activity (Mayo equal to or greater than 1), and if they were either on Vedolizumab or no biologic (5-ASA therapy) in the maintenace phase of treatment. Exclusion criteria included any contraindication to endoscopic forceps biopsy such as anticoagulation, dual antiplatelet agents that were not held prior to the procedure, or other high risk of bleeding. Subjects with anticipated procedural difficulty such as significant co-morbidities, prior difficult endoscopy, or any other planned high-risk endoscopic procedure were also excluded to minimize risk. All patients gave written informed consent and approval.
Ethics oversight	This study is approved by the Institutional Review Board of the University of California, San Francisco (19-27302).

Ethics oversight

Note that full information on the approval of the study protocol must also be provided in the manuscript.

Field-specific reporting

Please select the one below that is the best fit for your research. If you are not sure, read the appropriate sections before making your selection.

× Life sciences

Sample size

Behavioural & social sciences

Ecological, evolutionary & environmental sciences

For a reference copy of the document with all sections, see nature.com/documents/nr-reporting-summary-flat.pdf

Life sciences study design

All studies must disclose on these points even when the disclosure is negative.

The primary goal of this study was to determine the peripheral and mucosal cell subsets that contribute to ulcerative colitis (UC) and respond to Vedolizumab (VDZ). We wanted to compare healthy controls, patients with UC on aminosalicylates, and patients with UC on VDZ. Although we did not perform a detailed power calculation, we based our sample size on pilot scRNA-seq data showing that biopsy cryopreservation and batch processing facilitates inter-sample comparison,. We chose four patients per group as the minimum number to detect large effect sizes in relative cell subset frequency and differentially expressed genes. Our goal was to sequence >3000 cells per patient biopsy. For validation in longitudinal bulk transcriptomic data, all available pre- and post-treatment samples were included from GSE73661 to perform the indicated comparisons.

Data exclusions	Ribosomal and mitochondrial genes were not included in the final MAST output to prioritize differentially expressed genes that were most pertinent to the research question. For 12-plex RNA-ISH, MIBI, CODEX, and CosMx 960-plex RNA-ISH formalin-fixed, paraffin embedded (FFPE) analysis, low-quality cores or FOVs were excluded if they were grossly damaged or detached during processing, or if they had poor nuclei staining that precluded cell segmentation. For MIBI, FOVs that contained exclusively lymphoid aggregates, with no mucosal or submucosal cellular compartments, were also excluded.
Replication	Biopsy CITE-seq and blood scRNA-seq for all patients were performed on one day each, and both Right and Left colon biopsies were separately analyzed for each patient. The CITE-seq and scRNA-seq experiments were not replicated due to cost and limitations associated with small biospecimens. Additional research-only mucosal biopsies were obtained from two regions during endoscopic procedures for CITE-seq, CyTOF, bulk RNA-seq, and FFPE analysis. Given the number of biopsies obtained for the primary analysis, additional technical replicates could not be obtained without additional risk. CyTOF, 12-plex RNA-ISH, MIBI, CODEX, and 960-plex CosMx RNA-ISH on the same patient samples collected at the same time served as internal validation of major findings from CITE-seq analysis. The CyTOF, RNA-ISH, MIBI, CODEX, and CosMx experiments were not replicated due to cost and limitations associated with small biospecimens. However, the use of multiple, different methods on the same samples improved internal validation and served as a form of technical triangulation. For external validation, we then performed CyTOF and longitudinal CosMx spatial transcriptomics analysis in separate cohorts, and those analyses were not replicated due to cost and limitations associated with small biospecimens.
Randomization	This was a cross-sectional, case-control study design, so patients and corresponding biospecimens were not randomized. Patients were sequentially recruited with no additional pre-specified criteria other than disease and treatment status. Covariates and baseline demographics are highlighted in Supplementary Tables 1 & 2.
Blinding	All patient-derived biospecimens were de-identified with unique identifiers and researchers were blinded during all sample processing steps. Pathologists were blinded for histologic scoring. Samples were linked to disease and treatment status for data analysis. Blinding during data analysis was not employed due to the practical constraints of our case-control study design, which necessitated linking each sample to individual patients for accurate group assignment. However, to mitigate potential bias, we maintained blinding during the quantification of cell subset abundance in fields-of-view or regions-of-interest for our spatial multi-omics analyses.

Reporting for specific materials, systems and methods

We require information from authors about some types of materials, experimental systems and methods used in many studies. Here, indicate whether each material, system or method listed is relevant to your study. If you are not sure if a list item applies to your research, read the appropriate section before selecting a response.

terials & experimental systems	Methods	
Involved in the study	n/a Involved in the study	
X Antibodies	ChIP-seq	
Eukaryotic cell lines	Flow cytometry	
Palaeontology and archaeology	K MRI-based neuroimaging	
Animals and other organisms		
Clinical data		
Dual use research of concern		
Plants		
	Involved in the study Antibodies Eukaryotic cell lines Palaeontology and archaeology Animals and other organisms Clinical data Dual use research of concern	Involved in the study n/a Involved in the study Involved in the study Involved in the study Antibodies Involved in the study Eukaryotic cell lines Involved in the study Palaeontology and archaeology Involved in the study Animals and other organisms Involved in the study Clinical data Dual use research of concern

Antibodies

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Antibodies used

See Supplemental Table 3.

Markers for CyTOF:

CD45 (Hl30) at 1.5 µg/ml, BioLegend (# 304002); EpCAM (9C4) at 1.5 µg/ml, Biolegend (# 324202); CD7 (M-T701) at 1.5 µg/ml, BD Bioscience (#555359); CD15 (W6D3) at 1.5 µg/ml, Biolegend (# 323002); CD3 (UCHT1) at 0.4 µg/ml, Biolegend (# 300402); CD19 (HlB19) at 0.4 µg/ml, Biolegend (# 302202); CD163 (GHl61) at 3 µg/ml, Biolegend (# 333602); CD4 (RPA-T4) at 0.4 µg/ml, Thermo (# 14-0049-82); CD8a (RPA-T8) at 0.4 µg/ml, Biolegend (# 301002); CD11c (BU15) at 0.4 µg/ml, Thermo (#MA-82142); CD14 (M5E2) at 1.5 µg/ml, Biolegend (# 301802); CD127 (A019D5) at 1.5 µg/ml, Biolegend (# 351302); CD123 (6H6) at 0.8 µg/ml, Thermo (# 14-1239-82); gdTCR (5A6.E9) at 0.4 µg/ml, ATCC (#HB-9772); CD45RA (H100) at 1.5 µg/ml, Biolegend (# 304102); TIM3 (F38-2E2) at 3 µg/ml, Biolegend (#345002); TGIT (A15153G) at 1.5 µg/ml, Biolegend (# 372720); PD-L1 (29E.2A3) at 1.5 µg/ml, Biolegend (# 329702); Integrin a4 (9F10) at 1.5 µg/ml, Biolegend (# 304302); CD27 (O323) at 0.4 µg/ml, Biolegend (# 302802); CD137_4-1BB (4B4-1) at 1.5 µg/ml, Thermo (14-9056-82); Tbet (4B10) at 0.4 µg/ml, Thermo (# 14-5825-82); CTCLA-4 (14D3) at 0.4 µg/ml, Thermo (# 14-1529-82); FOXP3 (PCH101) at 0.8 µg/ml, Thermo (# 14-4776-82); CD31 (WM59) at 0.8 µg/ml, Biolegend (# 303102); Integrin b7 (FIB504) at 1.5 µg/ml, Biolegend (# 321202); CD141_BDCA3 (M80) at 0.8 µg/ml, Biolegend (# 304302); CCR7 (GO43H7) at 1.5 µg/ml, Biolegend (# 335202); KL6-7 (ki-67) at 0.8 µg/ml, Biolegend (# 30502); CD25 (M-A251) at 0.4 µg/ml, Biolegend (# 32502); CO38.4A) at 0.8 µg/ml, Biolegend (# 303502); ICOS (C398.4A) at 0.8 µg/ml, Biolegend (# 303502); ICOS (C398.4A) at 0.8 µg/ml, Biolegend (# 302049).

Markers for MIBI: dsDNA (3519 DNA) at 0.2 μ g/ml, Abcam (# ab27156); E-Cadherin (36) at 0.5 μ g/ml, Abcam (# ab287971); Histone H3 (D1H2) at 1:100, Ionpath (# 711501-100); FoxP3-AF488 (236A/E7) at 2 μ g/ml, BD Pharmigen (# 561181); anti-488 (polyclonal) at 3 μ g/ml, Thermofisher (# A-11094); CD163 (EPR14643-36) at 0.6 μ g/ml, Abcam (# ab215976); CD4 (EPR6855) at 1:100, Ionpath (# 714301-100); CD11c (EP1347Y) at 1:100, Ionpath (# 714401-100); CD14 (D7A27) at 0.6 μ g/ml, CST (# 42878); CD16 (D1N9L) at 0.6

µg/ml, CST (# 72204); LAG3 (17B4) at 5 µg/ml Novus (# NBP1-97657); PD-1 (D4W2J) at 1:100, lonpath (# 714801-100); PD-L1 (E1L3N) at 1:100, lonpath (# 714902-100); Granzyme B (D6E9W) at 1:100, lonpath (# 715002-100); CD56 (MRQ-42) at 1:100, lonpath (#7155101-100); CD31 (EP3095) at 1:100, lonpath (#715202-100); KI67 (D2H10) at 0.15 µg/ml, CST (# 44092); CD138 (EPR6454) at 5 µg/ml, Abcam (# ab226108); CD117 (YR145) at 1:100, lonpath (#715501-100); CD68 (D4B9C) at 1:100, lonpath (#715601-100); CD103 (EPR4166(2)) at 15 µg/ml, Abcam (# ab271889); CD8 (C8/144B) at 1:100, lonpath (# 715801-100); CD3 (D7A6E) at 1:100, lonpath (# 715903-100); Tbet (D6N8B) at 5 µg/ml, CST (# 27112); CD45RO (UCHL1) at 1:100, lonpath (716101-100); TIM3 (EPR22241) at 1:100, lonpath (# 716201-100); Vimentin (D21H3) at 1:100, lonpath (# 716301-100); CD27 (EPR8569) at 5 µg/ml, Abcam (ab256583); Pan-Keratin (AE1/AE3) at 1:100, lonpath (# 716501-100); aSMA (SP171) at 2.5 µg/ml, Abcam (# ab242395); CD20 (L26) at 1:100, lonpath (# 716701-100); CD11b (D6X1N) at 0.5 µg/ml, Abcam (#ab187537); BDCA3-CD141 (E7Y9P) at 10 µg/ml, CST (#34149); CD21 (EP3093) at 1:100, lonpath (# 717001-100); IDO1 (EPR20374) at 1:100, lonpath (# 717101-100); HLA-DR (EPR3692) at 1:100, lonpath (# 717201-100); EpCAM (D9S3P) at 0.5 µg/ml, CST (# 55725); CD45 (2B11/PD7/26) at 1:100, lonpath (# 717501-100); HLA-Class 1 (EMR8-5) at 1:100, lonpath (# 717602-100).

Markers for CODEX: HLA-A (EP1395Y) at 1:200, Abcam (# ab216653); CD34 (QBEND/10) at 1:150, Thermofisher (# MA1-10205); CD4 (EPR6885) at 1:200, Abcam (# ab133616); CD20 (L26) at 1:200, Thermofisher (# 14-0202-82); CD14 (EPR3653) at 1:500, Abcam (# ab133335); CD68 (KP1) at 1:200, Thermofisher (# MA5-13324); Vimentin (091D3) at 1:200, Biolegend (# 677802); CD8 (C8/144B) at 1:200, Biolegend (# 372908); CD11c (118/A5) at 1:200, Thermofisher (# 14-9761-82); CD31 (EP3095) at 1:100, Abcam (# ab134168); E-Cadherin (4A2C7) at 1:300, Thermofisher (# 33-4000); CD45 (D9M8I) at 1:300, CST (# 13917); SMA (1A4) at 1:200, Abcam (# ab7817); CD45RO (UCHL1) at 1:200, Biolegend (# 304202); CD3e (EP449E) at 1:200, Abcam (# ab52959); Pan-cytokeratin (AE1/AE3) at 1:200, Biolegend (# 914204); CD44 (156-3C11) at 1:400, CST (#3570); HLA-DR (EPR3692), at 1:200, Abcam (# ab92511); Granzyme B (D6E9W) at 1:50, CST (# 46890); CD16a (D11) at 1:200, Biolegend (# 30702); FOXP3 (259D/C7) AT 1:100, BD Pharmigen (# 560044); Ki67 (B56) at 1:300, BD Pharmigen (# 556003); CD163 (D6U1J) at 1:50, CST (#93498); CD19 (RM332) at 1:50, RevMab (# 31-1219-00); CD11b (EP1345Y) at 1:50, Abcam (# ab52478); CD21 (EP3093) at 1:400, Abcam (# ab75985); CD57 (HNK-1) at 1:100, Biolegend (# 359610).

Markers for CITE-seq:

NOTE dilutions are proprietary according BioLegen.

CD103 (Integrin aE) GACCTCATTGTGAAT A0145 Ber-ACT8 proprietary CD103 399907 BioLegend CD35 ACTTCCGTCGATCTT A0167 E11 proprietary CD35 399907 BioLegend CD25 TTTGTCCTGTACGCC A0085 BC96 proprietary CD25 399907 BioLegend CD62L GTCCCTGCAACTTGA A0147 DREG-56 proprietary CD62L 399907 BioLegend CD3 CTCATTGTAACTCCT A0034 UCHT1 proprietary CD3 399907 BioLegend CD33 TAACTCAGGGCCTAT A0052 P67.6 proprietary CD33.1 399907 BioLegend CD4 TGTTCCCGCTCAACT A0072 RPA-T4 proprietary CD4.2 399907 BioLegend CD58 (LFA-3) GTTCCTATGGACGAC A0174 TS2/9 proprietary CD58.1 399907 BioLegend CD8a GCTGCGCTTTCCATT A0080 RPA-T8 proprietary CD8a 399907 BioLegend CD56 TCCTTTCCTGATAGG 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) GGGCAATTAGCGAGT A0802 P44-8 proprietary CD336--NKp44 399907 BioLegend CD69 GTCTCTTGGCTTAAA A0146 FN50 proprietary CD69.1 399907 BioLegend CD335 (NKp46) ACAATTTGAACAGCG A0101 9E2 proprietary CD335--NKp46 399907 BioLegend CD49f TTCCGAGGATGATCT A0070 GoH3 proprietary CD49f 399907 BioLegend CD161 GTACGCAGTCCTTCT A0149 HP-3G10 proprietary CD161 399907 BioLegend CD45RO CTCCGAATCATGTTG A0087 UCHL1 proprietary CD45RO 399907 BioLegend CD20 TTCTGGGTCCCTAGA A0100 2H7 proprietary CD20 399907 BioLegend KLRG1 (MAFA) CTTATTTCCTGCCCT A0153 SA231A2 proprietary KLRG1--MAFA 399907 BioLegend CD19 CTGGGCAATTACTCG A0050 HIB19 proprietary CD19.1 399907 BioLegend CD32 GCTTCCGAATTACCG A0142 FUN-2 proprietary CD32 399907 BioLegend CD276 (B7-H3) GACTGGGAGGGTATT A0010 DCN.70 proprietary CD276--B7-H3 399907 BioLegend CD279 (PD-1) ACAGCGCCGTATTTA A0088 EH12.2H7 proprietary CD279--PD-1 399907 BioLegend CD197 (CCR7) AGTTCAGTCAACCGA A0148 G043H7 proprietary CD197--CCR7 399907 BioLegend CD252 (OX40L) TTTAGTGATCCGACT A0021 11C3.1 proprietary N/A 399907 BioLegend CD223 (LAG-3) CATTTGTCTGCCGGT A0152 11C3C65 proprietary CD223--LAG-3 399907 BioLegend HLA-DR AATAGCGAGCAAGTA A0159 L243 proprietary HLA-DR 399907 BioLegend CD45RA TCAATCCTTCCGCTT A0063 HI100 proprietary CD45RA 399907 BioLegend CD196 (CCR6) GATCCCTTTGTCACT A0143 G034E3 proprietary CD196--CCR6 399907 BioLegend CD2 TACGATTTGTCAGGG A0367 TS1/8 proprietary CD2.1 399907 BioLegend CD326 (Ep-CAM) TTCCGAGCAAGTATC A0123 9C4 proprietary CD326--EpCAM 399907 BioLegend CD169 (Sialoadhesin, Siglec-1) TACTCAGCGTGTTTG A0206 7-239 proprietary CD169--Sialoadhesin--Siglec-1 399907 BioLegend CD41 ACGTTGTGGCCTTGT A0353 HIP8 proprietary CD41 399907 BioLegend CD49b GCTTTCTTCAGTATG A0371 P1E6-C5 proprietary CD49b 399907 BioLegend CD303 (BDCA-2) GAGATGTCCGAATTT A0370 201A proprietary CD303--BDCA-2 399907 BioLegend CD29 GTATTCCCTCAGTCA A0369 TS2/16 proprietary CD29 399907 BioLegend CD184 (CXCR4) TCAGGTCCTTTCAAC A0366 12G5 proprietary CD184--CXCR4 399907 BioLegend CD98 GCACCAACAGCCATT A0374 MEM-108 proprietary CD98 399907 BioLegend CD124 (IL-4R) CCGTCCTGATAGATG A0363 G077F6 proprietary N/A 399907 BioLegend CD135 (Flt-3/Flk-2) CAGTAGATGGAGCAT A0351 BV10A4H2 proprietary CD135 399907 BioLegend CD370 (CLEC9A/DNGR1) CTGCATTTCAGTAAG A0207 8F9 proprietary CD370--CLEC9A--DNGR1 399907 BioLegend CD21 AACCTAGTAGTTCGG A0181 Bu32 proprietary CD21 399907 BioLegend

CD39 TTACCTGGTATCCGT A0176 A1 proprietary CD39 399907 BioLegend CD141 (Thrombomodulin) GGATAACCGCGCTTT A0163 M80 proprietary CD141--Thrombomodulin 399907 BioLegend CD64 AAGTATGCCCTACGA A0162 10.1 proprietary CD64 399907 BioLegend CD152 (CTLA-4) ATGGTTCACGTAATC A0151 BNI3 proprietary CD152 399907 BioLegend CD61 AGGTTGGAGTAGACT A0372 VI-PL2 proprietary CD61 399907 BioLegend CD163 GCTTCTCCTTCA A0358 GHI/61 proprietary CD163.1 399907 BioLegend CD357 (GITR) ACCTTTCGACACTCG A0360 108-17 proprietary CD357--GITR 399907 BioLegend CD137 (4-1BB) CAGTAAGTTCGGGAC A0355 4B4-1 proprietary CD137 399907 BioLegend CD81 (TAPA-1) GTATCCTTCCTTGGC A0373 5A6 proprietary CD81--TAPA-1 399907 BioLegend CD366 (Tim-3) TGTCCTACCCAACTT 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) CCAGCTCATTAGAGC A0156 DX2 proprietary CD95--Fas 399907 BioLegend CD80 ACGAATCAATCTGTG A0005 2D10 proprietary CD80.1 399907 BioLegend CD138 (Syndecan-1) ACTCTTTCGTTTACG A0055 MI15 proprietary CD138--Syndecan-1 399907 BioLegend CD70 CGCGAACATAAGAAG A0027 113-16 proprietary CD70.1 399907 BioLegend CD269 (BCMA) CAGATGATCCACCAT A0056 19F2 proprietary CD269--BCMA 399907 BioLegend IgM TAGCGAGCCCGTATA A0136 MHM-88 proprietary IgM 399907 BioLegend CD59 AATTAGCCGTCGAGA A0361 p282 (H19) proprietary CD59.1 399907 BioLegend CD194 (CCR4) AGCTTACCTGCACGA A0071 L291H4 proprietary CD194--CCR4 399907 BioLegend CD275 (B7-H2, ICOSL) GTGCATTCAACAGTA A0009 2D3 proprietary CD275--B7-H2--ICOSL 399907 BioLegend CD314 (NKG2D) CGTGTTTGTTCCTCA A0165 1D11 proprietary CD314--NKG2D 399907 BioLegend CD86 GTCTTTGTCAGTGCA A0006 IT2.2 proprietary CD86.1 399907 BioLegend Galectin-9 ACTCACTGGAGTCTC A0016 9M1-3 proprietary N/A 399907 BioLegend CD195 (CCR5) CCAAAGTAAGAGCCA A0141 J418F1 proprietary CD195--CCR5 399907 BioLegend CD1c GAGCTACTTCACTCG A0160 L161 proprietary CD1c 399907 BioLegend CD28 TGAGAACGACCCTAA A0386 CD28.2 proprietary CD28.1 399907 BioLegend TSLPR (TSLP-R) CAGTCCTCTCTGTCA A0387 1D3 proprietary TSLPR 399907 BioLegend CD38 TGTACCCGCTTGTGA A0389 HIT2 proprietary CD38.1 399907 BioLegend CD155 (PVR) ATCACATCGTTGCCA A0023 SKII.4 proprietary CD155--PVR 399907 BioLegend CD270 (HVEM, TR2) TGATAGAAACAGACC A0020 122 proprietary CD270--HVEM--TR2 399907 BioLegend CD178 (Fas-L) CCGGTCCTCTGTATT A0177 NOK-1 proprietary CD178--FasL 399907 BioLegend CD127 (IL-7R) GTGTGTTGTCCTATG A0390 A019D5 proprietary CD127--IL-7Ra 399907 BioLegend CD7 TGGATTCCCGGACTT A0066 CD7-6B7 proprietary CD7.1 399907 BioLegend CD117 (c-kit) AGACTAATAGCTGAC A0061 104D2 proprietary CD117--c-kit 399907 BioLegend CD10 CAGCCATTCATTAGG A0062 HI10a proprietary CD10 399907 BioLegend CD40 CTCAGATGGAGTATG A0031 5C3 proprietary CD40.1 399907 BioLegend CD48 CTACGACGTAGAAGA A0029 BJ40 proprietary CD48.1 399907 BioLegend CD154 GCTAGATAGATGCAA A0032 24-31 proprietary CD154 399907 BioLegend CD47 GCATTCTGTCACCTA A0026 CC2C6 proprietary CD47.1 399907 BioLegend CD112 (Nectin-2) AACCTTCCGTCTAAG A0024 TX31 proprietary CD112--Nectin-2 399907 BioLegend CD52 CTTTGTACGAGCAAA A0033 HI186 proprietary CD52.1 399907 BioLegend CD71 CCGTGTTCCTCATTA A0394 CY1G4 proprietary CD71 399907 BioLegend CD22 GGGTTGTTGTCTTTG A0393 S-HCL-1 proprietary CD22.1 399907 BioLegend CD15 (SSEA-1) TCACCAGTACCTAGT A0392 W6D3 proprietary CD15--BEA-1 399907 BioLegend XCR1 AAGACGCATGTCAAC A0208 S15046E proprietary XCR1.1 399907 BioLegend CD26 GGTGGCTAGATAATG A0396 BA5b proprietary CD26 399907 BioLegend B7-H4 TGTATGTCTGCCTTG A0395 MIH43 proprietary B7-H4 399907 BioLegend CD31 ACCTTTATGCCACGG 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) TGTTTACGAGAGCCC A0102 BM16 proprietary CD294--CRTH2 399907 BioLegend CD45 TCCCTTGCGATTTAC A0048 2D1 proprietary CD45 399907 BioLegend CD34 GCAGAAATCTCCCTT A0054 581 proprietary CD34.1 399907 BioLegend CD244 (2B4) TCGCTTGGATGGTAG A0189 C1.7 proprietary CD244--2B4 399907 BioLegend CD185 (CXCR5) AATTCAACCGTCGCC A0144 J252D4 proprietary CD185--CXCR5 399907 BioLegend CD18 TATTGGGACACTTCT A0385 TS1/18 proprietary CD18 399907 BioLegend CD119 (IFN-? R ? chain) TGTGTATTCCCTTGT 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 AGATTCCTTCGTGTT A0180 ML5 proprietary CD24.1 399907 BioLegend CTTCCGATTCATTCA A0139 B1 proprietary TCR-G--TCR-D 399907 BioLegend TCR / CD115 (CSF-1R) AATCACGGTCCTTGT A0398 9-4D2-1E4 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) TCCACTCATTCTGTA A0400 BV9 proprietary CD144--VE-cadherin 399907 BioLegend CD193 (CCR3) ACCAATCCTTTCGTC A0397 5E8 proprietary CD193--CCR3 399907 BioLegend CD55 GCTCATTACCCATTA A0383 JS11 proprietary CD55.1 399907 BioLegend

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CD79b (IgB) ATTCTTCAACCGAAG A0187 CB3-1 proprietary CD79b--Ig-B 399907 BioLegend CD123 CTTCACTCTGTCAGG A0064 6H6 proprietary CD123 399907 BioLegend CD11c TACGCCTATAACTTG A0053 S-HCL-3 proprietary CD11c 399907 BioLegend CD54 CTGATAGACTTGAGT A0217 HA58 proprietary CD54 399907 BioLegend CD268 (BAFF-R) CGAAGTCGATCCGTA A0215 11C1 proprietary CD268--BAFF-R 399907 BioLegend TIM-4 CGTCATATAGTATGG A0428 9F4 proprietary Tim-4 399907 BioLegend CD1a GATCGTGTTGTGTTA A0402 HI149 proprietary CD1a 399907 BioLegend CD301 (CLEC10A) ACCTAGAAATCAGCA A0401 H037G3 proprietary CD301--CLEC10A 399907 BioLegend CD63 GAGATGTCTGCAACT A0404 H5C6 proprietary CD63.1 399907 BioLegend CD44 AATCCTTCCGAATGT A0125 BJ18 proprietary CD44.1 399907 BioLegend IgD CAGTCTCCGTAGAGT A0384 IA6-2 proprietary IgD 399907 BioLegend CD72 CAGTCGTGGTAGATA A0419 3F3 proprietary CD72.1 399907 BioLegend CD93 GCGCTACTTCCTTGA A0446 VIMD2 proprietary CD93.1 399907 BioLegend CD85g (ILT7) TGTCAGTTCCTATGA A0409 17G10.2 proprietary CD85g--ILT7 399907 BioLegend CD172a (SIRP) CGTGTTTAACTTGAG A0408 15-414 proprietary CD172a--SIRPa 399907 BioLegend CD284 (TLR4) GCTTAGCTGTATCCG A0405 HTA125 proprietary CD284--TLR4 399907 BioLegend CD304 (Neuropilin-1) GGACTAAGTTTCGTT A0406 12C2 proprietary CD304--Neuropilin-1 399907 BioLegend CD36 TTCTTTGCCTTGCCA A0407 5-271 proprietary CD36.1 399907 BioLegend CD102 (ICAM-2) TGACCTTCCTCCT A0244 CBR-IC2/2 proprietary CD102 399907 BioLegend CGTAACGTAGAGCGA A0224 IP26 proprietary TCR-A--TCR-B 399907 BioLegend TCR/B CD122 (IL-2R?) TCATTTCCTCCGATT A0246 TU27 proprietary N/A 399907 BioLegend CD83 CCACTCATTTCCGGT A0359 HB15e proprietary CD83.1 399907 BioLegend CD267 (TACI) AGTGATGGAGCGAAC A0247 1A1 proprietary CD267--TACI 399907 BioLegend anti-human IgA AAGATGTCCGAGCAA A0186 #N/A proprietary N/A 399907 BioLegend CD107a (LAMP-1) CAGCCCACTGCAATA A0155 H4A3 proprietary CD107a--LAMP-1 399907 BioLegend CD146 CCTTGGATAACATCA A0134 P1H12 proprietary CD146 399907 BioLegend CD140b (PDGFR?) CAATGGTTCACTGCC A0129 18A2 proprietary CD140b 399907 BioLegend CD49a ACTGATGGACTCAGA A0575 TS2/7 proprietary CD49a 399907 BioLegend AACTTCTGTGGTAGC A0584 6B11 proprietary TCR-V-A-24-J-A-18--iNKT cell 399907 BioLegend CD9 GAGTCACCAATCTGC A0579 HI9a proprietary CD9.1 399907 BioLegend TCR V7.2 TACGAGCAGTATTCA A0581 3C10 proprietary TCR-V-A-7.2 399907 BioLegend 9 AAGTGATGGTATCTG A0583 B3 proprietary TCR-V-G-9 399907 BioLegend TCR V CD49d CCATTCAACTTCCGG A0576 9F10 proprietary CD49d 399907 BioLegend C5L2 ACAATTTGTCTGCGA A0572 1D9-M12 proprietary C5L2 399907 BioLegend CD338 (ABCG2) TAAGACTTGGCCGTC A0569 5D3 proprietary CD338--ABCG2 399907 BioLegend CD73 (Ecto-5'-nucleotidase) CAGTTCCTCAGTTCG A0577 AD2 proprietary CD73--Ecto-5-nucleotidase 399907 BioLegend anti-human CD79a (Ig?) CTTATCACCCGCTTT A0578 #N/A proprietary N/A 399907 BioLegend CD158b (KIR2DL2/L3, NKAT2) GACCCGTAGTTTGAT A0592 DX27 proprietary CD158b--KIR2DL2--KIR2DL3--NKAT2 399907 BioLegend CD226 (DNAM-1) AGACCAACTCATTCA A0805 TX25 proprietary CD226--DNAM-1.1 399907 BioLegend CD186 (CXCR6) GACAGTCGATGCAAC A0804 K041E5 proprietary CD186--CXCR6 399907 BioLegend CD158f (KIR2DL5) AAAGTGATGCCACTG A0600 UP-R1 proprietary CD158f--KIR2DL5 399907 BioLegend CD158e1 (KIR3DL1, NKB1) GGACGCTTTCCTTGA A0599 DX9 proprietary CD158e1--KIR3DL1--NKB1 399907 BioLegend CD140a (PDGFR?) ATGCGCCGAGAATTA A0128 16A1 proprietary CD140a 399907 BioLegend CD354 (TREM-1) TAGCCGTTTCCTTTG A0586 TREM-26 proprietary N/A 399907 BioLegend TCR V 2 TCAGTCAGATGGTAT A0582 B6 proprietary TCR-V-D-2 399907 BioLegend CD202b (Tie2/Tek) CGATCCCTTACCTAT A0588 33.1 (Ab33) proprietary CD202b--Tie2--Tek 399907 BioLegend CD96 (TACTILE) TGGCCTATAAATGGT A0175 NK92.39 proprietary CD96--TACTILE 399907 BioLegend CD158 (KIR2DL1/S1/S3/S5) TATCAACCAACGCTT A0420 HP-MA4 proprietary CD158--KIR2DL1--KIR2DS1--KIR2DS3--KIR2DS BioLegend CD337 (NKp30) AAAGTCACTCTGCCG A0801 P30-15 proprietary CD337--NKp30 399907 BioLegend CD253 (Trail) GCCATTCCTGCCTAA A0803 RIK-2 proprietary CD253--TRAIL 399907 BioLegend CD319 (CRACC) AGTATGCCATGTCTT A0830 162.1 proprietary CD319--CRACC 399907 BioLegend CD305 (LAIR1) ATTTCCATTCCCTGT A0590 NKTA255 proprietary CD305--LAIR1 399907 BioLegend CD325 (N-Cadherin) CCTTCCCTTTCCTCT A0433 8C11 proprietary CD325--N-Cadherin 399907 BioLegend mast cell tryptase ACTGATAGACCCGCT A0580 AA1 proprietary N/A 399907 BioLegend CLEC12A CATTAGAGTCTGCCA A0853 50C1 proprietary CD371--CLEC12A 399907 BioLegend CD90 (Thy1) GCATTGTACGATTCA A0060 5E10 proprietary CD90--Thy1 399907 BioLegend CD273 (B7-DC, PD-L2) TCAACGCTTGGCTAG A0008 24F.10C12 proprietary CD273--B7-DC--PD-L2 399907 BioLegend CD272 (BTLA) GTTATTGGACTAAGG A0170 MIH26 proprietary CD272--BTLA 399907 BioLegend CD5 CATTAACGGGATGCC A0138 UCHT2 proprietary CD5.1 399907 BioLegend CD23 TCTGTATAACCGTCT A0897 EBVCS-5 proprietary CD23 399907 BioLegend CD85j (ILT2) CCTTGTGAGGCTATG A0896 GHI/75 proprietary CD85j--ILT2 399907 BioLegend CD94 CTTTCCGGTCCTACA A0867 DX22 proprietary CD94 399907 BioLegend CD27 GCACTCCTGCATGTA A0154 O323 proprietary CD27.1 399907 BioLegend CD328 (Siglec-7) CTTAGCATTTCACTG A0902 6-434 proprietary CD328--Siglec-7 399907 BioLegend CD82 TCCCACTTCCGCTTT A0920 ASL-24 proprietary N/A 399907 BioLegend CD16 AAGTTCACTCTTTGC A0083 3G8 proprietary CD16 399907 BioLegend MERTK TCCTGCATGTACCCA A0423 590H11G1E3 proprietary MERTK.1 399907 BioLegend IgG Fc CTGGAGCGATTAGAA A0375 M1310G05 proprietary IgG-Fc 399907 BioLegend

FCERI CTCGTTTCCGTATCG A0352 AER-37 (CRA-1) proprietary FCERIa 399907 BioLegend CD274 (B7-H1, PD-L1) GTTGTCCGACAATAC A0007 29E.2A3 proprietary CD274--B7-H1--PD-L1 399907 BioLegend CD254 (TRANCE, RANKL) TCCGTGTTAGTTTGT A0356 MIH24 proprietary CD254--TRANCE--RANKL 399907 BioLegend CX3CR1 AGTATCGTCTCTGGG A0179 K0124E1 proprietary CX3CR1.1 399907 BioLegend CD137L (4-1BB Ligand) ATTCGCCTTACGCAA A0022 5F4 proprietary CD137L--4-1BB-Ligand 399907 BioLegend Siglec-8 CTTCTCCTCAGCAAT A0199 7C9 proprietary N/A 399907 BioLegend Validation CITE-seq antibodies from Biolegend have been validated by the manufacturer. CODEX antibodies were validated by Akoya Biosciences, and non-specific antibodies in the panel were excluded from phenotype analysis. MIBI and CyTOF antibodies were validated by their respective manufacturer, and then individually tested and titrated by serial dilution on relevant positive and negative controls by investigators to identify optimal concentrations. Validation of antibodies for CyTOF: https://www.biolegend.com/en-us/soluble-mhc/purified-anti-human-cd45-antibody-710 https://www.biolegend.com/ja-jp/products/purified-anti-human-cd326-epcam-antibody-3755 https://www.bdbiosciences.com/en-us/products/reagents/flow-cytometry-reagents/research-reagents/single-color-antibodies-ruo/ purified-mouse-anti-human-cd7.555359 https://www.biolegend.com/de-de/sean-tuckers-tests/purified-anti-human-cd15-ssea-1-antibody-3699 https://www.biolegend.com/en-gb/cell-health/purified-anti-human-cd3-antibody-867 https://www.biolegend.com/de-at/sean-tuckers-tests/purified-anti-human-cd19-antibody-721 https://www.biolegend.com/de-at/products/purified-anti-human-cd163-antibody-4789?GroupID=BLG10122 https://www.thermofisher.com/antibody/product/CD4-Antibody-clone-RPA-T4-Monoclonal/14-0049-82 https://www.biolegend.com/nl-be/cell-health/purified-anti-human-cd8a-antibody-839 https://www.thermofisher.com/antibody/product/CD11c-Antibody-clone-BU15-Monoclonal/MA1-82142 https://www.biolegend.com/en-us/cell-separation/purified-anti-human-cd14-antibody-797?GroupID=BLG4805 https://www.biolegend.com/en-ie/productstab/purified-anti-human-cd127-il-7ralpha-antibody-7093?GroupID=BLG9274 https://www.thermofisher.com/antibody/product/CD123-Antibody-clone-6H6-Monoclonal/14-1239-82 https://www.biolegend.com/en-ie/lyophilized-control-cells/purified-anti-human-cd45ra-antibody-689 https://www.biolegend.com/de-de/products/purified-anti-human-cd366-tim-3-antibody-6119?GroupID=BLG9937 https://www.biolegend.com/en-gb/soluble-mhc/ultra-leaf-purified-anti-human-tigit-vstm3-antibody-14287 https://www.biolegend.com/en-us/cell-health/purified-anti-human-cd274-b7-h1-pd-l1-antibody-4373 https://www.biolegend.com/nl-nl/products/apc-anti-human-cd49d-antibody-582?GroupID=BLG1167 https://www.biolegend.com/en-gb/productstab/purified-anti-human-cd27-antibody-812?GroupID=BLG10174 https://www.thermofisher.com/antibody/product/CD137-Ligand-4-1BB-Ligand-Antibody-clone-5G11-Monoclonal/14-9056-82 https://www.thermofisher.com/antibody/product/T-bet-Antibody-clone-eBio4B10-4B10-Monoclonal/14-5825-82 https://www.thermofisher.com/antibody/product/CD152-CTLA-4-Antibody-clone-14D3-Monoclonal/14-1529-82 https://www.thermofisher.com/antibody/product/FOXP3-Antibody-clone-PCH101-Monoclonal/14-4776-82 https://www.biolegend.com/de-de/products/purified-anti-human-cd31-antibody-883?Clone=WM59 https://www.biolegend.com/nl-nl/cell-health/purified-anti-human-mouse-integrin-beta7-antibody-2951 https://www.biolegend.com/nl-nl/antibodies-and-more/purified-anti-human-cd141-thrombomodulin-antibody-6106 https://www.biolegend.com/en-gb/clone-search/purified-anti-human-cd197-ccr7-antibody-7471?GroupID=BLG9610 https://www.biolegend.com/en-gb/search-results/purified-anti-human-ki-67-antibody-6967 https://www.biolegend.com/en-gb/cell-health/purified-anti-human-cd25-antibody-8323?GroupID=BLG10308 https://www.biolegend.com/en-gb/cell-health/purified-anti-human-cd38-antibody-748 https://www.biolegend.com/nl-nl/products/purified-anti-human-cd1c-antibody-4836?Clone=L161 https://www.biolegend.com/en-gb/cell-health/purified-anti-human-mouse-rat-cd278-icos-antibody-2477?GroupID=BLG3831 https://www.biolegend.com/en-us/soluble-mhc/purified-anti-human-hla-dr-antibody-792?GroupID=BLG11943 https://www.biolegend.com/fr-lu/cell-health/purified-anti-human-cd279-pd-1-antibody-4410 https://www.bdbiosciences.com/en-us/products/reagents/flow-cytometry-reagents/research-reagents/single-color-antibodies-ruo/ purified-mouse-anti-human-cd56.559043 https://www.biolegend.com/fr-lu/products/purified-anti-human-cd16-antibody-571?GroupID=BLG8436 Validation antibodies for MIBI: https://www.abcam.com/products/primary-antibodies/ds-dna-antibody-35i9-dna-bsa-and-azide-free-ab27156.html https://www.abcam.com/products/primary-antibodies/e-cadherin-antibody-36e-cadherin-bsa-and-azide-free-ab287971.html https://www.ionpath.com/histone-h3-antibody/ https://www.bdbiosciences.com/en-us/products/reagents/flow-cytometry-reagents/research-reagents/single-color-antibodies-ruo/ alexa-fluor-488-mouse-anti-human-foxp3.561181 https://www.thermofisher.com/antibody/product/Alexa-Fluor-488-Antibody-Polyclonal/A-11094 https://www.abcam.com/products/primary-antibodies/cd163-antibody-epr14643-36-low-endotoxin-azide-free-ab215976.html https://www.ionpath.com/cd4-antibody/ https://www.ionpath.com/cd11c-antibody/ https://www.cellsignal.com/products/primary-antibodies/cd14-d7a2t-rabbit-mab-bsa-and-azide-free/43878 https://www.cellsignal.com/products/primary-antibodies/cd16-d1n9l-rabbit-mab-bsa-and-azide-free/72204 https://www.novusbio.com/products/lag-3-antibody-17b4_nbp1-97657 https://www.ionpath.com/pd-1-antibody/ https://www.ionpath.com/pd-l1-antibody/ https://www.ionpath.com/granzyme-b-d6e9w-antibody/ https://www.ionpath.com/cd56-antibody-151eu/

https://www.ionpath.com/cd31-antibody-152sm/

https://www.cellsignal.com/products/primary-antibodies/ki-67-d2h10-rabbit-mab-bsa-and-azide-free/44092

https://www.ionpath.com/cd117-antibody-155gd/

https://www.ionpath.com/cd68-antibody/

https://www.abcam.com/en-za/products/primary-antibodies/cd103-antibody-epr41662-bsa-and-azide-free-ab271889 https://www.ionpath.com/cd8-antibody-158gd/

www.ionpath.com/wp-content/uploads/2021/03/TDS-715901_B-CD3e_D7A6E_159Tb_Human_FFPE.pdf https://www.ionpath.com/cd45ro-antibody-161dv/

https://www.ionpath.com/wp-content/uploads/2021/03/TDS-716201_B-TIM-3_EPR22241_162Dy_Human_FFPE.pdf https://www.ionpath.com/vimentin-antibody/

https://www.abcam.com/products/primary-antibodies/cd27-antibody-epr8569-bsa-and-azide-free-ab256583.html https://www.ionpath.com/keratin-antibody-165ho/

https://www.abcam.com/en-in/products/primary-antibodies/alpha-smooth-muscle-actin-antibody-sp171-bsa-and-azide-free-ab242395

https://www.ionpath.com/cd20-antibody-167er/

https://www.abcam.com/products/primary-antibodies/cd11b-antibody-ep1345y-bsa-and-azide-free-ab187537.html

https://www.cellsignal.com/products/primary-antibodies/cd141-thrombomodulin-e7y9p-xp-rabbit-mab-bsa-and-azide-free/34149 https://www.ionpath.com/cd21-antibody-170er/

https://www.ionpath.com/ido1-antibody-171yb/

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Plants

Seed stocks	Report on the source of all seed stocks or other plant material used. If applicable, state the seed stock centre and catalogue number. If plant specimens were collected from the field, describe the collection location, date and sampling procedures.
Novel plant genotypes	Describe the methods by which all novel plant genotypes were produced. This includes those generated by transgenic approaches, gene editing, chemical/radiation-based mutagenesis and hybridization. For transgenic lines, describe the transformation method, the number of independent lines analyzed and the generation upon which experiments were performed. For gene-edited lines, describe the editor used, the endogenous sequence targeted for editing, the targeting guide RNA sequence (if applicable) and how the editor
Authentication	was applied. Describe any authentication procedures for each seed stock used or novel genotype generated. Describe any experiments used to assess the effect of a mutation and, where applicable, how potential secondary effects (e.g. second site T-DNA insertions, mosiacism, off-target gene editing) were examined.

Flow Cytometry

Plots

Confirm that:

X The axis labels state the marker and fluorochrome used (e.g. CD4-FITC).

X The axis scales are clearly visible. Include numbers along axes only for bottom left plot of group (a 'group' is an analysis of identical markers).

X All plots are contour plots with outliers or pseudocolor plots.

X A numerical value for number of cells or percentage (with statistics) is provided.

Methodology

Sample preparation	A 37-parameter CyTOF panel was designed (Supplementary Table 3). All mass cytometry antibodies were conjugated in- house to their corresponding metal isotope. Metals were conjugated according to the manufacturer's instructions (Fluidigm, South San Francisco, CA, USA). In brief, this process consisted of loading the metal to a polymer for 1h at RT. The unconjugated antibody is transferred into a 50 kDA Amicon Ultra 500 V-bottom filter (Fisher Scientific, Hampton, NH, USA) and reduced for 30 min at 37°C with 1:125 dilution of Tris (2-carboxyethyl) phosphine hydrochloride (TCEP) (ThermoFisher, Waltham, MA, USA). Subsequently, the column was washed twice with C-buffer (Fluidigm) and the metal-loaded polymer was suspended in 200 µL of C-buffer in the 3 kDA Amicon Ultra 500 mLV-bottom filter. The suspension was transferred to the 50 kDA filter containing the antibody and incubated for 1.5h at 37°C. After incubation, antibodies were washed three times with W-buffer (Fluidigm) and quantified for protein content using Nanodrop. Once the concentration was determined, the antibodies were resuspended at a concentration of 0.2 mg/mL with Antibody Stabilizer (Boca Scientific, Dedham, Ma, USA) and stored at 4°C. Optimal concentrations for all antibodies were determined by different rounds of titrations. The staining protocol was optimized to use each antibody in aliquots of 6 million cells as previously described. CyTOF was performed on paired peripheral blood and colon biopsies obtained from the primary case control study (n=12) and from the secondary remission case control study (n=17). Single-cell suspensions of colon biopsies from the right and left colon were pooled to ensure sufficient cell counts per donor. Dead cells were labeled with Cisplatin (Fluidigm) according to the manufacturer's instructions, washed in wash buffer (PBS, 0.5% BSA, 10% DMSO) and stored at -80oC until staining. Samples were then thawed and washed with wash buffer. Prior to staining with the antibody panel (Supplementary Table 3), cells from each
	staining was performed following the manufacturer's institutions (Holding 1, South San Handsco, CA, OSA). Dieny, each sample was incubated for 15 min at RT on a shaker (200rpm) with a barcoding solution containing 10 µL of barcode in 1x Perm Buffer solution (Fluidigm, Cat#201068). Samples were then washed, centrifuged, resuspended in CSM, and pooled. Subsequently, extracellular staining was performed for 30 min at 4°C. After incubation, the samples were washed with CSM and centrifuged before resuspending in 1x Permeabilization Buffer (eBioscience™ Permeabilization Buffer Cat# 00-8333-56) for 10min at 4°C. The samples were then washed and incubated with Ir-intercalator (Biolegend CNS, San Diego, CA, USA) diluted 1:500 in 4% fresh PFA for 20 min at RT. After incubation, samples were washed and kept at 4°C overnight in EQTM bead solution (Fluidigm Cat#201078) diluted in MaxPar Water (Fluidigm Cat# 201069) at 1.2x106 cells/mL. Samples were analyzed on the CyTOF®2 instrument (Fluidigm).
Instrument	CyTOF®2
Software	CyTOF data were concatenated, normalized and de-barcoded using CyTOF software (Fluidigm); supervised analysis using FlowJo software (10.8.1); unsupervised analysis using an R-based Cytometry Clustering Optimization aNd Evaluation (Cyclone) pipeline developed by the UCSF Data Science CoLab (https://github.com/UCSF-DSCOLAB/cyclone).
Cell population abundance	See Supplemental Table 6
Gating strategy	See Supplemental Table 4 and Extended Figure 10

x Tick this box to confirm that a figure exemplifying the gating strategy is provided in the Supplementary Information.