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1 **Basophils promote barrier dysfunction and resolution in the atopic skin**

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24 **Declaration of interests**

25 The authors declare no competing interests

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28

29 **ABSTRACT**

- 30
- 31 • **Background:** Type 2 cytokines IL-4 and IL-13 promote atopic dermatitis (AD) but also
32 the resolution of inflammation. How type 2 cytokines participate in the resolution of
33 AD is poorly known.
 - 34 • **Objective:** To determine the mechanisms and cell types governing skin
35 inflammation, barrier dysfunction, and the resolution of inflammation in a model of
36 atopic dermatitis (AD)
 - 37 • **Methods:** Mice reporting the expression of IL-4, IL-13, and MCPT8, or that could be
38 deleted of basophils or eosinophils, or deficient in IL-4 or MHCII molecules, or in
39 which basophils lack M-CSF, were treated with MC903 as an acute model of AD.
40 Kinetics of the disease; keratinocyte differentiation; leukocyte accumulation,
41 phenotype, function and cytokine production were measured by histopathology,
42 molecular biology, or unbiased analysis of spectral flow cytometry.
 - 43 • **Results:** Basophils were activated systemically and were the initial and main source
44 of IL-4 in the skin in this model of AD. Basophil and IL-4 promoted epidermal
45 hyperplasia and skin barrier dysfunction by acting on keratinocytes differentiation
46 during inflammation. Basophil, IL-4 and basophil derived M-CSF inhibited the
47 accumulation of pro-inflammatory cells in the skin while promoting the expansion
48 and function of pro-resolution M2-like macrophages and the expression of pro-
49 barrier genes. Basophils kept their pro-resolution properties during AD resolution.
 - 50 • **Conclusion:** Basophils can display both beneficial and detrimental type 2 functions
simultaneously during atopic inflammation.

51 **Key Messages**

- 52 • Basophils and IL-4 drive skin barrier dysfunction and keratinocyte differentiation
- 53 • Basophils M-CSF and IL-4 promote M2 expansion, efferocytosis, and resolution
54 during atopic inflammation

55 **KEYWORDS:** Atopic dermatitis, basophils, M2, macrophages, efferocytosis, type 2

56 inflammation, resolution, IL-4, M-CSF.

57 Abbreviations: AD: Atopic dermatitis, ILC: innate lymphoid cell, TEWL: transepidermal water

58 loss, TSLP: thymic stromal lymphopoietin,

59

60 **Capsule Summary:**

61 Basophil promotes barrier dysfunction and the resolution during murine atopic skin
62 inflammation, partly by expressing interleukin 4 and M-CSF. Thus, basophil and interleukin 4
63 could play pathogenic and beneficial roles simultaneously in human atopic dermatitis.

64 INTRODUCTION

65 Atopic dermatitis (AD) is the most common chronic skin condition, affecting up to 20% of
66 the population. It is characterized by intense itch episodes and epidermal barrier
67 dysfunction, associated both epidemiologically and in mice models with the release of
68 thymic stromal lymphopoietin (TSLP), IL-33, IL-4, IL-13, and histamine(1). TSLP and IL-33 can
69 be secreted by the inflamed epidermis to directly induce the secretion of IL-4/13 by type 2
70 innate cells(2,3). The efficacy of Dupilumab (a monoclonal antibody targeting IL-4R α) in AD
71 underlines the critical functional role of IL-4 and/or IL-13 in this pathology(4).

72 Type 2 cytokines and histamine act on keratinocytes to prevent their timely differentiation
73 in an ordered epidermis(5). Alongside TSLP, they also promote chronic itch through direct
74 effects on sensory neurons(6). Keratinocyte differentiation and itch/scratch cycles are
75 critical contributors to the epidermal hyperplasia and barrier dysfunction observed in
76 AD(1,4). This leaky barrier enables the establishment of chronic inflammation and skin
77 dysbiosis, and allows entry of allergens which promote the development of the atopic
78 march(7).

79 Basophils are rare circulating granulocytes that can secrete type 2 cytokines and
80 histamine(8). They exacerbate Th2 cell differentiation(9), and infiltrate the skin to activate
81 innate lymphoid cell (ILC)2 expansion and IL-13 secretion, in AD-like disease through their
82 secretion of IL-4(10,11). Basophils have been considered both pro-(10,12) and anti-
83 inflammatory(13). Indeed, they infiltrate the dermis in diverse skin conditions(14), to induce
84 epidermal hyperplasia(15,16), or support the resolution of inflammation(16–18).

85 Besides their pathogenic role in atopic diseases, type 2 cytokines are involved in tissue
86 remodeling and resolution by inducing a M2 macrophage phenotype(19). M2-like
87 macrophages contain heterogeneous subsets specialized in efferocytosis, resolution of

88 inflammation, repair and remodeling(20–22). The dermis contains several subsets of
89 macrophages, including CD64⁺ MHCII^{hi} monocyte-derived macrophages and CD64⁺ CD206^{hi}
90 dermal resident self-renewed macrophages(23,24). The expression of CD206 by
91 macrophages is induced by type 2 cytokines and is critical for the resolution of inflammation
92 and tissue remodeling through pro-inflammatory molecule scavenging and the phagocytosis
93 of collagen fibers, respectively(25,26).

94 We used an established model of AD initiated by the topical application of calcipotriol
95 (MC903) for 4 days on the ear(27), known to be dependent on epidermal TSLP
96 expression(28). This led to an acute AD-like inflammation peaking at days 9-10, followed by
97 a spontaneous resolution. Unbiased high dimensional spectral flow cytometry analysis
98 revealed that basophils were the main source of IL-4 in the skin during both inflammation
99 and its resolution. Basophil depletion, or genetic deletion of IL-4, were sufficient to delay
100 and reduce epidermal hyperplasia and barrier dysfunction. Basophils also induced the
101 expansion of M2 macrophages, efferocytosis, and pro-barrier genes, key indicators of the
102 resolution phase. Thus basophils play both pathogenic and pro-homeostatic roles in a
103 mouse model of AD.

104

105 **Methods**

106 ***Mice***

107 6 to 12 weeks old, Basoph8 (“B8”, which express the Cre recombinase and the eYFP under
108 the control of MCPT8, specifically in basophils)(29,30), iDTR (expressing the simian Diphtheria
109 toxin receptor upon Cre activity) (31), dsRed.T3 (“dsRed”, ubiquitous expression) were from
110 Jackson Laboratories, C57BL/6J from Jackson Laboratories or Charles River Germany, iPhil
111 (which express the simian DTR specifically in eosinophils) were from James J. Lee (Mayo
112 Clinic Arizona)(32), IL-4^{G4/G4} (“IL-4KO”, express the GFP as a knockin, inactivating IL-4
113 expression)(33) and the 4C13R (Dual reporter of IL-4 and IL-13 expression)(30,34) were from
114 Bill Paul (NIH), CSF1^{fl/fl} were from Dr Jean X. Jiang (San Antonio University)(35), TSLPRKO
115 mice(36) from Dr Warren Leonard (NIH) and MHCIIKO mice were gifted by Dr. Horst
116 Bluethmann (Hoffman-La Roche, Basel)(37). All mice were bred, crossed, and housed in
117 specific pathogen-free conditions at the Malaghan Institute of Medical Research Biomedical
118 Research Unit or Novartis Institute for Biomedical Research Center. All experimental
119 protocols were approved by the Victoria University of Wellington Animal Ethics Committee
120 (Permit 24432) and performed according to local Institutional guidelines. Unless specified,
121 only female mice were used as leucocytes and type 2 cytokine expression showed different
122 kinetics in our models (data not shown). Littermates were used for all experiments
123 comparing groups issued from parents with the same genotype.

124 ***Treatments***

125 Mice were anaesthetized using intraperitoneal (i.p.) injection of ketamine/xylazine and
126 4nmoles of MC903 (Calcipotriol, Cayman Chemicals) in EtOH were applied daily topically for
127 4 days in 20µL while EtOH was applied on the contralateral ear (10µL per side). Diphtheria
128 toxin (Cayman Chemical) was injected i.p. (20ng/g/day) to B8xiDTR and every two days to

129 iPhil mice(30,38). AM156 was delivered intragastrically from D0 as described(38). Apoptosis
130 of dsRed mice splenocytes was induced in IMDM 10% FCS (Gibco) + 1 μ M dexamethasone at
131 10M/mL for 24h at 37°C 5% CO₂ (>90%) and washed 3 times in excess PBS before being
132 resuspended at 100x10⁶/mL. 20 μ L was then injected intradermally. 1mg of neutralizing anti-
133 mouse IL-4 (Novartis) or rat IgG1 isotype were injected i.p. every other day.

134 ***Measurements, Histology and Microscopy***

135 Ear thickness and TEWL were measured on sedated mice with a digital caliper (Capa System
136 or Kroeplin) and the DERMALAB™ TEWL probe (Cortex Technology Denmark)(38) or
137 Tewameter TM Nano (Courage+ Khazaka Electronics). Serum IgE was quantified using a
138 mouse IgE ELISA kit (Mybiosource.com, MBS564074). Ear tissue was fixed in 4% PFA 24h,
139 processed, paraffin embedded, sectioned at 4 μ m, and stained with Hematoxylin and eosin
140 (H&E) to measure epidermal and dermal thickness or Masson's trichrome (Sigma) to
141 measure the area of collagen as the percentage of the blue area from the total ear area
142 using ImageJ (NIH). Images were on a widefield microscope Olympus brightfield (BX51) with
143 a 20X NA 0.5 objective and Cell sens (Olympus). For immunofluorescence, whole ears were
144 incubated 1h at 4°C in 20% sucrose, rinsed, snap frozen in OCT in liquid nitrogen, sectioned
145 at 8 μ m, blocked in Superblock (ThermoFisher), and stained with rabbit anti-filaggrin
146 (Poly19058), chicken anti-keratin 14 (Poly9060), rat anti-Ki-67 (16A8, both from Biolegend),
147 and guinea pig anti-keratin 10 (Progen) overnight at 4°C. Sections were washed, stained
148 with goat anti-rabbit IgG AF488 (A27034, Thermofisher), goat anti-chicken IgY AF647
149 (Ab150171, Abcam) and anti-guinea pig AF568 (ab175714, Abcam) for 2h at 4°C before
150 being washed and mounted for confocal analysis(39). For the TUNEL assay, the Click-IT Plus
151 TUNEL assay AF647 kit (Thermofisher) was used as per the manufacturer's guidelines,
152 alongside DAPI. Images were recorded on an inverted microscope IX83 equipped with a

153 FV1200 Confocal head using a 20 X, N.A. 0.75 objective, and using FV10-ASW v4.2b
154 (Olympus) and analyzed with ImageJ v1.52n (NIH) or Cellprofiler 3.1.8 (Broad Institute).

155 ***Tissue digestion***

156 Blood was collected by cheek bleeding or intracardial puncture. Red blood, bone marrow,
157 and spleen cells were lysed in an ammonium chloride buffer and prepared as already
158 described(40). For skin cell preparations, ears were split into the dorsal and ventral layers,
159 minced and for digested 30 minutes at 37°C in a shaking incubator (150rpm) in IMDM 5%
160 FCS (Gibco) containing (2mg/mL) collagenase IV and (100µg/mL) DNase I (from Sigma).
161 Digestion was stopped by adding 5mM EDTA and a tissue filtered through a 70 µm nylon
162 mesh filter (BD). For experiments in Fig. 7B, ears were digested using (0.25mg/mL) Liberase
163 TM (Roche) for 20 minutes at 37°C and dissociated with the GentleMACS dissociator
164 (Miltenyi).

165 ***Flow cytometry***

166 Single cell suspensions were blocked for 15 min at 4°C in FACS buffer (1x PBS with 1%
167 bovine serum albumin and 0.05% NaN₃, (Sigma)) containing 0.5% of 2.4G2 hybridoma
168 supernatant. Cells were then stained in FACS buffer for 20 min at 4°C with an optimized
169 panel of fluorophore-conjugated antibodies , unless specified. Antibodies used are
170 presented in Table E1. A resume of gating strategies is represented in Table E2, and was
171 described previously(41). Compensation was performed using UltraComp eBeads
172 (Invitrogen) as single stained positive controls or dsRed or B8 splenocytes, and fluorescence
173 minus one (FMO) controls were used to set background expression when needed. B8xC57
174 mice were used to set B8x4C13R positivity gates as FMO. Flow cytometry was performed on
175 an Aurora spectral cytometer with 3 lasers (Cytek) or on a Fortessa X20 for Fig. 7B (BD).
176 FACS sorting was carried out on a BD Influx™ (Becton Dickinson). T-SNE analyses were done

177 as Opt-SNE on default settings from at least 4 concatenated samples per group using FlowJo
178 vX (Tree Star).

179 ***Molecular biology***

180 For cell specific gene expression, 2000 or 10000 sort purified cells were collected in RNA
181 lysis buffer and RNA was extracted using a Quick-RNA kit (Zymo Research). For whole skin
182 gene expression, ears were snap frozen, minced, homogenised using 5mm stainless steel
183 beads and a Tissue Lyzer II (Qiagen) and RNA was extracted using Trizol (Thermo Fisher) and
184 the Quick RNA kit (Zymo Research). cDNA was synthesized using the High Capacity RNA- to-
185 cDNA kit (Applied Biosystems). RT-qPCR was performed with a QuantStudio 7 (Applied
186 Biosystems) and following the manufacturer's guidelines.using SYBR Green Master Mix and
187 the following probes (*Gapdh*: F 5'- AATGGTGAAGGTCGGTGTGA, R 5'-
188 GCAACAATCTCCACTTTGCCA; *Furin*: F 5'- ACACACAGATGAATGACAAC, R 5'-
189 GCATTGTAAGCTACACCTAC; and *Dsg1a*: F 5'- AAGGCAGAAACGAGAATGGA, R 5' –
190 CGAGATGCGGTATGTCCTG) or Taqman Master Mix (all from Thermo Fisher) as a duplex
191 with probes described in Table E3. Transcript levels are expressed as the ratio of $2^{-\Delta CT}$ to
192 GAPDH.

193 ***Statistics***

194 Statistical analyses were performed using Prism 8 (GraphPad). Data from experimental
195 replicates was pooled if this led to a decreased variance, or representative results from an
196 individual experiment are represented. Parametric or non-parametric tests were used based
197 on the normality of the data distribution assessed with a D'Agostino-Pearson K2 omnibus
198 test. Individual dots and/or mean +/- SEM or median and the interquartile range or violin
199 plots are shown, depending on the distribution. Post-test adjusted p values are always

200 represented when used. In all cases, n represents an individual mouse and a two-

201 tailed p value < 0.05 was considered as threshold for significance.

202

203

204 RESULTS

205

206 1) MC903 induces an AD-like resolving epidermal pathology

207 After 4 days of daily MC903 treatment(27), ear thickness and transepidermal water loss

208 (TEWL, a measure of epidermal dysfunction) continued to increase until day 9-10 before

209 resolving, defining an inflammation (Day 0-9) and a resolution (Day 10-14) phase (Fig. 1A).

210 Skin presentation included a redness, stiffness, shrinking, dry skin and progressive scaling of

211 the ear (Fig. 1B). Although disease progression was associated with dermal and epidermal

212 thickening (Fig. 1C), only epidermal swelling was associated with disease kinetics (Fig. 1D).

213 The quantity of collagen in the dermis dropped during inflammation, before increasing

214 during the resolution phase (Fig. 1E, F). A basal layer of K10⁻ K14⁺ cells (*stratum basale*, Fig.

215 1G, H) and a suprabasal layer of K10⁺ K14⁺ keratinocytes (*stratum spinosum*, Fig. 1G, I)

216 expanded in the epidermis. Ki67⁺ proliferating keratinocytes were found mostly in the

217 *stratum basale* (Fig. 1J). The expression of profilaggrin (*Flg*), and an outer layer of FLG⁺

218 keratinocytes (*stratum granulosum*) expanded over time, revealing a peak of acanthosis at

219 D9(42). Hyperkeratosis was evident from D7 (Fig. 1C, E, G). These features were observed

220 alongside intercellular edemas (mild spongiosis) and focal parakeratosis (Fig. E1A).

221 Importantly, epidermal disease was more severe on the ear edges (Fig. E1B) Kinetics of

222 collagen deposition were confirmed using picosirius red staining (Fig. E1C, D). Thus 4 days

223 of induction with MC903 induced a skin barrier dysfunction with some histopathological

224 features of AD(1).

225

226 2) Basophils are the main source of type 2 cytokines in the skin

227 Unbiased neighbouring analysis of leukocyte kinetics measured by high dimensional
228 spectral flow cytometry from D0 to D12 after treatment revealed 12 time-dependent
229 clusters (Fig. 2A). The first cluster contained CD206^{hi} dermal resident and MHCII^{hi} monocyte
230 derived CD64^{hi} macrophages, and CD11b⁺, CD11b^{lo} (triple negative, TN) and CD206⁺CD301⁺
231 DCs (CD64⁻ CD11c⁺ MHCII⁺). Mast cells (3), ILC2s (7), dendritic epidermal T cells (DETCs) and
232 $\gamma\delta$ T cells (5) were identified in the naïve skin (D0). Resident cells were stable overtime, with
233 the exception of macrophages and DCs increasing steadily (Fig. 2A, Fig. E2A). The number of
234 infiltrating neutrophils (2), Ly6C⁺ monocytes and macrophages (9, CD64^{low/high}, respectively)
235 peaked around D7, while eosinophils (10), CD4⁺ and CD8⁺ T cells (6), natural killer (NK) and
236 NKT cells (8) and plasmacytoid DCs (11) peaked at D10. Basophils (4), CD4⁻CD8⁻ (DN) T cells
237 (6) and B cells continued to increase during the resolution phase (Fig. 2B, Fig. E2B). Cluster
238 12 represented highly fluorescent CD45 negative cells that were excluded from further
239 analyses. These results confirmed the kinetics of the inflammation and resolution phases
240 (Fig. 1), and highlighted a disappearance of neutrophils, but not basophils, at the beginning
241 of the resolution phase.

242 The expression of IL-4 and IL-13 in the skin was mainly restricted to basophils (4) and ILC2s
243 (7), respectively (Fig. 2C). Only minor subsets of CD4⁺ T cells were positive for IL-4 or IL-13
244 (Fig. 2D), contrary to the draining lymph nodes (LN) (27). Basophils were the initial and
245 main source of IL-4 in the skin (Fig. E2C, Fig. 2C, E), while eosinophils did not show a
246 significant expression of IL-4 or IL-13 (Fig. E2D). The proportion of IL-13⁺ cells was lower
247 than IL-4⁺ cells after D7 (Fig. 2E). IL-4 gene expression outweighed IL-13 by ~10 to 100 fold
248 from D9 to D12 (Fig. 2F). Thus, basophils were the main source of IL-4 in the skin during
249 both inflammation and the resolution.

250

251 **3) Cognate CD4⁺ T cells and eosinophils are redundant for the induction of skin**
252 **barrier dysfunction**

253 Upon MC903 treatment, MHCII^{ko} mice(43) showed normal skin inflammation and barrier
254 function (Fig. E3A), despite an ablation of CD4⁺ T cells, and an increase of CD8⁺ T cells and
255 ILC2s in the skin (Fig. E3B).

256 Specific conditional depletion of eosinophils(32) impaired the development of skin
257 inflammation but not barrier dysfunction (Fig. E3C). Eosinophils did not express IL-4 or IL-13
258 (Fig. 2B, D, E2D), or control the expression of IL-4 or IL-13 by basophils, ILC2s, CD4⁺ T cells or
259 CD206^{hi} macrophages (Fig. E3D, E3E). Overall, neither MHCII-dependent cognate CD4⁺ T
260 cells nor eosinophils played a non-redundant role in MC903-induced epidermal barrier
261 dysfunction during inflammation.

262

263 **4) Basophils are activated both systemically and locally in the skin**

264 Epidermal barrier dysfunction is critical for AD disease(1), and appeared between D5 and
265 D7 (Fig. 1A). From D4, peripheral blood and bone marrow (BM) basophils were more
266 activated: they were bigger and overexpressed CD11b, FcεRIα, and IL-4. This systemic
267 activation disappeared before D15 in the BM (Fig. E4A, B). Prostaglandin D2 (PGD2)
268 mediates eosinophil recruitment and AD-like pathology in a chronic MC903 model(38), and
269 can activate basophils systemically *in vivo*(40). However, in this acute model basophil
270 activation and IL-4/13 expression were not inhibited by AM156, a PGD2 receptor antagonist,
271 in the spleen or the bone marrow (Fig. E4C, D, E). TSLP is a major driver of MC903 induced
272 AD found at D4 in the serum in this model(27,28). TSLP is known to activate murine
273 basophils(2). MC903 did not induce basophils activation at D4 in TSLPR^{ko} mice (Fig. E4F).

274 Additionally, FACS-sorted TSLPR^{ko} basophils injected intradermally in MC903 treated ears
275 did not induce an optimal skin leukocyte recruitment (Fig. E4G).

276 TSLP and Cyp24a1 are expressed via a direct response of keratinocytes to MC903, which
277 ceases after treatment. Cyp24a1 codes for a member of the P450 cytochrome family
278 involved in vitamin D metabolism(27,28). IL-3 is the most potent cytokine that can activate
279 basophils in the skin(8). Interestingly, the expression of Cyp24a1, a gene known to be
280 induced by TSLP in the MC903 model, decreased from D7, and from D9 IL-3 expression was
281 dominant over TSLP, which was not detected anymore (Fig. E4H).

282

283 To explore skin and systemic basophil activation we sort-purified basophils from the spleen
284 and the skin at D4 or D15 after MC903 treatment. Spleen basophils showed an increased
285 expression of IL-4, IL-13, and IL-33R at D4 (but not at D15). At D4, skin basophils had a
286 distinct phenotype and expressed more GATA3, IL-6 and amphiregulin, but less CCR3. A
287 similar pattern was observed at D15 (Fig. E4I, J). Therefore, basophils seemed initially
288 systemically activated by TSLP (D4-D7), but their activation was imprinted by local signals in
289 the skin, which may include IL-3, during both the inflammation and resolution phases (D7-
290 15).

291

292 **5) Basophil and IL-4 control epidermal function and differentiation**

293 Basophil conditional depletion(30) or IL-4 deletion did not reduce ear swelling but delayed
294 and reduced the epidermal barrier dysfunction, which is critical to AD pathogenesis (Fig.
295 3A)(1). Basophil depletion reduced IL-4 gene expression by ~95% (Fig. 3B, C), confirming
296 that basophils are the main source of IL-4 in this model (Fig. 2), and suggesting that
297 basophil-derived IL-4 was important for the development of skin barrier dysfunction.

298 Skin barrier function is tightly controlled by keratinocyte proliferation, differentiation, and
299 cornification(1,5). Basophils and IL-4 induced epidermal hyperplasia (Fig. 3D), through an
300 expansion of Keratin 10⁺ and Filaggrin⁺ differentiated keratinocytes (*Stratum spinosum* and
301 *granulosum*) (Fig. 3E). Basophils also increased the number of proliferating Ki67⁺ K14⁺
302 keratinocytes, the area of the Keratin 14⁺ *stratum basale* (Fig. 3F), and keratinocyte
303 differentiation (Fig. 3G).

304 These results suggested that basophils and IL-4, among other mediators, can promote
305 keratinocyte differentiation, epidermal hyperplasia and skin barrier dysfunction,
306 cornerstone features of AD(1). Basophil also promoted the differentiation of keratinocytes
307 expressing the pro-barrier genes *Flg* and *Dsg1a*(44,45).

308

309 6) **Basophil and IL-4 promote a pro-resolution leucocyte landscape during** 310 **inflammation**

311 At D9 after MC903 treatment, during inflammation, we analyzed skin leucocytes after
312 basophil depletion by performing unbiased neighboring analyses and observed 11 different
313 clusters (Fig. 4A)(41). Specific basophil conditional depletion tended to decrease mast cells,
314 and CD206^{hi} and MHCII^{hi} macrophages proportions (Fig. 4B, E), but increased pro-
315 inflammatory cells such as neutrophils, CD4⁺ and CD8⁺ T cells (Fig. 4C), and Ly6C⁺
316 macrophages (Fig. 4D). CD206^{hi} macrophages expressed IL-4 at day 9, and this proportion
317 decreased after basophil depletion (Fig. 4F)(21,25,46–48). Basophils were also associated
318 with a decrease in ILC2 IL-13 expression (Fig. 4G)(10).

319 Expression of the M2 associated markers Relm α and Arginase 1 increased from the
320 inflammation phase (D7) to the resolution phase (D12) in the skin. Basophil depletion and
321 IL-4 deletion dampened the expression of Relm α but not Arginase 1 (Fig. 4H). Basophil

322 depletion tended to decrease skin Alox5 expression, but not Alox8, 12 and 15, the main
323 specialized pro-resolution mediators generating lipxygenases(49). The expression of IL-3,
324 IL-5, IL-6, and IL-33, but not other type 2 associated signals (IL-10, IL-18, TGF β , amphiregulin,
325 CCL17, M-CSF, galectin 3 and 9) also decreased after basophil depletion. Regulatory T cells
326 (Tregs) did not seem important in the resolution, as FoxP3 expression decreased from D0 to
327 D12 (Fig. E5A). Dermal macrophages seemed specifically biased by basophils to express
328 more of the pro-resolution markers Alox15 and Relm α at D9 (Fig. E5B).

329 Overall basophils and IL-4 limited the accumulation of pro-inflammatory cells, induced M2-
330 like macrophages and favored a pro-resolution landscape during skin inflammation.

331

332

333 **7) Basophils control dermal resident macrophage efferocytosis**

334 Efferocytosis is a critical step in the resolution. It is mainly carried out by CD206^{hi} dermal
335 resident macrophages, by IL-4 expressing macrophages, is induced by IL-4/13 and involves
336 CD206 expression(22,24,46,48). Indeed, the majority of dsRed⁺ apoptotic cells injected into
337 D9 treated ears were taken up by CD206^{hi} dermal resident macrophages (Fig. 5A). Basophil
338 depletion decreased efferocytosis by skin macrophages. This was due specifically to a defect
339 in the uptake of apoptotic cells by dermal resident CD206^{hi} macrophages (Fig. 5B). TUNEL⁺
340 apoptotic cells were observed in the epidermis in control mice (EtOH), but also in the dermis
341 at Day 9 after MC903 treatment. The number of apoptotic cells in the skin eventually
342 decreased during the resolution phase (Fig. 5C).

343 Thus, while apoptotic cells accumulated in the dermis during this AD-like inflammation,
344 basophils promoted the pro-resolution efferocytosis function of dermal resident CD206^{hi}
345 macrophages.

346

347 8) **Basophils show pro-homeostatic properties during the resolution phase**

348 Eosinophils express lipoxygenases, and can play a pro-resolution role(50). As we failed to
349 deplete skin-infiltrated eosinophils during the resolution phase, we couldn't explore a role
350 of eosinophils in the resolution of MC903 induced skin inflammation(Fig. E6A, B, C).

351 Basophil depletion during the resolution phase lead to a small reduction in TEWL(30).

352 Unexpectedly, this was associated with an accumulation of neutrophils and Ly6C⁺
353 monocytes, but not eosinophils (Fig. 6A). DETC, ILC2 and mast cell proportion, and IL-13
354 expression by DETCs but not by ILC2s decreased upon basophils depletion (Fig. 6B)(10).

355 Type 2 cytokine expression by CD4⁺ T cells was unaffected (data not shown). Proportions of
356 macrophages and of PDL2⁺ monocyte-derived M2 macrophages(51) were unchanged but
357 pro-inflammatory Ly6C⁺ macrophages tended to accumulate after basophil depletion.

358 Similarly, macrophage expression of the resolution associated markers CD206 and IL-4 were
359 decreased(24,25,46,48) (Fig. 6C), as well as the expression of *Il4* and *Retnla* in the skin (Fig.

360 6D). Interestingly *Flg* expression and FLG⁺ keratinocyte numbers decreased without any

361 change in dermal or epidermal hyperplasia (Fig. 6E, F).Collagen deposition increased in the

362 dermis, as measured after Masson's trichrome staining (Fig. 6G) or picosirius red staining

363 (data not shown). As dermal macrophages internalize and degrade collagen through CD206,

364 which is induced by IL-4, it is likely that basophils promoted this tissue remodeling through a
365 control of M2 expansion and function(26).

366 Collectively, basophils induced a M2-like macrophage bias, and a decrease of pro-

367 inflammatory cells during the resolution phase, associated with skin repair and remodeling.

368

369 9) **Both IL-4 and basophil derived M-CSF participate in the resolution**

370 IL-4 neutralization during the inflammatory phase did inhibit IgE secretion, as expected (Fig.
371 E7A). During the resolution phase, it did not inhibit the generation of IgE anymore (Fig. E7B),
372 nor did it lead to changes in ear swelling, TEWL (Fig. 7A) or total leucocyte infiltration.
373 However, IL-4 neutralization did inhibit the accumulation of neutrophils in the skin during
374 the resolution phase (Fig. 7B).

375 Basophil depletion during the inflammation phase revealed IL-4 independent effects such
376 as keratinocyte proliferation and the expansion of monocyte derived macrophages (Fig. E2,
377 E3). We looked for the genes expressed by basophils known to be associated with the
378 induction of M2 macrophages or the resolution of inflammation using a transcriptomic
379 database(52), and found lipoxygenases (*Alox5*, *Alox8*, *Alox12*, *Alox15*) and M-CSF were
380 potently expressed by peripheral basophils (Fig. E7C). At D9 after MC903 treatment,
381 basophils expressed similar or higher levels of pro-resolution genes as eosinophils or
382 macrophages. Besides *Il4*, they were specifically strong producers of *Alox8*, *Alox12*, *Areg*,
383 *Csf1*, *Il6* and *Tgfb1* (Fig. E7D). While skin basophil expression of *Il4* and *Il13* evolved time
384 dependently, *Csf1* expression stayed high from naïve conditions to D12 (Fig. E7E). As M-CSF
385 is important for both monocyte/macrophage and resident M2 macrophages, and shows
386 pro-repair and homeostatic functions(47,53,54), we generated the Basoph8.CSF1^{fl/fl} mice
387 (CSF1^{ΔMCPT8})(30,35), in which basophils specifically show a 90% decrease of M-CSF
388 expression (Fig. E7F). Topical application of MC903 induced an exacerbated pathology with
389 an increased TEWL and/or ear swelling (Fig. 7C, D), and an altered skin immune cells
390 content: they showed fewer MHCII^{hi} and CD206^{hi} macrophages, CD4⁺ T cells, $\gamma\delta$ T cells,
391 DETCs and ILC2s at D12 after MC903 treatment than their M-CSF-sufficient littermates.
392 CSF1^{ΔMCPT8} mice showed an accumulation of neutrophils in the skin (Fig. 7B).

393 CD206^{hi} dermal resident and infiltrating macrophage maintenance, M2 phenotype and
394 function rely on both IL-4 and M-CSF(24,47,54). Together, these results show that IL-4 and
395 basophil derived M-CSF control M2 homeostasis in an AD-like pathology. Basophils initiate
396 skin barrier dysfunction and control epidermal proliferation and differentiation after MC903
397 treatment, but also promote M2 macrophages expansion and function during skin
398 inflammation and its resolution.

399 **DISCUSSION**

400 Protective type 2 responses emerge following epithelial injury to expel parasites or noxious
401 substances and thus type 2 cytokines promote a characteristic tissue phenotype including
402 mucus secretion, remodeling, repair, and a return to homeostasis. Inappropriate or
403 excessive type 2 responses can become pathogenic if they are not resolved, and lead to
404 chronic itching or fibrosis(19). An intrinsic defect in the barrier function of the epidermis,
405 observed in filaggrin or desmoglein deficiencies, can lead to excessive type 2 inflammation
406 in the skin and to the development of AD(1,44,55). The type 2 cytokines IL-4 and/or IL-13
407 are critical for AD chronic inflammation, as shown by the clinical efficacy of Dupilumab, but
408 how type 2 effector cells impair or promote the resolution of AD inflammation has not been
409 studied(1,4).

410 The occurrence of AD is associated with industrialization, and environmental factors such as
411 pollutants can lead to chronic skin barrier dysfunction and scratching (56). Such an innate
412 chronic barrier dysfunction allows allergen entry and the development of specific adaptive
413 atopic responses. Indeed, MC903 induces a TSLP-dependent AD-like dermatitis in the
414 absence of functional adaptive immunity, but also promotes specific allergen sensitization
415 through the skin(11,12,55). TSLP has been found to be more elevated in the lesional skin of
416 pediatric cases compared to the adults(57). Most early mild pediatric cases enter remission
417 in the first years of life, and do not show high IgE levels or positive skin prick tests to
418 common allergens. Severe cases are more persistent and associated with allergen,
419 commensals, or autoantigen sensitization(58–60). Thus TSLP could be an important initiator
420 of allergen sensitization and T cell responses in infants, as in mice(9,27,61), but then
421 become redundant once allergen sensitization is driving the disease. Indeed Tezepelumab (a
422 TSLP neutralizing antibody) failed to improve moderate to severe adult AD over placebo

423 when associated with topical corticosteroids, with most patients showing high total IgE
424 titers >150kU/L(62). Epithelial derived cytokines such as TSLP or IL-33 can trigger IL-4 and IL-
425 13 expression by basophils, mast cells, eosinophils or ILC2s(63–65). These mechanisms
426 could be more important for AD initiation during infancy than for skin inflammation in
427 adulthood, when potent specific T cells and IgE responses are already established and
428 dominant.

429 Here we characterized a common model of TSLP-dependent dermatitis . Mice developed a
430 self-resolving AD-like disease histologically closer to acute than adult chronic AD. MC903
431 treatment was not associated with allergens(12,66), in order to study the mechanisms
432 leading to barrier dysfunction prior to allergen sensitization. In this regard, this model
433 differed from human AD, which is strongly associated with specific CD4+ T cell responses,
434 whereas we did not observe a MHCII dependent skin phenotype, despite the central
435 expansion of Th2/Th13 cells occurring in this model(27). CD4+ cells were also redundant to
436 induce skin inflammation in a chronic model of MC903 induced AD. Importantly, in this
437 chronic model, skin barrier function was determined by eosinophils and PGD2(38), which
438 differs from the acute situation described here. Another limitation of this study is that the
439 remission of human AD inflammation is usually controlled by weeks of corticosteroid
440 treatment, whereas we studied a model showing a spontaneous resolution after the
441 removal of the chemical offending the skin.

442 Immune cell recruitment to the inflamed skin tracked closely with keratinocyte
443 differentiation and skin pathology kinetics. Contrary to pro-inflammatory cells, IL-4
444 producing basophils remained in the skin during the resolution phase. Basophils were
445 activated systemically during the initiation of the disease, and were recruited to the dermis
446 to promote keratinocyte proliferation and differentiation, and to accelerate a skin barrier

447 dysfunction in a partially IL-4 dependent manner. However, their role could not be
448 described as pro-inflammatory or pathogenic: they also promoted the expansion and
449 function of M2-like macrophages, and the expression of pro-barrier genes such as *Flg* and
450 *Dsg1a*, during the inflammation phase. During the resolution phase, basophils supported a
451 pro-resolution macrophage phenotype, a decrease in pro-inflammatory cells, epidermal
452 repair and tissue remodeling. This late support seemed to rely on basophil IL-4 and M-CSF
453 expression. Both pro-inflammatory and pro-resolution roles have been attributed to
454 basophils during IgE mediated chronic and prurigo-like skin inflammation(16,18). However,
455 in a liver infection model dominated by type 1 inflammation, basophils showed only some
456 pro-resolution properties (17). Indeed, any basophil pro-homeostatic properties seem
457 hidden by the deleterious effects of type 2 inflammation during atopy or allergy.

458 This homeostatic role of basophils has been associated with resident dermal macrophages
459 in our current study, with monocyte derived macrophages during IgE mediated skin
460 inflammation(16,18), with Kupffer cells during liver infection(17), and with alveolar
461 macrophages during lung development(67). If basophils seem specialized to interact with
462 distinct macrophages subsets, they also express numerous ligands able to interact with non-
463 hematopoietic cells(67). Basophils can induce keratinocyte proliferation and
464 differentiation(15), but they also control the activation of fibroblasts(68,69) and endothelial
465 cells(70) via their secretion of IL-4 and histamine. As IL-4 has been shown to decrease
466 keratinocyte differentiation and *Flg* expression in vitro, it is likely that basophils and IL-4
467 promoted keratinocytes differentiation and *Flg* expression through indirect effects in the
468 current model in vivo, for example as a counterbalancing response to increased scratching.

469 Further investigation is needed to understand if basophils can directly contribute to tissue
470 remodeling and repair.

471 Cardinal immunological features of AD include an expansion of Th2 cells, eosinophils and
472 ILC2s in the skin, all of which have been observed in various MC903 induced AD-like
473 models(10,27,71). Surprisingly, eosinophils or cognate T cells did not represent a major
474 source of IL-4/13 in the current model. Hence, our data supports the notion that basophil
475 type 2 cytokine production can initiate an innate AD-like pathology.

476 Thus, basophils can promote a pleiotropic type 2 program inducing both a detrimental skin
477 barrier dysfunction, and beneficial efferocytosis, repair and remodeling through their
478 control of keratinocyte and dermal macrophage functional phenotypes.

479

480

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487 **Authors contribution**

488 Conceptualization, C.P. and G.L.G. ; Methodology, C.P., K.N., A.S. ; Investigation, C.P., K.N.,
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775

776 **Figure titles and Legends**

777 **Fig. 1: MC903 induces an AD-like resolving epidermal disease.** C57BL/6 mice ears were
778 treated with 4nmol MC903 or EtOH daily for 4 days. (A) Ear thickness and transepidermal
779 water loss (TEWL) were monitored over time (n=27). (B) Representative pictures of the ear
780 skin. (C) Representative H&E staining (scale = 50µm, 20X) and (D) quantification of dermal
781 and epidermal thickness (n=5, 5 Fields of view (FOV)/ear). (E) Representative Masson's

782 trichrome stained ear tissue sections at day 9 and day 12 (scale = 50 μ m, 20X) and (F)
783 quantification of the dermal area of collagen (n= 8, 6-8 FOV/ear 20X). (G) Representative 3D
784 reconstruction of Ki67, keratin 14 and 10 (K14, K10) and filaggrin (FLG) immunofluorescent
785 staining of the ear skin (40X) and quantification of the area of (H) K14, (I) K10, (J) K14+ Ki67+
786 keratinocytes, (K) the area of filaggrin and the (L) number of FLG+ cells (H-K: n=4, L: n=3; 5
787 FOV/ear). Results are from 5 (A, B), 2 (F-K) or (D, L) are representative of at least 2
788 independent experiments. Means +/- SEM are represented. (D, F, H-L) Statistics are one-way
789 ANOVA with Tukey's post-test. *: p<0.05; **: p<0.01; ***: p<0.001; ****: p<0.0001

790

791 **Fig. 2: Basophils are the main source of type 2 cytokines in AD skin.** B8x4C13R female mice
792 were treated with MC903 and skin leucocytes analysed by flow cytometry. (A) Contour and
793 pseudocolor plots of a representative t-SNE analysis of the kinetics of skin leucocytes
794 populations (10k CD45⁺/mouse, 4 mice/day). Clusters were numbered and immune cells
795 identified in each cluster. (B) Kinetics of selected leucocytes are depicted (n=10). (C) Overall
796 expression of IL-4 (AmCyan) and IL-13 (dsRed) from D0 to D12 (as in A) in heatmap of
797 intensity of expression. (D) Representative dot plots showing the expression of IL-4 and IL-
798 13 by basophils, ILC2s or CD4⁺ T cells at Day 9. (E) Quantification of the number of IL-4 or IL-
799 13+ cells (n=10) or (F) Il4 and Il13 transcripts in the ear (n= 4, 5, 5, 10) overtime. Results are
800 representative of at least two independent experiments showing similar results. (B) Means
801 +/- SEM or (E, F) medians and interquartile ranges are represented. (F) Statistics are a two
802 way ANOVA with Sidak's multiple comparison. *: p<0.05; ****: p<0.0001

803

804 **Fig. 3: Basophil derived IL-4 controls epidermal barrier function and differentiation.**
805 B8xiDTR or IL-4^{KO} mice were treated with MC903 or EtOH, and diphtheria toxin (DT) or PBS.
806 (A) Ear swelling and TEWL monitoring (n = 8, 12, 12, 14). (B) Gene expression of *Il4* (n=14,
807 17) and (C) *Il13* in the ear at D9 (n=10, 10, 12) ND: Not detected. (D) Dermal and epidermal
808 thickness assessed on H&E sections (n=3, 4, 4, 4, 6-7 Fields of view (FOV) / ear) at D0 or D9.
809 (E) Number of keratin 10+ (K10), Filaggrin+ (Flg+) cells assessed at 40X (n=4, 5-6 FOV/ear). (F)
810 K14+ Ki67+ cells (n=5, 5-6 FOV/ear) and keratin 14 area (n=4, 5-6 FOV/ear) at 20X at D9. (G)
811 C57BL/6 ear gene expression overtime (n=6-10) and at D9 (n=10-20). (A, E, F) Means +/-
812 SEM or (B, C, G) medians and violin plots are represented. (A, F) Representative of, or (B - E,
813 G) from at least two independent experiments. Statistics are (A) two-way ANOVA with a
814 Sidak's or (D) a Kruskal-Wallis or (E, F) one-way ANOVAs with a Dunnett's post-test and (B,
815 C, G) Mann Whitney tests. ns: non-significant. *: p<0.05; **: p<0.01; ***: p<0.001; ****:
816 p<0.0001

817 **Fig. 4: Basophil derived IL-4 promotes a pro-resolution leucocyte landscape during**
818 **inflammation.** B8.4C13RxiDTR mice were treated with MC903 with or without diphtheria
819 toxin (DT) until D9. Ear CD45+ cells were analysed by flow cytometry. (A) Representative t-
820 SNE analysis of skin leucocytes populations at D9 (20k CD45+ / mouse, n=5), showing
821 leucocyte clusters, (B) and overlay of DT treated (Red) and untreated mice (Black), revealing
822 depleted clusters. The proportion of (C) granulocytes and relevant lymphoid cells (n=13-17),
823 (D) Ly6C+ monocytes or macrophages (n=13-22), (E) CD206^{hi} or MHCII^{hi} macrophages (n=6 -
824 12) and (F) the expression of IL-4 (AmCyan) by CD206^{hi} macrophages (n=19-23), (G) CD4+ T
825 cells, or expression of IL-13 (dsRed) by CD4+ T cells and ILC2s (n=10-13) was analysed at Day
826 9 after EtOH (-) or MC903 treatment without (PBS) or with DT (DT) in B8.4C13RxiDTR or IL-

827 4KO (IL4^{ko}) mice. (H) Gene expression of Relm- α and Arginase 1 was analysed overtime
828 (n=6-17) and at D9 (n=10-12) by qPCR in the whole ear skin. Medians are represented. (A, B)
829 Representative of two or (C to H) from three independent experiments showing similar
830 results. Statistics are Kruskal Wallis with a Dunn's post-test. #: p<0.1; *: p<0.05; **: p<0.01;
831 ***: p<0.001; ****: p<0.0001

832

833 **Fig. 5: Basophils control dermal resident macrophage efferocytosis.** B8xiDTR mice were
834 treated with EtOH (-) or MC903 for 4 days. MC903 treated mice were depleted of basophils
835 by injections of diphtheria toxin (DT) or not (PBS). At D9 after MC903 treatment, 2M dsRed+
836 apoptotic cells were injected in the ear dermis. 2hr later mice were sacrificed and ear cells
837 analysed by flow cytometry. (A) Representative gating strategy of skin phagocytes and their
838 uptake of dsRed+ apoptotic cells. (B) Histograms representing the efferocytosis of
839 macrophages and relevant subsets as in (A), (n=6-8). (C) EtOH or MC903 treated ears were
840 analysed by immunofluorescence for macrophages and apoptotic cell content by TUNEL
841 assay (AF: autofluorescence, scale =50 μ m), and the TUNEL+ area was quantified (n=5-7 mice
842 with 10 Field of view (FOV) per mouse). Results are (A, B) representative of or (C) pooled
843 from two independent experiments showing similar results. Means +/- SEM are
844 represented. Statistics are unpaired t-tests. ns: non-significant. *: p<0.05; **: p<0.01; ****:
845 p<0.0001

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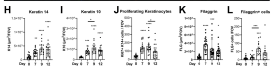
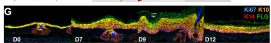
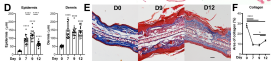
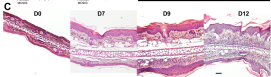
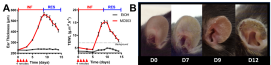
847 **Fig. 6: Basophils show anti-inflammatory and pro-remodeling properties during the**
848 **resolution phase.** B8.iDTRx4C13R mice were treated with MC903 or EtOH on the
849 contralateral ear and injected with diphtheria toxin (DT) or PBS (PBS) daily from D8. At D12
850 (A-C) ear skin leucocytes were analyzed by flow cytometry (n=5-9). (D) Whole skin

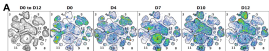
851 transcripts of relevant genes were analyzed by qPCR (n=8-16). (E) Dermal and epidermal
852 hyperplasia was quantified on H&E sections (n=8 ears with 5-7 sections /ear), (F) filaggrin
853 (FLG+), keratin 14 (K14+) and keratin 10 (K10+) cells counted from immunofluorescence
854 sections (n=3 mice, 8 Fields of view (FOV) / ear) and (G) collagen content was determined on
855 Masson's trichrome sections (n= 8 ears with 6-7 section /ear). Results are (B, C, F)
856 representative or (A, E, G) pooled from two to three independent experiments showing
857 similar results. (A-C) Medians and interquartile range or (D, G) in violin plots, or (E, F) means
858 +/- SEM are represented. Statistics are (A-D, G) Mann-Whitney or (E, F) unpaired t-tests. Ns:
859 non-significant. #: p<0.1; *: p<0.05; **: p<0.01; ***: p<0.001; ****: p<0.0001

860

861 **Fig. 7: Basophil derived IL-4 and M-CSF support the resolution of inflammation.** C57BL/6J
862 were treated with MC903 or EtOH on both ears and injected intraperitoneally with anti-IL-4
863 (a-IL4) or rat IgG1 (iso) or PBS at D9 and D11 and (A) their ear swelling and TEWL were
864 monitored overtime (n= 22, 26, 26, 26), and (B) skin leucocytes analysed at D12 by FACS
865 (n=17,19,19). B8^{tg/wt}.CSF1^{fl/fl} (CSF1^{ΔMCPT8}) and B8^{wt/wt}.CSF1^{fl/fl} (CSF1^{fl/fl}) littermates were
866 treated with MC903 and EtOH on the contralateral ear and (C) the ear swelling and TEWL of
867 males (n= 13 and 16) and (D) females (n= 20 and 19 for MC903, 15 and 9 for EtOH) was
868 monitored. (e) Skin leucocytes populations were quantified by FACS in female mice at D12
869 (n=12, 12, 17, 17). (A, C, D) Means +/- SEM or (B, E) medians and interquartile range are
870 represented. Statistics are (A, C, D) two-way ANOVAs with a Tukey's post-test comparing
871 each MC903 treated genotype and (B, E) Mann-Whitney tests. Results are from (A, B, E) two
872 or (C, D) three independent experiments. ns: non-significant. *: p<0.05; **: p<0.01; ***:
873 p<0.001; ****: p<0.0001

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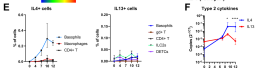
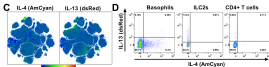
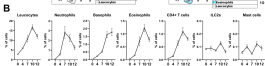


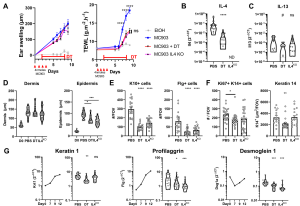


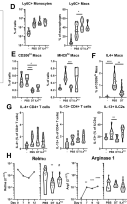
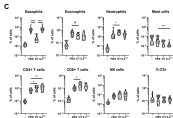
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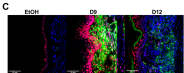
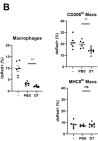
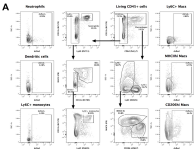


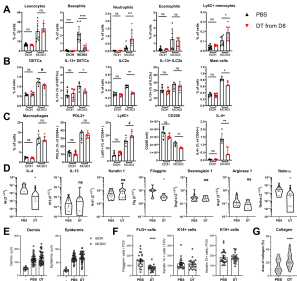
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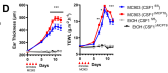
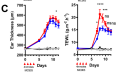
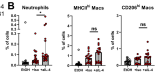
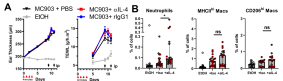




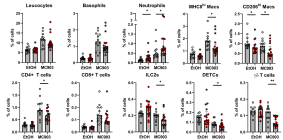






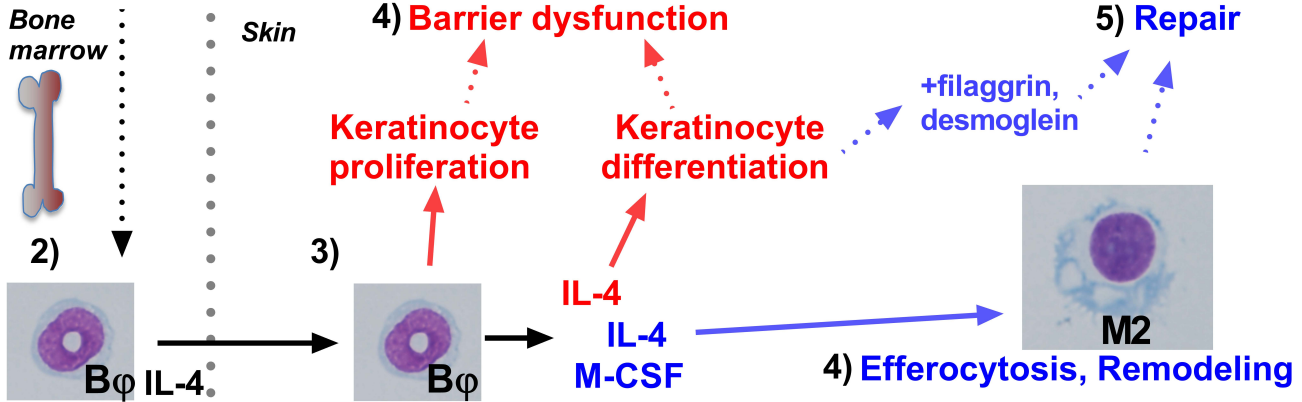
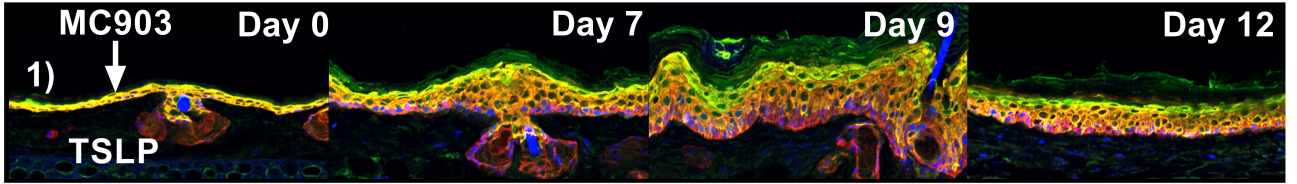


○ **CSP1^{EGFP}** ● **CSP1^{AMCP19}**





Basophil induces keratinocyte dysfunction and pro-resolution macrophages during atopic skin inflammation



B_{ϕ} : Basophils

M2: M2-like Macrophages

→ Direct evidence

··· Association

— Pro-Inflammation

— Pro-Resolution