



# Maturation-induced down-regulation of MFG-E8 impairs apoptotic cell clearance and enhances endotoxin response

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**Abstract.** In sepsis, phagocytosis and the killing of bacteria by phagocytes are important. Similarly, the clearance of accumulating apoptotic cells is critical in maintaining normal immunity. Upon maturation, peritoneal macrophages (PM) become a major source of proinflammatory cytokines, while losing their efficacy of phagocytosis. However, the underlying mechanism remains unknown. Here we investigated the differential effects of apoptotic thymocytes (AoTC) on TNF- $\alpha$  release in immature thioglycolate-elicited PM (TGPM) and mature resident PM (RPM) *in vitro* by culturing them with or without AoTC and/or LPS. MFG-E8 expression was assessed using Western blotting and the ability to engulf AoTC was determined histologically. Cytokine secretion was measured by ELISA. MAP kinase phosphorylation was assessed using Western blotting. Mature RPM express <50% of TGPM MFG-E8 levels and have a 30% lower capacity to clear AoTC. The proinflammatory response (TNF- $\alpha$  release) to LPS is 5 times higher, and the capability to phagocytose is decreased along with further down-regulation of MFG-E8 after LPS-stimulation. RPMs also lack phagocytosis-induced inhibition of TNF- $\alpha$  release after LPS stimulation. LPS-induced phosphorylation of ERK1/2, p38 and JNK is more enhanced in RPM compared to TGPM. MFG-E8-mediated apoptotic cell phagocytosis results in an inhibition of MAPK and NF $\kappa$ B signaling pathways. Differential MAPK activation may play a role in the enhanced LPS responsiveness of RPM and the lack of MFG-E8 impedes post-phagocytic suppression of LPS-response through the inhibition of those signaling pathways. These results provide a potential mechanistic insight into the benefit of promoting apoptotic cell clearance via MFG-E8 under inflammatory conditions.

## Introduction

Under normal conditions, resident peritoneal macrophages (RPMs) inhabiting the peritoneal cavity, are morphologically determined by numerous lamellipodia, subsurface vacuoles and an irregular eccentric nucleus with heterochromatin margination surrounded by a large number of lysosomes (1). Within 24 h after intraperitoneal injection of thioglycolate, two new subpopulations of macrophages emerge that can be distinguished morphologically as monocyte-like macrophages with a kidney-shaped nucleus and numerous lysosomes and the large irregular 'inflammatory macrophages' with many filopodia and cytoplasmic inclusions (1,2). Cytochemically these so-called exudate macrophages have a peroxidase activity limited to their lysosomes, while RPMs show abundant activity in the nuclear envelope and rough endoplasmic reticulum. Thioglycolate elicited peritoneal macrophages (TGPMs) have a higher antibody-dependent and -independent endocytotic and exocytotic activity but are less efficient than RPMs in antibody dependent cell-mediated cytotoxicity and resistance to *Listeria monocytogenes* infection (3-6). In 1994, Blasi *et al* discovered that cytokine mRNA transcription and activity are lower in TGPMs than those in RPMs after stimulation with either *Candida albicans* or LPS (7). Hanayama *et al* found that TGPMs as well as immature dendritic cells and marginal zone macrophages of the spleen contain abundant milk fat globule epidermal growth factor-factor VIII (MFG-E8)(8). MFG-E8 specifically binds to phosphatidylserine and acts as an opsonin for apoptotic cells by bridging phosphatidylserine on apoptotic cells to integrins on phagocytes (9,10). This aids to enhance the clearance of apoptotic cells, ultimately leading to suppression of proinflammatory responses that are crucial in autoimmunity (8) and sepsis (11). As mature RPMs have a lower capacity for phagocytosis than TGPMs, we speculated that this was also true for the engulfment of apoptotic cells. We therefore hypothesized that RPMs contain lower amounts of MFG-E8 and thus are less efficient to clear apoptotic cells. We further hypothesized that this is associated with an increased responsiveness to LPS.

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## Materials and methods

**Peritoneal macrophage culture.** Rat peritoneal macrophages (RPMs) were isolated by peritoneal lavage 72 h after i.p.

injection of 3% thioglycolate (TGPMs, 30 ml/kg) or without pretreatment and cultured in DMEM containing 10% FBS at 37°C. Cells were then incubated with LPS (*E. coli* 055:B5) with or without apoptotic thymocytes. All experiments were performed in accordance with the NIH Guidelines for the Use of Experimental Animals and approved by the IACUC of The Feinstein Institute for Medical Research.

**MFG-E8 Western blotting.** Macrophages were lysed and homogenized in 1 ml of lysis buffer (10 mM Tris-buffered saline, 1 mM EDTA, 1 mM EGTA, 2 mM sodium orthovanadate, 0.2 mM PMSF, 2 µg/ml leupeptin, 2 µg/ml aprotinin, and 1% Triton X-100) for 30 min on ice and tissue sample lysate was cleared by centrifugation at 400 × g for 15 min at 4°C. Samples were dissolved in 1% SDS and quantified using the DC Protein Assay (Bio-Rad, Hercules, CA). Plasma samples were ultrafiltered with Centricon 100 (Millipore) and elute was concentrated x30 using Centricon YM30 filters. Ten µg of protein or 5 µl of plasma concentrate was fractionated on a 4-12% Bis-Tris gel and transferred to 0.2-µm nitrocellulose membrane. Nitrocellulose blots were blocked by incubation in TBST (10 mM Tris-HCl, pH 7.5, 150 mM NaCl, 0.1% Tween-20) containing 10% bovine serum albumin for 1 h. Blots were incubated with goat anti-MFG-E8 IgG clone G-17 (1:100, Santa Cruz Biotechnology, Santa Cruz, CA; an antibody raised against a 17 amino acid peptide sequence shared by human, mouse and rat MFG-E8 in the C1 domain) overnight at 4°C and then washed vigorously 3 times in TBST 10 for min. Blots were incubated with horseradish peroxidase-labeled rabbit anti-goat IgG for 1 h at room temperature, washed as before, incubated with ECL (Amersham) and exposed on radiograph film.

**Phagocytosis assay.** Macrophages were cultured at  $2.5 \times 10^4$ /well in a 16-well chamber slide (Nalge Nunc, Rochester, NY). Thymocyte apoptosis was induced by 10 µM dexamethasone for 24 h (>99% AnnexinV<sup>+</sup>CD90<sup>+</sup>). After washing with PBS, macrophages were incubated with AoTC at a ratio of 4:1 (AoTC/macrophages) for 90 min. Adherent macrophages were washed with PBS, fixed with 4% paraformaldehyde, stained with TUNEL and analyzed by fluorescent microscopy using a Nikon Eclipse E600 microscope. Results presented as a ratio of apoptotic cells/macrophages (phagocytosis index).

**Cytokine assay.** TNF-α release was quantified in supernatants using a commercial ELISA kit (BD Pharmingen).

**Phosphoprotein and nuclear NFκB Western blots.** Macrophages were incubated with apoptotic cells for 90 min followed by washing with PBS and stimulation with LPS for 15 min. Cells were washed, lysed with whole cell lysate buffer (MAPK), or relaxation and nuclear extraction buffer (NFκB and IκBα). Protein (20 µg) was fractionated on a Bis-Tris gel and Western blotting was performed as described above, using specific antibodies against phospho-p38 and total p38, phospho-JNK and total JNK, phospho-ERK1/2 and total ERK1/2 (Cell Signaling, Danvers, MA), as well as anti-NFκB and IκBα antibodies (Santa Cruz Biotechnology, Santa Cruz, CA).

**Statistical analysis.** Data are expressed as mean ± SEM and compared by Student t-test or ANOVA and Student-Newman-Keuls (SNK) test. Statistical significance was set at  $P < 0.05$ .

## Results

We isolated resident peritoneal macrophages (RPM) from the peritoneal lavage fluid of normal rats or thioglycolate-induced peritoneal macrophages (TGPM) 3 days after injection of 10 ml 0.3% thioglycolate into the peritoneum. The initial lavage population was >90% RPMs and ~50% TGPM (the other half being ~50% neutrophils and sporadic eosinophils and basophils). After enrichment by adherence, both populations were >95% pure (Fig. 1A). MFG-E8 expression analysis by Western blot showed that RPMs contained <50% of TGPM MFG-E8 levels (Fig. 1B). Similarly, their ability to phagocytose apoptotic cells was reduced by 30% (Fig. 1C and D).

To investigate the changes of MFG-E8 expression in RPM after endotoxin challenge, we stimulated the cells *in vitro* with increasing doses of LPS and measured MFG-E8 expression using Western blotting. Ten ng/ml LPS was sufficient to significantly reduce MFG-E8 levels by 45% with no further reduction after stimulation with 100 ng/ml LPS (Fig. 2A). As expected, the ability to phagocytose apoptotic cells further decreased in RPMs and was consistently lower than that of TGPM under the same conditions (Fig. 2B).

To demonstrate the difference in cytokine response in TGPM and RPM, we measured TNF-α levels released by these cells 4 h after the stimulation with LPS. While TGPM were responsive to LPS, RPMs showed a 5-fold higher responsiveness to LPS (Fig. 3A). It has been previously demonstrated that apoptotic cell phagocytosis suppresses cytokine responsiveness in phagocytosing macrophages (12). To determine differences in RPM and TGPM-responsiveness after the phagocytosis of apoptotic cells, we measured the respective TNF-α levels. We showed here that the TNF-α release was suppressed in TGPM, here by 40%, and 32% after the coincubation with apoptotic thymocytes and stimulation with 10 and 100 ng/ml LPS, respectively (Fig. 3B). In RPM, however, this suppression was not seen (Fig. 3C).

Since this finding was very useful and novel, we sought to determine intracellular pathways involved in this non-responsiveness to apoptotic-cell mediated TNF-α suppression. We first investigated differences in MAP kinase activation in RPM and TGPM. TGPM showed a robust activation of the three major MAP kinases ERK1/2, p38 and JNK as demonstrated by their phosphorylation (Fig. 4A). By comparison, RPMs showed a stronger increase in ERK1/2, p38 and JNK activation 15 min after LPS stimulation, supporting the increased TNF-α responsiveness of these cells. We also explored whether MFG-E8 helps to suppress the pro-inflammatory pathway within RPMs. We hence incubated the macrophages with apoptotic cells, stimulated them with LPS and examined the activation of MAP kinase and NFκB pathways after the exogenous addition of recombinant MFG-E8 (rMFG-E8). While apoptotic cells alone had no effect on these pathways, the addition of rMFG-E8 to apoptotic cells prior to phagocytosis and LPS stimulation had a marked suppressive effect on each of the investigated pathways,

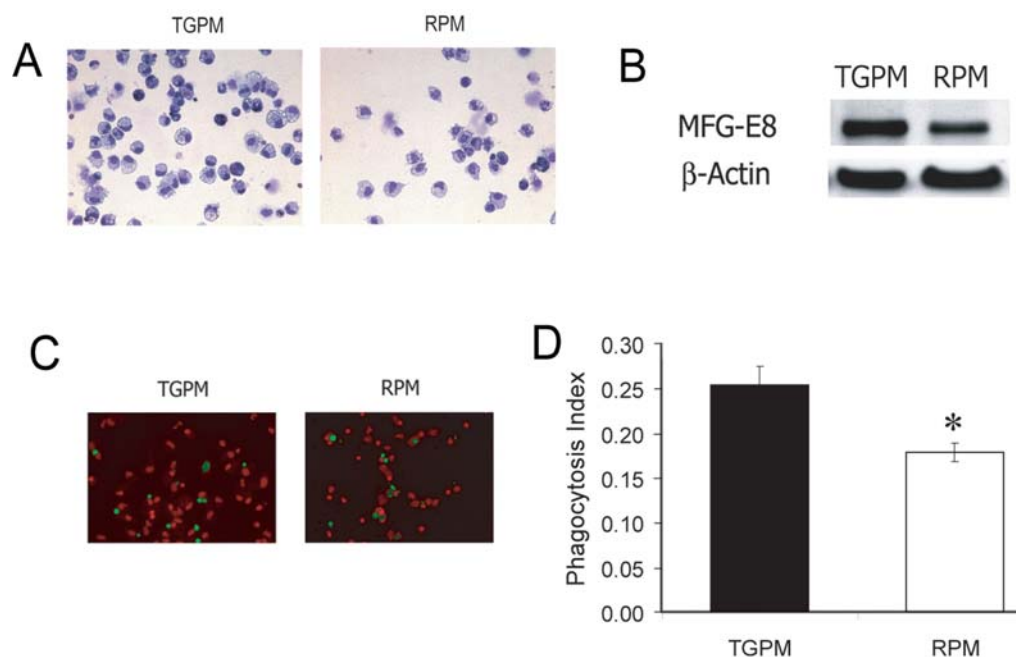


Figure 1. Mature resident peritoneal macrophages (RPM) contain less MFG-E8 than the immature thioglycolate-elicited (0.3% for 72 h) peritoneal (exsudate) macrophages (TGPM) resulting in a reduced efficacy in clearing apoptotic cells. (A) Giemsa-staining of freshly isolated peritoneal macrophages. (B) MFG-E8 Western blot of whole cell lysates. (C) Phagocytosis assay; macrophages were cultured on a chamber slide, fed with 4 apoptotic thymocytes for 90 min, fixed and stained using TUNEL. Red cells depict macrophages and green engulfed apoptotic cells. (D) Quantification of phagocytosis index by calculation engulfed apoptotic cells/macrophages ratio. \* $P < 0.05$  vs. TGPM, t-test,  $n = 6$ .

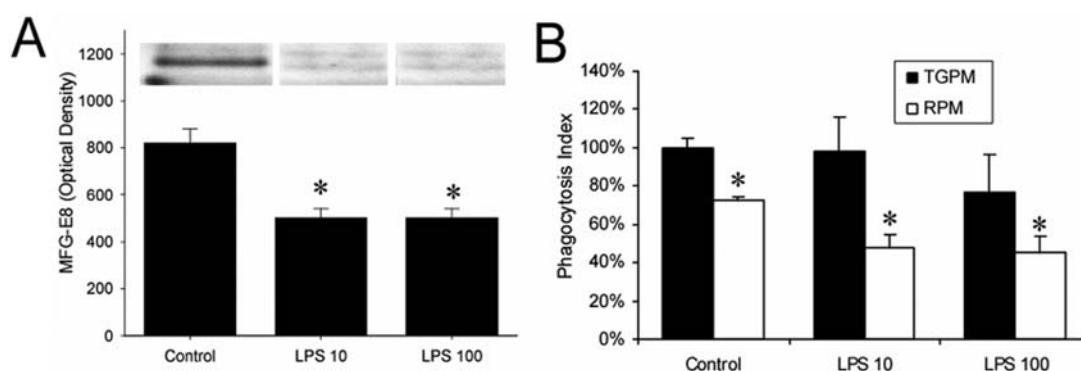


Figure 2. LPS down-regulates MFG-E8 in RPM and further deteriorates apoptotic cell clearance by RPMs compared with TGPMs. (A) RPMs were stimulated with 10 and 100 ng/ml LPS for 20 h, lysed and protein was used for Western blotting to detect MFG-E8. \* $P < 0.05$ , ANOVA using SNK test,  $n = 3$ . (B) TGPMs and RPMs were cultured with or without LPS for 20 h prior to an efferocytosis assay. Index was normalized to 100% for non-stimulated TGPM. \* $P < 0.05$  vs. TGPM, ANOVA using SNK test,  $n = 6$ .

particularly on the p38 and NF $\kappa$ B pathways that play a key role in the cytokine response of macrophages (Fig. 4B).

## Discussion

In our current study, we demonstrated that TGPMs have higher levels of MFG-E8 and are superior at phagocytosis of apoptotic cells. This leads to an effective suppression of post-phagocytic TNF- $\alpha$  release in these macrophages. While RPMs show a stronger activation of proinflammatory intracellular pathways after LPS-stimulation, MFG-E8-mediated apoptotic cell phagocytosis suppresses various

ERK1/2, p38, JNK, and NF $\kappa$ B activation, resulting in a lower TNF- $\alpha$  release (Fig. 5).

Macrophages are derived from a myeloid lineage through the differentiation of monoblasts and monocytes that undergo their final differentiation in different tissues, eventually becoming microglia, Kupffer cells, alveolar macrophages or osteoclasts, in the brain, liver, lungs and bone. In these tissues, they assume a central role in the recognition and activation phases of adaptive immune responses. The peritoneal cavity contains a similar cell population of tissue-fixed macrophages of which  $\sim 10^7$  cells can be retrieved by peritoneal lavage in a rat and  $\sim 10^6$  in a mouse. Over 25 years

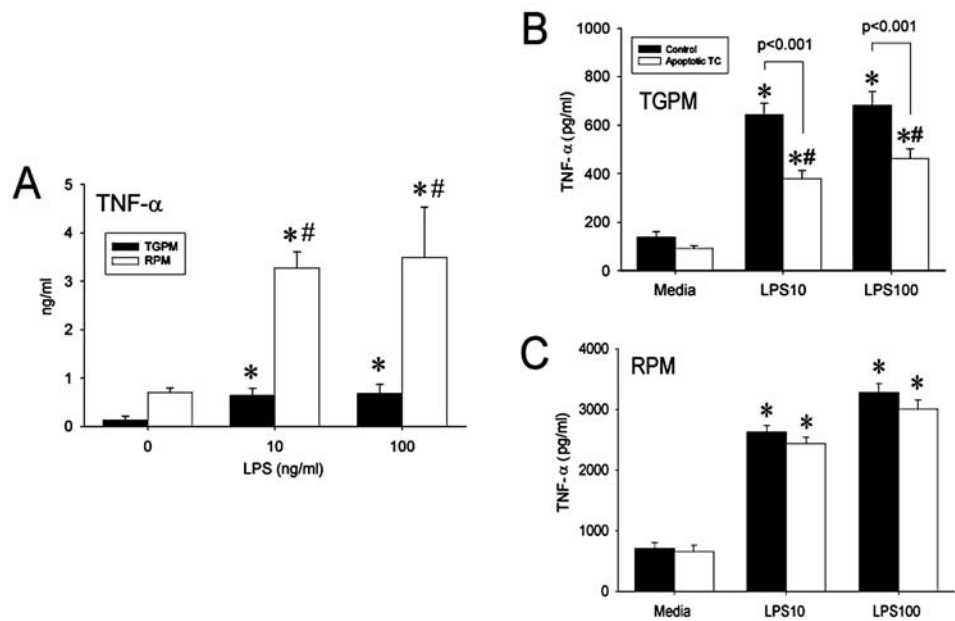


Figure 3. TGPM are less responsive to LPS-stimulated TNF- $\alpha$  release than RPM, which is further suppressed after apoptotic cell phagocytosis. (A) Macrophages were stimulated with LPS for 4 h and TNF- $\alpha$  release was assessed by measuring concentrations in the supernatant by ELISA. \* $P < 0.05$  vs. 0 ng/ml LPS, # $P < 0.05$  vs. TGPM, two-way ANOVA using SNK test,  $n = 10$ . (B and C) Macrophages were co-cultured for 90 min with apoptotic thymocytes. Non-phagocytosed apoptotic cells were removed by thorough washing followed by stimulation of the macrophages with LPS for 4 h. TNF- $\alpha$  released from TGPM (B) and RPM (C) was measured by ELISA. \* $P < 0.05$  vs. medium, # $P < 0.05$  vs. control, two-way ANOVA using SNK test,  $n = 10$ .

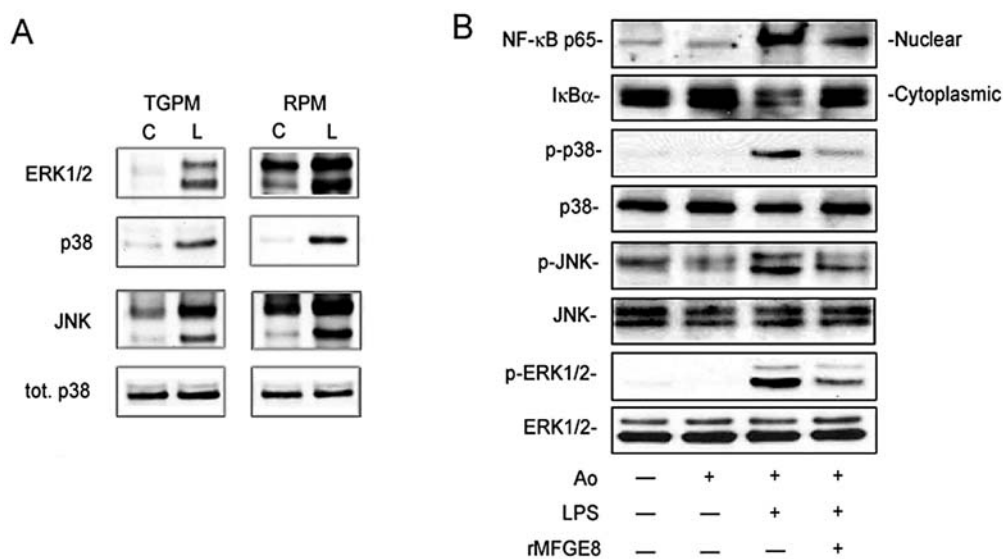


Figure 4. RPMs display a stronger MAPK activation by LPS (A) and apoptotic cell phagocytosis blocks multiple MAPK and the NF- $\kappa$ B pathways in macrophages (B). Macrophages were incubated with apoptotic cells for 90 min followed by thorough washing and stimulation with 100 ng/ml LPS for 15 min. Cells were immediately lysed total or separated into cytoplasmic and nuclear fractions, followed by Western blotting for NF- $\kappa$ B (nuclear), I $\kappa$ B $\alpha$  (cytoplasmic), phospho-ERK1/2, phospho-p38 and phospho-JNK. Total MAPK proteins served as internal controls. This figure is representative of three separate experiments with similar results. C, control; L, LPS.

ago researchers were interested in their development and discovered that injection of thioglycolate into the peritoneal cavity leads to an influx of immature monocyte-like (exudate) macrophages that undergo differentiation into mature resident tissue macrophages. This occurs over the course of 3-7 days after thioglycolate stimulation which can be

observed through morphological and biochemical changes in these cells (1). The exact mechanism of this influx of cells by an over 70-year old chemical, originally used for depilation and for permanent waves, is still poorly understood, although the involvement of advanced glycation end products (AGEs) are discussed (13,14). AGE has been reported to activate



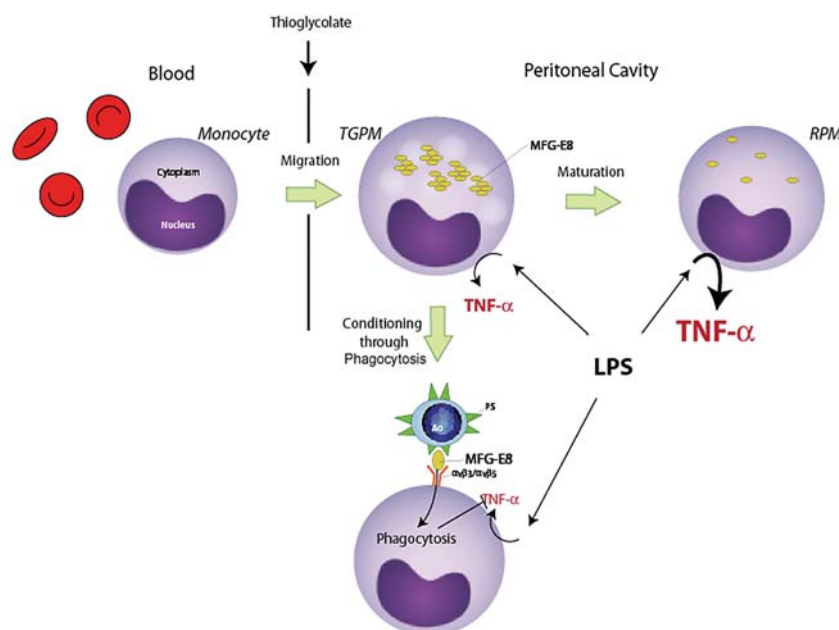


Figure 5. MFG-E8 and apoptotic cell clearance in immature and mature peritoneal macrophages. Thioglycolate recruits immature TGPMs to the peritoneal cavity that contain abundant MFG-E8. RPMs by comparison have less MFG-E8 and are more responsive to LPS stimulation. The apoptotic cell phagocytosis by TGPMs further suppresses LPS-mediated TNF- $\alpha$  release in these cells.

specific receptors, RAGE that induce proinflammatory responses through NF $\kappa$ B activation (15-17). Thus, AGEs themselves have been shown to elicit opposite effects from those have shown here, especially the increase of TNF- $\alpha$  release and suppression of phagocytic ability (18,19), indicating that AGEs may not be the crucial chemical component in the thioglycolate containing Brewer's medium.

Macrophages play a key role in the innate response and in the orchestration of an adaptive immune response, by releasing proinflammatory mediators that recruit other immune cells to the site of infection and by processing and presenting antigen to lymphocytes eventually mediating a specific host response. It appears, however, that these roles are not performed by a single phenotypic cell population but that these macrophages are, in a sense, specialized to perform certain tasks. For example, it was recognized that TGPMs were less effective in bacterial phagocytosis and killing, while RPMs were very effective to do so (3,20). Effective killing involves the activation of macrophages and bystander immune cells, which is associated with the release of proinflammatory cytokines such as IFN- $\gamma$  and TNF- $\alpha$ . We have demonstrated here that RPMs release five times the amount of TNF- $\alpha$  than the newly recruited and presumably more immature TGPMs. Furthermore, we have shown that the ability to clear apoptotic cells also differs between these two cell populations, with the more immature TGPMs being the major players in this respect. This effect is strongly associated with the expression levels of MFG-E8, a factor that may crucially influence the phagocytic ability of macrophages as previous reports have indicated (8,21). Even the exogenous administration of MFG-E8 is able to enhance the phagocytosis of apoptotic cells by RPMs as has been previously shown (8,21). In this respect, the role of MFG-E8 may be two-fold. It increases the clearance of apoptotic cells

and suppresses a proinflammatory response at the same time. We have shown that while TGPMs (abundant in MFG-E8) are very effective in this function, RPMs are not, which also reflects in the inability to suppress a proinflammatory response of RPMs after apoptotic cell exposure. MFG-E8-enhanced phagocytosis of apoptotic cells attenuates, as we have shown here, crucial intracellular pathways involved in the proinflammatory response of macrophages, such as p38 and NF $\kappa$ B. Immunosuppressive cytokines, such as TGF- $\beta$  and IL-10, released by the (non-phlogistic) phagocytic macrophages may play an additional role in this mechanism (8,12).

The impact of the difference in functionality is evident in acute inflammatory conditions such as sepsis. There is an entire dynamic orchestration of cell responses that regulate the host defense to intruding microbes. Resident macrophages wear off the pathogens by phagocytosing them and mounting an effective immune response, while newly migrating monocytic macrophages appear days later to 'clean up' i.e., remove cell debris and apoptotic cells. In this later phase an inflammatory response is inefficient and probably even detrimental. In sepsis, for example, the initial surge in inflammatory mediators does not subside but continues to stimulate immune responses that lead to self-inflicted damage, which is one of the reasons responsible for organ failure and morbidity in septic animals and possibly patients (22). We have previously shown that MFG-E8 is down-regulated in septic rats, which seriously compromised their ability to clear apoptotic cells and resulted in an increased systemic inflammatory response and morbidity (11). This happens usually within 72 h of septic onset, at the time point when immature monocytic macrophages, rich in MFG-E8, have not yet been recruited to the site of inflammation. Understanding and controlling the kinetics and dynamics of

this cellular response may be a future goal to attenuate an overzealous inflammatory response found in sepsis and chronic inflammatory diseases such as auto-immunity.

Our results provided new insight into the role of different macrophage subtypes in apoptotic cell clearance and a mechanism responsible for suppressed inflammation. We conclude that TGPMs, appearing to have an immature phenotype, play a major role in clearing apoptotic cells due to higher MFG-E8 levels, while RPM, which are tissue-fixed and mature, are responsible for a strong proinflammatory response.

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