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Inflammatory monocytes recruited after skeletal muscle injury switch into anti-inflammatory macrophages to support myogenesis

Ludovic ARNOLD¹, Adeline HENRY², Françoise PORON¹, Yasmine BABA-AMER¹, Nico van ROOIJEN³, Anne PLONQUET⁴, Romain K GHERARDI¹, Bénédicte CHAZAUD^{1,5}

¹ INSERM, Unité 841, IMRB, "Cell Interactions in the neuromuscular system" team, Créteil, F-94000, France; Université Paris 12 Val-de-Marne, Créteil, F-94000, France

² Plateforme cytométrie IM3, IFR10, Université Paris 12 Val-de-Marne, Créteil, F-94000, France

³ Vrije Universiteit, VUMC, Department of Molecular Cell Biology, Faculty of Medicine, Amsterdam, The Netherlands

⁴ AP-HP, Groupe Henri Mondor-Albert Chenevier, Service d'immunologie-Biologie, Créteil, F-94000, France

⁵ Corresponding author: Unité 841, Equipe 10, "Cellular Interactions in the neuromuscular system", Faculté de Médecine, 8 Rue du Général Sarrail, 94000 Créteil, France. Tel: 33-149813649, Fax: 33-149813642, email: benedicte.chazaud@creteil.inserm.fr

Abbreviations : 5-bromo-2-deoxy-uridine (BrdU); clodronate-encapsulated liposomes (clo-lip); cluster of differentiation (CD); dexamethasone (DEX); diphtheria toxin (DT); Diphtheria Toxin Receptor (DTR); latex beads (LX); macrophages (MPs); monocyte/MPs (MO/MPs); nitric oxide (NO); secretory leukocyte protease inhibitor (SLPI); tibialis anterior (TA)

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ABSTRACT

Macrophages are important for skeletal muscle regeneration in vivo and may exert beneficial effects on myogenic cell growth through mitogenic and anti-apoptotic activities in vitro. However, macrophages are highly versatile and may exert various, and even opposite, functions depending on their activation state. Here, we studied monocyte/macrophage phenotypes and functions during skeletal muscle repair. Selective labeling of circulating monocytes by latex beads in CX3CR1^{GFP/+} mice showed that injured muscle recruited only CX3CR1^{lo}/Ly-6C⁺ monocytes from blood that exhibited a non-dividing, F4/80^{lo}, proinflammatory profile. Then, within muscle, these cells switched their phenotype to become proliferating anti-inflammatory CX3CR1^{hi}/Ly-6C⁻ cells that further differentiated into F4/80^{hi} macrophages. In vitro, phagocytosis of muscle cell debris induced a switch of proinflammatory macrophages toward an anti-inflammatory phenotype releasing TGFβ1. In cocultures, inflammatory macrophages stimulated myogenic cell proliferation whereas anti-inflammatory macrophages exhibited differentiating activity, assessed by both myogenin expression and fusion into myotubes. Finally, depletion of circulating monocytes in CD11b-DTR mice at time of injury totally prevented muscle regeneration, whereas depletion of intramuscular F4/80^{hi} macrophages at later stages reduced diameter of regenerating fibers. In conclusion, injured skeletal muscle recruits monocytes exhibiting inflammatory profile that operate phagocytosis and rapidly convert into anti-inflammatory macrophages that stimulate myogenesis and fiber growth.

INTRODUCTION

Stromal cells are involved in organogenic processes by controlling parenchymal cell functions such as migration, proliferation, differentiation and programmed cell death (1). Among stromal cells, macrophages (MPs) play a major role in maintenance of tissue homeostasis (2). In addition to phagocytosis and antigen presentation, these cells play a supportive function in various tissues (2,3). However, MPs are renowned for their heterogeneity as reflected by the various specialized functions they adopt in different anatomical locations (2,4). Many of these activities appear to be opposing in nature: proinflammatory vs. anti-inflammatory, immunogenic vs. tolerogenic, and tissue-destructive vs. tissue repair activities (4).

Different activation states have been described in vitro for MPs, each being associated with a particular phenotype and function (2,5). Classical activation, obtained after LPS/IFN γ treatment, induces production of proinflammatory cytokines and reactive oxygen species. Alternative activation, obtained after IL-4 treatment, increases expression of the mannose receptor (CD206). Deactivation by IL-10 and glucocorticoids induces production of IL-10, TGF β 1 and expression of the scavenger receptor CD163. However, it is not known to what extent these distinct activation states exist in vivo and whether MP fate is determined for ever or whether it remains constantly malleable (2), although several in vitro studies have shown MP adaptation to changing microenvironment (4,6-8).

An additional degree of complexity is reached with the existence of several subsets of circulating monocytes. In both human and mouse, two main monocyte subsets are present in blood: the main CD14^{hi} CD16⁻ (CX3CR1^{lo}/Ly-6C^{hi} in mouse) monocyte subset that invades tissues during acute inflammation, and the CD14^{lo} CD16⁺ (CX3CR1^{hi}/Ly-6C^{lo} in mouse) monocyte subset, that is not -or in a less extent- recruited during inflammation (9-13) but is expanded in chronic diseases such as HIV infection, sepsis, tuberculosis or asthma (14). Both fate and function of this population remain unclear since, in mouse, it is present in

normal spleen and may correspond to MPs residing in normal tissues (9). However, there is growing evidence that they may result from maturation of Ly-6C^{hi} monocytes (11,15,16).

MPs are involved in skeletal muscle repair. Muscle damage induces massive MP infiltration at the injury site (17,18). Partial depletion of monocyte/MPs (MO/MPs) impairs muscle regeneration while reconstitution of bone marrow restores regeneration (19-23). Initially limited to phagocytosis of necrotic fibers, the pro-regenerative role of MPs was shown to involve a direct stromal support function (24,25). We have previously shown that monocyte-derived MPs enhance myogenic cell growth (26). MPs release mitogenic growth factors for myogenic precursor cells (mpcs) and establish cell-cell interactions that protect mpcs from apoptosis (26,27). However, little is known about both MO/MP subsets and phenotypes at work during muscle regeneration. Studies in rat have shown that ED1+ MPs are associated with muscle necrosis, whereas ED2+ (CD163) MPs, regarded as “resident MPs”, invade muscle once necrosis has been removed and are associated with regenerative myofibers (17,28). Although the nature of ED1 and ED2 MPs is not characterized, these results suggest that different MP subpopulations are associated with different stages of muscle repair.

In this study we used: *in vivo* tracing methods to analyze which monocyte subset was recruited after muscle injury; *ex-vivo* phenotyping analyses to characterize MO/MP profile during muscle regeneration; *in vitro* coculture experiments to identify which MP phenotype was associated with myogenesis; and sequential *in vivo* MO/MP depletion studies to evaluate the respective roles of both circulating and intramuscular MO/MPs in skeletal muscle repair.

RESULTS

Two MO/MP subsets are sequentially present during muscle regeneration. MO/MPs collected from skeletal muscle of CX3CR1^{GFP/+} mouse (9,15) were analyzed according to their GFP and Ly-6C expression during muscle regeneration. Normal muscle contained little CX3CR1^{hi}/Ly-6C⁻ MO/MPs (about 200 cells/mg of muscle). Notexin was injected in tibialis anterior (TA) muscle to induce the necrosis/regeneration process. As soon as 90 min after injury, CX3CR1^{lo}/Ly-6C⁺ MO/MPs were detected in muscle and rapidly invaded the tissue to reach a maximum point at 24 h post injury (25% of CD45⁺ cells). Then, CX3CR1^{lo}/Ly-6C⁺ MO/MP population quickly declined with time while CX3CR1^{hi}/Ly-6C⁻ MO/MPs appeared and continuously increased to reach a plateau at d 7 accounting for 35% of CD45⁺ cells extracted from muscle at this time point (Fig. 1A).

MO/MP percentage among CD45⁺ cells is a relative quantification that does not reflect the amount of MO/MPs actually present in muscle. Thus we calculated from these data the total number of extracted GFP⁺ cells per mg of muscle. Results indicated that the extent of the two MO/MP waves was different (Fig. 1B), and this was confirmed by histological examination (Fig. 1C). During the first 2 days, as injured myofibers became necrotic, 2000 to 3500 GFP⁺ cells were counted per mg of muscle. At 4 and 7 days, when myogenic cells proliferated and further differentiated (Fig. 1C), 5 to 6 fold more GFP⁺ MO/MPs were found in muscle (12000 to 16000 cells per mg) (Fig. 1B). Ten days after injury, when centrally nucleated fibers were visible throughout the injured area (Fig. 1C), the amount of GFP⁺ cells had dramatically declined to reach less than 2000 GFP⁺ cells per mg of tissue (Fig. 1B).

CX3CR1^{lo}/Ly-6C⁺ and CX3CR1^{hi}/Ly-6C⁻ MO/MP populations display distinct proliferative and inflammatory profiles. Between d 1 and d 4 post injury, the number of Ly-6C⁻ MO/MPs increased 20-fold while the number of Ly-6C⁺ MO/MPs increased 2-fold (Fig. 1B). We analyzed proliferation of both subsets that were isolated from regenerating muscle by cell sorting (Fig. 2A). Ki67 labeling showed that Ly-6C⁺ MO/MPs never proliferated

whereas Ly-6C⁻ MO/MPs partially entered into cell cycle from d 2, then the number of proliferating cells doubled at d 4 and further decreased by 2-fold at d 7 (Fig. 2B).

To compare the inflammatory profile of the two MO/MP populations, we analyzed cytokine expression of isolated Ly-6C⁻ and Ly-6C⁺ MO/MPs at d 4 post injury by RT-PCR. Results showed that Ly-6C⁺ MO/MPs expressed IL-1 β and, to a lesser extent, TNF α transcripts more strongly than Ly-6C⁻ MO/MPs (Fig. 2C). Conversely, Ly-6C⁻ MO/MPs expressed more strongly TGF β 1 and IL-10 than Ly-6C⁺ MO/MPs (Fig. 2C), indicating that Ly-6C⁺ MO/MPs had an inflammatory profile and Ly-6C⁻ MO/MPs had an anti-inflammatory profile.

Injured muscle only recruits CX3CR1^{lo}/Ly-6C⁺ MO/MPs that subsequently switch into anti-inflammatory CX3CR1^{hi}/Ly-6C⁻ MO/MPs and differentiate into MPs. We labeled each circulating subset with fluorescent latex beads (LX) before injury, as previously described (12,15). Labeling of monocytes does not alter their migratory capacities (12). After i.v. LX injection, circulating LX⁺ cells were CX3CR1^{hi}/Ly-6C⁻ monocytes (Fig. 3A upper panel, "blood"); LX⁺ cells were never detected into muscle after injury (Fig. 3A upper panel, "muscle"), indicating that Ly-6C⁻ MO/MPs appearing in muscle at d 2 were not recruited from blood. After liposome-encapsulated clodronate (clo-lip) injection followed by LX injection, circulating LX⁺ cells were CX3CR1^{lo}/Ly-6C⁺ monocytes, as previously described (15). 36 h after notexin injection, LX⁺ cells were still Ly-6C⁺ monocytes (95%) in the blood (Fig. 3A lower panel, "blood") while LX⁺ cells present within muscle were both Ly-6C⁺ and Ly-6C⁻ (38 ± 2 and 61 ± 3 %, respectively, on 3 experiments) (Fig. 3A right lower, "muscle"). These data indicate that muscle Ly-6C⁻ cells derived from early recruited Ly-6C⁺ MO/MPs.

Several markers have been associated with MP phenotype transition at time of resolution of inflammation. Among them, secretory leukocyte protease inhibitor (SLPI) is induced in MPs by proinflammatory signals to trigger down-regulation of proinflammatory cues (29-32). RT-PCR analysis showed that Ly-6C⁺ MO/MPs strongly expressed SLPI compared to Ly-6C⁻

MO/MPs (62% increase of band intensity on 3 experiments) (Fig. 3B) 4 days after injury. Inversely, PPAR γ is expressed by MPs that down-regulate inflammation (33-35). Our results showed that PPAR γ mRNA was strongly expressed by Ly-6C $^-$ compared to Ly-6C $^+$ MO/MPs (49% increase of band intensity on 3 experiments) (Fig. 3B).

Forward and side scatter of the two MO/MP subsets isolated at d 7 post injury showed that Ly-6C $^-$ MO/MPs had higher side scatter, and were larger than Ly-6C $^+$ MO/MPs (Fig. 3C), in accordance with a maturation of the Ly-6C $^-$ subset. Analysis of differentiation markers showed that Ly-6C $^-$ MO/MPs expressed CD11c and higher levels of F4/80 (Fig. 3C). Nearly all (> 90%) F4/80 $^+$ cells were CD11c $^+$ and I-A $^{b+}$, and no positivity was observed for the dendritic cell marker DEC-205 (data not shown). These results show that on the contrary to Ly-6C $^+$ MO/MPs, Ly-6C $^-$ MO/MPs exhibited high levels of MP differentiation markers.

Altogether, these results suggest that after muscle injury, inflammatory Ly-6C $^+$ monocytes are recruited from blood, convert into anti-inflammatory Ly-6C $^-$ MO/MPs that proliferate in situ and further differentiate into mature MPs.

Phagocytosis of muscle cell debris participates to MP phenotype transition. To study interactions between MPs and myogenic cells possibly involved in MP phenotype transition, we used in vitro human cell cocultures. Indeed, in vitro pro and anti-inflammatory MP activation states described in humans most likely correspond to Ly-6C $^+$ and Ly-6C $^-$ MO/MP profiles found in regenerating mouse muscle. Consistently, LPS/IFN γ treatment of human MPs induced TNF α (P<0.05) and IL-1 β (P<0.05) secretion (Fig. 4A) whereas dexamethasone (DEX)/IL-10 treatment induced IL-10 (P<0.05) secretion (Fig. 4A), as previously shown (36,37). Since differentiating muscle cells secrete IL-4 (38), a key inducer of alternative activation of MPs (5), we tested IL-4 treatment, that induced TGF β 1 secretion (Fig. 4A) as previously shown (39).

As a major role of MPs is to operate phagocytosis of dead cells, we evaluated the effect of phagocytosis of muscle cell debris on MP inflammatory profile. Upon phagocytosis of

necrotic mpcs, LPS/IFN γ treated MPs both decreased their TNF α secretion ($P<0.05$) and increased their TGF β 1 secretion ($P<0.03$) (Fig.4B). We attempted to inhibit phagocytosis of proinflammatory MPs by colchicine (40), cytochalasin D (41) and recombinant Annexin V (42). Colchicine failed to inhibit phagocytosis and MPs changed their cytokine secretion upon mpc debris phagocytosis, as in untreated cells (Fig. 4C). Conversely, phagocytosis was inhibited in the presence of cytochalasin D or recombinant Annexin V; in these conditions, TNF α and TGF β 1 secretion did not change in presence of mpc debris (Fig. 4C). These data indicate that phagocytosis of mpc debris induced a switch of MP phenotype toward an anti-inflammatory profile, as previously shown for other cell types (43-46).

We also evaluated whether interactions with living mpcs may influence MP phenotype. Neither addition of mpcs to previously activated MPs nor addition of mpcs prior to MP activation did significantly alter MP phenotype (data not shown). These data show that mpcs neither modified MP activation state nor prevented acquisition of any given MP phenotype.

Inflammatory MPs increase mpc growth whereas anti-inflammatory MPs stimulate their differentiation. In coculture experiments we analyzed mpc behavior depending on MP activation state. We have previously shown that untreated MPs stimulate mpc growth and proliferation (26). As shown in Fig. 5, mpc differentiation (Fig. 5C) and fusion (Fig. 5D) were also increased by untreated MPs ($P\leq 0.05$). LPS/IFN γ -treated MPs were more potent to enhance mpc growth (63%, $P<0.05$) (Fig. 5A), and accordingly, they enhanced mpc proliferation, as shown by increased 5-bromo-2-deoxy-uridine (BrdU) incorporation (25%, $P<0.05$) (Fig. 5B). Inversely, both myogenin expression (Fig. 5C) and myotube formation (Fig. 5D) were strongly reduced (58% and 68% respectively, $P<0.05$), indicating an inhibition of differentiation by inflammatory MPs. IL-4-treated MPs slightly reduced mpc proliferation (14%, $P<0.05$) (Fig. 5B) while they stimulated myogenin expression (28%, $P<0.05$) (Fig. 5C) and stimulated myotube formation (158%, $P<0.05$) (Fig. 5D) compared to untreated MPs. DEX/IL-10-treated MPs did not stimulate growth (Fig. 5A) and proliferation (Fig. 5B)

compared to untreated MPs, but presented a strong pro-differentiating activity, as assessed by increased myogenin expression (52%, $P < 0.05$) (Fig. 5C) and increased myotube formation (136%, $P < 0.01$) (Fig. 5D). These results strongly suggest that activation state of MPs may monitor the myogenic process.

MO/MPs presence is mandatory for skeletal muscle regeneration. We took advantage of the CD11b-diphtheria toxin receptor (DTR) mouse to study the respective roles of MO/MP subsets during muscle regeneration in vivo, i.m. injection of clo-lip being toxic for myogenic cells (unpublished data). In CD11b-DTR mice, monocytes - and possibly tissue MPs depending on the way of injection - are depleted after diphtheria toxin (DT) injection (47). After a single i.v. DT injection, circulating monocytes loss occurred from 6 h to reach a maximum at 12 h, that lasted 12 h more, with subsequent restoration and normalization at 48 h (Fig. 6A). At maximum of depletion, 75-80% of cells in the monocyte gate had disappeared (Fig. 6A) as did F4/80+, CD11b+ and Ly6C+ cells (data not shown). Time course of post injury muscle regeneration in untreated CD11b-DTR mice was similar to that of C57BL/6 mouse shown in figure 1 until d 4, but centrally nucleated myofibers appeared earlier in CD11b-DTR than in C57BL/6 (d 7 vs. d 10) (Fig. 6B). When DT was injected simultaneously to (Fig. 6C), or 12 h before (Fig. 6D) muscle injury, necrotic fibers were removed more slowly, and regeneration was impaired with appearance of adipose cells (Fig. 6C,D, Fig. 7). After double DT injection both 12 h before and simultaneously to notexin injection (that induced a monocyte depletion lasting 0-24 h post injury), almost no MO/MP infiltration was observed (Fig. 6E). None of the hallmarks of regeneration were detected and persistence of necrotic fibers was observed until d 7 (Fig. 6E, Fig. 7), although circulating monocytes reappeared in the circulation from d 2 post injury (data not shown). When DT was injected 4 days after notexin, no difference was observed compared to control (Fig. 6B, Fig. 7), indicating that circulating monocytes were no longer recruited into muscle at this time. Notably, i.v. injected DT did not significantly affect intramuscular MO/MPs, as shown by

unchanged expression of both F4/80 and CD11b by CD45⁺ cells isolated from muscle (Fig. 6F, insert).

We attempted to target intramuscular antiinflammatory MO/MPs, at a time when phagocytosis of necrotic myofibers is finished (Fig. 6B, Fig. 7A). To assess MP depletion in muscle, flow cytometric analysis of CD45⁺ cells was performed 24 h after a single i.m. DT injection. The total number of F4/80⁺ cells present in muscle was diminished by 25% after DT injection, especially that of F4/80^{hi} cells, by 75% (Fig. 8A) ($P < 0.05$), indicating that these differentiated FSC^{hi}/F4/80^{hi} MPs, corresponding to Ly-6C⁻ MPs, were targeted by DT. However, total number of CD45⁺ cells increased by 2 fold in DT vs. PBS injected muscle (Fig. 8A), indicating that i.m. DT injection induced a secondary inflammation accompanied by the recruitment of F4/80⁻ leukocytes (likely neutrophils and F4/80^{lo/-} monocytes). Consistently, histological examination 4-5 days later of whole TA muscle showed an area in which new cycles of regeneration were visible as assessed by the presence of necrotic and basophilic fibers (Fig. 8B, arrow). This area was restricted to the site of needle puncture, and was therefore excluded from the analysis to avoid counting of small regenerating myofibers coming from the secondary lesion (Fig. 8B). Diameter of centrally nucleated regenerating myofibers was evaluated in all the rest of TA and was found reduced by 41% in DT- vs. PBS-injected muscle (30.8 vs. 18.2 μm , respectively) (Fig. 8C) ($p < 0.001$).

DISCUSSION

In the present study, we have shown that: i) injured skeletal muscle selectively recruits Ly-6C⁺ circulating monocytes which exhibits an inflammatory profile; ii) within muscle, Ly-6C⁺ monocytes switch their phenotype to acquire an anti-inflammatory profile, phagocytosis of muscle cell debris being most likely involved in this transition; iii) as necrosis disappears from the injured area, anti-inflammatory Ly-6C⁻ MO/MPs proliferate and further differentiate into MPs; iv) in vitro, inflammatory MPs enhance myogenic cell proliferation and repress their differentiation, whereas anti-inflammatory MPs stimulate myogenesis and fusion; v) consistently, in vivo depletion of circulating monocytes at time of injury totally prevents muscle regeneration, whereas later depletion of intramuscular MPs decreases diameter of regenerating fibers.

It has been shown that the CX3CR1^{lo}/Ly-6C⁺ monocyte population is recruited in injured or inflamed tissues (9,13,48). Our experiments using LX labeled circulating monocytes showed that CX3CR1^{lo}/Ly-6C⁺ monocytes were selectively recruited rapidly after muscle injury. Mouse CX3CR1^{lo}/Ly-6C⁺ monocytes and their human analogs bear several chemokine receptors, including CCR2, CCR1, CCR4, CCR7, CXCR1 and CXCR2 (9). The corresponding chemokines are expressed by muscle tissue during the first days post injury (22,49,50) (cytokine expression profile is available at <http://pepr.cnmcresearch.org>). MCP-1 has been particularly involved in monocyte recruitment by injured muscle (19,20). We did not evidence recruitment of CX3CR1^{hi}/Ly-6C⁻ monocytes from blood to muscle. Moreover, transient depletion of circulating monocytes during the first 0-24 h after muscle injury showed no subsequent MP infiltration despite reconstitution of the pool of circulating monocytes. This suggests that CX3CR1^{lo}/Ly-6C⁺ MO/MP recruitment takes place early after injury, within the 2/3 first d post injury.

Our experiments showed that initially LX labeled CX3CR1^{lo}/Ly-6C⁺ monocytes converted into CX3CR1^{hi}/Ly-6C⁻ MO/MPs from d 2 after injury, indicating a phenotype transition of MO/MPs within muscle. Consistently, MP differentiation markers were weakly expressed by

CX3CR1^{lo}/Ly-6C⁺ MO/MPs and markedly expressed by CX3CR1^{hi}/Ly-6C⁻ MO/MPs (see below). We propose that CX3CR1^{lo}/Ly-6C⁺ cells, that exhibit a inflammatory phenotype (expressing TNF α and IL-1 β) progressively loose Ly-6C expression while converting into anti-inflammatory MO/MPs (expressing TGF β 1 and IL-10). These data are in keeping with the cytokine mRNA analysis of post injured muscle extract (51) (available at <http://pepr.cnmcresearch.org>): TNF α and IL-1 β expression peaks at d 2-3 post injury; IL-10 and TGF β 1 expression increases from d 2 post injury and is maintained throughout regeneration. MP phenotype conversion observed in skeletal muscle further supports previous in vitro evidence that MPs are capable of rapid adaptation to changing environments (4,6-8) and in vivo demonstration that MPs alter their phenotype according to their tissue environment (52).

MPs participate to both amplification of inflammation at time of injury and down-regulation of the inflammatory response to avoid excess tissue damage (31,53,54). MP stop-signals associated with resolution of inflammation include: i) induction of negative regulators of inflammation in MPs by environmental cues, and ii) non phlogistic phagocytosis. As shown herein, both likely operate during skeletal muscle regeneration. SLPI is considered as a brake on the response of MPs to inflammation (29,31). Its expression is induced by LPS and IL-6 and leads to inhibition of nitric oxide (NO) and TNF α production (30). In vivo, the highest SLPI synthesis occurs during the proinflammatory phase and thus anticipates increase of TGF β 1 and IL-10 production (32). Accordingly, we have shown that CX3CR1^{lo}/Ly-6C⁺ MO/MPs expressed more SLPI than CX3CR1^{hi}/Ly-6C⁻ cells. Another marker associated with the resolution of inflammation by MPs is PPAR- γ . Fatty acids and prostaglandin metabolites bind to PPAR- γ in activated MPs, which triggers inhibition of both inducible NO synthase and release of proinflammatory cytokines (33,35), thus conferring anti-inflammatory properties to PPAR- γ (34). Here we showed that antiinflammatory CX3CR1^{hi}/Ly-6C⁻ MO/MPs strongly expressed PPAR- γ . Non phlogistic phagocytosis by MPs is another signal for switching to healing (31). Binding and phagocytosis of apoptotic cells inhibit secretion of proinflammatory

mediators and stimulate secretion of TGF β 1 and IL-10 by inflammatory MPs (43-46). Induction of an anti-inflammatory process by phagocytosis of necrotic cells depends on the cell type ingested (42,43,55). We showed that both unstimulated and inflammatory MPs adopted an anti-inflammatory profile upon phagocytosis of necrotic muscle cell debris and that inhibition of MP phagocytosis prevented this phenotype switch. Altogether, these results strongly suggest that recruited inflammatory CX3CR1^{lo}/Ly-6C⁺ MO/MPs convert into anti-inflammatory CX3CR1^{hi}/Ly-6C⁻ MO/MPs upon both ingestion of muscle cell debris and expression of stop signals extinguishing inflammation.

Our results showed that once switched, CX3CR1^{hi}/Ly-6C⁻ MO/MPs actively proliferated, therefore contributing to the large amounts of MO/MPs observed in muscle at d 4 to d 7 post injury. In vitro, MP proliferation is inhibited by inflammatory mediators (LPS, IFN γ , NO, IL-6) and is stimulated by TGF β 1 (56-58). Thus, proliferation of recruited MO/MPs may be due to changes in their environment, although as yet unknown mechanisms related to the change of phenotype cannot be excluded. Later during muscle regeneration (d 7), CX3CR1^{hi}/Ly-6C⁻ MO/MP proliferation slowed down together with the fully differentiation into MPs, as shown by their strong F4/80 expression (59). Almost all F4/80⁺ cells were also positive for CD11c and MHCII, as previously observed after inflammation in various organs (60,61), and were DEC-205⁻. In the blood, Ly-6C⁻ monocytes, that come from Ly-6C⁺ cells, are also CD11c⁺ (12). Therefore, appearance of CD11c⁺ cells in the muscle may be a recapitulation of conversion from Ly-6C⁺/CD11c⁻ to Ly-6C⁻/CD11c⁺ monocytes within the tissue. Thus, proliferating CX3CR1^{hi}/Ly-6C⁻ MO/MPs and then differentiating CX3CR1^{hi}/Ly-6C⁻ MPs are associated with the regenerative phase of muscle repair.

Depending on the context, MPs may have supportive or deleterious effects on cells: in chronic diseases, including those affecting skeletal muscle, MPs are deleterious (62,63) while they support tissue repair in muscle and other various tissues including liver, brain, peripheral nerve and epithelium (19-22,64-67). If one except regulation of inflammation, studies documenting a direct role of MPs on cell behavior are scant and include intestinal

progenitor proliferation (65), erythroblast proliferation and maturation (68), oligodendrocytic differentiation and myelination (69). In vitro studies have shown that MP activation state may direct neural progenitor differentiation towards either neurogenesis or oligodendrogenesis (70). Our results substantiate the view that MP function may be related to an activation state. In coculture experiments, inflammatory MPs -the counterpart of inflammatory CX3CR1^{lo}/Ly-6C⁺ cells observed in vivo- stimulated mpc proliferation and inhibited their differentiation. Inversely, IL-4 treated and anti-inflammatory MPs exhibited a strong differentiating activity on mpc, assessed by both stimulation of myogenic program and increase of fusion into multinucleated cells. Molecular mechanisms involved in these processes are currently under investigation. It is likely that cytokines released by activated MPs influence myogenic cell behavior: TNF α is mitogenic for myoblasts and inhibits their differentiation (71,72); IL-1 β impairs myogenic differentiation through IGF-1 (73); in vitro effects of TGF β 1 are more controversial, although in vivo neutralization of TGF β 1 in regenerating muscle was shown to reduce the diameter of regenerating myofibers (74). Beyond cytokines, COX2 and its metabolites may also play a role, as they were shown to be fusogenic and necessary for good muscle repair (75-77).

Previous in vivo studies using different injury models have shown that partial reduction of MO/MP entry into injured muscle hinders muscle regeneration, as shown by a delay in the appearance of regenerating myofibers and the persistence of intermuscular adipocytes (19-22). We observed similar features upon partial inhibition of monocyte recruitment in toxic-induced muscle regeneration. Moreover, when nearly no MO/MP was allowed to enter damaged muscle, myofibers remained in the necrotic state, pointing out the indispensable role of MPs in muscle repair. Interestingly, despite normalized level of circulating monocytes from d 2 post injury, no delayed recruitment was observed suggesting that signals governing monocyte recruitment are only transiently expressed by the damaged tissue, as it was recently documented in ultraviolet-injured skin (48). Accordingly, depletion of circulating monocytes at d 4 post injury had no effect on muscle regeneration. On the other hand,

depletion of intramuscular anti-inflammatory F4/80^{hi} MO/MPs at time of regeneration, i.e. once phagocytosis of necrotic myofibers is achieved, reduced the diameter of the centrally nucleated regenerating myofibers, indicating that these MO/MPs were involved in fiber growth, in accordance with our in vitro data.

In conclusion, our data evidence a phenotype transition operated by recruited MO/MPs during resolution of inflammation and tissue repair, that is associated to changes in their functions. We propose sequential involvements of these two phenotypically distinct MP populations during skeletal muscle repair as follows: within the first 24-48 h after injury, skeletal muscle recruits inflammatory circulating monocytes, that stimulate myogenic cell proliferation and prevent their differentiation. While they are exposed to inflammatory environment and operate phagocytosis of muscle cell debris, inflammatory MO/MPs convert into anti-inflammatory MO/MPs. As phagocytosis of necrotic myofibers is finished, these anti-inflammatory MO/MPs actively proliferate and further differentiate into anti-inflammatory MPs, that sustain myogenic differentiation and myofiber growth, in addition to their protective effect on differentiating myotubes (27) and their effect on fiber membrane repair (78).

MATERIALS AND METHODS

Animals. C57BL/6, CX3CR1^{gfp/+} (9) and CD11b-DTR (47) mice were bred and used according to french legislation. Experiments were run at 4-8 weeks of age.

Muscle injury and muscle preparation. Notexin (10 μ l, 25 μ g/ml in PBS, Latoxan, Valence, France) was injected in the TA. For histological analysis, muscles were prepared as previously described (27). Quantitative analysis of muscle regeneration was performed on the whole injured area: about 7 fields (x20 objective) were analyzed in each mouse, representing 300-400 fibers per mouse. Myofiber diameter was evaluated after collagen IV immunolabeling (see below) on about 7 fields (x20 objective) in each mouse. The small diameter of only centrally nucleated myofibers was evaluated in late regenerating muscle (non hachured area in Fig. 8B) with Axiovision 4.6 software (Carl Zeiss SAS, LePecq, France), representing 250-350 fibers per mouse. In PBS-injected mice, the punctured fascicule was omitted from analysis.

Isolation of MO/MPs from muscle. Fascia of the TA was removed. Muscles were dissociated in DMEM containing collagenase B 0.2% (Roche Diagnostics GmbH, Mannheim, Germany) and trypsin-EDTA 0.2% at 37°C for 45 min twice, filtered and counted. CD45⁺ cells were isolated using magnetic sorting (Miltenyi Biotec GmbH, Bergisch Gladbach, Germany) and stained with PE- or PC5- conjugated anti-Gr1 Ab (reacts with Ly-6C and Ly-6G, but only Ly-6C is expressed by monocytes) (eBiosciences, San Diego, CA). Cells were sorted using an Epics Elite cell sorter (Beckman-Coulter, Fullerton, CA). Populations presenting over 90% of purity were used. In some experiments, PE-conjugated CD11c, I-A^b (BD Biosciences) and F4/80 (AbD Serotec, Oxford, UK) Ab were used. Analysis was performed with a FACSCalibur cytometer (BD Biosciences).

Labeling of blood monocytes. Labeling of circulating monocytes was performed exactly as described (15) with plain microspheres (Fluoresbrite polychromatic red 0.5- μ m microspheres 2.5% solids; Polysciences Inc., Warrington, PA) and clo-lip, that were prepared as previously described (79). Clodronate was a kind gift of Roche Diagnostics GmbH.

Depletion of circulating monocytes and intramuscular MPs. DT was i.v. injected into CD11b-DTR mice (12 ng/g). Blood was retroorbitally harvested at various times post-injection and cells were labelled with anti-CD11b, anti-Gr1, F4/80 Ab and analyzed by flow cytometry. Due to high interindividual variations (3 to 12% of PBMC), the number of circulating monocytes in control mice was normalized to 5% of PBMC for each series (mean calculated > 25 mice). To deplete infiltrated MO/MP, DT was i.m. injected (25 ng/g in less than 10 μ l) at d 5 and 6 post injury. Controls included i.v. injection of PBS.

Immunolabeling. Muscle slides were incubated with anti-collagen IV Ab (1/50, Chemicon International, Inc., Temecula, CA) revealed with Cy5-conjugated anti-rabbit Ab. Murine sorted cells were centrifugated on slides and labeled with anti-Ki67 Ab (1 μ g/ml) (Abcam plc, Cambridge, UK), revealed with FITC-conjugated anti-rabbit Ab. Cultured human mpcs were incubated with anti-desmin Ab (60 μ g/ml, Abcam) revealed by a Cy3-conjugated anti-rabbit Ab and with anti-myogenin Ab (10 μ g/ml, BD Biosciences) revealed by a biotinylated anti-mouse (Vector Laboratories Inc, Burlingame, CA) and by DTAF-streptavidin (Beckman Coulter). Controls included incubation with whole rabbit or mouse IgGs. Other secondary antibodies and IgGs were from Jackson ImmunoResearch Laboratories, Inc. West Grove, PA.

RT-PCR. Total RNA were prepared from sorted cells using the RNeasy mini kit (Qiagen GmbH, Hilden, Germany). 0.5 μ g of total RNA was reverse transcribed using Superscript II reverse transcriptase and then amplified with a platinum Taq DNA polymerase (Invitrogen Ltd, Paisley, UK) and specific primers (sens and antisens, respectively): β 2microglobulin: 5'-CAGTTCCACCCG-CCTCAC-3', 5'-CACATGTCTCGATCCCAG-3'; TNF α : 5'-TTCCAGATTCTTCCCTGAGGT-3', 5'-TAAGCAAAGAGGAGGCAACA-3'; IL-1 β : 5'-TGACGTTCCATTAGACAAGT-3', 5'-CCGTCTTTCATTACACAGGACA-3'; TGF β 1: 5'-GAGACGGAATACAGGGCTTTC-3', 5'-TCTCTGTGGAGCTGAAGCAAT-3'; IL-10: 5'-ACCAGCTGGACAACATACTGC-3', 5'-TCACTCTTCACCTGCTCCACT-3'; SLPI: 5'-CCTTAAGCTTGAGAAGCCACA-3', 5'-AGCACTTGTATTTGCCGTCAC; PPAR γ : 5'-

AAGAGCTGACCCAATGGTTG-3', 5'-GGATCCGGCAGTTAAGATCA-3'. Amplification was performed at 94, 60 and 72°C for 1 min each step for 30 cycles. Amplification products (10 µl) were subjected to electrophoresis on 2% agarose gel containing ethidium bromide for visualization. Quantification was performed by using Scion Image software.

Human mpc culture. Culture media components were from Invitrogen. Human mpcs were cultured from muscle samples as previously described (26).

Human MP cell culture. MPs were differentiated from monocytes isolated from human blood as previously described, at time of corrective surgery in conformity with french legislation (Code de la Santé Publique, livre II) (26) . Differentiated MPs were further cultured in either RPMI medium containing 15% FBS or advanced RPMI medium containing 0.5% FBS (supplemented with sodium pyruvate 1%, Hepes 10 mM, βmercaptoethanol 50 µM, non essential amino acids 1%, vitamins 100X 1%). MPs were treated, or not, for 48 h with either LPS (1 µg/ml, Sigma-Aldrich, St. Louis, MO) and IFN γ (10 ng/ml, PeproTech Inc., Rocky Hill, NJ), or IL-4 (10 ng/ml, PeproTech) or DEX (80 ng/ml, Sigma-Aldrich) and IL-10 (10 ng/ml, PeproTech).

Phagocytosis. Mpc necrosis was induced by H₂O treatment for 1 h at 37°C. 100% of cells were propidium iodide positive. Necrotic mpcs were seeded on MPs (5 dead mpcs for 1 MP) for 3 h at 37°C. MP cultures were washed 3 times to remove non-ingested material and further cultured in serum-free advanced RPMI supplemented medium during 24 h to make conditioned media. In some experiments, cells were treated, as previously described, with colchicine (10 µg/ml, Sigma) (40), cytochalasin D (1 µg/ml, Sigma) (41) or recombinant annexin V (40 µg/ml, BD Biosciences) (42). Phagocytosis was quantified in the same conditions after incubation with fluorescent microspheres (same as above). The number of LX⁺ cells was quantified under an inverted microscope and expressed as percentage of total cells.

Co-cultures. Mpcs were plated on previously prepared MP cultures (3:1 [MP:mpc] ratio) in advanced RPMI supplemented medium, except in some experiments in which MPs and mpcs were seeded together before MP treatment was applied as described above.

Mpc behavior. Mpc growth was evaluated as described before (26). Mpc proliferation was estimated by BrdU incorporation (Roche Diagnostics). Mpc differentiation was evaluated by counting the number of myogenin⁺ cells among desmin⁺ cells. Mpc fusion was estimated by counting the number of nuclei per myotube.

Statistical analyses. All experiments were performed using at least 3 different cultures or animals in independent experiments. The student t test was used for statistical analyses. P <0.05 was considered significant.

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FIGURE LEGENDS

Figure 1: Kinetics of MO/MP subsets during muscle regeneration in CX3CR1^{GFP/+} mouse. (A) CD45⁺ cells present into injured muscle were analyzed for Ly-6C and GFP (CX3CR1) expression by flow cytometry. Top: Results are expressed in percentage of CD45⁺ cells isolated from muscle and are means \pm SD of 3 experiments. Bottom: representative examples of FACS analysis at each time point. (B) Total number of MO/MPs was calculated from results obtained in (A) and the number of isolated CD45⁺ cells plotted to muscle weight. Results are means \pm SD of 3 experiments. (C) Representative hematoxylin-eosin staining of muscle sections at various times after notexin injection. Bar = 50 μ m.

Figure 2: Phenotype of Ly-6C⁺ and Ly-6C⁻ MO/MPs during muscle regeneration. Ly-6C⁺ and Ly-6C⁻ MO/MP subsets were isolated by cell sorting at various times post injury. (A) Representative example of cell sorting of Ly-6C⁻ (middle plot) and Ly-6C⁺ (right plot) cells from whole MO/MPs (left plot) at d 4 post injury. (B) Ki67 immunostaining is expressed in percentage of isolated MO/MPs. (C) Expression of TNF α , IL-1 β , TGF β 1 and IL-10 was analyzed by RT-PCR in isolated populations at d 4 post injury. Corresponding band intensities are given as means of 3 experiments.

Figure 3: Fate of Ly-6C⁺ and Ly-6C⁻ MO/MPs during muscle regeneration. (A) Circulating Ly-6C⁻ (upper panel) and Ly-6C⁺ (lower panel) monocytes were labeled with LX red microspheres and analyzed by flow cytometry. LX⁺ CD45⁺ GFP⁺ cells (gate in "all cells" plots) were analyzed for Ly-6C expression ("LX⁺ cells" plots) in both blood and muscle at d 3 (for Ly-6C⁻ labeling) and d 2 (for Ly-6C⁺ labeling) post injury. Results are representative of 3 experiments. (B) SLPI and PPAR γ expression was analyzed by RT-PCR in Ly-6C⁺ and Ly-6C⁻ MO/MPs sorted at d 4 post injury. (C) Flow cytometry analysis of FSC/SSC characteristics, F4/80 and CD11c expression of Ly-6C⁺ (blue label) and Ly-6C⁻ (red label)

MO/MPs sorted at d 7 post injury (black dotted line: isotypic control). Results are representative of 3 experiments.

Figure 4: Phagocytosis and cytokine secretion by activated MPs. Cytokine secretion was evaluated by ELISA in MP-conditioned medium. **(A)** Cytokine secretion by untreated (NT), LPS/IFN γ , IL-4 and DEX/IL-10 treated MPs. **(B)** Cytokine secretion by untreated (NT) and LPS/IFN γ treated MPs after phagocytosis of muscle cell debris. **(C)** Left: phagocytosis of LX by LPS/IFN γ treated MPs incubated or not (none) with colchicine (colc) cytochalasin D (cyt D) or recombinant annexin V (ann.V). Right: cytokine secretion by LPS/IFN γ treated MPs after phagocytosis of muscle cell debris in the presence of the same effectors. Results are means \pm SEM of 3 experiments.

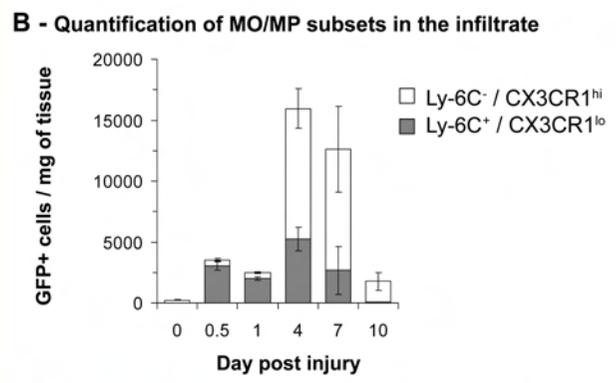
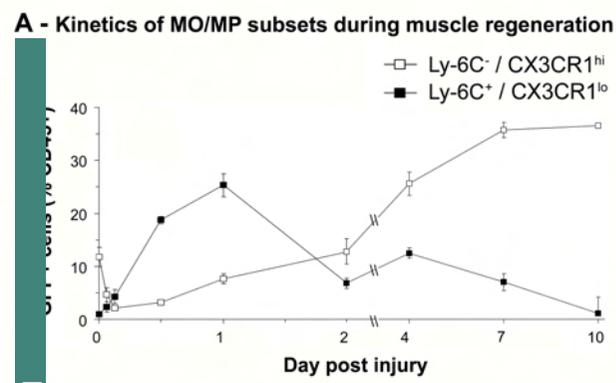
Figure 5: Effects of activated MPs on mpc fate. Mpc were cocultured with untreated (NT), LPS/IFN γ , IL-4 and DEX/IL-10 treated MPs and further analyzed for their: **(A)** growth, **(B)** proliferation, **(C)** differentiation and **(D)** fusion. All parameters were analyzed at d 3 of coculture except for BrdU incorporation that was monitored during 24 h of coculture. Results are means \pm SEM of 3 experiments. Bars bearing different symbols were significantly different.

Figure 6: Effects of circulating monocyte depletion on skeletal muscle regeneration. **(A)** Left: kinetics of circulating monocyte depletion after one i.v. DT injection, results are means \pm SD of at least 4 experiments. Right: examples of FACS analysis of blood before (-DT) and 18h after i.v. DT injection (+DT). **(B-F)** Notexin (Nx) was injected into TA of CD11b-DTR mice at d 0 and DT was injected i.v. at various times as indicated by the red arrows. Muscle histology was analyzed until d 7 after hematoxylin-eosin staining. Results are representative of at least 2 independent experiments. Bar = 50 μ m. Insert in (F): CD45 $^{+}$ cells

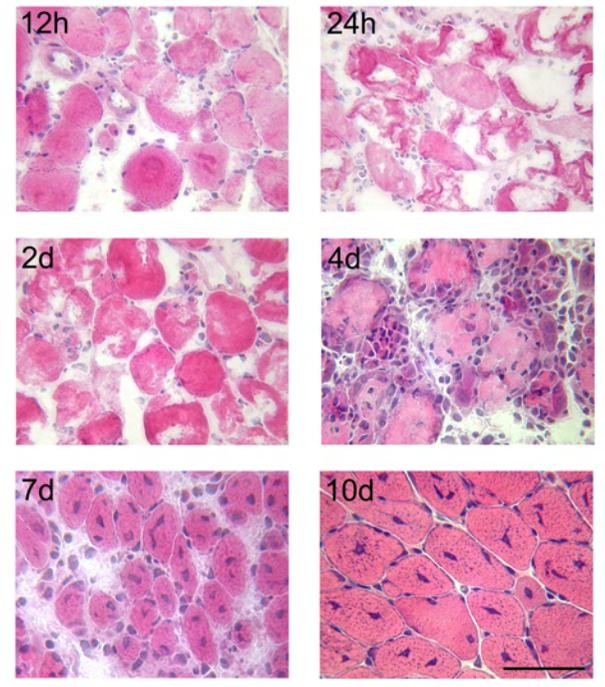
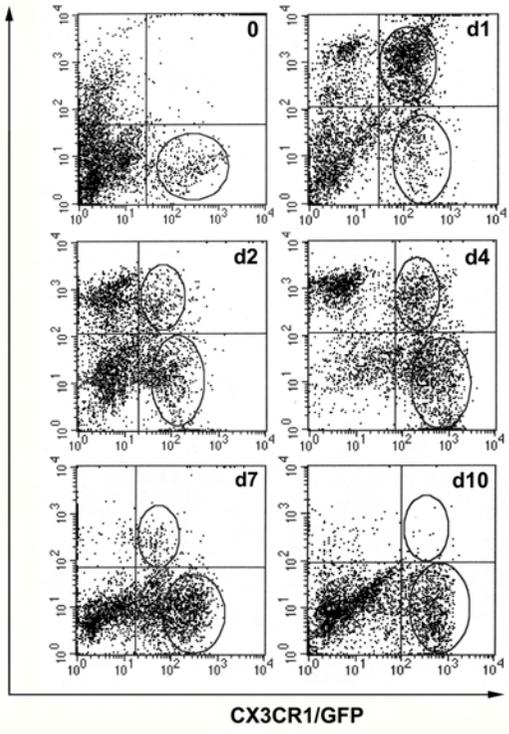
were isolated from muscle and analyzed for F4/80 and CD11b expression 24 h after an i.v. DT (blue line) or PBS (red line) injection at d 4 post injury (dotted line: isotypic control).

Figure 7: Quantitative analysis of muscle regeneration after circulating monocyte depletion. CD11b-DTR mice were treated as in figure 6 and in each case, the number of: **(A)** necrotic myofibers, **(B)** phagocytosed myofibers, **(C)** regenerating basic myofibers and **(D)** centrally nucleated regenerating myofibers was evaluated. **(closed squares)**: control corresponding to Fig. 6B; **(open diamonds)**: simultaneous DT and notexine injection corresponding to Fig. 6C; **(open triangles)**: DT injection 12 h before notexine injection corresponding to Fig. 6D; **(open squares)**: double DT injection corresponding to Fig. 6E; **(closed circles)**: DT injection 4 days after notexine injection corresponding to Fig. 6F. Results are expressed in percent of total counted myofibers and are means of at least 2 experiments.

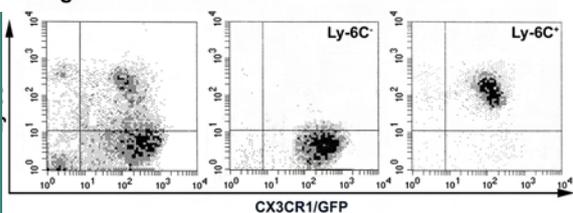
Figure 8: Effects of intramuscular DT injection on skeletal muscle regeneration. Notexin (Nx) was injected into TA of CD11b-DTR mice at d 0 and DT was injected in the same muscle at d 5 (A) and 5 and 6 (B,C). **(A)** 24 h after a single i.m. DT injection, CD45+ cells were isolated from muscle and analyzed for F4/80 expression. Red circles enclose $FSC^{hi}/F4/80^{hi}$ cells. **(B)** Reconstituted whole view of TA muscle at d 10 post Notexin injection (hematoxylin-eosine staining), presenting hallmarks of secondary regeneration restricted to the site of needle puncture (arrow). Hatched area on right picture represents the area excluded from analysis. Bar = 200 μ m. **(C)** PBS- and DT-injected muscles were analyzed at d 10 post injury for myofiber diameter evaluation (hematoxylin-eosine staining). Quantified results are given for 3 independent experiments, each point corresponding to a field and red bars representing means. Bar = 50 μ m.



C - Histological analysis of muscle regeneration



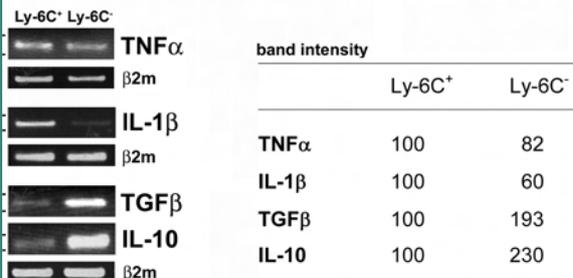
A - Sorting of MO/MP subsets



B Ki67 labeling of Ly-6C⁺ and Ly-6C⁻ MO/MP subsets

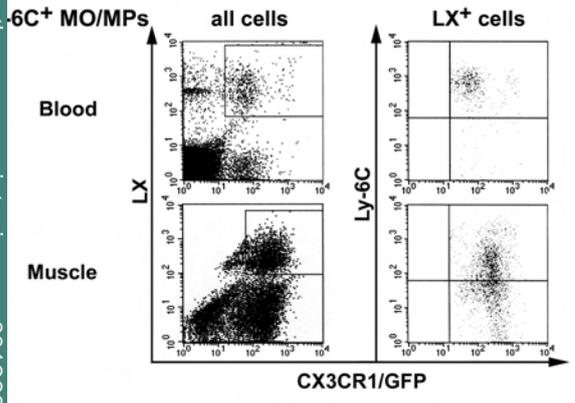
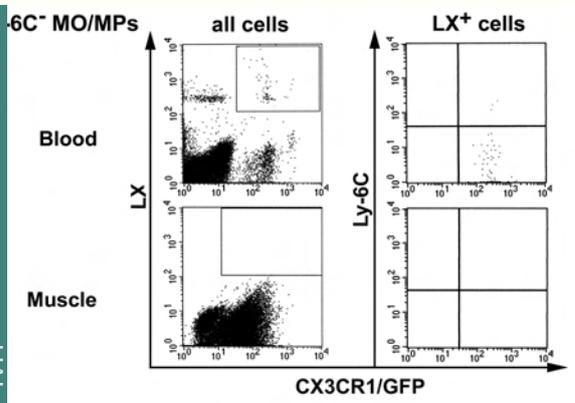
d post injury	15h	48h	4d	7d
CX3CR1 ^{hi} Ly-6C ⁻	0 ± 0	13.5 ± 2.9	27 ± 4.9	12.8 ± 4.9
CX3CR1 ^{lo} Ly-6C ⁺	0 ± 0	0 ± 0	0 ± 0	0 ± 0

C Cytokine expression of Ly-6C⁺ and Ly-6C⁻ MO/MP subsets

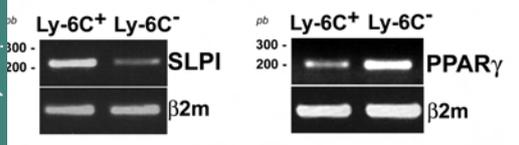


A - Recruitment of LX labeled circulating monocytes

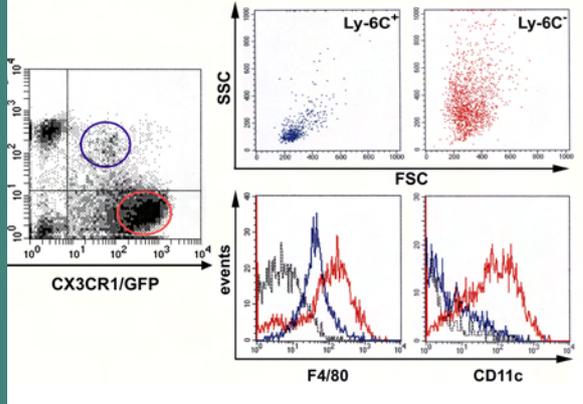
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Ly-6C+ MO/MP subset expression of SLPI and PPAR γ

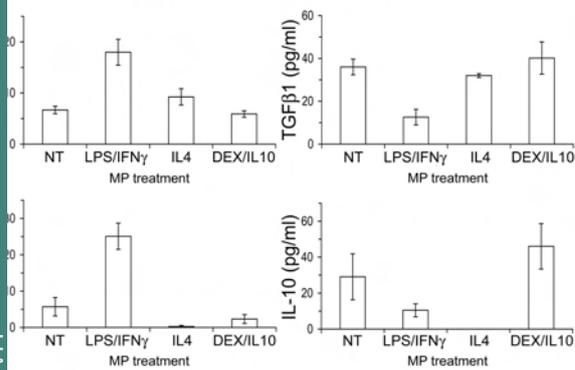


Differentiation of MO/MP subsets

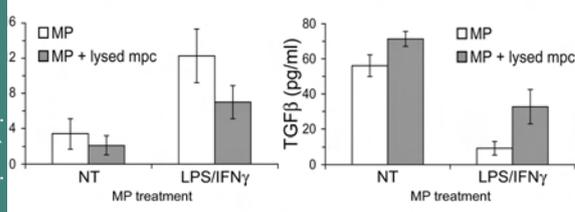


A - Cytokine secretion by MPs after activation

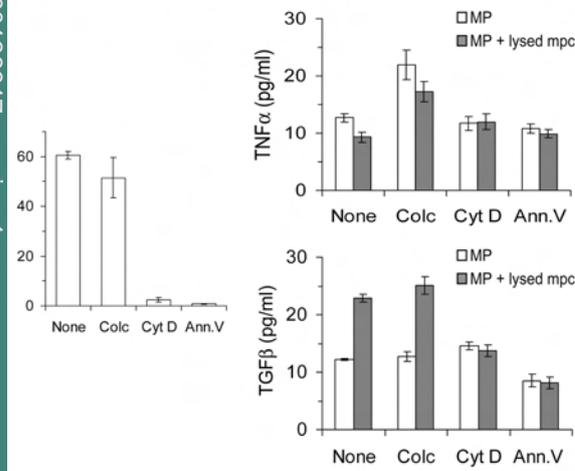
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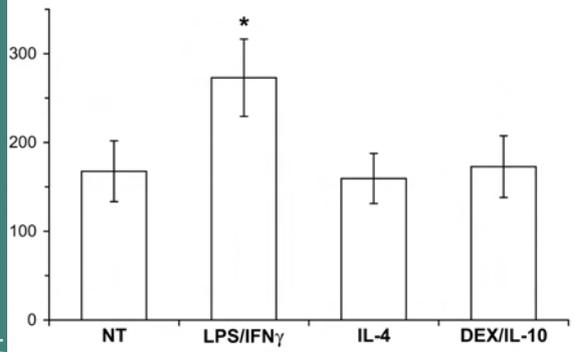
- Cytokine secretion by MPs after phagocytosis of muscle cell debris



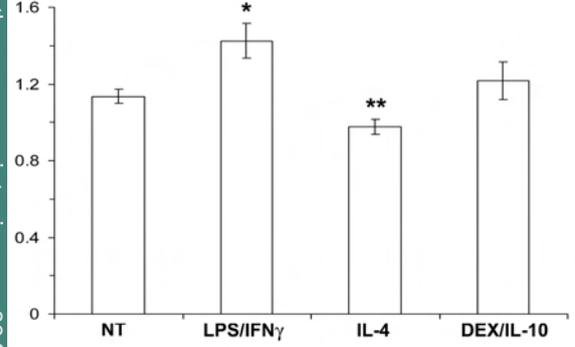
- Inhibition of phagocytosis in inflammatory MPs



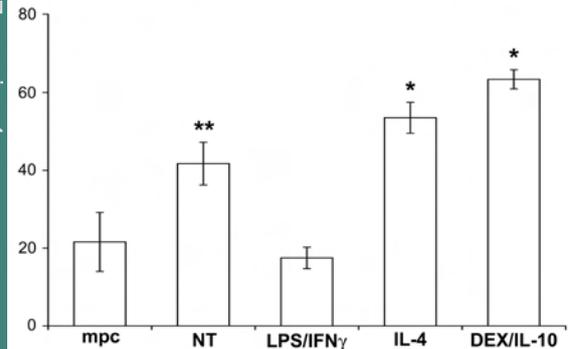
A - Mpc growth



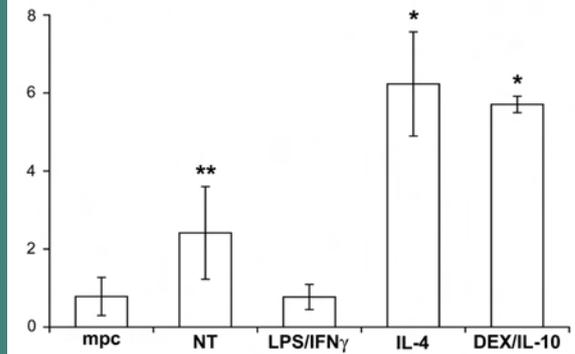
- Mpc proliferation



- Mpc differentiation

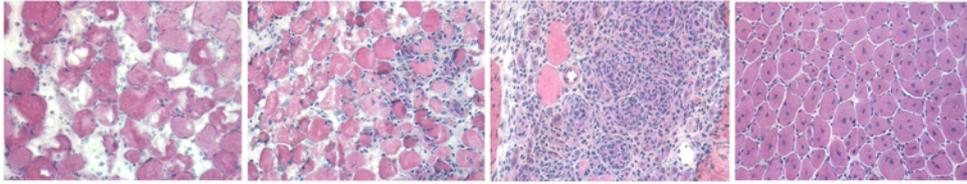
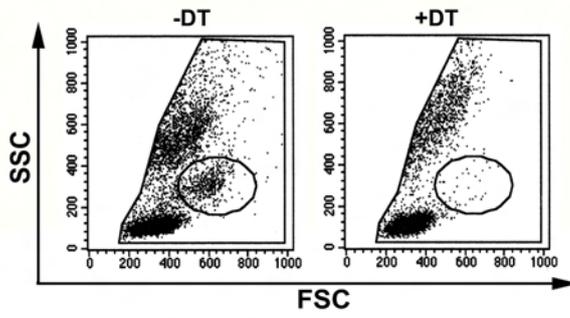
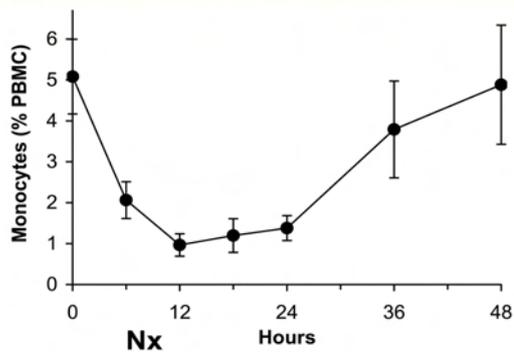


- Mpc fusion

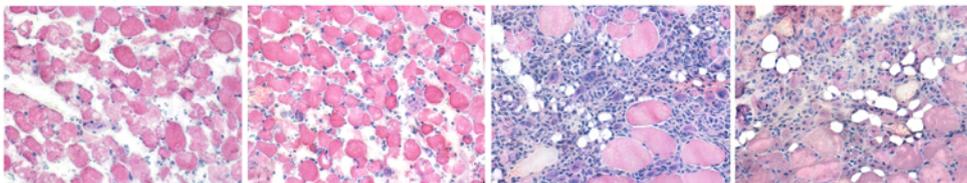


MP treatment

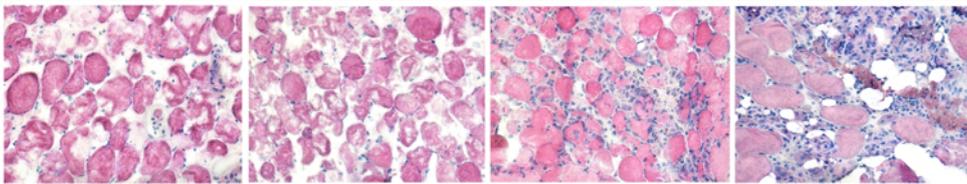
A - Circulating monocyte depletion



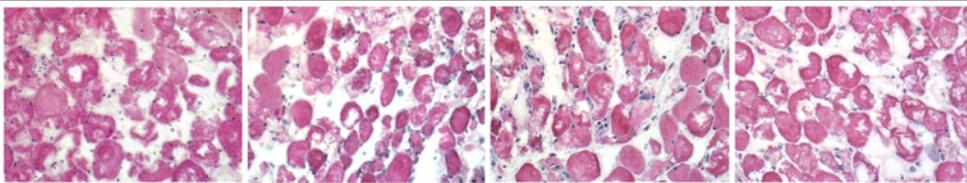
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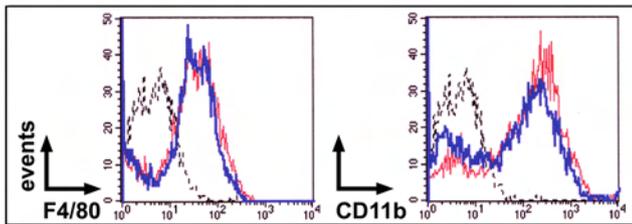
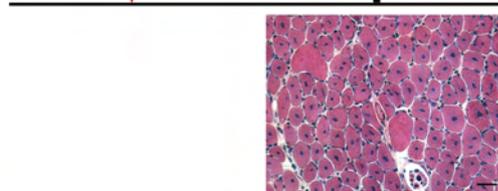
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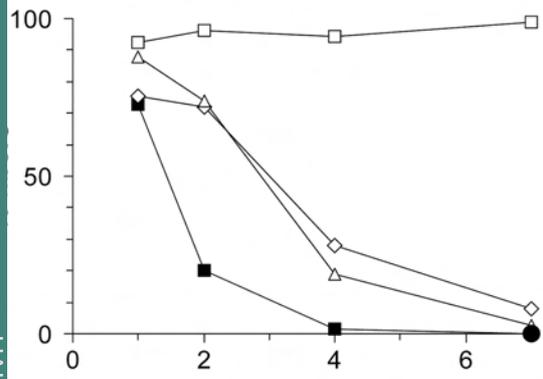
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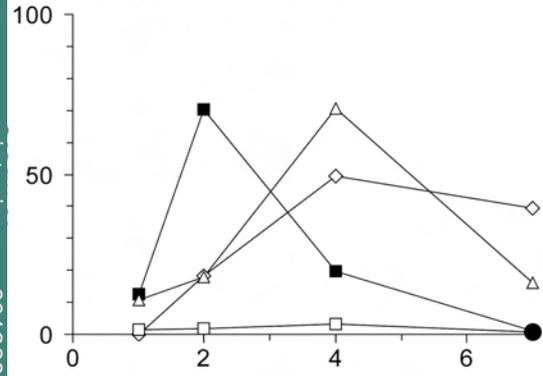
DT
 d4
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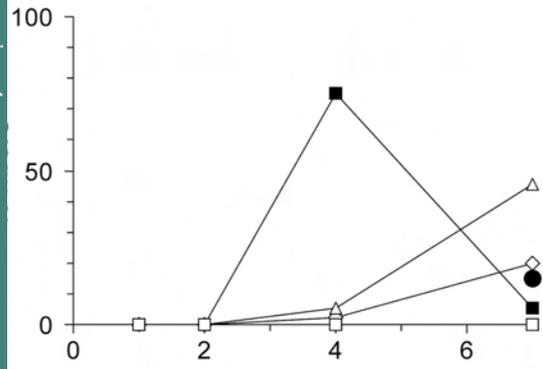
A - Necrotic myofibers



- Phagocytosed myofibers



- Regenerative basic myofibers



- Centrally nucleated myofibers

