

Protocol for evaluating in vivo the activation of the P2RX7 immunomodulator

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1	Pro	ptocol for evaluating <i>in vivo</i> the activation of
2		the P2RX7 immunomodulator
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1 Abstract

2 Background: P2RX7 is a purinergic receptor with pleiotropic activities that is activated by 3 high levels of extracellular ATP that are found in inflamed tissues. P2RX7 has immunomodulatory and anti-tumor proprieties and is therefore a therapeutic target for various 4 5 diseases. Several compounds are developed to either inhibit or enhance its activation. 6 However, studying their effect on P2RX7's activities is limited to in vitro and ex vivo studies 7 that require the use of unphysiological media that could affect its activation. Up to now, the only way to assess the activity of P2RX7 modulators on the receptor in vivo was in an indirect 8 9 manner.

10 **Results:** We successfully developed a protocol allowing the detection of P2RX7 activation *in* 11 *vivo* in lungs of mice, by taking advantage of its unique macropore formation ability. The 12 protocol is based on intranasal delivery of TO-PROTM-3, a non-permeant DNA intercalating 13 dye, and fluorescence measurement by flow cytometry. We show that ATP enhances TO-14 PROTM-3 fluorescence mainly in lung immune cells of mice in a P2RX7-dependant manner.

Conclusions: The described approach has allowed the successful analysis of P2RX7 activity directly in the lungs of WT and transgenic C57BL6 mice. The provided detailed guidelines and recommendations will support the use of this protocol to study the potency of pharmacologic or biologic compounds targeting P2RX7.

19

20 Keywords: P2RX7, P2X7 receptor, purinergic receptors, ATP, macropore, TOPRO-3, activity

1 Background

2 The purinergic P2X family of receptors is a family of 7 receptors that are all activated by 3 extracellular ATP (eATP) with various affinities. Unlike the other 6 receptors, P2RX7 requires 4 high levels of eATP (in the range of hundreds of micromolar) to be activated. Such levels of 5 eATP are found in inflamed tissues such as the tumor microenvironment or fibrotic sites [1].

6 P2RX7 is a receptor with pleiotropic activities. Its activation leads to calcium influx and 7 potassium efflux, a feature it shares with all members of the P2X family and that has been 8 shown to trigger activation of various signaling pathways [2]. However, P2RX7 has its own 9 particularities: its activation also leads to a macropore opening at the plasma membrane that 10 allows the non-selective entry of macromolecules (up to 900 Da) to the cell. Compromising 11 membrane integrity through macropore opening by a prolonged or sustained activation (over 12 1 hour) of P2RX7 could also lead to cell death[3]. Moreover, activation of P2RX7 leads subsequently to the assembly of the NLRP3 inflammasome and the release of mature IL-1 β 13 14 and IL-18 from the cell [4].

Due to its ability to induce cell proliferation and death but also to modulate the immune response, several approaches are undertaken to either inhibit or enhance its activation, by engineering chemical molecules [5,6] or nanobodies [7] targeting the receptor. Three assays are classically and routinely used to evaluate their ability to modulate P2RX7's activation, based on the 3 hallmarks of P2RX7:

<u>Calcium influx:</u> the goal is to measure intracellular calcium concentrations using cell
 permeant fluorescent calcium indicators such as Fluo-4 AM or Fura 2. Calcium
 concentration could be assessed by spectrophotometry, microscopy or flow cytometry.
 Calcium influx could also be detected by measuring ion fluxes by electrophysiology
 techniques such as patch-clamp.

<u>Macropore opening</u>: To evaluate the opening of the macropores, non-permeant DNA
 intercalating fluorescent dyes of high molecular weight (up to 900 Da) are used, such
 as TO-PROTM-3 or Ethidium bromide. Fluorescence is assessed by
 spectrophotometry, microscopy or flow cytometry.

NLRP3 inflammasome activation is determined by checking adaptor protein ASC
 oligomerization, caspase-1 cleavage or IL-1β/IL-18 release. It is assessed by
 immunofluorescence, western blot, flow cytometry or ELISA.

8 However, these assays are in vitro or ex vivo-based and rely on using unphysiological medias 9 that could interfere with P2RX7 activation. Indeed, the ionic composition of media is crucial. 10 ATP-induced cell death has been shown to be delayed in low-salt medias[8]. Moreover, macropore formation is impacted by Na⁺, iodide, thiocyanate and chloride-enriched medias, 11 since increasing concentration of these ions inhibited ethidium uptake by cells[9]. Channel 12 13 opening is also affected by external anions such as glutamate that potentiates human P2RX7 14 activation whereas chloride and iodide decreased it [10]. Since P2RX7 activation hangs in the 15 balance of ions concentrations in media, measurement of its activation in vitro and its translation in vivo can be not accurate. For example, under pathological conditions, glutamate 16 levels increase in the extracellular space during nerve injury, that could affect P2RX7 17 18 activation [11].

Since compounds targeting P2RX7 are tested *in vitro*, little is known on their impact on P2RX7 activities *in vivo*, i.e. in complete physiological conditions. Up to now, *in vivo* P2RX7 activation state is only seen indirectly through eATP levels or IL-1β/IL-18 production. Therefore, we aimed at setting up a way to measure its activation *in vivo* and developed a protocol based on intranasal delivery of TO-PROTM-3 and fluorescence measurement by flow cytometry.

24

1 Results

2 General strategy to measure P2RX7 activity in vivo

Since calcium influx is a common feature of P2X receptors and that NLRP3 assembly is a step
downstream of P2RX7 but also not specific to P2RX7, we opted to use macropore opening,
which directly represent the activation state of the receptor, as a readout for P2RX7 activity
(Fig. 1). Several dyes are available for measuring macropore opening. We opted to use the
TO-PROTM-3 dye (Invitrogen) since it is the most sensitive non-permeant DNA intercalating
dye and was shown to work best in flow cytometry [12].



9 Fig. 1. Activation of P2RX7 and macropore opening using TO-PRO[™]-3 dye

When the cell is in steady state (absence of eATP, membrane integrity) TO-PRO[™]-3 does not 10 enter the cell and the cell remains non fluorescent (Fig. 1A). In the presence of eATP, P2RX7 11 is activated and macropores are open, permitting the entry of TO-PRO[™]-3 into the cell. TO-12 13 PRO[™]-3 binds to DNA and renders the cell fluorescent (Fig. 1B). Macropore formation is a 14 reversible mechanism. However, once P2RX7 is activated, the cell remains fluorescent since 15 TO-PRO[™]-3 has already bound to DNA, even if no eATP remains or if P2RX7 is internalized [13,14] or cleaved [15] during a prolonged activation (Fig. 1C and D), allowing the detection 16 of P2RX7 activation at any point in time. 17

18 Channel opening and P2RX7-dependent calcium influx are detected few seconds after ATP 19 stimulation. However, detection of macropore formation is much slower due to the 20 incorporation of fluorescent dyes in DNA. and can take up to 1 hour to detect all activated

receptors [12]. We have previously shown in a time course experiment of ATP-induced TO-1 2 PRO[™]-3 uptake that a range between 20 and 30 minutes post stimulation is the ideal time point at which the TO-PRO-3 signal is well above background [5]. Moreover, since prolonged 3 activation of P2RX7 (over 1 hour) can also lead to cell death in a macropore-dependent 4 5 manner and since TO-PROTM-3 can also detect dead cells, it is crucial to add a Live/Dead 6 marker. Indeed, excluding dead cells to reduce background noise and macropore-unrelated 7 TO-PRO[™]-3 staining is important for accurate P2RX7-activation measurement. Therefore, to detect P2RX7-related TO-PRO[™]-3 fluorescence and to rule out detection of ATP-induced cell 8 death, we chose to activate P2RX7 for 30 minutes in vivo. 9

10 ATP activates P2RX7 in the lung and increases *in vivo* TO-PRO[™]-3 fluorescence.

We first analyzed TO-PRO[™]-3 fluorescence in the whole lung. We observed an increase of 11 12 fluorescence in WT mice receiving ATP compared to PBS (Fig. 2A and B), indicative of macropore opening and therefore activation of P2RX7. In order to know if this increase is 13 14 specific of P2RX7, the experiment was also done on p2rx7^{-/-} mice, where the increase of TO-PRO[™]-3 fluorescence in the ATP group was not observed and was similar to WT-PBS mice 15 (Fig. 2B). No difference was observed in the percentage of TO-PRO[™]-3 positive cells (supp 16 Fig. 1A-C). These results suggest that TO-PRO[™]-3 fluorescence after ATP administration 17 18 was specific to P2RX7 and indicative of its activation state.

To go further, we analyzed TO-PRO[™]-3 fluorescence in immune and non-immune cells. We
show that WT-ATP mice exhibit an enhanced fluorescence compared to WT-PBS mice only
in immune cells (Fig. 2C), and that was only observed in WT mice, demonstrating furthermore
the requirement of P2RX7 for the TO-PRO[™]-3 signal observed.

We also show that TO-PRO[™]-3⁺ cells express P2RX7. However, percentage of P2RX7⁺ cells
 among TO-PRO[™]-3⁺ cells was decreased in ATP-mice (Fig. 2D and E), but not in TO-PRO[™] 3⁺CD45⁻ cells (Fig. 2E). This is not surprising since it has been shown that sustained activation

- 1 of P2RX7 leads to the internalization of the receptor[14] or its cleavage [15] after its activation,
- 2 explaining the possibility of TO-PRO[™]-3⁺P2RX7⁻ cells.



3 Fig. 2. P2RX7 is activated in vivo

- 4 A Dotplot gated on live single cells of WT mice whole lungs
- 5 **B** GMFI of TO-PROTM- 3^+ cells in whole lung of WT and $p2rx7^{-/-}$ mice.
- 6 **C** GMFI of TO-PROTM-3⁺ cells in CD45⁺ cells (left) and CD45⁻ cells (right) in WT and *p2rx7^{-/-}* mice
- 7 D Percentage of P2RX7⁺ cells among TO-PROTM-3⁺ cells,
- 8 E Percentage of P2RX7⁺ cells among TO-PRO^{TM-3+}CD45⁺ cells (left) and TO-PRO^{TM-3+}CD45⁻
- 9 cells (right). One point represents one mouse, data is represented as mean±SEM. Two-tailed
- 10 unpaired *t*-test. **p* < 0.05. WT: Wildtype, GMFI: geomean fluorescence intensity
- 11

12 P2RX7 is mainly expressed and activated in lung immune cells

- 13 Since we observed a decrease of P2RX7 expression in TO-PRO-3⁺ cells (Fig. 2D and E), we
- 14 investigated furthermore its expression in whole lungs of mice. We show that the expression
- 15 of P2RX7 was decreased in all cells (Fig. 3A) but also in immune cells (Fig. 3B), supporting
- 16 furthermore the idea of an internalization of P2RX7 after its activation. Moreover, we show
- 17 that P2RX7 is more expressed by immune cells than non-immune cells (Fig. 3B). Even though

Commenté [VV1]: In cells positive for TO-PRO-3

- 1 WT-ATP mice show a decrease of P2RX7 expression, P2RX7⁺ cells show a higher TO-
- 2 PRO[™]-3 fluorescence in all cells (Fig. 3C) and in immune cells (Fig. 3D), meaning that
- 3 P2RX7⁺ cells are indeed activated by ATP.



5 Fig 3. TO-PRO[™]-3 fluorescence is increased in P2RX7⁺ cells in WT mice

- 6 A Percentage of P2RX7⁺ cells in the whole lung of WT mice
- 7 B Percentage of P2RX7⁺ cells in CD45⁺ and CD45⁻ cells
- 8 C GMFI of TO-PRO[™]-3 in P2RX7⁺ cells
- 9 D GMFI of TO-PROTM-3 in P2RX7⁺CD45⁺ and P2RX7⁺CD45⁻ cells
- 10 One point represents one mouse, data is represented as mean±SEM. Two-tailed unpaired *t*-test.
- 11 *p < 0.05, **p<0.01. WT: Wildtype, GMFI: geomean fluorescence intensity
- 12

Since TO-PROTM-3 fluorescence (Fig. 2C) and P2RX7 expression (Fig. 2E, Fig. 3B) were
 unchanged after ATP delivery in CD45⁻ cells, we can speculate that P2RX7-expressing
 immune cells are the main cells activated by ATP in the lung. However, we observed a
 decrease of TO-PROTM-3 fluorescence in CD45⁻ P2RX7⁺ cells (Fig. 3D).

1 Discussion

2 This protocol was set up to assess P2RX7 activation in vivo, by taking advantage of its 3 property of macropore formation. Among P2X receptors, a permeability for large cations was also described for P2RX2 and P2RX4. However, P2RX7 is activated by high concentrations 4 of eATP (0.5 to 1 mM), whereas P2RX2 and P2RX4 are activated by much lower 5 concentrations (3 to 10 µM). Unlike P2RX7, P2RX2 and P2RX4 receptors desensitize after 6 7 ATP stimulation. Even though P2RX4 is broadly expressed in most tissues of the body, it is 8 preferentially localized in lysosomes where the acidity prevents its activation [16]. Based on these properties, we hypothesized that the macropore activity of P2RX2 and P2RX4 will not 9 10 interfere with our assay. This is indeed the case since no increase in TO-PRO[™]-3 signal was 11 observed in $p2rx7^{-}$ mice (Fig. 2B).

12 Our protocol allows the measurement of P2RX7 activation in lungs and is based on TO-PRO[™]-3 uptake by cells and detection by flow cytometry. Flow cytometry was previously 13 shown as a sensitive tool to study P2RX7 activation in vitro [12] and is also efficient for analysis 14 15 of an in vivo activation of the receptor as shown in this study. Moreover, flow cytometry allows the reduction of background noise by excluding non P2RX7-dependent TO-PRO[™]-3 staining 16 17 of dead cells. The other advantage of using flow cytometry is coupling TO-PRO[™]-3 with other 18 surface markers to distinguish P2RX7 activation at the single cell level. We only studied its 19 activation in immune versus non-immune cells, however, one can easily increase the number 20 of markers during the surface staining step to precisely identify the subtypes of immune and 21 non-immune cells activated by ATP.

To our knowledge, we show for the first time in a direct manner that P2RX7 is mainly activated
in lung immune cells of mice, as per an increase of TO-PROTM-3 fluorescence. P2RX7 has a
pro-inflammatory role since it releases IL-1β and is at the etiology of inflammatory diseases.
Nonetheless, direct activation of the receptor i.e. calcium influx or macropore formation was
never assessed *in vivo*. However, expression of P2RX7 on hematopoietic cells [17,18], ATP

levels [19,20] and IL-1β release [21,22] were shown in vivo to be important for airway 1 2 inflammation in mouse models. Even so, these experiments require the use of transgenic and chimeric mice and do not evaluate the receptor activation status. Our protocol could be used 3 instead for accurate and easier assessment of the activation of P2RX7. We previously showed 4 5 using human tumor lung adenocarcinoma (LUAD) tissues that P2RX7 is expressed in tumor 6 associated immune cells whereas non-immune cells do not express P2RX7. Further, we 7 showed that the overall activity of P2RX7 is impaired by expression of the C-terminal domain truncated P2RX7B receptor in immune cells [23]. Accordingly, we also show that ATP did not 8 enhance TO-PRO[™]-3 fluorescence in non-immune cells. Lung CD45⁻ cells comprise 9 10 endothelial cells, fibroblasts, and alveolar epithelial cells of type I (AEC I) and type II (AEC II). 11 They all express P2RX7 except for AEC II. It has been shown in vitro and ex vivo studies that 12 they express a functional receptor (macropore opening, calcium influx, IL-1β release) [24,25] but we do not seem to detect an activation in vivo with our protocol by measuring TO-PRO[™]-13 14 3 fluorescence in CD45⁻ cells. This could be due to the lower levels of expression of P2RX7 15 in CD45⁻ cells than in CD45⁺ cells (Fig. 3B, supp Fig. 2B) and the heterogeneity of the 16 amplitude of its activation in these cells that is lost when looking at overall CD45⁻ cells. One 17 possible way to overcome this is to analyze TO-PRO[™]-3 fluorescence in each CD45⁻ cell type 18 by including specific markers. However, we could not exclude that the overall macropore activity of P2RX7 is impaired by the expression of truncated splice variants as previously 19 described in mouse astrocytes [26]. Surprisingly, we observed that TO-PROTM-3 fluorescence 20 21 was decreased in P2RX7⁺CD45⁻ cells in ATP-mice (Fig 3D). It should be noted that the intensity of TOPRO[™]-3 fluorescence in P2RX7-expressing CD45⁻ cells (Fig. 3D) is identical 22 to the background value observed in $p2rx7^{\prime -}$ mice (Fig 2C) accompanied by unchanged 23 24 P2RX7 expression (Fig 3B, supp Fig 2B). Thus, these observations support the background 25 signal of TO-PRO-3 fluorescence rather than an ATP-related effect.

We have also shown that the expression of P2RX7 was decreased in WT-ATP mice. P2RX7
activation by ATP stimulates membrane internalization [27] that results to the loss of

expression of many surface proteins[28,29] including P2RX7 [13,14]. Indeed, prolonged 1 2 stimulation with high levels of ATP results in decreased surface expression and internalization 3 of the receptor in RAW cells [14], Caski cells and HEK 293 transfected with the human P2RX7 [13], at least 15 to 30 minutes after its activation depending on the cell type. Decreased 4 5 expression of the receptor was also reported on human and mouse monocytes after sustained 6 stimulation with bzATP [30]. Moreover, it has been shown that sustained activation of P2RX7 7 induces its cleavage at the plasma membrane by MMP-2, starting 15 minutes after its 8 activation [15]. Altogether, these observations support the decrease of P2RX7 surface expression in lungs of mice due to high levels of ATP administration and duration of P2RX7 9 10 activation in this study.

Even though this protocol successfully detects P2RX7 activation in the lungs, it can be adapted for studying its activation in other organs by giving TO-PROTM-3 to the mice by intravenous (i.v.) route. YO-PRO-1 (an analog of TO-PROTM-3) and propidium iodide (PI) are also used to study macropore formation. They have been given i.v. in mice and rats to study *in vivo* cell death in lungs [31,32], brain [33,34] and liver [35]. However, TO-PROTM-3's time to access and penetrate the organ of interest should be carefully assessed in order to study the activation of P2RX7.

To go further, this protocol could also be helpful to determine the *in vivo* efficacy of compounds targeting P2RX7, to potentially adapt their administration route or their dosage or even to identify the targeted cell type of the compounds. HEI3090 is a novel positive modulator of P2RX7 that enhances calcium influx, macropore opening and IL-18 release, as determined by *in vitro* and *ex vivo* experiments. Although we showed that HEI3090 enhances IL-18 release *in vivo* [5], it is of interest to study its impact on P2RX7 activation *in vivo*, especially since HEI3090 is a promising molecule for treatment of lung cancer [5].

25 Conclusions

- 1 We described a protocol assessing P2RX7 activation in vivo, by taking advantage of its unique
- 2 property of macropore formation. Given the immunomodulatory functions of P2RX7 and the
- 3 increasing number of molecules aiming to modulate its activity, this protocol represents an
- 4 important advance in the field of purinergic signaling.

5

1 Methods

The described methods include the description of the work, along with general guidelines that
can be used for implementing the strategy with various compounds described to modulate
P2RX7 activity.

5 Animals: 8 weeks old WT C57BL/6J Olahsd mice are from Envigo, (Gannat, France), and 8 weeks old p2rx7^{-/-} (B6.129P2-P2rx7tm1Gab/J) backcrossed more than 10 times with WT 6 7 C57BL/6J Olahsd and are inbreeded in our animal facility. Mice were housed under standardized light-dark cycles in a temperature-controlled air-conditioned environment under 8 9 specific pathogen-free conditions at IRCAN, Nice, France, with free access to food and water. 10 All experiments were approved by the committee for Research and Ethics of the local authorities (CIEPAL #772, protocol number MESRI APAFIS #33150-2021091413316813) and 11 followed the European directive 2010/63/UE, in agreement with the ARRIVE guidelines. 12 13 Experiments were performed in accord with animal protection representative at IRCAN.

- 14 Protocol description is summarized in Fig. 4. Measuring P2RX7 activity in vivo requires 5
- 15 sequential and mandatory steps as illustrated below



16

17 Fig. 4: Schematic protocol assessing P2RX7 activity in vivo

18 Time 0 - Intranasal administration

- 19 1. Anesthetize mouse i.p. with a mixture of 25 mg/kg ketamine and 2.5 mg/kg xylazine
- 20 using 1 ml syringe with a 26G needle.
- 2. Place mouse in a box with 5% isoflurane for 1 minute.

1	3.	When asleep, mouse is held upright and was given 25 μI of TO-PRO^TM-3 (1 $\mu M/kg)$ in	
2		one nostril then 25 μI of ATP (100 mM) or 1X PBS in another using a micropipette.	
3		/!\ Intranasal administration should be slow and follow the respiration rate of the mouse.	
4		/!\ Mice should be anesthetized and processed for intranasal delivery one by one.	
5		/!\ As soon as the 25 μ l+25 μ l are completely taken up by the mouse, start the 30	
6		minutes timer.	
7	Time	30 minutes – Mouse sacrifice and lung harvesting	
8	1.	Sacrifice mouse by cervical elongation	
9	2.	Spray mouse thorax with 70% ethanol. Harvest lungs using fine and sharp scissors	
10		and pliers.	
11	3.	Separate the lobes. Transfer the lobes into a GentleMACS $^{\mbox{\scriptsize TM}}$ C-tube (Miltenyi Biotech)	
12		containing the enzymatic buffer for dissociation (Lung dissociation kit, Miltenyi Biotech)	
13	4.	Dissociate lungs using the 37_m_LDK1 program on a GentleMACS $^{\rm TM}$ Octo dissociator	
14		with heaters (Miltenyi Biotech)	
15	5.	At the end of the program, resuspend cells and pass them through a 100 μm cell	
16		strainer on top of a 50 ml tube for single cell suspensions. Wash the GentleMACS $^{\rm TM}$	
17		C-tube with 1X Buffer S (Lung dissociation kit, Miltenyi Biotech) for maximum retrieval	
18		of cells and pass them though the 100 μm cell strainer.	
19	6.	Discard the cell strainer and centrifuge the tube at 1200 rpm 4°C for 5 minutes.	
20		/!\ Keep cells in the dark at all times, and on ice except during the dissociation program.	
21	Time	60 minutes – Red blood lysis and live/dead staining	
22	1.	Aspirate completely the supernatant and resuspend pellet gently with 1 ml of ice-cold	
23		ACK lysis buffer (Gibco) for 30 seconds to lyse red blood cells.	
24	2.	Add 30 ml of 1X PBS for stopping the reaction.	

- 25 3. Centrifuge at 1200 rpm 4°C for 5 minutes
- 26 4. Count cells

1	5.	Add 1 million of lung single cells in 150 μI 1X PBS in a 5 ml polypropylene tube	
2	6.	Add 150 μI of Green Live/Dead stain (2X) (Invitrogen) or 1X PBS (unstained control)	
3		into the 5 ml polypropylene tube	
4		/!\ Prepare appropriate number of samples without Live/Dead staining for single-	
5		stained controls	
6	7.	Vortex and incubate in the dark at room temperature for 30 minutes	
7	8.	Add 3 ml of FACS buffer (PBS 5% FBS 0.5% EDTA) into the 5 ml polypropylene tube	
8	9.	Centrifuge at 1200 rpm 4°C for 5 minutes	
9	10	. Aspirate supernatant completely and resuspend cells in 100 μI FACS Buffer	
10		/!\ Keep cells in the dark at all times, and on ice unless otherwise stated.	
11	Time	90 minutes – Surface staining	
40			
12	1.		
13	2.	Centriluge plate at 1300 rpm 4°C for 3 minutes	
14	3.	Flick the plate	
15	4.	Block FC receptor using anti-CD16/32 (dil 1/100, BD Blosciences) diluted in FACS	
16		Buffer (1X PBS, 5% FBS, 0.5% EDTA) in the dark under agitation for 10 minutes	
17	5.	Centrifuge plate at 1300 rpm 4°C for 3 minutes	
18	6.	Flick plate	
19	7.	Add antibody mix diluted in FACS buffer and homogenize immediately using a	
20		multichannel pipette.	
21		Antibodies: CD45 BUV395 (dil 1/100, BD Biosciences), P2RX7 PE (dil 1/8, Biolegend)	
22		or isotype control rat IgG2b κ PE (dil 1/8, Biolegend)	
23		/!\ Prepare appropriate single-stained controls	
24	8.	Incubate at 4°C in the dark under agitation for 20 minutes	
25	9.	Add 100 μI of FACS buffer per well using a multichannel pipette	
26	10	. Centrifuge plate at 1300 rpm 4°C for 3 minutes	
27	11	. Flick the plate	

1	12	. Resuspend cell in 100 μl of FACS buffer
2		/!\ Keep cells in the dark at all times, and on ice.
3	Time '	110 minutes – Flow cytometry analysis
4	1.	Cells were acquired using the CytoFLEX LX (Beckman Coulter)
5	2.	Samples were analyzed using FlowJo (LLC)
6		
7	Materi	ials
8	1.	1 ml syringe omnifix-F (Braun, catalog number 9161406V)
9	2.	26G needle (BD Microlance 3, catalog number 304300)
10	3.	Pipette
11	4.	Pipette tips
12	5.	Fine scissors
13	6.	Sharp scissors
14	7.	Pliers
15	8.	GentleMACS [™] C-tube (Miltenyi Biotech, catalog number 130-093-237)
16	9.	100 μm cell strainer (Falcon, catalog number 352360)
17	10	. Polypropylene 50 ml tube (Falcon, catalog number 352070)
18	11	. Polypropylene 5 ml tube (Falcon, catalog number 352096)
19	12	. 96-well clear V bottom plates (Greiner, catalog number 651101)
20	13	. Serological pipette (Falcon, catalog number 357543)
21	14	. Serological pipette gun
22	15	. Multi-channel pipette (Starlab, catalog number 57112-330)
23	Reage	ents

24 1. Ketamine (Virbac, catalog number 03597132111010)

- 1 2. Xylazine (Sedaxylan, Dechra, catalog number 08714225151523) 2 1. isofluorane / Aerrane (Baxter, catalog number DAGG9223) 2. TO-PRO[™]-3 (Life Technologies, catalog number T3605) 3 3. ATP (Sigma-Aldrich, catalog number A6419) 4 4. 1X Dulbecco's PBS (Gibco, catalog number 14190144) 5 6 5. 70% ethanol 7 6. Double distilled water 7. 20X Buffer S, enzyme D and enzyme A from the mouse lung dissociation kit (Miltenyi 8 Biotech, catalog number 130-095-927) 9 8. ACK lysis buffer (Gibco, catalog number A1049201) 10 11 9. LIVE/DEAD[™] fixable green dead cell staining kit (Invitrogen, catalog number L23101) 12 10. Fetal bovine serum (FBS) 11. Ethylenediaminetetraacetic acid (EDTA) (Invitrogen, catalog number 15575020) 13 14 12. CD16/32 (BD Biosciences, catalog number 553141) 13. CD45 BUV395 (BD Biosciences, catalog number 564279) 15 16 14. P2RX7 PE (Biolegend, catalog number 148703) 17 15. Rat IgG2b κ PE (Biolegend, catalog number 400607) 18 Equipment 19 1. Anesthesia machine and box (Anesteo) 2. GentleMACS[™] Octo dissociator with heaters (Miltenyi Biotech, catalog number 130-20 21 096-427)
- 22 3. Tube and plate centrifuge (Thermofischer scientific, catalog number 75009527)
- 23 4. Plate agitator
- 24 5. CytoFLEX LX (Beckman Coulter)
- 25 Reagent preparation
- 26 /!\ Prepare all reagents in sterile conditions using a PSM II tissue culture hood

1	1.	Anesthetics
2		Prepare 25 mg/kg of ketamine and 2.5 mg/kg of xylazine in 1X Dulbecco's PBS as per
3		250 μl of mix per 25 g of mouse.
4	2.	Lung dissociation buffer
5		Dilute 20X Buffer S in double distilled water and store at 4°C. Resuspend enzyme A $$
6		and enzyme D with 1X Buffer S according to manufacturer's instructions. Add 2.4 ml $$
7		of 1X Buffer S, 15 μI of enzyme A and 100 μI of enzyme D per GentleMACS^M C-tube
8		(Miltenyi Biotech)
9	3.	ATP
10		Prepare 100 mM stock of ATP in sterile 1X PBS, pH 6.8, aliquot and store at -80°C
11		/!\ Avoid freeze/thaw cycles
12	4.	<u>TO-PRO[™]-3</u>
13		Dilute TO-PRO [™] -3 in 1X PBS to 1 µM/kg
14		/!\ Manipulate TO-PRO TM -3 in the dark
15		/!\ Aliquot TO-PRO TM - 3 stock and avoid freeze/thaw cycles
16	5.	LIVE/DEAD [™] fixable green dead cell staining kit
17		Dilute LIVE/DEAD^{TM} in 1X PBS to 2X working concentration per tube. Add 0.4 μL
18		LIVE/DEAD [™] in 150µL per tube.
19		/!\ Manipulate LIVE/DEAD [™] green in the dark
20	6.	/!\ Prepare mix of appropriate volume
21	7.	FACS Buffer
22		Prepare 500 ml of FACS Buffer containing 1X Dulbecco's PBS, 5% FBS, 0.5% EDTA
23		and store at 4°C
24		/!\ Keep on ice at all times
25	8.	Antibody mix cocktail
26		Dilute CD45 BUV395 (dil 1/100) and P2RX7 PE (dil 1/8) or isotype control rat IgG2b κ
27		PE (dil 1/8) in FACS Buffer
28		/!\ Keep on ice at all times and manipulate in the dark

1 Software

- 2 1. CytExpert (Beckman Coulter)
- 3 2. FlowJo (LLC) or any other flow cytometry software
- 4 3. Graphpad prism

5 Statistical analyses:

- 6 All analyses were carried out using Prism software (GraphPad). Mouse experiments were
- 7 performed on at least n = 5 individuals, as indicated. Mice were equally divided for treatments
- 8 and controls. Data is represented as mean values and error bars represent SEM. Two-tailed
- 9 Mann–Whitney and unpaired *t*-test were used to evaluate the statistical significance between
- 10 groups.
- 11

1 **Declarations**

2 Author's contribution:

- 3 S.J.H. and V.V.-C. conceived and designed the study. S.J.H. developed the methodology.
 4 S.J.H, T.J. and V.V.-C. acquired the data (provided animals, provided facilities, and so on).
- 5 S.J.H, P.H. and V.V.-C. analyzed and interpreted the data. S.J.H. and V.V.-C. wrote the 6 manuscript. All authors reviewed the manuscript.
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- 17 Ethics Approval: All experiments were approved by the committee for Research and Ethics
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- 24

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fig3





SUPP







