# Immunophenotyping HMGULA\_IMM\_001

# 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 of animals : 3M + 3F
- Age at test: Week 63
- Sex: We would expect the results of this test to show sexual dimorphism

# 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

#### Supplies

- 96-well V-bottomed plates (Falcon #353263)
- Petri dishes
- Dispensing troughs
- Extra long 10 µl 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 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 Mg2+, with Ca2+ (for <u>enzyme buffer</u> used for DNAse and Collagenase D digestions)
- PBS without Mg2+, without Ca2+ (for <u>FACS buffer</u> to be used in all steps subsequent to enzymatic digest)
- EDTA (final concentration 2mM)
- Digestion enzyme (Collagenase D from Roche, #11088858001) stock solution in <u>enzym</u> <u>e buffer</u> (see below), aliquoted and stored at –20°C
- DNAse I stock solution (Sigma, #DN25) in <u>enzyme buffer</u> (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)

## 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 Ca2+ and with Mg2+, another without Ca2+ and without Mg2+.

- Collection buffer:
  - (*if spleens are to be processed on the same day*) HBSS with Ca2+/Mg2+ and phenol red (Life Technologies 14170161; check if it has phenol red) **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 <u>without</u> Ca2+/Mg2+ OR
  - HBSS 1X <u>without</u> Ca2+/Mg2+
  - EDTA 2mM
  - 2% CS (v/v)
  - 10mM HEPES
- **Enzyme buffer** (for DNAse and Collagenase D digestions; Stable for up to 1 month in the fridge):
  - PBS <u>with</u> Ca2+ and Mg2+ OR
  - HBSS 1X <u>with</u> Ca2+/Mg2+
  - 2% CS (v/v);
  - 10mM HEPES
- **RBC Lysis buffer**: Prepare a 1X solution in ddH<sub>2</sub>0 from lysis buffer.
- **Stopping buffer** (require 300 µl per sample):
  - 1x PBS without Ca2+ and without Mg2+ or HBSS
  - 0.1 M EDTA (37.5 g/L)
- Antibody cocktails for Panels 1 & 2
  - Protect antibodies and prepared cocktails from direct light.
  - Mastermix concentration, storage temperature and stability to be determined after panels 1 and 2 have been finalised and tested.

- Each sample will require 50 µl (or up to 100 µl) of diluted 1X antibody cocktail.
- Antibody cocktails should be gently but thoroughly mixed or quickly vortexed 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.
- 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 per each spleen, containing final concentrations of:
  - DNAse I: 30 g
  - Collagenase D: 600 Mandl Units

*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 m Nylon mesh filter into a 50 mL Falcon tube OR

- directly from C-tubes pour splenocyte suspension through 30 mm 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 flowthrough 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 Ca2+ and Mg2+ (0.17-0.2 Wünsch unit/ml)
- Mince into fine pieces using small scissors, place on ice until all samples are minced.
- Add 2ml collagenase (0.17-0.2 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 down the cell count, correct for dilution and calculate the concentration in cells per µl.
- Cell count:
  - <u>If performed before RBC lysis</u>, 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 ml 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 ml 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 ml 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 ml <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 ml of 1:100 Fc block and incubate on ice for 10 min. Top up to 200 ml 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.
- Centrifuge plate, discard supernatant and resuspend in 50 to 100 ml 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 ml with <u>FACS buffer</u> after incubation. Centrifuge, discard supernatant and resuspend in 200 ml <u>FACS buffer</u>.
  - When ready to read plate, centrifuge again and discard supernatant. Resuspend the pellet in 200 ml 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 ml of PBS (RT) to all samples
    - Spin at 2000 rpm for 1 minute 8°C
    - Add 100 ml/well of Zombie Near-IR Live/Dead dye (1/2000) made up in PBS incubate at room temperature for 10 mins, add 200 ml <u>FACS buffer</u>.

#### 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 of low frequencies (macrophages, DCs).

#### Gating Panel 1

Parameters	Gating steps			
Panel A live leukocyte count				Γ
T cells (panel A)	number of live leukocytes	CD5+	CD161-	
NKT cells (panel A)	number of live leukocytes	CD5+	CD161+	
NK cells (panel A)	number of live leukocytes	CD5-	CD161+	
Others	number of live leukocytes	CD5-	CD161-	
CD4 T cells	number of live leukocytes	CD5+	CD161-	CD
CD8 T cells	number of live leukocytes	CD5+	CD161-	CD
DN T cells	number of live leukocytes	CD5+	CD161-	CD
DP T cells	number of live leukocytes	CD5+	CD161-	CD
CD4 NKT cells	number of live leukocytes	CD5+	CD161+	CD
CD8 NKT cells	number of live leukocytes	CD5+	CD161+	CD
		1		1

DN NKT cells	number of live leukocytes	CD5+	CD161+	CD
CD4 CD25+ T cells		number of CD5+	CD161-	CD
CD4 CD25- T cells		number of CD5+	CD161-	CD
CD8 CD25+ T cells		number of CD5+	CD161-	CD
CD8 CD25- T cells		number of CD5+	CD161-	CD
DN CD25+ T cells		number of CD5+	CD161-	CD
DN CD25- T cells		number of CD5+	CD161-	CD
CD4 CD25+ NKT cells		number of CD5+	CD161+	CD
CD4 CD25- NKT cells		number of CD5+	CD161+	CD
CD8 CD25+ NKT cells		number of CD5+	CD161+	CD
CD8 CD25- NKT cells		number of CD5+	CD161+	CD
DN CD25+ NKT cells		number of CD5+	CD161+	CD
DN CD25- NKT cells		number of CD5+	CD161+	CD
CD4 CD44+CD62L- T cells		number of CD5+	CD161-	CD
CD4 CD44+CD62L+ T cells		number of CD5+	CD161-	CD
CD4 CD44-CD62L+ T cells		number of CD5+	CD161-	CD
CD4 CD44-CD62L- T cells		number of CD5+	CD161-	CD
CD8 CD44+CD62L- T cells		number of CD5+	CD161-	CD
CD8 CD44+CD62L+ T cells		number of CD5+	CD161-	CD
CD8 CD44-CD62L+ T cells		number of CD5+	CD161-	CD
CD8 CD44-CD62L- T cells		number of CD5+	CD161-	CD
DN CD44+CD62L- T cells		number of CD5+	CD161-	CD
DN CD44+CD62L+ T cells		number of CD5+	CD161-	CD
DN CD44-CD62L+ T cells		number of CD5+	CD161-	CD
DN CD44-CD62L- T cells		number of CD5+	CD161-	CD
CD4 CD44+CD62L- NKT cells		number of CD5+	CD161+	CD
CD4 CD44+CD62L+ NKT cells		number of CD5+	CD161+	CD
CD4 CD44-CD62L+ NKT cells		number of CD5+	CD161+	CD
CD8 CD44+CD62L- NKT cells		number of CD5+	CD161+	CD
CD8 CD44+CD62L+ NKT cells		number of CD5+	CD161+	CD
CD8 CD44-CD62L+ NKT cells		number of CD5+	CD161+	CD
DN CD44+CD62L- NKT cells		number of CD5+	CD161+	CD
DN CD44+CD62L+ NKT cells		number of CD5+	CD161+	CD
DN CD44-CD62L+ NKT cells		number of CD5+	CD161+	CD

#### Gating Panel B

Parameters	Gating steps				
Panel B live leukocyte count					
Neutrophils	Live	CD11b+	Ly6G+		
Monocytes	Not Granulocytes	CD11b+	Ly6C High		
Eosinophils	Not Monocytes	CD11b+	SSC-H High		
NK Cells (panel B)	Not Eosinophils	CD161+	CD19-	CD5-	
NK Subsets (Q1)	Not Eosinophils	CD161+	CD19-	CD5-	С
NK Subsets (Q2)	Not Eosinophils	CD161+	CD19-	CD5-	С
NK Subsets (Q3)	Not Eosinophils	CD161+	CD19-	CD5-	С

NK Subsets (Q4)	Not Eosinophils	CD161+	CD19-	CD5-	С
NKT Cells (panel B)	Not Eosinophils	CD161+	CD19-	CD5+	Т
NKT Subsets (Q1)	Not Eosinophils	CD161+	CD19-	CD5+	С
NKT Subsets (Q3)	Not Eosinophils	CD161+	CD19-	CD5+	С
T Cells (panel B)	Not Eosinophils	CD161-	CD5+		
T Subset	Not Eosinophils	CD161-	CD5+	Ly6C+	
B Cells	Not Eosinophils	MHCII+	CD19+		
B1B Cells	Not Eosinophils	MHCII+	CD19+	CD5+	
B2B Cells	Not Eosinophils	MHCII+	CD19+	CD5-	
Follicular B Cells	Not Eosinophils	MHCII+	CD19+	CD5-	С
pre-B Cells	Not Eosinophils	MHCII+	CD19+	CD5-	С
MZB	Not Eosinophils	MHCII+	CD19+	CD5-	С
cDCs	Not Eosinophils	MHCII+	CD19-	CD11c+	
cDCs CD11b Type	Not Eosinophils	MHCII+	CD19-	CD11c+	С
pDCs	Not Eosinophils	Not T Cells	Ly6C+	CD317+	
RP Macrophage (F4/80+)	Not Eosinophils	MHCII+	F4/80+		Τ
or					Τ
RP Macrophage (CD19- CD11c-)	Not Eosinophils	MHCII+	CD19-	CD11c-	

#### **Parameters and Metadata**

#### Spleen weight HMGULA\_IMM\_001\_001 | v1.0

simpleParameter

Req. Analysis: false Req. Upload: true

Is Annotated: false

Unit Measured: g

## Percentage of live gated events in Panel A HMGULA\_IMM\_002\_0 01 | v1.0

**Unit Measured:** %


T cells (panel A) HMGULA_IMM_003_001   v1.0 simpleParameter			
Req. Analysis: false	Req. Upload: false	Is Annotated: true	
NKT cells (panel A) simpleParameter	HMGULA_IMM_004_001	v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: true	
NK cells (panel A)	HMGULA_IMM_005_001  v1	.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: true	
Others HMGULA_IMM_C	006_001  v1.0		

simpleParameter

Req. Analysis: falseReq. Upload: falseIs Annotated: true

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# CD4 T cells HMGULA\_IMM\_007\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true
CD8 T cells HMGULA_ simpleParameter	_IMM_008_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
DN T cells HMGULA_IN simpleParameter	MM_009_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false
<b>DP T cells</b> HMGULA_IN simpleParameter	/IM_010_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false

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## CD4 NKT cells HMGULA\_IMM\_011\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true
CD8 NKT cells HMGU simpleParameter	JLA_IMM_012_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: false
DN NKT cells HMGUL simpleParameter	.A_IMM_013_001   v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
CD4 CD25+ T cells simpleParameter	HMGULA_IMM_014_001   v	1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: true

## CD4 CD25- T cells HMGULA\_IMM\_015\_001 | v1.0

Req. Analysis: false	Req. Upload: false	Is Annotated: true
CD8 CD25+ T cells simpleParameter	HMGULA_IMM_016_001   v	1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: true
CD8 CD25- T cells	HMGULA_IMM_017_001   v1	.0
Req. Analysis: false	Req. Upload: false	Is Annotated: true
DN CD25+ T cells H simpleParameter	IMGULA_IMM_018_001  v1.	0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
DN CD25- T cells HI simpleParameter	MGULA_IMM_019_001   v1.0	)
Req. Analysis: false	Req. Upload: false	Is Annotated: false

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## CD4 CD25+ NKT cells HMGULA\_IMM\_020\_001 | v1.0

simpleParameter

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

#### CD4 CD25- NKT cells HMGULA\_IMM\_021\_001 | v1.0

simpleParameter

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

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## CD8 CD25+ NKT cells HMGULA\_IMM\_022\_001 | v1.0

simpleParameter

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

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#### CD8 CD25- NKT cells HMGULA\_IMM\_023\_001 | v1.0

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

#### DN CD25+ NKT cells HMGULA\_IMM\_024\_001 | v1.0

simpleParameter

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

#### DN CD25- NKT cells HMGULA\_IMM\_025\_001 | v1.0

simpleParameter

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

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## Total number of acquired events in Panel A HMGULA\_IMM\_026\_

001 | v1.0 simpleParameter

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

## Total number of acquired events in Panel B HMGULA\_IMM\_027\_

001 | v1.0

simpleParameter

Req.	Analy	/sis:	false
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Req. Upload: false

Is Annotated: false

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#### **CD4 CD44+CD62L- T cells** HMGULA\_IMM\_028\_001 | v1.0

simpleParameter

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

#### CD4 CD44+CD62L+ T cells HMGULA\_IMM\_029\_001 | v1.0

simpleParameter

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

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#### CD4 CD44-CD62L+ T cells HMGULA\_IMM\_030\_001 | v1.0

simpleParameter

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

#### CD4 CD44-CD62L- T cells HMGULA\_IMM\_031\_001 | v1.0

simpleParameter

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

## CD8 CD44+CD62L- T cells HMGULA\_IMM\_032\_001 | v1.0

#### CD8 CD44+CD62L+ T cells HMGULA IMM 033 001 | v1.0

simpleParameter

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

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#### CD8 CD44-CD62L+ T cells HMGULA\_IMM\_034\_001 | v1.0

simpleParameter

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

#### CD8 CD44-CD62L- T cells HMGULA\_IMM\_035\_001 | v1.0

simpleParameter

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

#### DN CD44+CD62L- T cells HMGULA\_IMM\_036\_001 | v1.0

simpleParameter

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

#### DN CD44+CD62L+ T cells HMGULA\_IMM\_037\_001 | v1.0

simpleParameter

Req. Analysis: falseReq. Upload: falseIs Annotated: false

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#### DN CD44-CD62L+ T cells HMGULA\_IMM\_038\_001 | v1.0

simpleParameter

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

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#### DN CD44-CD62L- T cells HMGULA\_IMM\_039\_001 | v1.0

simpleParameter

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

#### CD4 CD44+CD62L- NKT cells HMGULA\_IMM\_040\_001 | v1.0

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

## CD4 CD44+CD62L+ NKT cells HMGULA\_IMM\_041\_001 |v1.0

simpleParameter

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

#### CD4 CD44-CD62L+ NKT cells HMGULA\_IMM\_042\_001 | v1.0

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#### CD8 CD44+CD62L- NKT cells HMGULA\_IMM\_043\_001 | v1.0

simpleParameter

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

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#### CD8 CD44+CD62L+ NKT cells HMGULA IMM 044 001 | v1.0

simpleParameter

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

## CD8 CD44-CD62L+ NKT cells HMGULA\_IMM\_045\_001 | v1.0

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#### DN CD44+CD62L- NKT cells HMGULA\_IMM\_046\_001 | v1.0

simpleParameter

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

#### DN CD44+CD62L+ NKT cells HMGULA\_IMM\_047\_001 | v1.0

simpleParameter

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

#### DN CD44-CD62L+ NKT cells HMGULA\_IMM\_048\_001 | v1.0

simpleParameter

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

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## Percentage of live gated events in Panel B HMGULA\_IMM\_049\_0

01 | v1.0 simpleParameter

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# NK Cells (panel B) HMGULA\_IMM\_053\_001 | v1.0

# NK Subsets (Q1) HMGULA\_IMM\_054\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true NK Subsets (Q2) HMGULA\_IMM\_055\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true \_\_\_\_\_ NK Subsets (Q3) HMGULA\_IMM\_056\_001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true

#### NK Subsets (Q4) HMGULA\_IMM\_057\_001 | v1.0

Req. Analysis: false	Req. Upload: false	Is Annotated: true
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# NKT Cells (panel B) HMGULA\_IMM\_058\_001 | v1.0

simpleParameter

	Req. Upload: false	Is Annotated: true
NKT Subsets (Q1)	HMGULA_IMM_059_001   v1	.0
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true
NKT Subsets (Q3)	HMGULA_IMM_060_001  v1	.0
Req. Analysis: false	Req. Upload: false	Is Annotated: true
T Cells (panel B) HM simpleParameter	/IGULA_IMM_061_001  v1.0	
Req. Analysis: false	Req. Upload: false	Is Annotated: true

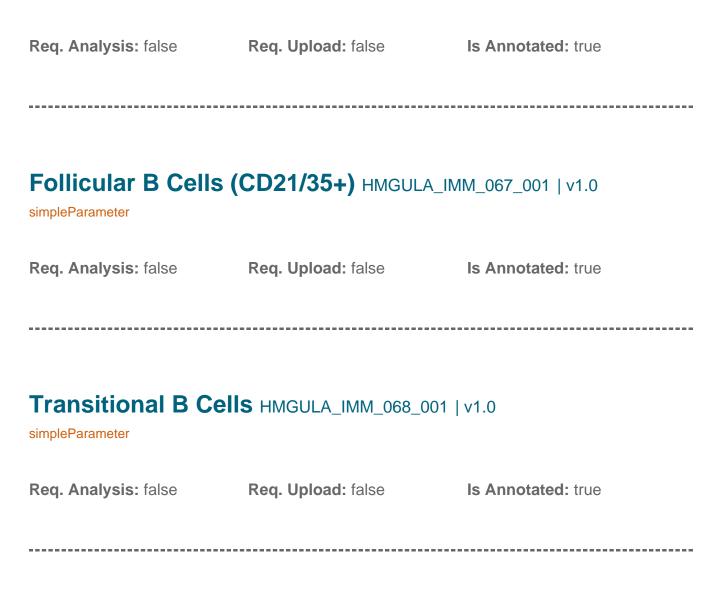
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#### T Subset HMGULA\_IMM\_062\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true		
<b>B Cells</b> HMGULA_IMM_( simpleParameter	063_001  v1.0			
Req. Analysis: false	Req. Upload: false	Is Annotated: true		
<b>B1B Cells</b> HMGULA_IN simpleParameter	B1B Cells HMGULA_IMM_064_001   v1.0 simpleParameter			
Req. Analysis: false	Req. Upload: false	Is Annotated: true		
B2B Cells HMGULA_IMM_065_001   v1.0 simpleParameter				
Req. Analysis: false	Req. Upload: false	Is Annotated: true		

# Follicular B Cells HMGULA\_IMM\_066\_001 | v1.0



## Transitional B Cells (CD21/35 low) HMGULA\_IMM\_069\_001 | v1.0

simpleParameter

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

#### MZB HMGULA\_IMM\_070\_001 | v1.0

simpleParameter

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

# MZB (CD21/35 high) HMGULA\_IMM\_071\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true	
cDCs HMGULA_IMM_072 simpleParameter	2_001   v1.0		
	Req. Upload: false		
cDCs CD11b Type HMGULA_IMM_073_001   v1.0 simpleParameter			
Req. Analysis: false	Req. Upload: false	Is Annotated: true	
pDCs HMGULA_IMM_074_001   v1.0 simpleParameter			
Req. Analysis: false	Req. Upload: false	Is Annotated: true	

# **RP Macrophage (F4/80+)** HMGULA\_IMM\_075\_001 | v1.0

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true		
RP Macrophage (C simpleParameter	<b>:D19- CD11с-)</b> нмgui	LA_IMM_076_001   v1.0		
Req. Analysis: false	Req. Upload: false	Is Annotated: true		
Equipment name	HMGULA_IMM_077_001   v1	.0		
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
<b>Options:</b> FACS, Fortessa_1, LSR II, Flow cytometer,				
Equipment manufacturer HMGULA_IMM_078_001   v1.0				
Req. Analysis: true	Req. Upload: true	Is Annotated: false		
Options: BD Biosciences, Be	eckman Coulter, IntelliCyt,			

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#### Equipment model HMGULA\_IMM\_079\_001 | v1.0

procedureMetadata

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

Options: BD LSRFortessa Cell Analyzer, H47100123, Gallios, FACSAria III, BD LSR-II, CANTO-II, iQue Screener PLUS,

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#### CS&T Bead lot HMGULA\_IMM\_080\_001 | v1.0

procedureMetadata

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

#### Anesthesia HMGULA IMM 081 001 | v1.0

procedureMetadata

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

Options: Injection narcosis with Sodium Pentobarbital (Somnopentyl), none, Injection narcosis with Ketamine (100mg/kg)/Xylazine (10mg/kg), Injection narcosis with Tribromoethanol (Avertin), Isoflurane,

#### Cell digestion HMGULA\_IMM\_082\_001 | v1.0

Req. Analysis: true	Req. Upload: true	Is Annotated: false		
Options: GentleMACS, manual,				
Cell digestion agent HMGULA_IMM_083_001   v1.0 procedureMetadata				
Req. Analysis: false	Req. Upload: true	Is Annotated: false		
Options: Collagenase D, Collagenase II,				

## Cell digestion agent manufacturer HMGULA\_IMM\_084\_001 | v1.0

procedureMetadata

Req.	Analysis: fal	se Req.	Upload: true	Is Annotated: false
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Options: Roche, Worthington, Gibco, Sigma,

## Cell digestion agent catalog number HMGULA\_IMM\_085\_001 | v1.0

procedureMetadata

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

Options: #11088858001, CLS2LS004176, 17101-015, C6885,

.....

#### Cell counting HMGULA\_IMM\_086\_001 | v1.0

procedureMetadata

Req. Analysis: false	Req. Upload: true	Is Annotated: false
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**Options:** pre-lysis, post-lysis,

# Cell counting equipment manufacturer HMGULA\_IMM\_087\_001 | v1

.0

procedureMetadata

Req. Analysis: false Req. Upload: true	Is Annotated: false
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**Options:** Life Technologies, American Optical, Beckman Coulter, BD Biosciences, Merck Millipore, Orflo, Nexcelom Bioscience, IntelliCyt,

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#### Cell counting equipment model HMGULA\_IMM\_088\_001 | v1.0

procedureMetadata

Req. Analysis: false

Req. Upload: true

Is Annotated: false

**Options:** Countess Automated Cell Counter, Reichert Brightline, Gallios, BD LSR-II, Scepter, Attune, Moxi Z, 4468770, Cellometer Auto T4, iQue Screener PLUS,

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.....
```

#### Cell counting equipment name HMGULA\_IMM\_089\_001 | v1.0

procedureMetadata

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

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## Cell lysis buffer manufacturer HMGULA\_IMM\_090\_001 | v1.0

procedureMetadata

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

Options: eBioscience, BD PharmLyse, Jax, JMC, LONZA,

#### Cell lysis buffer catalog number HMGULA\_IMM\_091\_001 | v1.0

procedureMetadata

Req. Analysis: falseReq. Upload: trueIs Annotated: false

Options: 00-4300-54, 555899, home brew, 10-548E,

.....

#### Date and time of sacrifice HMGULA\_IMM\_092\_001 | v1.0

procedureMetadata

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

#### Date and time of sample preparation HMGULA\_IMM\_093\_001 | v1.0

procedureMetadata

Req. Analysis: falseReq. Upload: trueIs Annotated: false

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## Sample storage temperature until analysis (in Celsius) HM

GULA\_IMM\_094\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Req. Analysis: false Req. Req. Req. Req. Req. Req. Req. Req	eq. Upload: true	Is Annotated: fa	alse
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Unit Measured: C

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#### FCS repository reference (URL/ID) HMGULA\_IMM\_095\_001 | v1.0

procedureMetadata

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

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#### Balanced salt solution type HMGULA\_IMM\_096\_001 | v1.0

procedureMetadata

Req. Analysis: false	Req. Upload: true	Is Annotated: false
Options: HBSS, PBS,		

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## Balanced salt solution manufacturer HMGULA\_IMM\_097\_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Req. Analysis:	eq. Upload: true	Is Annotated: false
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Options: Sigma, Life Technologies, Wisent, Wako, Gibco, Biochrom,

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## Balanced salt solution catalog number HMGULA\_IMM\_098\_001 | v1

.0

procedureMetadata

Req. Analysis: falseReq. Upload: trueIs Annotated: false

**Options:** D1408, H6136-1L, 041-20211, 14190-144, L 182-10, HBSS 1X 14170-088, 14175-095, 14190169,

#### **RPMI manufacturer** HMGULA\_IMM\_099\_001 | v1.0

procedureMetadata

Req. Analysis: false	Req. Upload: true	Is Annotated: false
Options: Sigma, Life Technolo	ogies, Jax, Wako, Gibco, none	used,
RPMI catalog numb	<b>Der</b> HMGULA_IMM_100_00	1   v1.0
Req. Analysis: false	Req. Upload: true	Is Annotated: false
<b>Options:</b> R8758, 11875-101, H	home brew, 189-02145, 31800-	-022, none used, 11875-093,
DNAse I manufacturer HMGULA_IMM_101_001   v1.0 procedureMetadata		
Req. Analysis: false	Req. Upload: true	Is Annotated: false

Options: Sigma,

## DNAse I catalog number HMGULA\_IMM\_102\_001 | v1.0

procedureMetadata

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

Options: DN25, D8764,

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#### Dead cell exclusion dye HMGULA\_IMM\_103\_001 | v1.0

procedureMetadata

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

Options: Sytox Blue, Sytox Green, Zombie NIR, DAPI, Propidium Iodide,

#### Dead cell exclusion dye manufacturer HMGULA\_IMM\_104\_001 | v1 .0

procedureMetadata

Req. Analysis: false	Req. Upload: true	Is Annotated: false
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Options: Life Technologies, Biolegend, Sigma, home brew,

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#### Dead cell exclusion dye catalog number HMGULA\_IMM\_105\_001 | v1.0

procedureMetadata

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

Options: S34857, S-34860, 423106, D9542, S11348, home brew, R37606, P4170,

# Cell digestion temperature (in Celsius) HMGULA\_IMM\_106\_001 | v1

procedureMetadata   Req. Analysis: false Req. Upload: true Is Annotated: false	
Reg. Analysis: false Reg. Upload: true Is Annotated: false	
Reg. Analysis: false Reg. Upload: true Is Annotated: false	
Options: 37, RT,	
Panel A FCS file(s) HMGULA_IMM_107_001   v1.0	

seriesMediaParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Increments: Minimum 1		
Panel B FCS file(s) seriesMediaParameter	HMGULA_IMM_108_001   v	1.0

Req. Analysis: false	Req. Upload: false	Is Annotated: false
	ited. opiedai laide	
Increments: Minimum 1		

#### Automated analysis HMGULA\_IMM\_109\_001 | v1.0

procedureMetadata

Req. Analysis: false	Req. Upload: true	Is Annotated: false
Options: Yes, No,		

## Collection buffer manufacturer HMGULA\_IMM\_110\_001 | v1.0

procedureMetadata

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

**Options:** Life Technologies,

# Collection buffer catalog number number HMGULA\_IMM\_111\_001

| v1.0 procedureMetadata

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

**Options:** 24020,

## FACS buffer manufacturer HMGULA\_IMM\_112\_001 | v1.0

procedureMetadata

Req.	Ana	lysis:	false
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Options: Life Technologies,

#### FACS buffer catalog number HMGULA\_IMM\_113\_001 | v1.0

procedureMetadata

Req. Analysis: false	Req. Upload: false	Is Annotated: false
<b>Options:</b> 14175,		

#### Enzyme buffer manufacturer HMGULA\_IMM\_114\_001 | v1.0

procedureMetadata

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

**Options:** Life Technologies,

#### Enzyme buffer catalog number HMGULA\_IMM\_115\_001 | v1.0

procedureMetadata

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

**Options:** 14025,