17beta-estradiol modulates monocyte heterogeneity and macrophage polarization

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Mononuclear phagocytes respond to environmental cues with the acquisition of distinct functional phenotypes, M1 (classical) or M2 (alternative), which in turn characterize different pathological conditions (1). 17 β -estradiol (E₂) is known to mediate profound effects on monocyte and macrophage immune function (2, 3) acting through estrogen receptors (ER). Data from our laboratory indicate that ER α mediates most of the beneficial effects of estrogens in the cardiovascular system (4). We hypothesized that estrogen-dependent effects on the monocyte/macrophage system protect postmenopausal women from cardiovascular disease (5). Therefore, we investigated the effects of E₂ on mouse circulating monocytes and human monocyte-derived macrophage subsets in resting state (M0) and after M1 or M2 polarized activation.

METHODS: Monocytes from ERα-knockout and wild-type mice were stained with antibodies against surface antigens for classical (Ly6C^{high}/CCR2⁺) and alternative (Ly6C^{low}/CX3CR1⁺) phenotypes, respectively, and analysed by flow cytometry. Macrophages were obtained by culturing monocytes isolated from buffy coats of healthy donors. After 7 days, macrophages were polarized to M1 or M2 phenotypes for 48 h using LPS (1 µg/ml)/IFNg (10 ng/ml) or IL-4 (20 ng/ml)/IL-13 (5 ng/ml), respectively, in the presence or absence of 17β-estradiol or dexamethasone (both 100 nM). M1 and M2 phenotypes were characterized by flow cytometry as percentage of CD68⁺/CCR2⁺ or CD163⁺/CD206⁺/CX3CR1⁺ cells, respectively. In selected experiments cells were further analyzed for CD14 and CD16 expression. ERα expression was assessed by quantitative real-time PCR and Western blot analysis in M0, M1 and M2 polarized macrophages. Intracellular TNF-α and IL-10 production was measured by flow cytometry.

RESULTS: Genetic deletion of ERα in mice was associated with a decrease in the anti-inflammatory (M2) monocyte subset, suggesting that ER-mediated pathways are involved in the development of inflammatory phenotypes. Resting macrophages displayed a predominant M2 phenotype ($36.8\pm7.1\%$ vs $8.1\pm2.2\%$, p<0.05). M1 polarization significantly increased the % of CD68⁺ macrophages compared with M0 cells ($17.8\pm3.3\%$ vs $8.9\pm2.1\%$, p<0.05). The same activation pattern was observed for CD14^{high}/CD16^{low}/CD68⁺ cells (27.8 ± 5.3 vs $7.4\pm1.4\%$, p<0.05). Interestingly, following pro-inflammatory stimulus, M2 cells dropped from $36.8\pm7.1\%$ to $20.5\pm4.7\%$ (p<0.05), suggesting a dynamic M2-to-M1 switch. Overnight pretreatment with dexamethasone enhanced CD163 surface expression in resting (M0: $83.3\pm5.5\%$ vs $49.6\pm9.5\%$) and M2-polarized macrophages ($78.8\pm8.8\%$ vs $50.7\pm11.2\%$, p<0.05), and reversed LPS/IFNγ-induced CD163 down-regulation ($64.4\pm10.3\%$ vs $23.4\pm8.3\%$, p<0.05). Similarly, treatment with 17β-estradiol reversed M1-induced down-regulation of the alternative phenotype through a 25% increase in CD163⁺/CD206⁺ cells. After 48 h, ERα was down-regulated in M1 compared with M0 macrophages, as assessed at both protein (-23%, p<0.05) and mRNA levels. By contrast, no differences in ER β expression were observed between macrophage subsets. Finally, overnight pre-incubation with 17β -estradiol reduced LPS-induced TNF α production compared with untreated macrophages ($13.8\pm2.3\%$ vs $24.4\pm1.7\%$, p<0.05).

CONCLUSIONS: Overall we demonstrated that M1 activation with LPS/IFN γ down-regulate M2 markers and ER α expression, whereas E_2 and dexamethasone promote M2 macrophage signature counteracting the effect of pro-inflammatory stimuli on both M2 surface marker expression and cytokine production.

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