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 80
- 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 enzyme buffer 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 enzyme buffer (see below), aliquoted and stored at -20°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)

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 with Ca2+ and Mg2+ OR
 - HBSS 1X with Ca2+/Mg2+
 - 2% CS (v/v):
 - 10mM HEPES
- RBC Lysis buffer: Prepare a 1X solution in ddH₂0 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 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 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 1x10⁸ 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 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 of low frequencies (macrophages, DCs).

Gating Panel 1

| Parameters | Gating steps | | | |
|------------------------------|---------------------------|------|--------|--------|
| Panel A live leukocyte count | | | | Τ |
| T cells (panel A) | number of live leukocytes | CD5+ | CD161- | \top |
| 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- | П |
| 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 |
| | i | | İ | |

| 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

| | 1 | | | | |
|------------------------------|------------------|--------|------------|------|---|
| 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+ | |
| 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+ | \mathbf{L} |
| cDCs CD11b Type | Not Eosinophils | MHCII+ | CD19- | CD11c+ | С |
| pDCs | Not Eosinophils | Not T Cells | Ly6C+ | CD317+ | \mathbf{L} |
| RP Macrophage (F4/80+) | Not Eosinophils | MHCII+ | F4/80+ | | \mathbf{L} |
| or | | | | | \mathbf{L} |
| RP Macrophage (CD19-CD11c-) | Not Eosinophils | MHCII+ | CD19- | CD11c- | \prod |

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

simpleParameter

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

| Unit Measured: % | | |
|--------------------------------------|--------------------------|--------------------|
| | | |
| T cells (panel A) HN simpleParameter | MGULA_IMM_003_001 v1.0 | |
| 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) simpleParameter | HMGULA_IMM_005_001 v1 | 1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| Others HMGULA_IMM_0 simpleParameter | 006_001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |

| CD4 T cells HMGULA_simpleParameter | _IMM_007_001 v1.0 | |
|-------------------------------------|---------------------|---------------------|
| | Req. Upload: false | |
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| CD8 T cells HMGULA_simpleParameter | _IMM_008_001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| DN T cells HMGULA_INsimpleParameter | MM_009_001 v1.0 | |
| | Req. Upload: false | |
| | | |
| DP T cells HMGULA_INsimpleParameter | MM_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 HMG simpleParameter | GULA_IMM_012_001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |
| DN NKT cells HMGU simpleParameter | LA_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 |
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| | | |
| CD8 CD25+ T cells simpleParameter | HMGULA_IMM_016_001 v | 1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
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| CD8 CD25- T cells simpleParameter | HMGULA_IMM_017_001 v1 | 1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
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| DN CD25+ T cells H simpleParameter | HMGULA_IMM_018_001 v1. | 0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
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| DN CD25- T cells H simpleParameter | MGULA_IMM_019_001 v1.0 |) |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |

| CD4 CD25+ NKT ce simpleParameter | ells HMGULA_IMM_020_00 | 1 v1.0 |
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| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| CD4 CD25- NKT ce | IIS HMGULA_IMM_021_001 | v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| CD8 CD25+ NKT ce | PIIS HMGULA_IMM_022_00 | 1 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| CD8 CD25- NKT ce simpleParameter | IIS 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 | |
|---|----------------------|------------------------|
| DN CD25- NKT cell simpleParameter | S HMGULA_IMM_025_001 | v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
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| Total number of ac 001 v1.0 simpleParameter | quired events in Pa | anel A HMGULA_IMM_026_ |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |
| Total number of ac 001 v1.0 simpleParameter | quired events in Pa | nel B HMGULA_IMM_027_ |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |

CD4 CD44+CD62L- T cells HMGULA_IMM_028_001 | v1.0

simpleParameter

| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
|--------------------------------|-------------------------------|---------------------|
| | | |
| CD4 CD44+CD62L simpleParameter | + T cells hmgula_imm | _029_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| CD4 CD44-CD62L-simpleParameter | + T cells hmgula_imm_ | .030_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| CD4 CD44-CD62L-simpleParameter | - T cells HMGULA_IMM_0 | 031_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |

CD8 CD44+CD62L- T cells HMGULA_IMM_032_001 | v1.0

| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
|--------------------------------|------------------------|---------------------|
| | | |
| CD8 CD44+CD62L-simpleParameter | + T cells hmgula_imm_ | _033_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| CD8 CD44-CD62L+simpleParameter | - T cells hmgula_imm_o | 034_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| CD8 CD44-CD62L-simpleParameter | T cells HMGULA_IMM_0 | 35_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |
| DN CD44+CD62L- | T cells HMGULA_IMM_03 | 6_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |

| DN CD44+CD62L+ simpleParameter | - T cells hmgula_imm_0 | 037_001 v1.0 |
|--------------------------------|------------------------|---------------------|
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |
| DN CD44-CD62L+ simpleParameter | T cells HMGULA_IMM_0 | 38_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |
| DN CD44-CD62L- | T cells hmgula_imm_03 | 39_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |
| CD4 CD44+CD62L simpleParameter | NKT cells hmgula_i | MM_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 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false CD8 CD44+CD62L- NKT cells HMGULA IMM 043 001 Lv1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: false 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

| Req. Analysis: false | | Is Annotated: false |
|----------------------------------|------------------------|---------------------|
| | NKT cells hmgula_imn | |
| simpleParameter | THE COME THE COLA_INIT | VI_040_001 V1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| DN CD44+CD62L+ simpleParameter | NKT cells HMGULA_IM | M_047_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| DN CD44-CD62L+ I simpleParameter | NKT cells hmgula_imm | M_048_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |

Percentage of live gated events in Panel B HMGULA_IMM_049_0 01 | v1.0

simpleParameter

| Req. Analysis: false | Req. Upload: false | Is Annotated: false |
|------------------------------------|---------------------|---------------------|
| Unit Measured: % | | |
| | | |
| Neutrophils HMGULA simpleParameter | _IMM_050_001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| Monocytes HMGULA_simpleParameter | IMM_051_001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| Eosinophils HMGULA simpleParameter | _IMM_052_001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |

NK Cells (panel B) HMGULA_IMM_053_001 | v1.0

simpleParameter

| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
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| | | |
| NK Subsets (Q1) H simpleParameter | MGULA_IMM_054_001 v1.0 |) |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| NK Subsets (Q2) H simpleParameter | MGULA_IMM_055_001 v1.0 | 0 |
| | Req. Upload: false | Is Annotated: true |
| | | |
| NK Subsets (Q3) H | MGULA_IMM_056_001 v1.0 |) |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| NK Subsets (Q4) H simpleParameter | MGULA_IMM_057_001 v1.0 | 0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |

| NKT Cells (panel EsimpleParameter | 3) HMGULA_IMM_058_001 | v1.0 |
|-------------------------------------|------------------------------|--------------------|
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| NKT Subsets (Q1) simpleParameter | HMGULA_IMM_059_001 v | 1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| NKT Subsets (Q3) simpleParameter | HMGULA_IMM_060_001 v | 1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| T Cells (panel B) H simpleParameter | MGULA_IMM_061_001 v1.0 |) |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |

simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true B Cells HMGULA IMM 063 001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true B1B Cells HMGULA IMM 064 001 | v1.0 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true

T Subset HMGULA_IMM_062_001 | v1.0

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

Follicular B Cells HMGULA_IMM_066_001 | v1.0

B2B Cells HMGULA_IMM_065_001 | v1.0

simpleParameter

| | Req. Upload: false | |
|------------------------------------|----------------------------|----------------------------|
| | | |
| simpleParameter | CD21/35+) HMGULA_IM | M_067_001 v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| Tues all D Oall | _ | |
| simpleParameter | S HMGULA_IMM_068_001 | v1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| Transitional R Cells | s (CD21/35 low) HMG | LII A IMM 060 001 Lv1 0 |
| simpleParameter | S (ODZ 1/33 IOW) TIME | OLA_IIVIIVI_009_001 V1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| MZB HMGULA_IMM_070_simpleParameter | _001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |

| MZB (CD21/35 high simpleParameter |) HMGULA_IMM_071_001 | v1.0 |
|-------------------------------------|-------------------------|--------------------|
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| cDCs HMGULA_IMM_072 simpleParameter | 2_001 v1.0 | |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| cDCs CD11b Type simpleParameter | HMGULA_IMM_073_001 v1 | 1.0 |
| Req. Analysis: false | Req. Upload: false | Is Annotated: true |
| | | |
| pDCs HMGULA_IMM_074 simpleParameter | 4_001 v1.0 | |
| 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 (CD19- CD11c-) HMGULA_IMM_076_001 | v1.0

simpleParameter

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

Equipment name HMGULA_IMM_077_001 | v1.0

procedureMetadata

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

Options: FACS, Fortessa_1, LSR II, Flow cytometer,

.....

Equipment manufacturer HMGULA_IMM_078_001 | v1.0

procedureMetadata

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

Options: BD Biosciences, Beckman Coulter, IntelliCyt,

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, CS&T Bead lot HMGULA_IMM_080_001 | v1.0 procedureMetadata Reg. Analysis: false Reg. Upload: true Is Annotated: false Anesthesia HMGULA IMM 081 001 | v1.0 procedureMetadata Reg. Analysis: true Reg. 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,

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: false Req. Upload: true Is Annotated: false

Cell digestion agent catalog number HMGULA_IMM_085_001 | v1.0

procedureMetadata

Options: Roche, Worthington, Gibco, Sigma,

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 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 Options: Life Technologies, American Optical, Beckman Coulter, BD Biosciences, Merck Millipore, Orflo, Nexcelom Bioscience, IntelliCyt,

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, | | |
|---|---------------------------|-----------------------|
| | | |
| Cell counting equiposedureMetadata | oment name HMGULA | _IMM_089_001 v1.0 |
| Req. Analysis: false | Req. Upload: true | Is Annotated: false |
| | | |
| Cell lysis buffer ma | anufacturer HMGULA_ | IMM_090_001 v1.0 |
| Req. Analysis: false | Req. Upload: true | Is Annotated: false |
| Options: eBioscience, BD Pha | armLyse, Jax, JMC, LONZA, | |
| | | |
| Cell lysis buffer car | talog number нмвог | _A_IMM_091_001 v1.0 |
| Req. Analysis: false | Req. Upload: true | Is 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 sa procedureMetadata | ample preparation ⊢ | MGULA_IMM_093_001 v1.0 |
| Req. Analysis: false | Req. Upload: true | Is Annotated: false |
| Sample storage ter GULA_IMM_094_001 v1.0 | nperature until anal | lysis (in Celsius) нм |
| procedureMetadata Req. Analysis: false | Req. Upload: true | Is Annotated: false |
| Unit Measured: C | | |
| FCS repository refe | erence (URL/ID) нмс | ULA_IMM_095_001 v1.0 |
| procedureMetadata Req. Analysis: false | Req. Upload: false | Is Annotated: false |
| | | |

Balanced salt solution type HMGULA_IMM_096_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: true Is Annotated: false Options: HBSS, PBS, Balanced salt solution manufacturer HMGULA_IMM_097_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: true Is Annotated: false Options: Sigma, Life Technologies, Wisent, Wako, Gibco, Biochrom, Balanced salt solution catalog number HMGULA_IMM_098_001 | v1 .0 procedureMetadata Req. Analysis: false Req. Upload: true **Is 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 Technologies, Jax, Wako, Gibco, none used,

RPMI catalog number HMGULA_IMM_100_001 | v1.0

procedureMetadata

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

Options: R8758, 11875-101, 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

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

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

Options: Life Technologies, Biolegend, Sigma, home brew,

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

.0

procedureMetadata

Req. Analysis: false Req. 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) HMGULA_IMM_108_001 | v1.0

seriesMediaParameter

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

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 I v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false **Options:** 24020,

FACS buffer manufacturer HMGULA_IMM_112_001 | v1.0

Req. Analysis: false **Req. Upload:** false Is Annotated: false Options: Life Technologies, FACS buffer catalog number HMGULA_IMM_113_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: false Is Annotated: false **Options:** 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,