# Immunophenotyping UCDLA\_IMM\_002

## **Purpose**

This test differentiates immune cell sub-populations via flow cytometry.

Description: increased CD4-positive T cell number (MP:0008074), decreased CD4-positive T cell number (MP:0008075), etc., ...

# **Experimental Design**

• Minimum number: 3M + 3F

Age at test: Week 59

• **Sex:** Both (sexually dimorphic)

# **Equipment**

### **Equipment**

- Scissors and forceps for biopsy
- Precision balance
- Calibrated single and multichannel pipettes
- Plate shaker
- Refrigerated centrifuge
- Flow Cytometer (capable of distinguishing a minimum of 8 colours per well)
- Tissue dissociator:
  - GentleMACS tissue dissociator OR
  - Equipment for manual dissociation
- Cell counter equipment:
  - Orflo Moxi-Z Cell counter OR
  - Coulter Vicell XR OR Life Technologies Attune® Flow Cytometer OR
  - Haemocytometer

### **Supplies**

- 96-well V-bottomed plates (Falcon #353263)
- Petri dishes
- Dispensing troughs
- Low retention pipette tips for antibody solutions
- (if using GentleMACS for dissociation) C Tubes. It is acceptable to re-use these once.
- 50ml Falcon tubes
- Cell strainers e.g. 70m cell strainers that fit 50ml Falcon tubes (BD Falcon, #352350) OR 70-80µM Nytex
- Cell counter recipients (i.e., slides/cassettes/etc. for cell counter)
- (if sample processing delayed) RPMI 1640

- (if sample processing on same day) HBSS, with phenol red
- CS (calf serum)
- PBS with Mg<sup>2+</sup>, with Ca<sup>2+</sup> (for <u>enzyme buffer</u> used for DNAse and Collagenase D digestions)
- PBS without Mg<sup>2+</sup>, without Ca<sup>2+</sup> (for <u>FACS buffer</u> to be used in all steps subsequent to enzymatic digest)
- EDTA (0.5M stock; final concentration 2mM)
- Digestion enzyme (Collagenase D from Roche #11088858001), stock solution in enzym e buffer (see below), aliquoted and stored at -2 0°C
- DNAse I stock solution (Sigma, #DN25) in enzyme buffer (see below), aliquoted and stored at -20°C
- RBC lysis buffer (eBioscience #00-4300-54 or BD Biosciences #555899, both 10X from manufacturer)
- HEPES (pH 7.2-7.4)

### **Procedure**

This protocol requires several steps in the collection, preparation and analysis of the samples. Each one is detailed separately below.

### Reagent preparation

Note that two different PBS solutions are required for the protocol below, one with  $Ca^{2+}$  and  $Mg^{2+}$ , another without  $Ca^{2+}$  and  $Mg^{2+}$ .

- Collection buffer:
  - (*if spleens are to be processed on the same day*) HBSS with Ca<sup>2+</sup> and Mg<sup>2+</sup> and phenol red (e.g. Life Technologies 14170161) *OR*
  - (*if analysis will be delayed*) RPMI medium with 2% CS added.
- FACS buffer (for all steps subsequent to enzymatic digest; stable for up to 1 month in the fridge):
  - PBS 1X without Ca<sup>2+</sup>/Mg<sup>2+</sup> OR
  - HBSS 1X without Ca<sup>2+</sup>/Mg<sup>2+</sup>
  - EDTA 2mM
  - 2% (v/v) CS
  - 10mM HEPES, pH 7.2-7.4
- **Brilliant Stain Buffer** (BD 563794; for all steps when two or more brilliant violet antibodies are used to prevent non-specific dye-to-dye interaction)
- Enzyme buffer (for DNAse and Collagenase D digestions; Stable for up to 1 month in the fridge):
  - PBS with Ca<sup>2+</sup> and Mg<sup>2+</sup> OR
  - HBSS 1X with Ca<sup>2+</sup>/Mg<sup>2+</sup>
  - 2% (v/v) CS:
  - 10mM HEPES, pH 7.2-7.4
- RBC Lysis buffer: Prepare a 1X solution in ddH<sub>2</sub>0 from 10X stock lysis buffer.
- **Stopping buffer** (require 300 µl per sample):
  - 1x PBS without Ca<sup>2+</sup> and Mg<sup>2+</sup> or 1X HBSS without Ca<sup>2+</sup> and Mg<sup>2+</sup>
  - 0.1 M EDTA (37.5 g/L)

### Antibody cocktails for Panels 1 & 2

- Protect antibodies and prepared cocktails from direct light.
- Final concentration of antibodies should be determined by titration to ensure saturating amounts of antibody are used. Appropriate amounts of antibodies can be mixed together from the manufacturer's stock solutions and stored for 1 week at 4°C prior to dilution in FACS buffer immediately before use. Do NOT pre-mix BV antibodies. These should be added fresh to the diluted staining mixure.
- Each sample will require 50 μl (or up to 100 μl) of diluted 1X antibody cocktail.
- Antibody cocktails should be gently but thoroughly mixed to ensure homogeneity of the solutions.
- In order to eliminate aggregated antibodies from your mix, centrifuge each antibody cocktail for 8 min at 20,000xg and 8°C prior to staining cells.

### Antibody Panels

 Recommended antibody (marker) panels, Panel A for T, NKT and NK cells, Panel B for B, myeloid and NK cells are shown below, along with optional markers that may be used by some centres. Core antibodies are required for upload of data; optional markers are not and are listed in alphabetical order. Clones and fluorochromes used should be uploaded for required and optional markers. Where not indicated, clone and fluorochrome choice is dependent on available detectors and filters on the cytometer used at each centre.

### Panel A

Туре	Antibody (Marker)	Clone	Fluorochrome
Required	CD5	53-7.3	BV421
Required	CD4	RM4-5	FITC
Required	CD44	IM7	PE
Required	CD8a	53-6.7	PE-CF594
Required	CD25	PC61	PE-Cy7
Required	CD161	PK136	APC
Required	CD62L	MEL-14	APC-Cy7
Required	Live/Dead	-	SytoxBlue

Optional	CD3e	145-2C11	
Optional	CD24	M1/69	
Optional	CD27	LG.3A10	
Optional	CD357/GITR	DTA-1	
Optional	CD45	30-F11	
Optional	KLRG1	2F1	
Optional	Ly6c	AL-21	
Optional	TCRd	GL-3	

### Panel B

Туре	Antibody (Marker)	Clone	Fluorochrome
Required	CD5	53-7.3	BV421
Required	Ly6G	1A3	BV421
Required	CD19	1D3	BV510
Required	Ly6C	AL-21	FITC
Required	CD21/CD35	7G6	PE
Required	CD11b	M1/70	PE-CF594

Required	CD11c	HL3	PE-Cy7
Required	CD161	PK136	APC
Required	MHCII	M5/114.15.2	APC-Cy7 or A700
Required	Live/Dead	-	SytoxBlue
Optional	CD23	B3B4	
Optional	CD27	LG3.A10	
Optional	CD43	S7	
Optional	CD44		
Optional	CD45	30-F11	
Optional	CD317	927	
Optional	F4/80	BM8	
Optional	IgD		
Optional	KLRG	2F1	

### • Read buffer / dead cell exclusion dye

- SytoxBlue at 1:10000 concentration in FACS buffer *OR*
- SytoxGreen at 1:20000 concentration in FACS buffer
- Zombie Near Infra-Red live dead from Biolegend at 1:2000 concentration
- Require 200 I per well (i.e. 400 I for each spleen).
- Enzyme cocktail (working solution): 3 ml for each spleen, containing final concentrations of:
  - DNAse I: 30-100 g (from 10 mg/ml stock in enzyme buffer stored in single experiment aliquots at -20°C, do not freeze-thaw stock)

 Collagenase D: 600 Mandl Units (from 30 U/µl stock in enzyme buffer stored in single experiment aliquots at -20°C, do not freeze-thaw stock)

**NOTE:** To top up to the 3ml use enzyme buffer; any intermediate dilutions of the enzyme stock solutions should be prepared with <u>enzyme buffer</u>.

### Other preparations on the day

- Bring RBC lysis buffer and stop solution to room temperature.
- Prepare wet ice box, label tubes, etc.

### Note all centrifuge steps are: 5 min, 400 x g at 8°C

### Spleen collection

- Collect the spleen from euthanized mice.
- Remove all fat from the spleen and weigh the organ on a petri dish (do not hydrate the organ before weighing it as this would lead to substantial errors in measurement).
- Place the spleen in a 1.5ml eppendorf tube with 1 mL of sample collection buffer on ice.
   Use:
  - (if spleens are to be processed on the same day) HBSS without calcium, without magnesium but with phenol red OR
  - (if analysis will be delayed) RPMI with 2% CS buffer.

### Spleen dissociation / digests

### If using a GentleMacs tissue dissociator:

- Add the spleen to a GentleMACS C tube containing 3 ml of 1X enzyme cocktail.
- Clip the tube on GentleMACS dissociator and run programme spleen\_2.
- Incubate cell suspension for 30 minutes with gentle mixing at least every 5 minutes. Register incubation temperature.
- Run programme spleen 3.
- Add 300 L of stopping buffer and mix by inversion to block enzymatic digestion and dissociate T cell-dendritic cell interactions.
- Filter cell suspension:
  - through 70-80 m Nylon mesh filter into a 50 mL Falcon tube OR
  - directly from C-tubes pour splenocyte suspension through 30  $\mu$ m CellTrics Partec filters (#04-0042-2316) into 15 ml tubes.
- (optional) Wash the GentleMACS C tube with 5ml <u>FACS buffer</u>, filter and pool with flow-through from previous step.
- Centrifuge for 5 minutes, 400 x g at 8°C and discard supernatant.
- Resuspend total splenocytes in 1 mL cold <u>FACS buffer</u> and keep on ice (this step is not required if counting is performed on the attune).

### OR, if performing manual digests:

- Place weighed spleen in 12x75mm tube containing 1ml of collagenase solution in 1X HBSS with Ca<sup>2+</sup> and Mg<sup>2+</sup> (17-0.2 Wünsch unit/ml)
- Mince into fine pieces using small scissors, place on ice until all samples are minced.

- Add 2ml collagenase (17-0.2 Wünsch unit/ml) to each tube and place in a 37°C water bath for 30 minutes.
- Tricturate (pipetting vigorously up and down using a 1 mL pipetman) the mixture to break up clumps.
- Spin at 500 x g in a swing bucket rotor for 5 min at 10°C. Decant the supernatant, rack the tubes or vortex to resuspend the pellet. Add 2ml <u>FACS buffer</u>, mix well by vortexing, take 10 µl for the counting step.
- Dilutions for counting: 2 serial 1:10 dilutions (10μl cells + 90μl <u>FACS buffer</u>, then 10μl of the 1:10 dilution + 90μl buffer.)
- Spin for 5min, 500 x g at 10°C, decant supernatant, blot the top of the tube, resuspend pellet at 1x10<sup>8</sup> cells/ml.

### Cell counting

- Perform a cell count on an aliquot of the re-suspended cells (adjust concentration according to the cell counter method used).
- Note the cell count, correct for dilution and calculate the concentration in cells per µl.
- Cell count:
  - If performed before RBC lysis, pipette the volume containing approximately 4
    million cells/well to a 96 well plate in horizontal fashion starting from A1 onwards
    for panel 1 staining.
  - <u>If performed after RBC lysis</u>, pipette the volume containing approximately 1-2 million cells/well to a 96 well plate in horizontal fashion starting from A1 onwards for panel 1 staining.
- Do the same for panel 2 staining in separate wells leaving a few empty rows between the panels to avoid cross contamination.
- Top up to final volume of 100  $\mu$ l using <u>FACS buffer</u>, centrifuge, discard supernatant and keep plate on wet ice.

### Red blood cell lysis, blocking & staining

- Remove plate from ice and add 30 to 100 μl of 1X RBC lysis buffer (at room temperature) to each cell pellet from the previous step.
- Pipette up and down 2-3 times to break up the pellet and ensure complete lysis. Alternatively, vortex the edges of the plates, then pipet quickly once to ensure resuspension is ideal for optimal lysis.
- Incubate for 1 minute at room temperature and then return to ice and add 100 to 200  $\mu$ l of <u>FACS buffer</u> (to stop lysis) to each well.

**Note**: Following RBC lysis, every centrifugation step can be performed at 2000rpm for 1 minute in a 96 well plate, which significantly speeds up the protocol. Do take care to resuspend the cells very well to prevent HTS clumping.

- Centrifuge, discard supernatant and resuspend in 200 μl <u>FACS buffer</u> (this step is not required if lysis was performed in 30 μl, since there will be enough volume left in the well for a bigger wash of 200 μl; saves time on a spin).
- Again centrifuge and discard supernatant and resuspend in 50 μl of 1:100 Fc block and incubate on ice for 10 min. Top up to 200 μl using <u>FACS buffer</u> after incubation.

- Take antibody (AB) cocktails from the fridge. In order to eliminate aggregated ABs from your mix before use, centrifuge each AB cocktail for 8 min at 20,000 x g and 4°C. Dilute antibody cocktail to final working concentration with FACS buffer, or Brilliant stain buffer when two or more brilliant violet antibodies are used, to make the AB mix.
- Centrifuge plate, discard supernatant and resuspend in 50 to 100 μl 1X AB mix in appropriate wells for individual panels followed by incubation on ice and in the dark for 20 min.
- If using Sytox Blue/Sytox Green as live/dead discriminator:
  - Top up to 200 μl with <u>FACS or Brilliant Stain buffer</u> after incubation. Centrifuge, discard supernatant and resuspend in 200 μl <u>FACS or Brilliant Stain buffer</u>.
  - When ready to read plate, centrifuge again and discard supernatant. Resuspend the pellet in 200 μl of read buffer (Sytox Blue diluted 1:10000 in <u>FACS buffer</u>; Sytox Green diluted 1:20000 in <u>FACS buffer</u>).
- If using Zombie NIR dye as live/dead discriminator:
  - Add 200 μl of PBS (RT) to all samples
  - Spin at 2000 rpm for 1 minute 8°C
  - Add 100 μl/well of Zombie Near-IR Live/Dead dye (1/2000) made up in PBS incubate at room temperature for 10 mins, add 200 μl FACS buffer.

### **General Recommendations for Setting up Cytometer**

Set up the analyser to aim acquire 300,000 viable events (live cells) for each of Panels 1 and 2. 500,000 are recommended for panel 2 in order to increase robustness of myeloid population assessment for low frequency populations (macrophages, DCs).

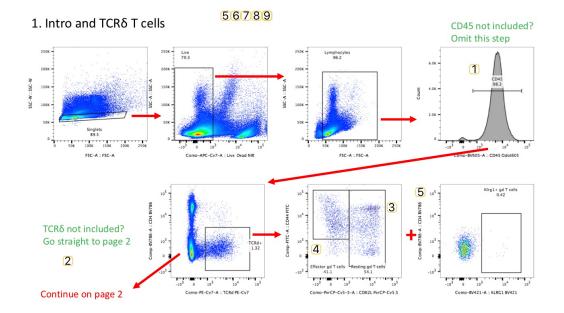
### **Notes**

**Visual help for Gating** 



### **APPENDIX 1. GATING HIERARCHIES**

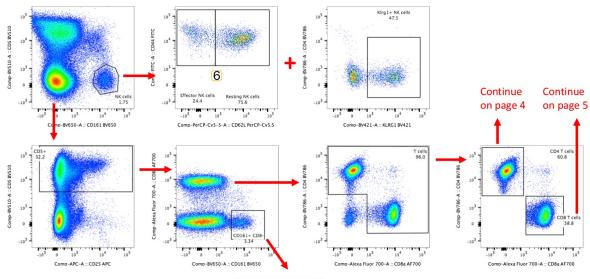
### Panel A. Page 1





### Panel A. Page 2.

### 2. NK cells and further gating

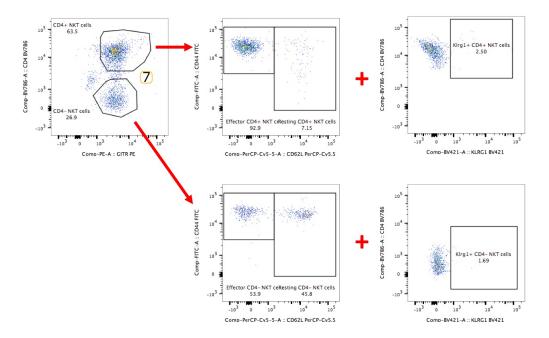


Continue on page 3



### Panel A. Page 3.

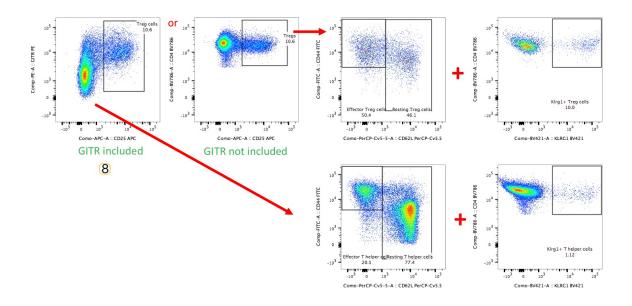
### 3. NKT cells





### Panel A. Page 4.

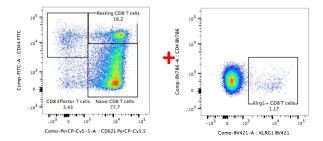
### 4. Tregs and T helper cells





### Panel A. Page 5.

### 5. CD8 T cells and notes

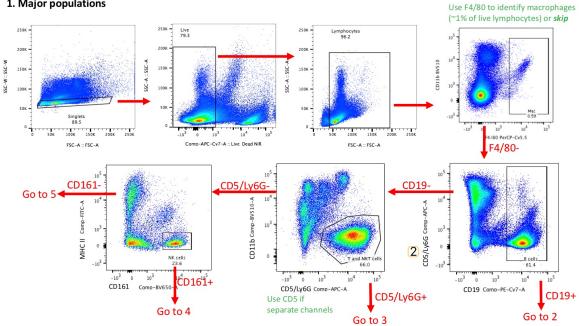


- 1 If there is no CD45 in the panel, omit this step.
- 2 Approximately 50% of γδ T cells are CD5-, so they will drop out when gating on CD5 later on. Of the remaining cells, approximately 90% are CD4- CD8- and will drop out of the T cell gate. Only 5% (approximately 0.2% of lymphocytes) will end up in the CD8 T cell gate which is negligible.
- 3 Please note that each cell type requires different thresholds for both CD44 and CD62L.
- CD44- CD62L- cells do not occur naturally and show up when CD62L is shed from resting cells during sample preparation.
- I have chosen CD4 for the y axis because gives a nice compact population for almost all cell types which makes it easy to see the Klrg1+ cells. However, if CD4 doesn't work for you because of your fluorochrome combinations, it can be substituted by any other marker.
- The name effector is fine for CD4 and CD8 T cells, it is a bit unusual for yδ T cells, NKT cells and NK cells. We settled for this term in the end and also added these population names (with a more detailed description) to the MGI ontology, so MP terms that we use now carry these names.
- These need to be added up to give the counts of total NKT cells. Use any fluorochrome on the y axis that gates out the non-specific autofluorescent population between the two distinct populations
- 8 If you don't have GITR, use CD4 on the y axis instead. It works almost as well.



### Panel B. Page 1.

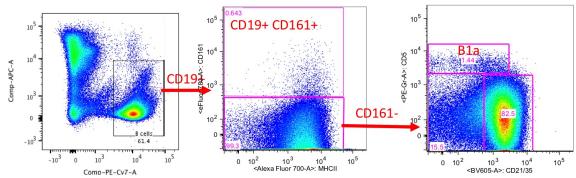
### 1. Major populations





Panel B. Page 2A.

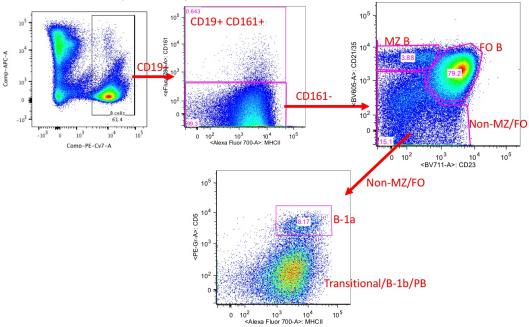
### 2A. B cells – core panel only (CD21/35 & CD5/Ly6G)





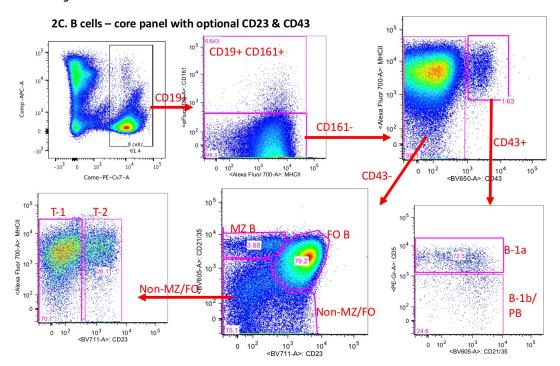
Panel B. Page 2B.







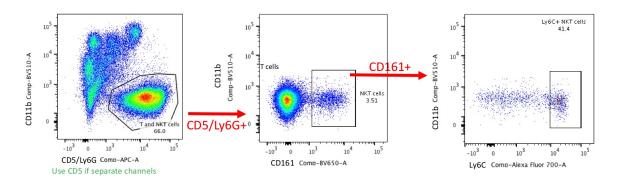
### Panel B. Page 2C.





### Panel B. Page 3.

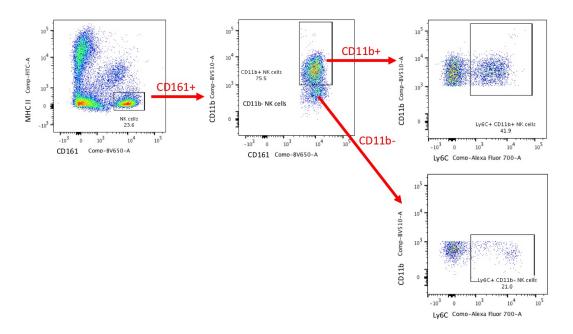
### 3. T cells and NK T cells





### Panel B. Page 4.

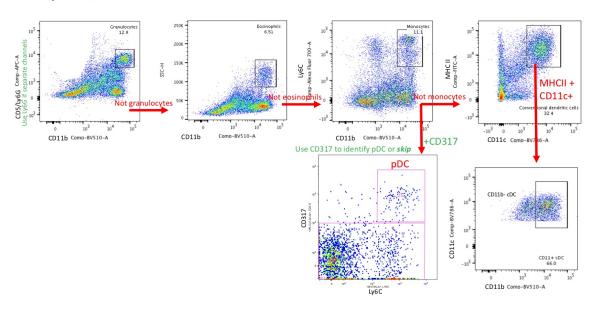
### 4. NK cells





### Panel B. Page 5.

### 5. Myeloid cells



### **Parameters and Metadata**

# Spleen weight UCDLA\_IMM\_001\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

\_\_\_\_\_

# T cells (Panel A) - % of live leukocytes (Panel A) UCDLA\_IMM\_

003\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: true

**Unit Measured:** %

.....

# NKT cells (panel A) - % of live leukocytes (Panel A) UCDLA\_I

MM\_004\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: true

**Unit Measured:** %

NK cells (Panel A) - % of live leukocytes (Panel A) UCDLA\_IM

M\_005\_002 | v2.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true
Unit Measured: %		
CD4 T cells - % of I	live leukocytes (Par	nel A) ucdla_imm_007_002
simpler arameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Unit Measured: %		
CD8+ T cells - % of	live leukocytes (Pa	anel A) ucdla_imm_008_
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Unit Measured: %		

# CD4+ NKT cells - % of live leukocytes (Panel A) UCDLA\_IMM\_0

11\_002 | v2.0

simpleParameter

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Unit Measured: %		
CD4- NKT cells - %	of live leukocytes	(Panel A) UCDLA_IMM_0
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
Treg cells - % of liv	e leukocytes (Pane	A) UCDLA_IMM_014_002
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Unit Measured: %		

CD4+ T helper cells - % of live leukocytes (Panel A) UCDLA\_I

MM\_015\_002 | v2.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true
Unit Measured: %		
Total events (Pane simpleParameter	<b>I A)</b> UCDLA_IMM_026_002	v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Total events (Pane simpleParameter	IB) UCDLA_IMM_027_002	v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Effector CD4+ T he A) UCDLA_IMM_028_002 simpleParameter	•	e leukocytes (Panel
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Unit Measured: %		

# Resting CD4+ T helper cells - % of live leukocytes (Panel A) UCDLA\_IMM\_029\_002 | v2.0

simpleParameter

Reg. Analysis: false Reg. Upload: false Is Annotated: true Unit Measured: % Effector CD8+ T cells - % of live leukocytes (Panel A) UCDL A\_IMM\_032\_002 | v2.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true Unit Measured: % NaÃÂ-ve CD8+ T cells - % of live leukocytes (Panel A) UCDL A\_IMM\_033\_002 | v2.0 simpleParameter Is Annotated: true Req. Analysis: false Req. Upload: false **Unit Measured:** %

# Resting CD8+ T cells - % of live leukocytes (Panel A) UCDLA

\_IMM\_034\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: true
Unit Measured: %

# Effector CD4+ NKT cells - % of live leukocytes (Panel A) U

CDLA\_IMM\_040\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: true

**Unit Measured:** %

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# Resting CD4+ NKT cells - % of live leukocytes (Panel A) UC

DLA\_IMM\_041\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

# Effector CD4- NKT cells - % of live leukocytes (Panel A) UC

DLA\_IMM\_046\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false
Unit Measured: %

# Resting CD4- NKT cells - % of live leukocytes (Panel A) UC

DLA\_IMM\_047\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

**Unit Measured:** %

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# Live leukocytes (Panel B) - % of total events (Panel B) UCD

LA\_IMM\_049\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

# Granulocytes - % of live leukocytes (Panel B) UCDLA\_IMM\_050 \_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false Unit Measured: % Monocytes - % of live leukocytes (Panel B) UCDLA\_IMM\_051\_002 | v2.0 simpleParameter Req. Analysis: false Req. Upload: true Is Annotated: false Unit Measured: % Eosinophils - % of live leukocytes (Panel B) UCDLA\_IMM\_052\_0 02 | v2.0 simpleParameter Req. Analysis: false Req. Upload: true Is Annotated: false **Unit Measured:** %

# NK cells (Panel B) - % of live leukocytes (Panel B) UCDLA\_IM

M\_053\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

Unit Measured: %

Ly6C+ CD11b- NK cells - % of live leukocytes (Panel B) uc

simpleParameter

DLA\_IMM\_054\_002 | v2.0

Req. Analysis: false Req. Upload: true Is Annotated: false

Unit Measured: %

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# Ly6C+ CD11b+ NK cells - % of live leukocytes (Panel B) UC

DLA\_IMM\_055\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

# CD11b- NK cells - % of live leukocytes (Panel B) UCDLA\_IMM\_

056\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false
Unit Measured: %

# CD11b+ NK cells - % of live leukocytes (Panel B) UCDLA\_IMM

\_057\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

Unit Measured: %

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# NKT cells (panel B) - % of live leukocytes (Panel B) UCDLA\_I

MM\_058\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

# Ly6C+ NKT cells - % of live leukocytes (Panel B) UCDLA\_IMM\_ 059\_002 | v2.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: %

# T cells (panel B) - % of live leukocytes (Panel B) UCDLA\_IMM\_

061\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

Unit Measured: %

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# B cells - % of live leukocytes (Panel B) UCDLA\_IMM\_063\_002 | v2.

0

simpleParameter

Req. Analysis: false Req. Upload: true Is Annotated: false

.0 simpleParameter		
Req. Analysis: false	Req. Upload: true	Is Annotated: false
Unit Measured: %		
Marginal zone B c 002   v2.0 simpleParameter	ells - % of B cells (F	Panel B) UCDLA_IMM_071_
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
Conventional DC072_002   v2.0 simpleParameter	· % of live leukocyte	es (Panel B) ucdla_imm
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		

Follicular B cells - % of B cells (Panel B) UCDLA\_IMM\_067\_002 | v2

# Plasmacytoid DC- % of live leukocytes (Panel B) UCDLA\_IMM

\_074\_002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

.....

# Macrophages- % of live leukocytes (Panel B) UCDLA\_IMM\_075\_

002 | v2.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

.....

### Equipment name UCDLA\_IMM\_077\_002 | v2.0

procedureMetadata

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Options: FACS, Flow cytometer, Fortessa\_1, LSR II, Fortessa\_I Custom Build,

\_\_\_\_\_

# Equipment manufacturer UCDLA\_IMM\_078\_002 | v2.0

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: BD Biosciences, Beckman Coulter, IntelliCyt, Cytek,

# Equipment model UCDLA\_IMM\_079\_002 | v2.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: BD LSR-II, BD LSRFortessa Cell Analyzer, CANTO-II, FACSAria III, Gallios,

H47100123, iQue Screener PLUS, Aurora,

.....

# CS&T Bead lot UCDLA\_IMM\_080\_002 | v2.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

# Anesthesia UCDLA\_IMM\_081\_002 | v2.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: Injection narcosis with Ketamine (100mg/kg)/Xylazine (10mg/kg),

Injection narcosis with Sodium Pentobarbital (Somnopentyl),

Injection narcosis with Tribromoethanol (Avertin), Isoflurane, none, Injection narcosis with Medetomidine/Midazolam/Butorphanol,			
Cell digestion UCDL procedureMetadata	_A_IMM_082_002   v2.0		
	Req. Upload: false		
Cell digestion age procedureMetadata	<b>nt</b> ucdla_imm_083_002	v2.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Cell digestion age procedureMetadata	nt manufacturer ∪ci	DLA_IMM_084_002   v2.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false	

# Cell digestion agent catalog number UCDLA\_IMM\_085\_002 | v2.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Cell counting performance procedure Metadata	ormed ucdla_imm_086_i	002   v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Cell counting equi	oment manufacture	<b>I</b> UCDLA_IMM_087_002   v2.
0 procedureMetadata		• 00BE/(_IWIWI_00/_002   V2.
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Cell counting equiporcedureMetadata	oment model ucdla_	IMM_088_002   v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Cell counting equiports procedureMetadata	oment name ucdla_ii	MM_089_002   v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false

Options: Cellometer,

Cell lysis buffer manufacturer UCDLA\_IMM\_090\_002 | v2.0 procedureMetadata Reg. Analysis: false Reg. Upload: false Is Annotated: false Options: BD PharmLyse, eBioscience, Jax, JMC, LONZA, In house, Cell lysis buffer catalog number UCDLA\_IMM\_091\_002 | v2.0 procedureMetadata Reg. Analysis: false Reg. Upload: false Is Annotated: false Date and time of sacrifice UCDLA\_IMM\_092\_002 | v2.0 procedureMetadata Reg. Analysis: false Reg. Upload: false Is Annotated: false

#### Date and time of sample preparation UCDLA\_IMM\_093\_002 | v2.0

procedureMetadata

Sample storage ter LA_IMM_094_002   v2.0 procedureMetadata	nperature until ana	lysis (in Celsius) ucd
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: C		
FCS repository referencedureMetadata	erence (URL/ID) ucd	LA_IMM_095_002   v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Balanced salt solut procedureMetadata	tion type ucdla_imm_0	096_002   v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: HBSS, PBS, KDS B	SS,	

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Balanced salt solution  0  procedureMetadata	tion catalog numbe	<b>I</b> UCDLA_IMM_098_002   v2.
Req. Analysis: false	Req. Upload: false	Is Annotated: false
RPMI manufacture procedureMetadata	<b>r</b> ucdla_imm_099_002   v2	2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: Gibco, Jax, Life Tec	hnologies, none used, Sigma, \	Wako, Thermo Fisher Scientific,
RPMI catalog numl	ber ucdla_imm_100_002	v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false

Req. Analysis: false Req. Upload: false Is Annotated: false Options: Sigma, Spleen Dissociation Kit, N/A, DNAse I catalog number UCDLA\_IMM\_102\_002 | v2.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Dead cell exclusion dye UCDLA\_IMM\_103\_002 | v2.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Options: DAPI, Propidium Iodide, Sytox Blue, Sytox Green, Zombie NIR, LIVE/DEAD Fixable Aqua stain, Ghost Dye UV450, Trypan Blue,

### Dead cell exclusion dye manufacturer UCDLA\_IMM\_104\_002 | v2.0

procedureMetadata

<b>Options:</b> Biolegend, home brew, Life Technologies, Sigma, Invitrogen by Thermo Fisher, Tonbo biosciences,			
	n dye catalog numb	<b>DET</b> UCDLA_IMM_105_002   v2	
procedureMetadata			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
<b>Options:</b> 423106, D9542, hon 13-0868-T500, T8154-100ML,		1348, S34857, P4170, L34966,	
Cell digestion temp 0 procedureMetadata	perature (in Celsius	<b>)</b> UCDLA_IMM_106_002   v2.	
	Req. Upload: false		
Panel A FCS file(s) seriesMediaParameter	UCDLA_IMM_107_002   v2	.0	
	Req. Upload: false		

### Panel B FCS file(s) UCDLA\_IMM\_108\_002 | v2.0

seriesMediaParameter

v2.0

procedureMetadata

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Automated analysi procedureMetadata	<b>S</b> UCDLA_IMM_109_002   v	2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Collection buffer m procedureMetadata	nanufacturer ucdla_ii	MM_110_002   v2.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: Life Technologies, B		
		<b>ber</b> ucdla_imm_111_002
	3	

<b>Options:</b> 24020, 563503, 084-08965,				
FACS buffer manuf procedureMetadata	facturer ucdla_imm_11	2_002   v2.0		
Req. Analysis: false	Req. Upload: false	Is Annotated: false		
Options: Life Technologies, Ir				
FACS buffer catalo procedureMetadata	g number ucdla_imm	_113_002   v2.0		
Req. Analysis: false	Req. Upload: false	Is Annotated: false		
<b>Options:</b> 14175, In house, 048-29805,				
Enzyme buffer mar	nufacturer ucdla_imm	_114_002   v2.0		
Req. Analysis: false	Req. Upload: false	Is Annotated: false		
Options: Life Technologies, N	I/A, Wako,			

# Enzyme buffer catalog number UCDLA\_IMM\_115\_002 | v2.0

procedureMetadata

Req. Analysis: false	Req. Upload: false	Is Annotated: false	
<b>Options:</b> 14025, N/A, 084-08	965,		
Total spleen leuko simpleParameter	cyte count ucdla_imi	M_116_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Clog- events (PanesimpleParameter	PIA) UCDLA_IMM_117_00	1   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
FSC/SSC Singlets (Panel A) UCDLA_IMM_118_001   v1.0			
simpleParameter			
Des Analysis folos			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	

#### Effector NK cells - % of live leukocytes (Panel A) UCDLA\_IMM

\_119\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false
Unit Measured: %

# Effector Treg cells - % of live leukocytes (Panel A) UCDLA\_IM

M\_120\_001 | v1.0

simpleParameter

Reg. Analysis: false Reg. Upload: false Is Annotated: false

**Unit Measured:** %

.....

### Effector T cells - % of live leukocytes (Panel A) UCDLA\_IMM\_1

21\_001 | v1.0

simpleParameter

Reg. Analysis: false Reg. Upload: false Is Annotated: false

Unit Measured: %

#### KIrg1+ CD4- NKT cells - % of live leukocytes (Panel A) UCD

LA\_IMM\_122\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false
Unit Measured: %

### Klrg1+ CD4+ NKT cells - % of live leukocytes (Panel A) UCD

LA\_IMM\_123\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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# Klrg1+ CD4+ T helper cells - % of live leukocytes (Panel A) UCDLA\_IMM\_124\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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#### KIrg1+ CD8 T cells - % of live leukocytes (Panel A) UCDLA\_IM

M\_125\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false
Unit Measured: %

# Klrg1+ NK cells - % of live leukocytes (Panel A) UCDLA\_IMM\_1

26\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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## Klrg1+ Treg cells - % of live leukocytes (Panel A) UCDLA\_IMM

\_127\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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# Klrg1+ T cells - % of live leukocytes (Panel A) UCDLA\_IMM\_12 8\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false
Unit Measured: %

### Resting NK cells - % of live leukocytes (Panel A) UCDLA\_IMM

\_129\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: %

#### Resting Treg cells - % of live leukocytes (Panel A) UCDLA\_IM

M\_130\_001 | v1.0

simpleParameter

Reg. Analysis: false Reg. Upload: false Is Annotated: false

**Unit Measured:** %

# Resting T cells - % of live leukocytes (Panel A) UCDLA\_IMM\_1 31\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
T cells - % of live	leukocytes (Panel A	<b>4)</b> UCDLA_IMM_132_001   v1.
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
	leukocytes (Panel A	<b>4)</b> UCDLA_IMM_133_001   v1.
0 simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		

# CD4- NKT cells - % of NKT cells (Panel A) UCDLA\_IMM\_134\_001 | v1.0 simpleParameter Reg. Analysis: false Reg. Upload: false Is Annotated: false Unit Measured: % CD4+ NKT cells - % of NKT cells (Panel A) UCDLA\_IMM\_135\_001 I v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: % CD4+ T cells - % of T cells UCDLA IMM 136 001 | v1.0 simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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#### CD4+ T helper cells - % of CD4 T cells UCDLA\_IMM\_137\_001 | v1.0

Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: % CD8+ T cells - % of T cells UCDLA\_IMM\_138\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: % Effector CD4- NKT cells - % of CD4- NKT cells UCDLA\_IMM\_139 \_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: % Effector CD4+ NKT cells - % of CD4+ NKT cells ucdla imm 1

Req. Analysis: false Req. Upload: false Is Annotated: false

40 001 | v1.0

simpleParameter

Unit Measured: %
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#### Effector CD4+ T helper cells - % of CD4+ T helper cells ucd

LA\_IMM\_141\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: %

#### Effector CD8+ T cells - % of CD8+ T cells UCDLA\_IMM\_142\_001 |

v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

#### Effector NK cells - % of NK cells (Panel A) UCDLA\_IMM\_143\_001

| v1.0

simpleParameter

ĺ	Jn	iŧ	M	ea	SI	ıre	he	%
•				vu	9	413		70

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#### Effector Treg cells - % of Treg cells UCDLA\_IMM\_144\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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#### Effector T cells - % of T cells UCDLA\_IMM\_145\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: %

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### Kirg1+ CD4- NKT cells - % of CD4- NKT cells UCDLA\_IMM\_146\_

001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

KIrg1+ CD4+ NKT ( _001   v1.0 simpleParameter	cells - % of CD4+ NI	KT cells ucdla_imm_147
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
KIrg1+ CD4+ T help A_IMM_148_001   v1.0 simpleParameter	per cells - % of CD4	+ T helper cells ucdl
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
KIrg1+ CD8 T cells simpleParameter	- % of CD8+ T cells	6 UCDLA_IMM_149_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		

### Kirg1+ NK cells - % of NK cells (Panel A) UCDLA\_IMM\_150\_001 |

v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

#### Kirg1+ Treg cells - % of Treg cells UCDLA\_IMM\_151\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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#### Kirg1+ T cells - % of T cells UCDLA\_IMM\_152\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

.....

Naïve CD8+ T cells - % of CD8+ T cells UCDLA\_IMM\_153\_001 | v1.

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

# Resting CD4- NKT cells - % of CD4- NKT cells UCDLA\_IMM\_154 \_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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# Resting CD4+ NKT cells - % of CD4+ NKT cells UCDLA\_IMM\_15 5\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

#### Resting CD4+ T helper cells - % of CD4+ T helper cells UCD

LA\_IMM\_156\_001 | v1.0

Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: % Resting CD8+ T cells - % of CD8+ T cells UCDLA\_IMM\_157\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: % Resting NK cells - % of NK cells (Panel A) UCDLA\_IMM\_158\_001 | v1.0 simpleParameter Reg. Analysis: false Reg. Upload: false Is Annotated: false Unit Measured: %

#### Resting Treg cells - % of Treg cells UCDLA\_IMM\_159\_001 | v1.0

simpleParameter

Unit Measured: %		
Resting T cells - %	of T cells ucdla_imi	M_160_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
Treg cells - % of Cl	D4 T cells ucdla_imm_	_161_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
Clog- events (Pane simpleParameter	B) UCDLA_IMM_162_00	I   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false

#### FSC/SSC Singlets (Panel B) UCDLA\_IMM\_163\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

#### B1a cells - % of B cells (Panel B) UCDLA\_IMM\_164\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

#### B1b cells - % of B cells (Panel B) UCDLA\_IMM\_165\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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#### CD11b-high cDC - % of conventional DC (Panel B) UCDLA\_IM

M\_166\_001 | v1.0

simpleParameter

Unit Measured: %		
CD11b-low cDC - % _167_001   v1.0 simpleParameter	% of conventional D	C (Panel B) UCDLA_IMM
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
CD161+ B cells - % 68_001   v1.0 simpleParameter	of live leukocytes	(Panel B) ucdla_imm_1

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

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# Transitional 1 B cells - % of B cells (Panel B) UCDLA\_IMM\_169\_ 001 | v1.0

simpleParameter

Unit Measured: %	nit N	/leasured:	%	
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# Transitional 2 B cells - % of B cells (Panel B) UCDLA\_IMM\_170\_

001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: %

#### CD11b- NK cells - % of NK cells (Panel B) UCDLA\_IMM\_171\_001 |

v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: %

#### CD11b+ NK cells - % of NK cells (Panel B) UCDLA\_IMM\_172\_001

| v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

**Unit Measured:** %

CD161+ B cells - % of B cells UCDLA\_IMM\_173\_001 | v1.0 simpleParameter **Reg. Analysis:** false **Reg. Upload:** false **Is Annotated:** false Unit Measured: % Follicular B cells - % of B cells UCDLA\_IMM\_174\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false **Unit Measured:** % Ly6C+ CD11b- NK cells - % of NK cells (Panel B) UCDLA\_IMM\_ 175\_001 | v1.0 simpleParameter Reg. Analysis: false Reg. Upload: false Is Annotated: false Unit Measured: %

# Ly6C+ CD11b+ NK cells - % of NK cells (Panel B) UCDLA\_IMM \_176\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: false			
Unit Measured: %					
Ly6C+ NKT cells -   v1.0   simpleParameter	% of NKT cells (Par	nel B) ucdla_imm_177_001			
Req. Analysis: false	Req. Upload: false	Is Annotated: false			
Unit Measured: %					
Marginal zone B cells - % of B cells UCDLA_IMM_178_001   v1.0 simpleParameter					
Req. Analysis: false	Req. Upload: false	Is Annotated: false			
Unit Measured: %					

Transitional 1 Bcells - % of B cells UCDLA\_IMM\_179\_001 | v1.0

Req. Analysis: false Req. Upload: false Is Annotated: false **Unit Measured:** % Transitional 2 B cells - % of B cells UCDLA\_IMM\_180\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false **Unit Measured:** % T cells (Panel A) - cell count UCDLA\_IMM\_181\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented()

T cells - cell count UCDLA\_IMM\_182\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() CD8+ T cells - cell count UCDLA\_IMM\_183\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() Effector CD8+ T cells - cell count UCDLA\_IMM\_184\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented()

**Req. Analysis:** false **Req. Upload:** false **Is Annotated:** false **Unit Measured:** count **Derivation:** unimplemented() Naïve CD8+ T cells - cell count UCDLA\_IMM\_186\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() Kirg1+ CD8 T cells - cell count UCDLA\_IMM\_187\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented()

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		
CD4+ T helper cells	s - cell count ucdla_	IMM_189_001   v1.0
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		
Effector CD4+ T he	elper cells - cell cou	<b>nt</b> UCDLA_IMM_190_001   v1
.0		·
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		

# Resting CD4+ T helper cells - cell count UCDLA\_IMM\_191\_001 | v1

.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

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# Kirg1+ CD4+ T helper cells - cell count UCDLA\_IMM\_192\_001 | v1.

0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

#### Treg cells - cell count UCDLA\_IMM\_193\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

#### Effector Treg cells - cell count UCDLA\_IMM\_194\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

#### Resting Treg cells - cell count UCDLA\_IMM\_195\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

#### Kirg1+ Treg cells - cell count UCDLA\_IMM\_196\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

#### T cells - cell count UCDLA\_IMM\_197\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

#### Effector T cells - cell count UCDLA\_IMM\_198\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

#### Resting T cells - cell count UCDLA\_IMM\_199\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

#### Kirg1+ T cells - cell count UCDLA\_IMM\_200\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

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#### NKT cells (panel A) - cell count UCDLA\_IMM\_201\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

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### CD4+ NKT cells - cell count UCDLA\_IMM\_202\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

#### Effector CD4+ NKT cells - cell count UCDLA\_IMM\_203\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

#### Resting CD4+ NKT cells - cell count UCDLA\_IMM\_204\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

#### Kirg1+ CD4+ NKT cells - cell count UCDLA\_IMM\_205\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

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#### CD4- NKT cells - cell count UCDLA\_IMM\_206\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

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#### Effector CD4- NKT cells - cell count UCDLA\_IMM\_207\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

**Derivation:** unimplemented()

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### Resting CD4- NKT cells - cell count UCDLA\_IMM\_208\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

Derivation: unimplemented()		
KIrg1+ CD4- NKT of simpleParameter	cells - cell count ucd	LA_IMM_209_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
Derivation: unimplemented()		
NK cells (Panel A) simpleParameter	- cell count ucdla_im	M_210_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		

# Effector NK cells - cell count UCDLA\_IMM\_211\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false

Unit Measured: count

Derivation: unimplemented()			
Resting NK cells - cell count UCDLA_IMM_212_001   v1.0 simpleParameter			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Unit Measured: count  Derivation: unimplemented()			
Kirg1+ NK cells - cell count UCDLA_IMM_213_001   v1.0 simpleParameter			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Unit Measured: count  Derivation: unimplemented()			

# T cells (panel B) - cell count UCDLA\_IMM\_214\_001 | v1.0

simpleParameter

Unit Measured: count		
<b>Derivation:</b> unimplemented()		
NKT cells (panel B) simpleParameter	) - cell count ucdla_i	MM_215_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		
	cell count ucdla_imm	_216_001   v1.0
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		

# NK cells (Panel B) - cell count UCDLA\_IMM\_217\_001 | v1.0

simpleParameter

Unit Measured: count		
<b>Derivation:</b> unimplemented()		
CD11b- NK cells - 0 simpleParameter	cell count ucdla_imm_	_218_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		
Ly6C+ CD11b- NK simpleParameter	cells - cell count uc	DLA_IMM_219_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		

# CD11b+ NK cells - cell count UCDLA\_IMM\_220\_001 | v1.0

simpleParameter

Unit Measured: count		
Derivation: unimplemented()		
Ly6C+ CD11b+ NK simpleParameter	cells - cell count uc	DLA_IMM_221_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
Derivation: unimplemented()		
B cells - cell count simpleParameter	UCDLA_IMM_222_001   v1.0	)
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
Derivation: unimplemented()		

B1a cells - cell count UCDLA\_IMM\_223\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		
	I <b>nt</b> UCDLA_IMM_224_001	v1.0
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		
Follicular B cells - simpleParameter	cell count ucdla_imm	_225_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		

**Req. Analysis:** false **Req. Upload:** false **Is Annotated:** false **Unit Measured:** count **Derivation:** unimplemented() Transitional 1 B cells - cell count UCDLA\_IMM\_227\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() Transitional 2 B cells - cell count UCDLA\_IMM\_228\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented()

Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() Conventional DC - cell count UCDLA\_IMM\_230\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() CD11b-low cDC - cell count UCDLA\_IMM\_231\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented()

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		
Plasmacytoid DC - simpleParameter	cell count ucdla_imm	I_233_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count  Derivation: unimplemented()		
Macrophages - cell simpleParameter	Count UCDLA_IMM_234	_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: count		
<b>Derivation:</b> unimplemented()		

### Monocytes - cell count UCDLA\_IMM\_235\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() Granulocytes - cell count UCDLA\_IMM\_236\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented() Eosinophils - cell count UCDLA\_IMM\_237\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false Unit Measured: count **Derivation:** unimplemented()

# Panel A anti-CD5 clone UCDLA\_IMM\_238\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

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# Panel A anti-CD5 fluorochrome UCDLA\_IMM\_239\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: BV421, eF450, PE-Gr-A,

.....

### Panel A anti-CD5 RRID UCDLA\_IMM\_240\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: AB\_2737758, AB\_1603250,

.....

#### Panel A anti-CD4 clone UCDLA\_IMM\_241\_001 | v1.0

procedureMetadata

Options: RM4-5, GK1.5,		
Panel A anti-CD4 f	<b>luorochrome</b> ucdla_	IMM_242_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: FITC, PO, PE-CF59	)4,	
Panel A anti-CD4 F procedureMetadata	RRID UCDLA_IMM_243_00	01   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
<b>Options:</b> AB_394583, AB_14	74250, AB_396633,	
Panel A anti-CD44 procedureMetadata	clone ucdla_imm_244_	001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false

# Panel A anti-CD44 fluorochrome UCDLA\_IMM\_245\_001 | v1.0

Options: PE, BV650, PE-Cy7,

### Panel A anti-CD44 RRID UCDLA\_IMM\_246\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: AB\_394649, AB\_2562600, AB\_10895375,

.....

# Panel A anti-CD8a clone UCDLA\_IMM\_247\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

# Panel A anti-CD8a fluorochrome UCDLA\_IMM\_248\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: PE-CF594, APCeF780, eFluor 450,

# Panel A anti-CD8a RRID UCDLA\_IMM\_249\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: AB\_11152075, AB\_1272185,

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# Panel A anti-CD25 clone UCDLA\_IMM\_250\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

**Options:** PC61, PC61.5,

.....

### Panel A anti-CD25 fluorochrome UCDLA\_IMM\_251\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: PE-Cy7, APC, FITC,

Options: AB\_394509, AB\_398623, AB\_10562035,

### Panel A anti-CD161 clone UCDLA\_IMM\_253\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

### Panel A anti-CD161 fluorochrome UCDLA\_IMM\_254\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: APC, PE, PE-Cy7, eFluor 780,

.....

# Panel A anti-CD161 RRID UCDLA\_IMM\_255\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

**Options:** AB\_398463, AB\_394677, AB\_394507,

Panel A anti-CD62L clone UCDLA\_IMM\_256\_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Panel A anti-CD62L fluorochrome UCDLA\_IMM\_257\_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Options: APC-Cy7, PE-Cy7, APC, Panel A anti-CD62L RRID UCDLA IMM 258 001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false

Options: AB\_10611861, AB\_469633,

#### Panel A Live/Dead stain UCDLA IMM 259 001 | v1.0

Options: PI, Aqua, Sytox Blue,

.....

#### Panel A additional maker 1 name UCDLA\_IMM\_260\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: TCRd, CD3,

### Panel A additional marker 1 clone UCDLA\_IMM\_261\_001 | v1.0

procedureMetadata

Reg. Analysis: false Reg. Upload: false Is Annotated: false

Options: GL3, eBio500A2,

.....

### Panel A additional marker 1 fluorochrome UCDLA\_IMM\_262\_001

| v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: FITC, Alexa 700,

Panel A additional procedureMetadata	marker 1 RRID ucdl	A_IMM_263_001   v1.0
	Req. Upload: false	Is Annotated: false
	maker 2 name ucdl	
procedureMetadata	maker 2 mame ocol	A_IIVIIVI_264_001   V1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: CD45, TCRd,		
Panel A additional procedureMetadata	marker 2 clone ucdi	_A_IMM_265_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
<b>Options:</b> 30-F11, GL3,		

Panel A additional marker 2 fluorochrome UCDLA\_IMM\_266\_001 | v1.0

**Options:** BV785, BV711,

### Panel A additional marker 2 RRID UCDLA\_IMM\_267\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

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### Panel A additional maker 3 name UCDLA\_IMM\_268\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: CD3, KLRG1,

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# Panel A additional marker 3 clone UCDLA\_IMM\_269\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: ebio500A2, 2F1,

Panel A additional marker 3 fluorochrome UCDLA\_IMM\_270\_001 | v1.0 procedureMetadata **Reg. Analysis:** false **Reg. Upload:** false Is Annotated: false Options: Af700, BV605, Panel A additional marker 3 RRID UCDLA IMM 271 001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Panel A additional maker 4 name UCDLA IMM 272 001 | v1.0 procedureMetadata Req. Analysis: false **Req. Upload:** false **Is Annotated:** false Options: GITR,

Panel A additional marker 4 clone UCDLA\_IMM\_273\_001 | v1.0

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: DTA-1,		
Panel A additional   v1.0   procedureMetadata	marker 4 fluorochro	ome ucdla_imm_274_001
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: BV786,		
Panel A additional procedureMetadata	marker 4 RRID UCDL	A_IMM_275_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Panel A additional procedureMetadata	maker 5 name ucdl/	A_IMM_276_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false

### Panel A additional marker 5 clone UCDLA\_IMM\_277\_001 | v1.0

procedureMetadata

Panel A additional marker 5 fluorochrome UCDLA\_IMM\_278\_001

| v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

# Panel A additional marker 5 RRID UCDLA\_IMM\_279\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

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#### Panel B anti-CD5 clone UCDLA\_IMM\_280\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

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Panel B anti-CD5 fluorochrome UCDLA\_IMM\_281\_001 | v1.0

Options: BV421, eF450, PE-Gr-A,

### Panel B anti-CD5 RRID UCDLA\_IMM\_282\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

### Panel B anti-Ly6G clone UCDLA\_IMM\_283\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

**Options:** 1A8, RB6-8C5,

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## Panel B anti-Ly6G fluorochrome UCDLA\_IMM\_284\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: BV421, BV785, violetFluor 450,

Panel B anti-Ly6G RRID UCDLA\_IMM\_285\_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Options: AB\_2737756, AB\_2566317, Panel B anti-CD19 clone UCDLA\_IMM\_286\_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Panel B anti-CD19 fluorochrome UCDLA\_IMM\_287\_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Options: BV510, PE-Cy7, PE-CF594,

#### Panel B anti-CD19 RRID UCDLA\_IMM\_288\_001 | v1.0

Options: AB\_2737915, AB\_394495,

.....

### Panel B anti-Ly6C clone UCDLA\_IMM\_289\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: AL-21, HK1.4,

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### Panel B anti-Ly6c fluorochrome UCDLA\_IMM\_290\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: FITC, PerCP Cy5.5, BV785,

#### Panel B anti-Ly6c RRID UCDLA\_IMM\_291\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: AB\_394628, AB\_2723343,

Panel B anti-CD21/35 clone UCDLA\_IMM\_292\_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Panel B anti-CD21/35 fluorochrome UCDLA\_IMM\_293\_001 | v1.0 procedureMetadata Reg. Analysis: false Reg. Upload: false Is Annotated: false Options: PE, BV605, Panel B anti-CD21/35 RRID UCDLA\_IMM\_294\_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false Options: AB\_394532, AB\_2738048,

Panel B anti-CD11b clone UCDLA\_IMM\_295\_001 | v1.0

procedureMetadata

Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Panel B anti-CD11 procedureMetadata	<b>b fluorochrome</b> ucd	LA_IMM_296_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Options: PE-CF594, PerCP-	Cy5.5, FITC,		
Panel B anti-CD11b RRID UCDLA_IMM_297_001   v1.0 procedureMetadata			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
<b>Options:</b> AB_11154216, AB_	_2033995,		
Panel B anti-CD11c clone UCDLA_IMM_298_001   v1.0 procedureMetadata			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Options: HL3, N418,			

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#### Panel B anti-CD11c fluorochrome UCDLA\_IMM\_299\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: PE-Cy7, APC-Cy7, APC,

### Panel B anti-CD11c RRID UCDLA\_IMM\_300\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: AB\_647251, AB\_10611727,

### Panel B anti-CD161 clone UCDLA\_IMM\_301\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

#### Panel B anti-CD161 fluorochrome UCDLA IMM 302 001 | v1.0

procedureMetadata

Options: APC, PE, eFluor 780,		
Panel B anti-CD16 <sup>2</sup> procedureMetadata	1 RRID UCDLA_IMM_303	_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Panel B anti-MHCII procedureMetadata	clone ucdla_imm_304_	_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Panel B anti-MHCII procedureMetadata	fluorochrome ucdl	A_IMM_305_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: APC-eFluor(R) 780,	BV650, APC-Cy7, Alexa 700,	

# Panel B anti-MHCII RRID UCDLA\_IMM\_306\_001 | v1.0

Req. Analysis: false	Req. Upload: false	Is Annotated: false			
<b>Options:</b> AB_1548783, AB_2565975, AB_2069377,					
Panel B Live/Dead stain UCDLA_IMM_307_001   v1.0 procedureMetadata					
Req. Analysis: false	Req. Upload: false	Is Annotated: false			
Options: PI, Aqua, Sytox Blue,					
Panel B additional maker 1 name UCDLA_IMM_308_001   v1.0 procedureMetadata					
Req. Analysis: false	Req. Upload: false	Is Annotated: false			
Panel B additional marker 1 clone UCDLA_IMM_309_001   v1.0 procedureMetadata					
Req. Analysis: false	Req. Upload: false	Is Annotated: false			

### Panel B additional marker 1 fluorochrome UCDLA\_IMM\_310\_001

| v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: FITC, BV711,

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# Panel B additional marker 1 RRID UCDLA\_IMM\_311\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

#### Panel B additional maker 2 name UCDLA IMM 312 001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

**Options:** F4/80, CD317,

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### Panel B additional marker 2 clone UCDLA\_IMM\_313\_001 | v1.0

procedureMetadata

Options: BM8, eBio927,		
Panel B additional	marker 2 fluorochro	<b>ome</b> ucdla_imm_314_001
v1.0 procedureMetadata		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: APC, PE-Cy7,		
Panel B additional procedureMetadata	marker 2 RRID UCDL	_A_IMM_315_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Panel B additional procedureMetadata	maker 3 name ucdl	A_IMM_316_001   v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
<b>Options:</b> CD45, F4/80,		

# Panel B additional marker 3 clone UCDLA\_IMM\_317\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

Options: 30-F11, BM8,

Panel B additional marker 3 fluorochrome UCDLA\_IMM\_318\_001

| v1.0 | procedureMetadata

Req. Analysis: false Req. Upload: false Is Annotated: false

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# Panel B additional marker 3 RRID UCDLA\_IMM\_319\_001 | v1.0

procedureMetadata

Options: BV510, Cy5PE,

Req. Analysis: false Req. Upload: false Is Annotated: false

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# Panel B additional maker 4 name UCDLA\_IMM\_320\_001 | v1.0

procedureMetadata

Panel B additional procedureMetadata	marker 4 clone ucpi	LA_IMM_321_001   v1.0	
	Req. Upload: false		
		ome ucdla_imm_322_001	
procedureMetadata			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Panel B additional procedureMetadata	marker 4 RRID UCDL	_A_IMM_323_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Panel B additional maker 5 name UCDLA_IMM_324_001   v1.0 procedureMetadata			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	

# Panel B additional marker 5 clone UCDLA\_IMM\_325\_001 | v1.0

procedureMetadata

Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Panel B additional   v1.0   procedureMetadata	marker 5 fluorochr	ome ucdla_imm_326_001	
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Panel B additional marker 5 RRID UCDLA_IMM_327_001   v1.0 procedureMetadata			
Req. Analysis: false	Req. Upload: false	Is Annotated: false	
Analysis results fil	• UCDLA_IMM_328_001   v	1.0	
	<b>e</b> UCDLA_IMM_328_001   v	1.0  Is Annotated: false	