

1 **IL-17 induced inflammation modulates mPGES-1/PPAR- $\gamma$  pathways in**  
2 **monocytes/macrophages**

3

4 **Running title:** IL-17A and mPGES-1/PPAR- $\gamma$  pathway

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41This article has been conducted and written in the absence of any commercial or financial  
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43

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45Data available on request from the authors The data that support the findings of this study are  
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47available because of privacy or ethical restrictions.

48

#### 49ABSTRACT

50**Background and Purpose:** Recent biochemical and pharmacological studies have reported that in  
51several tissues and cell types, microsomal prostaglandin E<sub>2</sub> synthase (mPGES) and peroxisome  
52proliferator-activated receptor- $\gamma$  (PPAR- $\gamma$ ) expression are modulated by a variety of inflammatory  
53factors and stimuli Considering that very little is known about the biological effects promoted by  
54IL-17 in the context of mPGES-1/PPAR- $\gamma$  modulation, we sought to investigate the contribution of  
55this unique cytokine on these integrated pathways during the onset of inflammation.

56**Experimental Approach:** We evaluated PF 9184 (mPGES-1 antagonist) and Troglitazone (PPAR-  
57 $\gamma$  agonist) activity utilising integrated *in vitro* and *in vivo* approaches. The dorsal air pouch model  
58was employed, and inflammatory infiltrates were analysed by flow cytometry. Locally produced  
59cyto-chemokines and prostaglandins were assessed using ELISA assays. Western blots were also  
60employed to determine the activity of various enzymes involved in downstream signalling  
61pathways.

62**Key Results:** PF 9184 and Troglitazone, in a time and dose-dependent manner, were shown to  
63significantly modulate leukocyte infiltration, myeloperoxidase activity, and the expression of COX-  
64/mPGES-1, NF- $\kappa$ B/I $\kappa$ B- $\alpha$  and mPGES-1/PPAR- $\gamma$  induced by IL-17. Moreover, both compounds  
65were found to modulate prostaglandins (PGE<sub>2</sub>, PGD<sub>2</sub>, and PGJ<sub>2</sub>) production, the expression of  
66different pro-inflammatory cyto-chemokines and the recruitment of inflammatory monocytes in  
67response to IL-17.

68**Conclusions and Implications:** Collectively, the data presented suggests that IL-17 may constitute  
69a specific modulator of inflammatory monocytes during later phases of the inflammatory response.  
70Therefore, the results of this study show, for the first time, that IL-17/mPGES-1/PPAR- $\gamma$  “axis”  
71could represent a potential therapeutic target for inflammatory-based and immune-mediated  
72diseases.

73**Keywords:** IL-17A, Inflammation, mPGES-1, PGE<sub>2</sub>, PPAR- $\gamma$ .

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75**Abbreviations.** BLC, B lymphocyte chemoattractant; C5a, complement component 5a; CD,  
76Crohn's disease; CMC, carboxymethyl cellulose; COX-, cyclooxygenase-; cPGES-, cytosolic  
77prostaglandin E<sub>2</sub> synthase; CTRL, control; 15d-PGJ<sub>2</sub>, 15-deoxy- $\Delta$ 12,14-prostaglandin J<sub>2</sub>; DMEM,  
78Dulbecco's modified Eagle's medium; FBS, fetal bovine serum; FCS, fetal calf serum; I $\kappa$ B- $\alpha$ ,  
79nuclear factor kappa-B inhibitor  $\alpha$ ; IL-, interleukin-; IMIDs, immune-mediated inflammatory  
80diseases; INF- $\gamma$ , interferon- $\gamma$ ; IP-10, interferon  $\gamma$ -induced protein-10; ITAC, interferon-inducible T-  
81cell  $\alpha$  chemoattractant; JE, junctional epithelium; KC, keratinocyte chemoattractant; LPS,  
82lipopolysaccharides; LT, leukotriene; MCP-, monocytes chemoattractant protein-; MCSF,  
83macrophage colony-stimulating factor; MIG, monokine induced by interferon  $\gamma$ ; MIPs, macrophage  
84inflammatory proteins; mPGES-, microsomal prostaglandin E<sub>2</sub> synthase-; MPO, myeloperoxidase;  
85NF- $\kappa$ b, nuclear factor kappa-B; NKT, natural killer T; NSAID, nonsteroidal anti-inflammatory  
86drugs; OA, osteoarthritic; PBS-T, PBS Tween; PF, PF 9184; PG, prostaglandin; PMN,  
87polymorphonuclear leukocytes; PPAR- $\gamma$ , peroxisome proliferator-activated receptor- $\gamma$ ; PsA,  
88psoriatic arthritis; PUFA, polyunsaturated fatty acid; RA, rheumatoid arthritis; RT, room  
89temperature; SDF-1, stromal cell-derived factor-1; sICAM-1, soluble intercellular adhesion  
90molecular-1; SpA, spondylarthritis; Th, T-helper; TIMP-1, metalloproteinase inhibitor-1; TNF- $\alpha$ ,  
91tumor necrosis factor- $\alpha$ ; TREM-1, triggering receptor expressed on myeloid cells-1; TRO,  
92Troglitazone.

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## 941. INTRODUCTION

95Inflammation is a complex defence mechanism characterised by leukocyte extravasation from the  
96vasculature to local tissue damage resulting from injurious and noxious agents/stimuli (Serhan,  
972014). Neutrophils dominate the initial influx of leukocytes followed by monocytes and  
98macrophages. The recruitment of inflammatory monocytes is correlated with a transient increase of  
99pro-inflammatory mediators including cytokines, chemokines, prostaglandins (PGs), and  
100leukotrienes (LTs) (D'Acquisto et al., 2010; Perretti et al., 2017). Indeed, inappropriate  
101monocyte/macrophage survival or overactivation perpetuate inflammatory pathways and strengthen  
102disease activity/duration (Zhou et al., 2009). Therefore, regulating the function of  
103monocytes/macrophages during inflammation is critical to promote resolution and healing. PGs and  
104LTs are potent bioactive lipid mediators involved not only in the onset of inflammation but also in  
105numerous homeostatic functions (Funk, 2001). Their biosynthesis is initialized by cyclooxygenases  
106(COXs) isoenzymes (COX-1 and COX-2) that convert arachidonic acid to PGH<sub>2</sub> (Hawkey, 1999)  
107and subsequently isomerized, by three different PGE<sub>2</sub> synthases (situated downstream of COXs in  
108the prostaglandin synthesis pathway), to PGE<sub>2</sub> (Koeberle et al., 2015). The cytosolic prostaglandin  
109E<sub>2</sub> synthase (cPGES) and the microsomal prostaglandin E<sub>2</sub> synthase (mPGES)-2 are constitutive  
110enzymes, whereas mPGES-1 is an inducible isoform (Samuelsson et al., 2007). PGE<sub>2</sub> production  
111can also be indirectly modulated by the alternative pathway of peroxisome proliferator-activated  
112receptor- $\gamma$  (PPAR- $\gamma$ ), a member of the nuclear receptor superfamily of ligand-dependent  
113transcription factors activated by 15-deoxy- $\Delta$ 12,14-prostaglandin J<sub>2</sub> (15d-PGJ<sub>2</sub>) (Cuzzocrea et al.,  
1142002). Anti-inflammatory effects related to PPAR- $\gamma$  activation have previously been demonstrated  
115in several studies (Ricote et al., 1998; Tsai et al., 2010) through its ability to increase the expression  
116of nuclear factor kappa-B inhibitor  $\alpha$  (I $\kappa$ B- $\alpha$ ), an endogenous inhibitor which interferes with the  
117activation of p65 nuclear factor kappa-B (NF- $\kappa$ B) (Korbecki et al., 2019), via its natural ligand 15d-  
118PGJ<sub>2</sub>. Although a large body of work has been carried out to elucidate the biological function of

119PPAR- $\gamma$  activation, this nuclear receptor's role remains poorly defined in the context of monocytes/  
120macrophages activation via the arachidonic pathway.

121Recently, several biochemical and pharmacological studies demonstrated a molecular interaction  
122between COXs and PGES isoenzymes, which results in preferential functional coupling activity.  
123Specifically, mPGES-2 was shown to utilize COX-1 to generate PGH<sub>2</sub> in contrast to mPGES-1  
124which uses COX-2 (Samuelsson et al., 2007). Moreover, studies with PGES deficient mice have  
125shown that induced PGE<sub>2</sub> synthesis is largely and preferentially dependent on mPGES-1 enzyme  
126(Inada et al., 2006). Consistently, the upregulation of mPGES-1 expression and the involvement of  
127COX-2/mPGES-1/PGE<sub>2</sub> cascade in terms of PGs production has been extensively reported in  
128pathological settings in which PGE<sub>2</sub> is implicated, such as fever, pain and inflammatory-based  
129diseases (Uematsu et al., 2002; Trebino et al., 2003). In line with these observations, it has been  
130reported that in several tissues and different cell types including fibroblasts, osteoblasts,  
131chondrocytes and osteoarthritic cartilage, mPGES-1 expression is enhanced by a variety of  
132inflammatory factors including lipopolysaccharides (LPS), interleukin-(IL-) $1\beta$ , tumor necrosis  
133factor- $\alpha$  (TNF- $\alpha$ ) and IL-17A (commonly known as IL-17), the latter of which was used in this  
134study (Stichtenoth et al., 2001; Li et al., 2005).

135In this context IL-17, has received much attention as a significant driver of autoimmune and  
136autoinflammatory conditions (Maione, 2016; Maione et al., 2020; Raucci et al., 2020) This cytokine  
137is mainly produced by T-helper (Th)-17 lymphocytes, but it also released by natural killer T (NKT)  
138cells, macrophages, neutrophils, monocytes, CD8<sup>+</sup> T cells,  $\gamma\delta$  T cells and innate lymphoid cells  
139(Onishi & Gaffen, 2010). We have previously demonstrated that IL-17 sustains chronic  
140inflammation and tissue remodelling rather than initiating it, through its ability to prime monocytes/  
141macrophages towards an inflammatory phenotype (Maione et al., 2009; Maione et al., 2011;  
142Maione, 2016). Considering that very little is known about the biological effects promoted by IL-17  
143both *in vitro* and *in vivo* in the context of mPGES-1/PPAR- $\gamma$  modulation, here we sought to  
144investigate and characterize the role of IL-17 on these pathways during an ongoing inflammatory  
145response.

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## 1472. METHODS

1482.1 **Materials:** PGE<sub>2</sub> Elisa kit, proteome profiler mouse cytokine array kit and recombinant mouse  
149IL-17 were purchased from R&D System (Milan, Italy). PF 9184 and Troglitazone were purchased  
150from Tocris (Milan, Italy), whereas FACS buffer and all conjugated antibodies from BioLegend  
151(London, UK). PGD<sub>2</sub> Elisa kit and the primary antibodies for western blot analysis were obtained  
152from Elabscience (Milan, Italy) whereas the HRP-conjugated IgG secondary antibodies from Dako  
153(Copenhagen, Denmark). 15d-PGJ<sub>2</sub> Elisa kit was purchased from Abcam (Cambridge, UK). Unless  
154otherwise stated, all the other reagents were from BioCell (Milan, Italy).

1552.2 **Murine cell isolation and culture:** Mouse macrophage cell line (J774A.1, ATCC® TIB-67™)  
156was cultured in Dulbecco's modified Eagle's medium (DMEM) supplemented with fetal bovine  
157serum (FBS) (ATCC® 30-2020™) to a final concentration of 10%. Cells were seeded in petri  
158culture dishes (100 × 20 mm, Falcon®) at a density of  $5 \times 10^5$  cells/dish and allowed growing for 24  
159h. The medium was then replaced, and cells were treated for 4 and 24 h with recombinant mouse  
160IL-17 (0.5-500 ng/ml). In another set of experiments, cells were treated with IL-17 (50 ng/ml) in  
161presence or absence of PF 9184 (50  $\mu$ M) or Troglitazone (50  $\mu$ M) according to previous *in vitro*  
162studies (Li et al., 2005). Following incubations for 4 and 24 h, cells were collected with a cell

163scraper and pellets were lysed at 4°C for 10 min with a buffer containing 1g/100 ml Triton X-100, 5  
164mM EDTA in PBS (pH 7.4) containing protease inhibitors. After centrifugation at 14000×g for 10  
165min at 4°C, the supernatant was collected and stored at -80°C for future western blot analysis.  
166Protein concentration was determined by the Bio-Rad protein assay (Bio-Rad, Milan, Italy).

167**2.3 Animals:** Experimental procedures were carried out in 8-12-week-old male CD1 mice in  
168compliance with the international and national law and policies and approved by Italian Ministry of  
169Health. Animal studies were performed in compliance with the ARRIVE guidelines and with the  
170recommendations made by the British Journal of Pharmacology (EU Directive 2010/63/EU for  
171animal experiments, ARRIVE guidelines, and the Basel declaration including the 3Rs concept;  
172Kilkenny et al., 2010; McGrath & Lilley, 2015). Mice, purchased from Charles River (Milan, Italy),  
173were housed with *ad libitum* access to food and water and maintained on a 12 h light/dark cycle.  
174Experimental study groups were randomized and blinded. All procedures were carried out to  
175minimize the number of animals used (n=7 per group) and their suffering.

176**2.4 Air Pouch:** Dorsal air pouches were induced by injection of 2.5 ml of air on day 0 and day 3 as  
177previously described (Maione et al., 2018a). On day 6, mice received the following treatments: i)  
178Control (CTRL) 0.25 ml of 0.5% carboxymethyl cellulose (CMC); ii) IL-17 (1 µg) in 0.25 ml of  
1790.5% CMC; iii) IL-17 (1 µg) in 0.25 ml of 0.5% CMC co-administrated with PF 9184 (1-9  
180µg/pouch); iv) IL-17 (1 µg) in 0.25 ml of 0.5% CMC co-administrated with Troglitazone (1-9  
181µg/pouch) accordingly to previous studies (Li et al., 2005; Maione et al., 2009). Mice were  
182sacrificed after 4 and 24 h from the injection and air pouches washed thoroughly with 2 ml of PBS  
183containing 50 U/ml heparin and 3 mM EDTA. Lavage fluids were centrifuged at 220×g for 10 min  
184at 4°C to separate the exudates from the recruited cells. Inflammatory exudates were collected and  
185measured to evaluate the level of inflammatory cyto-chemokines, whereas cell pellet subjected to  
186western blot and FACS analysis as described below. Cell number was determined by TC20  
187automated cell counter (Bio-Rad, Milan, Italy) using Bio-Rad's automated cell counter uses  
188disposable slides, TC20 trypan blue dye (0.4% trypan blue dye w/v in 0.81% sodium chloride and  
1890.06% potassium phosphate dibasic solution) and a CCD camera to count cells based on the  
190analyses of capture images (Maione et al., 2018a; Bellavita et al., 2020).

191**2.5 Myeloperoxidase assay:** Leukocytes myeloperoxidase (MPO) activity was assessed by  
192measuring the H<sub>2</sub>O<sub>2</sub>-dependent oxidation of 3,3',5,5'-tetramethylbenzidine as previously reported  
193(Maione et al., 2009). Cellular lysate, from air pouch experiments, were homogenized in a solution  
194composed of hexadecyltrimethylammonium bromide (0.5% w/v) in 50 mM sodium phosphate  
195buffer at pH 5.4. After homogenization, samples were centrifuged at 1008×g for 10 min and the  
196supernatant used for the assay. Aliquots of 20 µl were incubated with 160 µl of 3,3',5,5' -  
197tetramethylbenzidine and 20 µl of H<sub>2</sub>O<sub>2</sub> (in 80 mM phosphate buffer, pH 5.4) in 96-well plates.  
198Plates were incubated for 5 min at room temperature (RT), and OD was read at 620 nm using a  
199plate-reader (Biorad, Italy). Assay was performed in duplicates and normalized for protein content.

200**2.6 Elisa and ElisaSpot assay:** Enzyme-linked immunosorbent assays for PGE<sub>2</sub>, PGD<sub>2</sub> and 15d-  
201PGJ<sub>2</sub> were carried out on pouch inflammatory exudates. Briefly, 100 µl of samples, diluted  
202standards, quality controls and dilution buffer (blank) were added to a pre-coated plate with  
203monoclonal anti-PGE<sub>2</sub>, PGD<sub>2</sub> or 15d-PGJ<sub>2</sub> for 2 h. After washing, 100 µl of biotin labelled antibody  
204was added for 1 h. The plate was washed and 100 µl of streptavidin-HRP conjugate was added and  
205the plate was incubated for a further 30 min period in the dark. The addition of 100 µl of the  
206substrate and stop solution represented the last steps before the reading of absorbance (measured at  
207450 and 405 nm for PGE<sub>2</sub>, PGD<sub>2</sub> and 15d-PGJ<sub>2</sub> respectively) on a microplate reader. Antigen levels

208in the samples were determined using a standard curve and expressed as pg/pouch (Raucci et al.,  
2092019). For cyto-chemokines protein array equal volumes (1.5 ml) of pouch inflammatory fluids in  
210all described experimental conditions were incubated with the pre-coated proteome profiler array  
211membranes according to the manufacturer's instructions. Dot plots were detected by using the  
212enhanced chemiluminescence detection kit and Image Quant 400 GE Healthcare software (GE  
213Healthcare, Italy) and successively quantified using GS 800 imaging densitometer software (Biorad,  
214Italy) as previously described (Cristiano et al., 2019).

215**2.7 Flow Cytometry:** Cells collected from the pouch cavities, at 4 and 24 h, were first washed with  
216PBS and then re-suspended in FACS buffer (PBS containing 1% FCS and 0.02% NaN<sub>2</sub>) containing  
217CD16/CD32 FcγIIIR blocking antibody (clone 93; eBioscience, London, UK) for 30 min at 4°C.  
218Thereafter, cells were labelled for 30 min at 4°C with the following conjugated antibodies (all from  
219BioLegend, London, UK): CD45 (1:100; clone 30-F11), LY6C (1:100; clone HK1.4), LY6G  
220(1:100; clone 1A8), CD115 (1:100; clone AFS98), CD11b (1:100; clone M1/70), F4/80 (1:100;  
221clone BM8), CD206 (1:100; clone C068C2), prior to analysis by FACS calibre using CellQuest  
222software (Becton Dickinson, Franklin Lakes, NJ). Neutrophils, macrophage and  
223resident/inflammatory monocytes were defined according to the flow cytometry procedure  
224previously described (Maione et al., 2018a; Kapellos et al., 2019; Podaru et al., 2019). At least  
2251×10<sup>4</sup> cells were analysed per sample, and determination of positive and negative populations was  
226performed based on the staining obtained with related IgG isotypes. Flow cytometry was performed  
227on BriCyte E6 flow cytometer (Mindray Bio-Medical Electronics, Nanshan, China) using MRFlow  
228and FlowJo software operation (Raucci et al., 2019b)

229**2.8 Western blot analysis:** Whole cellular pellets homogenates (50 µg of protein) from *in vitro* and  
230*in vivo* experiments were subjected to SDS-PAGE (10 and 12% gel) using standard protocols as  
231previously described (Maione et al., 2018b). The proteins were transferred to nitrocellulose  
232membrane (0.2 µm nitrocellulose membrane, Trans-Blot® Turbo™, Transfer Pack, Bio-Rad  
233Laboratories, Hercules, CA, USA) in transfer buffer (25 mM Tris-HCl pH 7.4 containing 192 mM  
234glycine and 20% v/v methanol) at 400 mA for 2 h. The membranes were saturated by incubation for  
2352 h with non-fat dry milk (5% wt/v) in PBS supplemented with 0.1% (v/v) Tween 20 (PBS-T) for 2  
236h at RT and then incubated with 1:1000 dilution of primary antibodies over-night at 4°C such as:  
237mouse monoclonal anti-actin (E-AB-20094), mouse monoclonal anti-COX-2 (E-AB-27666), mouse  
238monoclonal anti-IκB-α (4814), rabbit polyclonal anti-IL-17 Receptor (E-AB-63080), mouse  
239monoclonal anti-NFκB (MAB3026), rabbit polyclonal anti-mPGDS-1 (TA301420), rabbit  
240polyclonal anti-mPGES-1 (E-AB-32563), rabbit polyclonal anti-PPAR-γ (NBP2-22106), and then  
241washed 3 times with PBS-T. In all cases, blots were then incubated with a 1:3000 dilution of related  
242horseradish peroxidase-conjugated secondary antibody for 2 h at RT and finally washed 3 times  
243with PBS-T. Protein bands were detected by using the enhanced chemiluminescence method  
244(Clarify™ Western ECL Substrate, Bio-Rad Laboratories, Hercules, CA, USA) and Image Quant  
245400 GE Healthcare software (GE Healthcare, Italy). Finally, protein bands were quantified using the  
246GS 800 imaging densitometer software (Biorad, Italy) and normalized with respective actin.

247**2.9 Statistical analysis:** In this study, statistical analysis complies with the recommendations of the  
248British Journal of Pharmacology on experimental design and analysis in pharmacology (Curtis et  
249al., 2018) and data sharing and presentation in preclinical pharmacology (Alexander et al., 2018;  
250George et al., 2017). All data are presented as means ± S.D. and were analysed using Student's t-  
251test (two groups) or one-way ANOVA followed by Bonferroni's for multiple comparison test (more  
252than two groups). GraphPad Prism 8.0 software (San Diego, CA, USA) was used for analysis.

253Differences between means were considered statistically significant when  $P \leq 0.05$  was achieved.  
254Sample size was chosen to ensure alpha 0.05 and power 0.8. Animal weight was used for  
255randomization and group allocation. No animals were excluded from the analysis.

256

## 2573. RESULTS

258**3.1 PF 9184 and Troglitazone, in a time- and dose-dependent manner, revert leukocytes**  
259**accumulation and activation at site of inflammation:** We knew from our previous studies  
260(Maione et al., 2009; Maione et al., 2018a) that single administration of IL-17 (1  $\mu\text{g}/\text{pouch}$ ) into a  
2616-day old air pouch causes a transient infiltration of leukocytes by 4 h, which peaks at 24 h and then  
262declines by 48 h. To test the potential dose-responsive effect of an mPGES-1 antagonist and a  
263PPAR- $\gamma$  agonist at this local site of inflammation, we administered PF 9184 (PF, 1-9  $\mu\text{g}/\text{pouch}$ ) and  
264Troglitazone (TRO, 1-9  $\mu\text{g}/\text{pouch}$ ) concomitantly with IL-17. All mice were sacrificed at 4, 24 and  
26548 h time point. Consistent with our previous findings, 4 and 24 h after IL-17 administration mice  
266showed significant differences in the number of inflammatory leukocytes compared to vehicle-  
267treated (Fig. 1A-B). No significant differences were observed at 48 h (Fig. 1C). Interestingly, mice  
268receiving PF at a dose of 9  $\mu\text{g}/\text{pouch}$  showed a marked decrease (~40%) in inflammatory infiltrates,  
269compared to IL-17-treated mice, at both 4 and 24 h (Fig. 1A-B). A similar anti-inflammatory profile  
270was observed after TRO administration (9  $\mu\text{g}/\text{pouch}$ ), with a marked reduction in infiltrating  
271leukocytes at 4 and 24 h (Fig. 1A-B). No biological effects were observed at 48 h (Fig. 1C). Based  
272on these results, we selected the most active PF and TRO (9  $\mu\text{g}/\text{pouch}$ ) dose for all subsequent  
273analyses. The level of myeloperoxidase (MPO, Fig. 1D-F), a peroxidase enzyme most abundantly  
274expressed in polymorphonuclear leukocytes, and different prostaglandins such as PGE<sub>2</sub>, PGD<sub>2</sub> and  
27515d-PGJ<sub>2</sub>, were measured in pouch fluid. Administration with IL-17 correlated with increased  
276levels of MPO (Fig. 1D-E) and PGE<sub>2</sub> (Fig. 1G-H) at both 4 and 24 h. Conversely a reduction in  
277PGD<sub>2</sub> (Fig. 1I-J) and 15d-PGJ<sub>2</sub> (Fig. 1K-L) at both 4 and 24 h was observed. When co-injected in  
278the presence of PF and TRO the opposite profile was observed with a marked reduction in MPO  
279(Fig. 1E) activity and PGE<sub>2</sub> (Fig. 1G,H), and reversal in PGD<sub>2</sub> (Fig. 1I,J) and 15d-PGJ<sub>2</sub> levels (Fig.  
2801K,L). Collectively these data indicate time- and dose-dependent protective effects of a selective  
281mPGES-1 antagonist and a PPAR- $\gamma$  agonist on inflammatory cell recruitment and activation in  
282response to IL-17 induced inflammation.

283**3.2 mPGES-1 and PPAR- $\gamma$  enzymatic pathway modulation on the onset of ongoing**  
284**inflammation:** Previous studies have shown in various tissues and cell types, both COX-2 and  
285mPGES-1 expression is enhanced in response to a range of inflammatory mediators including LPS,  
286IL-1 $\beta$ , TNF- $\alpha$ , and IL-17 (Raucci et al., 2019a). We, therefore, carried out western blot analysis on  
287recruited cells and found that COX-2 (Fig. 2B) and mPGES-1 (Fig. 2H) were upregulated following  
288IL-17-induced inflammation and were both significantly reduced at 4 and, in particular, 24 h after  
289PF and TRO treatment. Moreover, we found that IL-17 receptor (IL-17R) expression (increased in  
290IL-17-treated animals compared to CTRL group at 24 h) were not influenced by either PF or TRO  
291administration (Fig. 2A). We also observed a substantial increase in NF- $\kappa$ B expression in mice  
292treated with IL-17 (Fig. 2C). The natural inhibitor of NF- $\kappa$ B (I $\kappa$ B- $\alpha$ ) was also measured and the  
293converse was observed in mice treated with PF or TRO at 4, and 24 h (Fig. 2F). Notably, mPGES-1  
294and PPAR- $\gamma$  expression were weakly present in IL-17 group, whereas it significantly increased in  
295PF (only at 24 h) and TRO-treated mice (at both 4 and 24 h), confirming the hypothesis that  
296activation of an alternative molecular pathway following thiazolidinedione administration (Fig.  
2972D,G respectively). Densitometric values (at the bottom of the Fig. 2) are expressed as OD Ratio

298with actin (Fig. 2E) for both 4 and 24 h. Uncropped and original western blots are presented in  
299Supplementary Figure 1-8.

**3003.3 PF 9184 and Troglitazone selectively modulate the recruitment of inflammatory**  
301**monocytes:** We went on to further characterize the phenotype of recruited cells at different time  
302points. Flow cytometry was employed to determine neutrophil and monocyte/macrophage subsets.  
303Specifically, to identify potential differences in leukocytes sub-populations total cells were gated,  
304(Fig. 3A, 4A, and 4N, *gate* R1) followed by single cells (Fig. 3B, 4B, and 4O, *gate* R2). CD45 (pan  
305leukocyte/immune cell marker; Fig. 3C) and CD11b (myeloid marker; Fig. 4C, P) were selected  
306(<sup>+</sup>, *gate* R3). Neutrophils were identified as CD45<sup>+</sup>/Ly6G<sup>Hi</sup>/Ly6C<sup>Hi</sup> as shown both at 4 (Fig. 3D-  
307G) and 24 h (Fig. 3I-L). Monocytes and macrophages were further delineated based upon a range of  
308markers. CD11b<sup>+</sup> cells were plotted for Ly6C and CD115 at 4 (Fig. 4D-G) and 24 h (Fig. 4I-L) to  
309distinguish CD11b<sup>+</sup>/CD115<sup>+</sup>/Ly6C<sup>lo</sup> patrolling monocytes (*gate* R4) from  
310CD11b<sup>+</sup>/CD115<sup>+</sup>/Ly6C<sup>hi</sup> inflammatory monocytes (*gate* R5) (Raucci et al., 2019a; Kapellos et al.,  
3112019; de Brito et al., 2019), and for CD206 and F4/80 (Fig. 4Q-T and 4V-Y at 4 and 24 h  
312respectively) to identify CD11b<sup>+</sup>/F480<sup>+</sup>/CD206<sup>+</sup> reparative macrophages (Podaru et al., 2019).  
313Our results show that in IL-17-injected mice, at 4 h, the majority of cells recovered were neutrophils  
314(Fig. 3H) which were largely replaced by inflammatory monocytes (Fig. 4M) and lower proportion  
315of reparative macrophages (Fig. 4Z) at 24 h. Interestingly, both PF and TRO treatments  
316significantly inhibited this selective and time-dependent recruitment of neutrophils and  
317inflammatory monocytes (Fig. 3H,M and 4H,M). Moreover, TRO administration at 24 h maintained  
318similar levels of reparative macrophages as the control group (Fig. 4Z). In all experimental  
319conditions, no significant differences were found in resident monocyte recruitment (*gate* R4).  
320Reported values were strengthened by a low percentage of positive cells found in the staining for  
321the isotype control antibodies (Supporting Information 1). These results suggest that PF and TRO  
322treatment significantly disrupts the recruitment of inflammatory cells in the first phase of the  
323response and that TRO selectively promotes resolution in the subsequent reparative phase.

**3243.4 Co-administration of PF 9184 and Troglitazone with IL-17 into the air pouch decreases**  
325**the release of cyto-chemokines in the inflammatory fluids:** To gain some insights into other  
326possible differences in the inflammatory response caused by co-administration of an mPGES-1  
327antagonist or a PPAR- $\gamma$  agonist with a IL-17 inflammatory stimulus, we used an unbiased approach  
328(pre-made protein array) based on profiling cytokines and chemokines present in the inflammatory  
329fluids. As shown in Figure 5, the pouch fluid obtained from IL-17 administered mice showed a  
330large increase, at 4 (Fig. 5A-D) and, in particular, 24 h (Fig. 5F-I), of cyto-chemokines profile  
331compared to vehicle (CTRL group). When comparing pouch fluids from PF and TRO treated  
332groups with IL-17 (alone), we observed a selective decrease in a range of mediators (Fig. 5A-D and  
333Fig. 5F-I at 4 and 24 h respectively). Densitometric analysis revealed that the PF treated group had  
334a specific modulation, at 4 h (Fig. 5E), in the following factors: B lymphocyte chemoattractant  
335(BLC), soluble intercellular adhesion molecular-1 (sICAM-1), IL-16, interferon  $\gamma$ -induced protein-  
33610 (IP-10), keratinocyte chemoattractant (KC), macrophage colony-stimulating factor (MCSF),  
337junctional epithelium (JE), monocytes chemoattractant protein-5 (MCP-5), monokine induced by  
338interferon  $\gamma$  (MIG), macrophage inflammatory proteins (MIPs), metalloproteinase inhibitor-1  
339(TIMP-1) and triggering receptor expressed on myeloid cells-1 (TREM-1) compared to IL-17 group  
340(Fig. 5E). This profile was confirmed at the 24 h time-point (Fig. 5J) in addition to complement  
341component 5a (C5a), interferon- $\gamma$  (INF- $\gamma$ ), IL-1 $\alpha$ , IL-1 $\beta$ , IL-7, interferon-inducible T-cell  $\alpha$   
342chemoattractant (ITAC) and stromal cell-derived factor-1 (SDF-1) (Fig. 5J). A similar inhibitory  
343profile was found in TRO injected mice, but surprisingly we observed a more prominent

344modulation of BLC, C5a, IL-17, IP-10, SDF-1 and TIMP-1 at 4 h (Fig. 5E) and of BLC, sICAM-1,  
345IL-16, IL-17, JE, MCP-5 and MIG at 24 h time-point (Fig. 5J).  $\Delta$  increase/decrease of cito-  
346chemokines modulation in all experimental conditions are reported in Supporting Information 2.

347**3.5 Protective effect of PF 9184 and Troglitazone on murine isolated, and IL-17-stimulated,**  
348**macrophage:** To further confirm the validity of our *in vivo* findings, we carried out *in vitro* studies  
349using a murine macrophage cell line J774. Stimulation of macrophages with increasing  
350concentrations of IL-17 (0.5-500 ng/ml) induced a significant increase in IL-17R expression at both  
3514 and 24 h (Fig. 6A). Notably, pre-treatment of IL-17 (50 ng/ml)-stimulated macrophages with PF  
352or TRO (50  $\mu$ M) reverted the expression of COX-2 and mPGES-1 at 4 and 24 h (Fig. 6B).  
353Moreover, PPAR- $\gamma$  expression was weakly present in IL-17 group, whereas it significantly  
354increased in PF (only at 24 h) and TRO-treated mice (at both 4 and 24 h), further strengthening our  
355hypothesis of activation of an alternative pathway following thiazolidinedione administration (Fig.  
3566B). Densitometric values (at the bottom of the Fig. 6A, B) are expressed as OD Ratio with actin  
357for both 4 and 24 h. Original western blots are presented in Supplementary Fig 9-14.

358

#### 3594. DISCUSSION AND CONCLUSIONS

360The integration of inflammatory signals is paramount to controlling the intensity and duration of the  
361immune response. Eicosanoids, particularly PGE<sub>2</sub>, are critical molecules in the initiation of  
362inflammation and transition from innate to acquired immune responses. mPGES-1, an integral  
363membrane enzyme functionally coupled to COX-2, whose regulated expression controls PGE<sub>2</sub>  
364levels at sites of inflammation (Park et al., 2006; Kalinski, 2012), has pleiotropic effects on many  
365cells of the immune system, influencing both the innate and acquired immune responses  
366(Sreeramkumar et al., 2012; Sreeramkumar et al., 2016). PGE<sub>2</sub> can promote the influx and  
367activation of neutrophils, macrophages and mast cells (Westman et al., 2004; Kojima et al., 2008),  
368but can also suppress NKT cytolytic and granulocyte functions (Jakobsson et al., 1999). *In vitro*  
369evidence from several groups have shown that the induction of mPGES-1 is increased in response  
370to the pro-inflammatory cytokines IL-1 $\beta$ , TNF, or LPS and that its expression, in certain immune-  
371mediated inflammatory diseases (IMIDs), maybe upregulated by a wide range of stimuli  
372(Korotkova et al., 2008). Li and colleagues (Li et al., 2005) have shown that expression and  
373regulation of microsomal prostaglandin E synthase-1 and peroxisome proliferator-activated  
374receptor-gamma in human osteoarthritic cartilage and chondrocytes are regulated by pro-  
375inflammatory stimuli such as IL-1 $\alpha$  and TNF- $\alpha$  and that the concomitant presence of IL-17  
376displayed a synergistic effect reversed by troglitazone or exogenous PGE<sub>2</sub>. These data suggest that  
377mPGES-1 may prove to be an interesting therapeutic target for controlling PGE<sub>2</sub> synthesis in certain  
378IMIDs such as rheumatoid arthritis (RA), psoriatic arthritis (PsA) and spondylarthritis (SpA)  
379(Jouzeau et al., 2008; Navarini et al., 2020).

380Self-regulation of the COXs pathway does not consist solely of positive feedback but also involves  
381mechanisms that inhibit the inflammatory response. At the centre of the COXs auto-inhibitory  
382pathway, are PPARs. Activation of inflammatory responses causes an increase in the expression of  
383PPAR- $\alpha$  and a decrease in PPAR- $\gamma$  (Kapoor et al., 2007; García-Alonso et al., 2013). PPARs are  
384nuclear receptors activated by oxidised and nitrated fatty acid derivatives and cyclopentenone  
385prostaglandins (PGA<sub>2</sub> and 15d-PGJ<sub>2</sub>) during the inflammatory response. Other activators include  
386nonsteroidal anti-inflammatory drugs (NSAID), fatty acids, especially polyunsaturated fatty acid  
387(PUFA) (arachidonic acid, ALA, EPA, and DHA) and thiazolidinedione derivates (Yousefnia et al.,

3882018). The primary function of PPARs during the inflammatory reaction is to promote the  
389inactivation of NF- $\kappa$ B (by direct binding of p65 NF- $\kappa$ B) or the proteolytic degradation of p65 NF-  
390 $\kappa$ B. PPARs also cause an increase in the expression of I $\kappa$ B- $\alpha$ , SIRT1, and PTEN, which interfere  
391with the activation and function of NF- $\kappa$ B in inflammatory reactions (Thommesen et al., 1998).  
392From a pathological point of view, it should also be taken into account that IL-1 $\alpha$ , TNF- $\alpha$ , IL-17  
393and PGE<sub>2</sub>, which are involved in the pathogenesis and progression of certain IMIDS, also  
394downregulate PPAR- $\gamma$  expression in a dose- and time-dependent manner (Afif et al., 2007;  
395Korbecki et al., 2014). Several lines of evidences suggest that PPAR- $\gamma$  activation may have  
396therapeutic benefits in RA, PsA and possibly other chronic articular diseases (Murakami, 2011;  
397Korbecki et al., 2019).

398IL-17 is an archetype molecule for the entire family of IL-17 cytokines. Currently believed to be  
399produced by a specific subset of CD4<sup>+</sup> T cells, named Th17 cells, but also by many innate cell  
400components (D'Acquisto et al., 2010), this cytokine is functionally located at the interface of innate  
401and adaptive immunity (Maione, 2016). Specifically, IL-17 ability to induce the release of a range  
402of cyto-chemokines and growth factors, has led to its emergence as an essential co-ordinator of  
403local inflammatory reactions due to its ability to modulate neutrophil and monocyte accumulation in  
404inflamed tissues. Furthermore, there is growing evidence that suggests targeting IL-17 signalling  
405may prove useful in a variety of inflammatory-based diseases including RA, OA, asthma, Crohn's  
406disease (CD), psoriatic like-disease and PsA. Here, using a well-established preclinical model of  
407ongoing inflammation, the dorsal air pouch (Maione et al., 2009), we tested the hypothesis that pre-  
408treatment with an mPGES-1 antagonist or PPAR- $\gamma$  agonist could retard the process of IL-17-  
409induced inflammation. We confirmed the pro-inflammatory action (at 4 and 24 h) of IL-17, but also  
410found a novel protective role for PF and TRO, as exemplified by a reduction in both PMN  
411(polymorphonuclear leukocytes) recruitment (at both 4 and in particular 24 h), MPO activity and  
412modulation of crucial lipid mediators strictly related to mPGES-1 and PPAR- $\gamma$  enzymatic activity  
413(PGE<sub>2</sub>, PGD<sub>2</sub>, and PGJ<sub>2</sub>). From a mechanistic basis, we demonstrated that IL-17 increased in a time-  
414dependent manner, the levels of its receptor (IL-17R), COX-2, mPGES-1, and NF- $\kappa$ B and  
415decreased mPGDS-1, PPAR- $\gamma$  and I $\kappa$ B- $\alpha$  expression, in pouch-recruited leukocytes. Moreover, co-  
416administration with PF and TRO was shown to significantly revert PMN recruitment, MPO activity,  
417and reduce the level of lipid mediators and, most importantly, to modulate mPGES-1, mPGDS-1,  
418PPAR- $\gamma$  and NF- $\kappa$ B/I $\kappa$ B- $\alpha$  expression, leaving unchanged IL-17R expression. Another exciting  
419aspect of our study is that we provide the first reported evidence of an indirect, co-ordinated  
420functional regulation in both neutrophils and monocytes by mPGES-1 and PPAR- $\gamma$ . This is most  
421likely related to the presence of complex mechanisms regulating COX-2/mPGES-1 *in vivo* which  
422go onto to impact PPAR- $\gamma$  activity (Maione et al., 2020).

423This hypothesis is consistent with previous studies showing that pre-adipocytes stably transfected  
424with either COX-1 or COX-2 had lower PPAR- $\gamma$  expression (Chu et al., 2009) and that mice  
425genetically deficient for mPGES-1 had basal elevations in PPAR- $\gamma$  expression and transcriptional  
426activity (Kapoor et al., 2007). The mechanisms by which the COX-2/mPGES-1/PGE<sub>2</sub> axis and the  
427nuclear receptor PPAR- $\gamma$  interact during the inflammatory process are not entirely delineated but  
428may linked to PGE<sub>2</sub> ability (mainly produced by neutrophils at the early stage of inflammation) to  
429decrease the amount of well-known lipid mediators (such as PGD<sub>2</sub> and PGJ<sub>2</sub>) which are implicated  
430in the induction of PPAR- $\gamma$  (Maione et al., 2020). This biological event is interconnected with a  
431cellular shift from neutrophils to monocytes as exemplified by the release of specific neutrophil  
432(KC, C5a) and monocytes/macrophages (INF- $\gamma$ , IL-16, IL-17, IP-10, JE, MIPs, TREM-1, ICAM,  
433IL-1 $\alpha$ / $\beta$ , MCPs, MIG, TIMP-1) cyto-chemokines. It is well-established in the literature that i)

434 PPAR- $\gamma$  acts as a negative regulator of macrophage activation (Ricote et al., 1998) due to its ability  
435 to control polarization of monocyte differentiation between pro-inflammatory (M1) and alternative  
436 anti-inflammatory (M2) macrophage phenotypes (Tontonoz & Spiegelman, 2008) and to reduce  
437 neutrophil migration to sites of injury (Napimoga et al., 2008); ii) neutrophils are a principle  
438 cellular source of mPGES-1, with activated M1, rather than alternatively activated M2  
439 macrophages, a secondary source (Posadas et al., 2000; Mosca et al., 2007). It is clear from this  
440 current study that the release of PGE<sub>2</sub> and PGD<sub>2</sub>/PGJ<sub>2</sub> related to COX-2/mPGES-1 and  
441 PPAR- $\gamma$ /mPGDS-1 expression follows the temporal shift from neutrophils to monocytes that are  
442 implicated in the potential resolution of inflammatory response. This was confirmed by our  
443 experiments performed in the presence of either PF or TRO's where both compounds were shown to  
444 revert the inflammatory response (Fig. 7). To further support this association between IL-17 and  
445 mPGEs-1/PPAR $\gamma$  expression and modulation, and in light of the detrimental role of macrophages  
446 and macrophage-derived cytokines in RA and SpA synovium (Kabala et al., 2020), we performed *in*  
447 *vitro* experiments where we used J774 cell line. Interestingly, we confirmed our *in vivo* results  
448 demonstrating a direct involvement of mPGES-1/PPAR- $\gamma$  axis in IL-17 stimulated-macrophages.

449 In conclusion, our results reveal a novel interaction between IL-17 and mPGES-1/PPAR- $\gamma$   
450 generated by macrophages/inflammatory monocytes during inflammation. Therefore, we believe  
451 that the IL-17/mPGES-1/PPAR- $\gamma$  "axis" could represent a potential therapeutic target for  
452 inflammatory-based and immune-mediated diseases.

453

#### 454 Author contributions

455 FR, AS, GMC, MGR, AAM, MP and MGF performed the experiments. EP, VV, CI, FC, RS, NM,  
456 MA, AJI and FM performed data analysis and wrote the manuscript. FR, AJI and FM revised the  
457 final version of the manuscript for intellectual contents. All authors gave final approval to the  
458 publication.

459

#### 460 Declaration of transparency and scientific rigour

461 This Declaration acknowledges that this paper adheres to the principles for transparent reporting  
462 and scientific rigour of preclinical research as stated in the BJP guidelines for Design & Analysis,  
463 Immunoblotting, and Animal Experimentation, and as recommended by funding agencies,  
464 publishers and other organisations engaged with supporting research.

465

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662

## 663FIGURE LEGENDS

664**Fig. 1 PF and TRO, in a time- and dose-dependent manner, revert leukocytes accumulation** 665**and activation at site of inflammation:** Mice were treated with IL-17 vehicle (CTRL), IL-17 (1 666 $\mu\text{g/pouch}$ ) alone (IL-17) or co-administrated with PF 9184 (PF, 1-9  $\mu\text{g/pouch}$ ) and Troglitazone 667(TRO, 1-9  $\mu\text{g/pouch}$ ) and thereafter total cell number from pouches inflammatory exudates was 668evaluated at 4 (A), 24 (B) and 48 (C) h. At the same time-point (D-F), supernatants from cell pellet 669lysate were tested for myeloperoxidase activity. Subsequently, inflammatory fluids from all 670experimental conditions were assayed by Elisa for PGE<sub>2</sub> (G-H), PGD<sub>2</sub> (I-J), and PGJ<sub>2</sub> (K-L) levels 671at both 4 and 24 h. Data were expressed as millions of cells for pouch (A-C), units/mg of protein

672(D-F) or pg/pouch (G-L) and presented as means  $\pm$  S.D. of n=7 mice per group. Statistical analysis  
673was conducted by one-way ANOVA followed by Bonferroni's for multiple comparisons.  $^{\#}P\leq 0.05$ ,  
674 $^{\#\#}P\leq 0.01$ ,  $^{\#\#\#}P\leq 0.005$  vs CTRL group;  $^*P\leq 0.05$ ,  $^{**}P\leq 0.01$  vs IL-17 group.

675

676**Fig. 2 mPGES-1 and PPAR- $\gamma$  enzymatic pathway modulation on the onset of ongoing**  
677**inflammation:** Whole cellular pellets homogenates from air pouch experiments in all experimental  
678conditions were assayed by western blot for IL-17 Receptor (IL-17R) (A), COX-2 (B), NF-Kb (C),  
679PPAR- $\gamma$  (D), Ikb- $\alpha$  (F), mPGDS-1(G), mPGES-1 (H) expression at 4 and 24 h. Western blot  
680images are representative of three separate experiments with similar results. Cumulative  
681densitometric values (at the bottom of the Figure) are expressed as OD Ratio with actin (E) for both  
6824 and 24 h. Values are presented as means  $\pm$  S.D. of three experiments with n=7 mice per group.  
683Statistical analysis was conducted by one-way ANOVA followed by Bonferroni's for multiple  
684comparisons.  $^{\#}P\leq 0.05$ ,  $^{\#\#}P\leq 0.01$ ,  $^{\#\#\#}P\leq 0.005$  vs CTRL group;  $^*P\leq 0.05$ ,  $^{**}P\leq 0.01$ ,  $^{***}P\leq 0.005$  vs IL-  
68517 group.

686

687**Fig. 3 Flow cytometry strategy applied to identify the modulation of neutrophils after IL-17**  
688**stimulation:** Mice were treated with IL-17 vehicle (CTRL), IL-17 (1  $\mu$ g/pouch) alone (IL-17) or in  
689co-administrated with PF 9184 (PF, 9  $\mu$ g/pouch) and Troglitazone (TRO, 9  $\mu$ g/pouch). Cells  
690collected from the pouch cavities were washed, gated in their totality (A, *gate* R1) and singlet (B,  
691*gate* R2) before the identification of CD45 positive (C, CD45<sup>+</sup>) population (*gate* R3). CD45<sup>+</sup>  
692cells (C) were then plotted for Ly6C and Ly6G expression at 4 (D-G) and 24 h (I-L) to identify  
693CD45<sup>+</sup>/Ly6C<sup>+</sup>/Ly6G<sup>+</sup> as neutrophils. Histograms values (expressed as million for pouch)  
694indicate the total positive cells, in the different experimental conditions, of  
695CD45<sup>+</sup>/Ly6C<sup>+</sup>/Ly6G<sup>+</sup> (H,M) at 4 and 24 h. FACS pictures are representative of independent  
696experiments with similar results. Values are presented as means  $\pm$  S.D. of three separate  
697experiments with n=7 mice per group. Statistical analysis was conducted by one-way ANOVA  
698followed by Bonferroni's for multiple comparisons.  $^{\#\#}P\leq 0.01$ ,  $^{\#\#\#}P\leq 0.005$  vs CTRL group;  $^*P\leq 0.05$ ,  
699 $^{**}P\leq 0.01$  vs IL-17 group.

700

701**Fig. 4 PF and TRO selectively modulate the recruitment of inflammatory monocytes and**  
702**reparative macrophages:** Mice were treated with IL-17 vehicle (CTRL), IL-17 (1  $\mu$ g/pouch) alone  
703(IL-17) or in co-administrated with PF 9184 (PF, 9  $\mu$ g/pouch) and Troglitazone (TRO, 9  $\mu$ g/pouch).  
704Cells collected from the pouch cavities were washed, gated in their totality (A,N, *gate* R1) and  
705singlet (B,O, *gate* R2) before the identification of CD11b positive (C,P CD11b<sup>+</sup>) population (*gate*  
706R3). CD11b<sup>+</sup> (C) cells were then plotted for Ly6C and CD115 expression at 4 (D-G) and 24 h (I-  
707L) to distinguish CD11b<sup>+</sup>/CD115<sup>+</sup>/Ly6C<sup>low</sup> patrolling monocytes (*gate* R4) from  
708CD11b<sup>+</sup>/CD115<sup>+</sup>/Ly6C<sup>hi</sup> inflammatory monocytes (*gate* R5). Therefore, CD11b<sup>+</sup> cells were then  
709plotted for CD206 and F480 expression at both 4 (Q-T) and 24 h (V-Y) to identify  
710CD206<sup>+</sup>/F480<sup>+</sup> population. Histograms values (expressed as million for pouch) indicate the total  
711positive cells, in the different experimental conditions, of CD11b<sup>+</sup>/CD115<sup>+</sup>/Ly6C<sup>hi</sup> (H,M) and  
712CD11b<sup>+</sup>/CD206<sup>+</sup>/F480<sup>+</sup> (U,Z) at 4 and 24 h. FACS pictures are representative of independent  
713experiments with similar results. Values are presented as means  $\pm$  S.D. of three separate  
714experiments with n=7 mice per group. Statistical analysis was conducted by one-way ANOVA

715 followed by Bonferroni's for multiple comparisons. <sup>#</sup>P≤0.05, <sup>###</sup>P≤0.005 vs CTRL group; \*P≤0.05,  
716 \*\*P≤0.01 vs IL-17 group.

717

718 **Fig. 5 Co-administration of PF and TRO with IL-17 into the air pouch decreases the release of**  
719 **cyto-chemokines in the inflammatory fluid:** Inflammatory supernatants obtained from the pouch  
720 cavities were assayed using a Proteome Profiler cytokine array at both 4 and 24 h for CTRL  
721 (respectively **A, F**), IL-17 (1 µg/pouch, respectively **B, G**), IL-17 + PF (9 µg/pouch, respectively **C,**  
722 **H**) and IL-17 + TRO (9 µg, respectively **D, I**). Densitometric analysis is presented as heatmap at 4  
723 (**E**) and 24 h (**J**). Data (expressed as INT/mm<sup>2</sup>) are presented as means ± S.D. of positive spots of  
724 three independent experiments with n=7 mice per group.

725

726 **Fig. 6 Protective effect of PF and TRO on murine isolated, and IL-17-stimulated,**  
727 **macrophage:** Whole cellular pellets homogenates from murine macrophages stimulated with  
728 increasing concentration of IL-17 (0.5-500 ng/ml) (**A**) or with IL-17 at 50 ng/ml in presence or  
729 absence of PF 9184 (PF, 50 µM) and Troglitazone (TRO, 50 µM) at 4 and 24 h (**B**) were analysed,  
730 by western blot, for IL-17R, COX-2, PPAR-γ, mPGES-1 and actin expression. Western blot  
731 images are representative of three separate experiments with similar results. Cumulative  
732 densitometric values (at the bottom of the Figure **A,B**) are expressed as OD Ratio with actin for  
733 both 4 and 24 h. Values are presented as means ± S.D. of three experiments. Statistical analysis was  
734 conducted by one-way ANOVA followed by Bonferroni's for multiple comparisons. <sup>#</sup>P≤0.05,  
735 <sup>##</sup>P≤0.01, <sup>###</sup>P≤0.005 vs CTRL group; \*P≤0.05, \*\*P≤0.01, \*\*\*P≤0.005 vs IL-17 group.

736

737 **Fig. 7 Schematic representation of involvement of IL-17/mPGES-1/PPAR-γ axis on the onset**  
738 **and resolution of inflammation:** IL-17 injection into air pouch recruits neutrophils and, more  
739 specifically, inflammatory monocytes, producing a massive release of pro-inflammatory cyto-  
740 chemokines at both 4 and 24 h. However, the co-administration of IL-17 with an mPGES-1  
741 antagonist (PF) or PPAR-γ agonist (TRO) shifts the *equilibrium* between COX-2/mPGES-1 and  
742 PPAR-γ/mPGES-1 pathways, downregulating PGE<sub>2</sub> and upregulating PGD<sub>2</sub>/PGJ<sub>2</sub> levels via NF-κB.