FACS ICS 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 16
- 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+		\mathbf{I}
T Subset	Not Eosinophils	CD161-	CD5+	Ly6C+	\mathbf{I}
B Cells	Not Eosinophils	MHCII+	CD19+		\mathbf{I}
B1B Cells	Not Eosinophils	MHCII+	CD19+	CD5+	\mathbf{I}
B2B Cells	Not Eosinophils	MHCII+	CD19+	CD5-	\perp
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+	\perp
cDCs CD11b Type	Not Eosinophils	MHCII+	CD19-	CD11c+	С
pDCs	Not Eosinophils	Not T Cells	Ly6C+	CD317+	\perp
RP Macrophage (F4/80+)	Not Eosinophils	MHCII+	F4/80+		\perp
or					\perp
RP Macrophage (CD19-CD11c-)	Not Eosinophils	MHCII+	CD19-	CD11c-	

Parameters and Metadata

Spleen weight IMPC_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 IMPC_IMM_002_001 |

v1.7

simpleParameter

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Unit Measured: %		
T cells (panel A) IMI simpleParameter	PC_IMM_003_001 v1.5	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
NKT cells (panel A) simpleParameter) IMPC_IMM_004_001 v1.5	
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NK cells (panel A) simpleParameter	IMPC_IMM_005_001 v1.5	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Others IMPC_IMM_006_simpleParameter	_001 v1.4	
Req. Analysis: false	Req. Upload: false	Is Annotated: true

0D4 T		
CD4 T cells IMPC_IMN simpleParameter	/I_007_001 v1.4	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
CD8 T cells IMPC_IMM	/I_008_001 v1.4	
simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true
DN T cells IMPC_IMM_ simpleParameter	_009_001 v1.5	
•		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
DP T cells IMPC_IMM_ simpleParameter	010_001 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: false

CD4 NKT cells IMPC_IMM_011_001 | v1.4

simpleParameter

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CD8 NKT cells IMPC_IMM_012_001 | v1.5

simpleParameter

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DN NKT cells IMPC_IMM_013_001 | v1.4

simpleParameter

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CD4 CD25+ T cells IMPC_IMM_014_001 | v1.4

simpleParameter

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CD4 CD25- T cells IMPC_IMM_015_001 | v1.4

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CD8 CD25+ T cells simpleParameter	IMPC_IMM_016_001 v1.4	
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CD8 CD25- T cells simpleParameter	IMPC_IMM_017_001 v1.4	
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DN CD25+ T cells II simpleParameter	MPC_IMM_018_001 v1.5	
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DN CD25- T cells IN simpleParameter	MPC_IMM_019_001 v1.5	
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CD4 CD25+ NKT cosimpleParameter	ells IMPC_IMM_020_001	v1.4
	Req. Upload: false	
CD4 CD25- NKT ce	ells	
simpleParameter Req. Analysis: false	Req. Upload: false	Is Annotated: true
	ells IMPC_IMM_022_001	
	Req. Upload: false	
CD8 CD25- NKT ce	EIIS IMPC_IMM_023_001 v	1.5
	Req. Upload: false	

DN CD25+ NKT cells IMPC_IMM_024_001 | v1.4

simpleParameter

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DN CD25- NKT cel	 S MPC_IMM_025_001 v1	.2
Req. Analysis: false	Req. Upload: false	Is Annotated: true
CD4 CD44+CD62L simpleParameter	T cells IMPC_IMM_028	_001 v1.2
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CD4 CD44+CD62L simpleParameter	+ T cells IMPC_IMM_029	9_001 v1.2
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CD4 CD44-CD62L+ T cells IMPC_IMM_030_001 | v1.2

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CD4 CD44-CD62LesimpleParameter	- T cells IMPC_IMM_031	_001 v1.2
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CD8 CD44+CD62L simpleParameter	T cells IMPC_IMM_032	2_001 v1.2
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CD8 CD44+CD62L simpleParameter	.+ T cells IMPC_IMM_03	3_001 v1.2
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CD8 CD44-CD62LesimpleParameter	+ T cells IMPC_IMM_034	001 v1.2
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CD8 CD44-CD62L-simpleParameter	T cells IMPC_IMM_035_0	001 v1.2
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DN CD44+CD62L- simpleParameter	Cells IMPC_IMM_036_0	01 v1.3
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DN CD44+CD62L+ simpleParameter	T cells IMPC_IMM_037_0	001 v1.3
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DN CD44-CD62L+ simpleParameter	Cells IMPC_IMM_038_0	01 v1.3
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DN CD44-CD62L- T cells IMPC_IMM_039_001 | v1.3

simpleParameter

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CD4 CD44+CD62L simpleParameter	- NKT cells IMPC_IMM	_040_001 v1.2
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CD4 CD44+CD62L simpleParameter	+ NKT cells IMPC_IMM	1_041_001 v1.2
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CD4 CD44-CD62L-simpleParameter	+ NKT cells IMPC_IMM	_042_001 v1.3
Req. Analysis: false	Req. Upload: false	Is Annotated: false

CD8 CD44+CD62L- NKT cells IMPC_IMM_043_001 | v1.3

Req. Analysis: false	Req. Upload: false	Is Annotated: false
CD8 CD44+CD62L	.+ NKT cells IMPC_IMN	/ 044 001 Lv1 3
simpleParameter	T IVIT CONS IVIT C_IVIV	1_044_001 V1.5
Req. Analysis: false	Req. Upload: false	Is Annotated: false
CD8 CD44-CD62L simpleParameter	+ NKT cells IMPC_IMM	_045_001 v1.3
Req. Analysis: false	Req. Upload: false	Is Annotated: false
DN CD44+CD62L-simpleParameter	NKT cells IMPC_IMM_0	046_001 v1.2
Req. Analysis: false	Req. Upload: false	Is Annotated: true
DN CD44+CD62L+ NKT cells IMPC_IMM_047_001 v1.2 simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true

DN CD44-CD62L+ I simpleParameter	NKT cells IMPC_IMM_04	48_001 v1.3
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Percentage of live	gated events in Par	nel B IMPC_IMM_049_001
v1.5 simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Unit Measured: %		
Neutrophils IMPC_IMI	M_050_001 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Monocytes IMPC_IMM simpleParameter	1_051_001 v1.2	

Req. Upload: false Is Annotated: true

Req. Analysis: false

Eosinophils IMPC_IM simpleParameter	M_052_001 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
NK Cells (panel B) simpleParameter	IMPC_IMM_053_001 v1.3	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
NK Subsets (Q1) IM simpleParameter	PC_IMM_054_001 v1.3	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
NK Subsets (Q2) IM simpleParameter	PC_IMM_055_001 v1.2	
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NK Subsets (Q3) IMPC_IMM_056_001 | v1.2

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true
NK Subsets (Q4) I simpleParameter	MPC_IMM_057_001 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
NKT Cells (panel B) IMPC_IMM_058_001 v1.2 simpleParameter		
Req. Analysis: false	Req. Upload: false	
. ,	Neq. Opioau. Taise	Is Annotated: true
	Req. Opload. laise	Is Annotated: true
		Is Annotated: true
	IMPC_IMM_059_001 v1.2	Is Annotated: true
NKT Subsets (Q1)		Is Annotated: true

NKT Subsets (Q3) IMPC_IMM_060_001 | v1.2

Req. Analysis: false Req. Upload: false Is Annotated: true T Cells (panel B) IMPC_IMM_061_001 | v1.2 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true T Subset IMPC_IMM_062_001 | v1.2 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true **B Cells** IMPC IMM 063 001 | v1.2 simpleParameter Req. Analysis: false Req. Upload: false Is Annotated: true

B1B Cells IMPC_IMM_064_001 | v1.2

simpleParameter

Req. Analysis: false	Req. Upload: false	Is Annotated: true
B2B Cells IMPC_IMM_simpleParameter	_065_001 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Follicular B Cells III simpleParameter	MPC_IMM_066_001 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Follicular B Cells (simpleParameter	CD21/35+) IMPC_IMM_	067_001 v1.1
Req. Analysis: false	Req. Upload: false	Is Annotated: true
Transitional B Cell simpleParameter	S IMPC_IMM_068_001 v1.	3
Req. Analysis: false	Req. Upload: false	Is Annotated: true

Transitional B Cells (CD21/35 low) IMPC_IMM_069_001 v1.2 simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true
MZB IMPC_IMM_070_00 simpleParameter	01 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
MZB (CD21/35 high) IMPC_IMM_071_001 v1.1 simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true
cDCs IMPC_IMM_072_001 v1.2 simpleParameter		
Req. Analysis: false	Req. Upload: false	Is Annotated: true

cDCs CD11b Type IMPC_IMM_073_001 | v1.2

simpleParameter

	Req. Upload: false	Is Annotated: true
pDCs IMPC_IMM_074_00 simpleParameter	01 v1.2	
Req. Analysis: false	Req. Upload: false	Is Annotated: true
DD Magraphaga /F	4/00 - \	
simpleParameter	4/80+)	01 v1.1
Req. Analysis: false	Req. Upload: false	Is Annotated: true
DD Maaranhaga (C	D40 CD44e \	
simpleParameter	D19- CD11c-) IMPC_IN	ИМ_076_001 v1.1
Req. Analysis: false	Req. Upload: false	Is Annotated: true

Equipment name IMPC_IMM_077_001 | v1.0

Req. Analysis: false Req. Upload: true Is Annotated: false Options: FACS, Fortessa_1, LSR II, Flow cytometer, Equipment manufacturer IMPC_IMM_078_001 | v1.0 procedureMetadata Reg. Analysis: true Reg. Upload: true Is Annotated: false Options: BD Biosciences, Beckman Coulter, IntelliCyt, Equipment model IMPC_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, BD FACSVerse, iQue Screener PLUS,

CS&T Bead lot IMPC_IMM_080_001 | v1.0

procedureMetadata

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

Anesthesia IMPC 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 IMPC_IMM_082_001 | v1.0

procedureMetadata

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

Options: GentleMACS, manual, manual with needles,

Cell digestion agent IMPC_IMM_083_001 | v1.0

procedureMetadata

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

Options: Collagenase D, Collagenase II, Spleen Dissociation Kit,

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Cell digestion agent manufacturer IMPC_IMM_084_001 | v1.1

procedureMetadata

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

Options: Roche, Worthington, Gibco, Miltenyi Biotec,

Cell digestion agent catalog number IMPC_IMM_085_001 | v1.2

procedureMetadata

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

Options: #11088858001, CLS2LS004176, 17101-015, 130-095-926,

Cell counting IMPC_IMM_086_001 | v1.1

procedureMetadata

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

Options: pre-lysis, post-lysis,

Cell counting equipment manufacturer IMPC_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,		
Cell counting equiports procedure Metadata	oment model IMPC_IM	IM_088_001 v1.0
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,		
	oment name IMPC_IMN	
Req. Analysis: false	Req. Upload: true	Is Annotated: false
Cell lysis buffer manufacturer IMPC_IMM_090_001 v1.2 procedureMetadata		
Req. Analysis: false	Req. Upload: true	Is Annotated: false
Options: eBioscience, BD PharmLyse, Jax, JMC, LONZA, home brew,		

Cell lysis buffer catalog number IMPC_IMM_091_001 | v1.2

procedureMetadata

Req. Analysis: false Req. Upload: true Is Annotated: false **Options:** 00-4300-54, 555899, home brew, 10-548E, Date and time of sacrifice IMPC IMM 092 001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: true Is Annotated: false Date and time of sample preparation IMPC_IMM_093_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: true Is Annotated: false

Sample storage temperature until analysis (in Celsius) IMP

C_IMM_094_001 | v1.0

procedureMetadata

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

Unit Measured: C		
FCS repository ref	erence (URL/ID) IMPO	C_IMM_095_001 v1.0
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Total number of ac v1.4 simpleParameter	equired events in Pa	anel A IMPC_IMM_026_001
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Total number of ac v1.2 simpleParameter	equired events in Pa	anel B IMPC_IMM_027_001
Req. Analysis: false	Req. Upload: false	Is Annotated: false

Balanced salt solution type IMPC_IMM_096_001 | v1.0

procedureMetadata

Req. Analysis: false Req. Upload: true Is Annotated: false Options: HBSS, PBS, Balanced salt solution manufacturer IMPC IMM 097 001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: true Is Annotated: false **Options:** Sigma, Life Technologies, Wisent, Wako, Gibco, Biochrom, home brew, Balanced salt solution catalog number IMPC_IMM_098_001 | v1.1 procedureMetadata Reg. Analysis: false Reg. Upload: true Is Annotated: false Options: D1408, H6136-1L, 041-20211, 14190-144, L 182-10, HBSS 1X 14170-088, 14175-095, home brew, RPMI manufacturer IMPC IMM 099 001 | v1.0 procedureMetadata

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

Options: Sigma, Life Technologies, Jax, Wako, Gibco, none used,

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RPMI catalog number IMPC_IMM_100_001 | v1.1

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 IMPC_IMM_101_001 | v1.1

procedureMetadata

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

Options: Sigma, Spleen Dissociation Kit,

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DNAse I catalog number IMPC_IMM_102_001 | v1.2

procedureMetadata

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

Options: DN25, D8764, 130-095-926,

Dead cell exclusion dye IMPC_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 IMPC_IMM_104_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: true Is Annotated: false **Options:** Life Technologies, Biolegend, Sigma, home brew, Dead cell exclusion dye catalog number IMPC_IMM_105_001 | v1. 1 procedureMetadata Req. Analysis: false Req. Upload: true Is Annotated: false **Options:** S34857, S-34860, 423106, D9542, S11348, home brew, R37606,

Cell digestion temperature (in Celsius) IMPC_IMM_106_001 | v1.1

procedureMetadata

Req. Analysis: false Req. Upload: true Is Annotated: false Options: 37, RT, Panel A FCS file(s) IMPC_IMM_107_001 | v1.0 seriesMediaParameter Req. Analysis: false Req. Upload: false Is Annotated: false **Increments:** Minimum 1 Panel B FCS file(s) IMPC_IMM_108_001 | v1.0 seriesMediaParameter Reg. Analysis: false Reg. Upload: false Is Annotated: false **Increments:** Minimum 1 Automated analysis IMPC_IMM_109_001 | v1.0 procedureMetadata Req. Analysis: false Req. Upload: true Is Annotated: false

Options: Yes, No,

Collection buffer m	nanufacturer імрс_імі	M_110_001 v1.2
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: Life Technologies,		
Collection buffer ca.2 procedureMetadata	atalog number num	ber IMPC_IMM_111_001 v1
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: 24020,		
FACS buffer manuf procedureMetadata	facturer IMPC_IMM_112	_001 v1.1
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: Life Technologies,		

FACS buffer catalog number IMPC_IMM_113_001 | v1.1

procedureMetadata

Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: 14175,		
Enzyme buffer ma	nufacturer IMPC_IMM_	114_001 v1.1
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: Life Technologies,		
	alog number імрс_ім	
Req. Analysis: false	Req. Upload: false	Is Annotated: false
Options: 14025,		