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## Chronic risperidone normalizes elevated pro-inflammatory cytokine and C-reactive protein production in omega-3 fatty acid deficient rats

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### Abstract

Prior clinical and preclinical studies suggest that omega-3 fatty acids negatively regulate pro-inflammatory signaling cascades, and that the atypical antipsychotic risperidone up-regulates omega-3 fatty acid biosynthesis. In the present study, we investigated the effects of chronic (40 d) risperidone treatment (3 mg/kg/day) on basal pro-inflammatory cytokine (interleukin-6, IL-6; tumor necrosis factor-alpha, TNF $\alpha$ ) and C-reactive protein (CRP) production in control and *n*-3 fatty acid deficient rats. Relationships with erythrocyte polyunsaturated fatty acid composition were determined. Compared with untreated controls, untreated *n*-3-deficient rats exhibited significantly greater basal IL-6, TNF $\alpha$ , and CRP production. Following chronic risperidone treatment there were trends for greater IL-6, TNF $\alpha$ , and CRP production in controls, but these did not reach significance. In *n*-3-deficient rats, chronic risperidone normalized elevated IL-6, TNF $\alpha$ , and CRP levels. Erythrocyte arachidonic acid (20:4*n*-6) composition was positively correlated, and erythrocyte eicosapentaenoic (20:5*n*-3) and docosahexaenoic acid (22:6*n*-3) inversely correlated, with plasma IL-6, TNF $\alpha$ , and CRP levels in untreated control and *n*-3-deficient rats, and these associations were not observed among risperidone-treated rats. The adrenic acid (22:4*n*-6)/arachidonic acid ratio, an index of elongase-mediated arachidonic acid biosynthesis, was reduced by risperidone in controls and elevated in *n*-3-deficient rats. These preclinical data demonstrate that chronic risperidone treatment normalizes constitutively elevated pro-inflammatory cytokine and CRP production in *n*-3 fatty acid deficient rats but not in controls, and that the mechanism is dissociable from *n*-3 fatty acid biosynthesis.

### Keywords

Risperidone; Antipsychotic; Cytokine; Interleukin-6; Tumor necrosis factor-alpha; Inflammation; C-reactive protein; Eicosapentaenoic acid; Docosahexaenoic acid; Arachidonic acid; Erythrocyte; Rat

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## 1. Introduction

Emerging evidence from case-control studies suggest that a dysregulation in the inflammatory immune response may be a pathophysiological feature associated with schizophrenia (Watanabe et al., 2010). Although some case-control studies have found that antipsychotic medications have immunosuppressive effects in schizophrenic patients, this has not been consistently observed across different studies (Drzyzga et al., 2006). Multiple factors that influence inflammatory immune activity, including diet, smoking, and medications, may contribute to discrepant findings in clinical studies (Drexhage et al., 2010). Preclinical studies have found that atypical antipsychotic medications, including risperidone, significantly attenuate elevated production of pro-inflammatory cytokines, including interleukin-6 (IL-6) and tumor necrosis factor-alpha (TNF $\alpha$ ), in microglia cells following interferon- $\gamma$  exposure (Bian et al., 2008; Kato et al., 2007), and in mice following peripheral lipopolysaccharide administration (Sugino et al., 2009). These preclinical data demonstrate that atypical antipsychotic medications suppress inflammatory immune activation in response to exogenous stimuli. However, the mechanisms mediating the effects of atypical antipsychotics on constitutive inflammatory immune activity have not been systematically investigated.

Emerging data from clinical and preclinical studies suggest that atypical antipsychotic medications up-regulate long-chain polyunsaturated fatty acid biosynthesis (McNamara, 2009). Importantly, the long-chain omega-6 (*n*-6) fatty acid arachidonic acid (AA, 20:4*n*-6) is a primary substrate for cyclooxygenase-2 (COX-2)-mediated prostaglandin E<sub>2</sub> (PGE<sub>2</sub>) synthesis, and PGE<sub>2</sub> up-regulates IL-6 biosynthesis at the level of transcription via nuclear factor (NF)- $\kappa$ B (Portanova et al., 1996; Wang et al., 2010). In contrast, long-chain *n*-3 fatty acids, including eicosapentaenoic acid (20:5*n*-3) and docosahexaenoic acid (22:6*n*-3), and their lipid metabolites have robust immunosuppressive and anti-inflammatory properties (Calder, 2008; Groeger et al., 2010; Khalfoun et al., 1997; Lu et al., 2010; Mingam et al., 2008; Serhan, 2005). Importantly, we found that chronic risperidone treatment significantly increased erythrocyte membrane long-chain *n*-3 fatty acid composition in rats, and this effect was only observed in rats maintained on a diet fortified with the short-chain *n*-3 fatty acid  $\alpha$ -linolenic acid (ALA, 18:3*n*-3)(McNamara et al., 2009). Together these data support the hypothesis that up-regulation of *n*-3 fatty acid biosynthesis may contribute to the anti-inflammatory and immunosuppressive effects of risperidone.

In a companion paper we reported that *n*-3 fatty acid deficient rats exhibit elevated basal production of pro-inflammatory cytokines (IL-6 and TNF $\alpha$ ) and the acute phase protein C-reactive protein (CRP) compared with *n*-3 fatty acid adequate controls (McNamara et al., 2010). We additionally demonstrated that elevated basal cytokine and CRP production in *n*-3 fatty acid deficient rats was significantly attenuated by prior dietary-induced normalization of *n*-3 fatty acid status. In the present study, we determined the effect of chronic risperidone treatment on constitutive cytokine (IL-6 and TNF $\alpha$ ) and CRP production in *n*-3 fatty acid adequate controls and *n*-3 fatty acid deficient rats. Our specific prediction was that risperidone would significantly reduce cytokine and CRP production in rats maintained on diet ALA-fortified diet in association with increased *n*-3 fatty acid biosynthesis, but not in rats maintained on diet ALA-free diet.

## 2. Materials and methods

### 2.1. Diets

Diets were either  $\alpha$ -linolenic acid (ALA, 18:3*n*-3)-fortified (ALA+, TD.04285) or ALA-free (ALA-, TD.04286)(Harlan-TEKLAD, Madison, WI). Both diets were matched for all non-fat nutrients, and both diets contained *n*-3 fatty acid-free hydrogenated coconut (45 g/kg)

and safflower (19 g/kg) oils. The ALA+ diet additionally contained ALA-containing flaxseed oil (6 g/kg). Analysis of diet fatty acid composition by gas chromatography confirmed that the ALA- diet did not contain ALA, but was matched with the ALA+ diet in saturated fatty acids (C8:0, C10:0, C12:0, C14:0, C16:0, C18:0), monounsaturated fatty acids (18:1*n*-9), and *n*-6 fatty acids (18:2*n*-6, linoleic acid, 22% of total fatty acid composition) (see Table 1 in McNamara et al., 2008). ALA represented 4.6% of total fatty acid composition in the ALA+ diet. Neither diet contained long-chain *n*-3 or *n*-6 fatty acids including DHA and arachidonic acid, respectively.

## 2.2. Animals

For perinatal ALA deficiency, nulliparous Long-Evans hooded dams (Harlan Farms, Indianapolis, IN) were fed the ALA- diet for 1 month prior to mating through weaning, and male offspring were maintained on the ALA- diet from P21-P100. Controls were born to nulliparous dams maintained on the ALA+ diet, and received the ALA+ diet from P21-P100. Rats were housed 2 per cage with food and water available *ad libitum*, and maintained under standard non-barrier *vivarium* conditions (i.e., not specific pathogen-free) on a 12:12 h light:dark cycle. Rats were sacrificed by decapitation on P100 during the light portion of the cycle in a counterbalanced manner. Trunk blood was collected into EDTA-coated tubes, plasma was isolated by centrifugation (4° C), and erythrocytes washed 3× with 4° C 0.9% NaCl. All samples were immediately stored at -80°C.

## 2.3. Drug administration

On P60, one-half of rats in the control (n=10) and *n*-3-deficient (n=10) diet groups were randomly assigned to receive chronic treatment with drug vehicle (0.1 M acetic acid diluted in deionized water) or risperidone (3.0 mg/kg/day; Ortho-McNeil Janssen Pharmaceuticals) through their drinking water, as previously described (McNamara et al., 2009). This dose was selected based on our prior studies finding that it produces therapeutically-relevant plasma risperidone and 9-OH-RSP concentrations and significantly increases *n*-3 fatty acid biosynthesis in rats following chronic oral administration (McNamara et al., 2009). For three days prior to drug delivery, 24 h water consumption was determined for each cage using bottle weights (1 g water = 1 ml water), and ml water intake/mean kg body weight calculated. Risperidone was dissolved and diluted in 0.1 M acetic acid to prepare a stock solution (stored at 4 deg) which was added to tap water in a volume required to deliver the targeted daily dose. Fresh solutions were prepared, and drug concentrations adjusted to mean body weight, every 3 days. Red opaque drinking bottles were used to protect drug from light degradation. Rats were maintained on their respective drug and dose until being sacrificed on P100 (40 days of treatment).

## 2.4. Plasma cytokine and CRP levels

Plasma cytokine (pg/ml) and CRP (ng/ml) concentrations were determined with a multiplexing suspension array and flow-cytometry based analyzer Luminex™ 100 IS (MiraiBio, South San Francisco, CA) using a LINCoplex Cytokine/Chemokine Luminex® Bead immunoassay Kit according to manufacturer's protocol (LINCO Research, St. Charles, MO). All analyses were performed by a technician blinded to group identity.

## 2.5. Erythrocyte fatty acid composition

The gas chromatography procedure used to determine erythrocyte composition (mg fatty acid/100 mg fatty acids) has been described in detail previously (McNamara et al., 2009). Briefly, total fatty acid composition was determined with a Shimadzu GC-2014 (Shimadzu Scientific Instruments Inc., Columbia MD). Analysis of fatty acid methyl esters was based on area under the curve calculated with Shimadzu Class VP 4.3 software. Fatty acid

identification was based on retention times of authenticated fatty acid methyl ester standards (Matreya LLC Inc., Pleasant Gap PA). We focused our analysis on fatty acids implicated in the regulation of immune and inflammatory signaling, including *n*-3 fatty acids, DHA (22:6*n*-3), EPA (20:5*n*-3), and EPA+DHA ('omega-3 index'), and the *n*-6 fatty acids arachidonic acid (20:4*n*-6) and its elongase product adrenic acid (ADA, 22:4*n*-6).

## 2.6. Statistical analysis

Group differences in erythrocyte fatty acid composition and plasma inflammatory markers were evaluated with a two-factor ANOVA, with diet (ALA+, ALA-) and drug treatment (vehicle, RSP) as the main factors. Pairwise comparisons were made with unpaired *t*-tests (2-tail,  $\alpha=0.05$ ). Homogeneity of variance was first confirmed using Bartlett's test. Parametric (Pearson) correlation analyses were performed to determine relationships between erythrocyte fatty acid composition and inflammatory markers (2-tail,  $\alpha=0.05$ ). Statistical analyses were performed with GB-STAT (V.10, Dynamic Microsystems, Inc., Silver Springs MD).

## 3. Results

### 3.1. Erythrocyte fatty acid composition

There was a significant main effect of Diet on erythrocyte EPA+DHA composition,  $F(1,39)=1859$ ,  $P\leq 0.0001$ , and the main effect of Treatment and the Diet  $\times$  Treatment interaction were not significant (Fig. 1A). For erythrocyte arachidonic acid composition, the main effect of Diet was significant,  $F(1,39)=98.6$ ,  $P\leq 0.0001$ , and the main effect of Treatment and the Diet  $\times$  Treatment interaction were not significant (Fig. 1B). For the erythrocyte AA/EPA+DHA ratio, the main effect of Diet was significant,  $F(1,39)=1202$ ,  $P\leq 0.0001$ , and the main effect of Treatment and the Diet  $\times$  Treatment interaction were not significant (Fig. 1C). For the erythrocyte ADA/AA ratio, the main effect of Diet,  $F(1,39)=1037$ ,  $P\leq 0.0001$ , and the Diet  $\times$  Treatment interaction,  $F(1,39)=9.3$ ,  $P=0.004$ , were significant, and the main effect of Treatment was not significant (Fig. 1D).

### 3.2. Plasma inflammatory markers

For plasma IL-6 concentrations (pg/ml), the Diet  $\times$  Treatment interaction was significant,  $F(1,39)=8.8$ ,  $P=0.006$ , and the main effects of Diet and Treatment were not significant (Fig. 2A). For plasma TNF $\alpha$  concentrations (pg/ml), the Diet  $\times$  Treatment interaction was significant,  $F(1,39)=9.6$ ,  $P=0.004$ , and the main effects of Diet and Treatment were not significant (Fig. 2B). For plasma CRP (ng/ml), the Diet  $\times$  Treatment interaction,  $F(1,39)=11.5$ ,  $P=0.002$ , and the main effects of Diet,  $F(1,39)=9.4$ ,  $P=0.004$ , were significant, and the main effect of Treatment approached significance,  $F(1,39)=3.6$ ,  $P=0.07$  (Fig. 2C). Among all rats ( $n=40$ ), IL-6 was positively correlated with CRP ( $r = +0.56$ ,  $P=0.001$ ) and TNF $\alpha$  ( $r = +0.53$ ,  $P=0.002$ ) concentrations.

### 3.3. Correlations with erythrocyte fatty acid composition

Among untreated control and *n*-3 fatty acid deficient rats ( $n=20$ ), erythrocyte arachidonic acid composition was positively correlated with IL-6 ( $r = +0.54$ ,  $P=0.01$ ), TNF $\alpha$  ( $r = +0.69$ ,  $P=0.003$ ) and CRP ( $r = +0.54$ ,  $P=0.02$ ) concentrations, and erythrocyte EPA+DHA composition was inversely correlated with IL-6 ( $r = -0.66$ ,  $P=0.003$ ), TNF $\alpha$  ( $r = -0.65$ ,  $P=0.006$ ), and CRP ( $r = -0.62$ ,  $P=0.008$ ). The arachidonic acid/EPA+DHA ratio was positively correlated with IL-6 ( $r = +0.60$ ,  $P=0.008$ ), TNF $\alpha$  ( $r = +0.66$ ,  $P=0.005$ ), and CRP ( $r = +0.74$ ,  $P=0.001$ ). The ADA/arachidonic acid ratio was not correlated with plasma IL-6 ( $r = -0.32$ ,  $P=0.18$ ), TNF $\alpha$  ( $r = +0.02$ ,  $P=0.92$ ), or CRP ( $r = +0.02$ ,  $P=0.94$ ). Among risperidone-treated control and *n*-3 fatty acid deficient rats ( $n=20$ ), erythrocyte arachidonic

acid composition was not significantly correlated with IL-6 ( $r = -0.17$ ,  $P=0.53$ ), TNF $\alpha$  ( $r = -0.24$ ,  $P=0.33$ ), or CRP ( $r = -0.09$ ,  $P=0.71$ ), and erythrocyte EPA+DHA composition was not significantly correlated IL-6 ( $r = +0.18$ ,  $P=0.46$ ), TNF $\alpha$  ( $r = +0.07$ ,  $P=0.77$ ), or CRP ( $r = +0.06$ ,  $P=0.82$ ). The ADA/AA ratio was inversely correlated with TNF $\alpha$  ( $r = -0.49$ ,  $P=0.03$ ), and similar trends were found for IL-6 ( $r = -0.38$ ,  $P=0.09$ ) and CRP ( $r = -0.35$ ,  $P=0.16$ ).

#### 4. Discussion

A principal finding of the present study is that chronic risperidone treatment normalized constitutive elevations in pro-inflammatory cytokine (IL-6, TNF $\alpha$ ) and CRP production in *n*-3 fatty acid deficient rats. The differences in pro-inflammatory cytokine and CRP levels observed in risperidone-treated versus drug-free *n*-3 fatty acid deficient rats is similar to those previously observed between *n*-3 fatty acid deficient and *n*-3 fatty acid repleted rats (McNamara et al., 2010). In controls, chronic risperidone treatment did not significantly alter IL-6, TNF $\alpha$ , or CRP production, though trends for greater concentrations were observed. Consistent with our prior study (McNamara et al., 2009), chronic risperidone treatment increased erythrocyte membrane long-chain *n*-3 fatty acid composition in rats maintained on a diet containing the short-chain *n*-3 fatty acid precursor (ALA, 18:3*n*-3), but not in rats maintained on the ALA-free diet. Contrary to our hypothesis, however, risperidone decreased pro-inflammatory cytokine and CRP production in rats maintained on the ALA-free diet in the absence of corresponding increases in *n*-3 fatty acid composition, and did not significantly alter cytokine and CRP production in rats maintained on the ALA-fortified diet despite increased *n*-3 fatty acid composition. Nevertheless, these data are in agreement with prior preclinical reports that risperidone has immunosuppressive and anti-inflammatory properties (Kato et al., 2007; Sugino et al., 2009), and further indicate that this effect is not mediated by augmentation of *n*-3 fatty acid biosynthesis.

Consistent with prior studies finding that *n*-3 fatty acid deficiency is associated with reciprocal increases in *n*-6 fatty acid biosynthesis (Igarashi et al., 2007), we found that erythrocyte membrane arachidonic acid (20:4*n*-6) composition was significantly elevated in untreated *n*-3 fatty acid deficient rats. Prior studies indicate that elevated arachidonic acid composition in immune cell membranes is positively correlated with PGE<sub>2</sub> production (Calder, 2008). In the present study, erythrocyte membrane arachidonic acid composition was positively correlated with pro-inflammatory cytokine and CRP levels in untreated control and *n*-3 deficient rats. However, positive correlations were not observed in risperidone treated rats which exhibited similar erythrocyte arachidonic acid composition to untreated rats. This finding suggests that the immunosuppressant and anti-inflammatory effects of risperidone are dissociable from its effects on arachidonic acid membrane composition, and implicate involvement of down-stream signaling events potentially including COX-2-mediated arachidonic acid→PGE<sub>2</sub> production. It is relevant, therefore, that combined treatment with the COX-2 inhibitor celecoxib and risperidone was found to be superior to risperidone alone in reducing symptom severity in schizophrenic patients (Akhondzadeh et al., 2007).

Chronic risperidone treatment differentially altered erythrocyte composition of adrenic acid (ADA, 22:4*n*-6) and the ADA/AA ratio, an index of arachidonic acid→ADA elongation, in controls and *n*-3 fatty acid deficient rats. Specifically, risperidone-treated controls exhibited a lower ADA/AA ratio relative to untreated controls, consistent with risperidone shunting arachidonic acid away from elongation to ADA to alternate pathways including COX-2 mediated arachidonic acid→PGE<sub>2</sub> production. In contrast, risperidone-treated *n*-3 fatty acid deficient rats exhibited a higher erythrocyte ADA/AA ratio relative to untreated *n*-3 fatty acid deficient rats, and the ADA/AA ratio was inversely correlated with plasma TNF $\alpha$



concentrations. The latter findings are consistent with risperidone shunting arachidonic acid to arachidonic acid→ADA fatty acid elongation and away from arachidonic acid→PGE<sub>2</sub> production. This significant Drug by Diet interaction suggests that *n*-3 fatty acid status is an important determinant of the immunosuppressant and anti-inflammatory effects of risperidone, and may contribute in part to the variability in immune responses observed in schizophrenic patients following treatment with atypical antipsychotic medications including risperidone (Drzyzga et al., 2006).

In conclusion, the present study found that chronic risperidone treatment has immunosuppressant and anti-inflammatory properties in *n*-3 fatty acid deficient rats, but not in *n*-3 adequate rats. These findings add to a growing body of preclinical data demonstrating that *n*-3 fatty acid deficiency (Calder, 2008; Groeger et al., 2010; Khalfoun et al., 1997; Lu et al., 2010; McNamara et al., 2010; Mingam et al., 2008; Serhan, 2005) and atypical antipsychotic medications including risperidone (Bian et al., 2008; Kato et al., 2007; Sugino et al., 2009) have opposing effects on inflammatory immune signaling pathways. Moreover, the present data support our prior finding that chronic risperidone augments long-chain *n*-3 fatty acid biosynthesis (McNamara et al., 2009), and demonstrate that this mechanism does not contribute to its immunosuppressive effects. The present data therefore suggest that the anti-inflammatory and immunosuppressive actions of risperidone may be mediated by alternate signaling pathways, including down-regulation of arachidonic acid→PGE<sub>2</sub> production. Future studies will be required to determine whether other antipsychotic medications also exhibit anti-inflammatory and immunosuppressive actions in this model, and whether the effects are correlated with receptor binding profiles. Lastly, in conjunction with previous findings, the present data support the view that elevations in inflammatory and immunological signaling may contribute to the pathophysiology of schizophrenia.

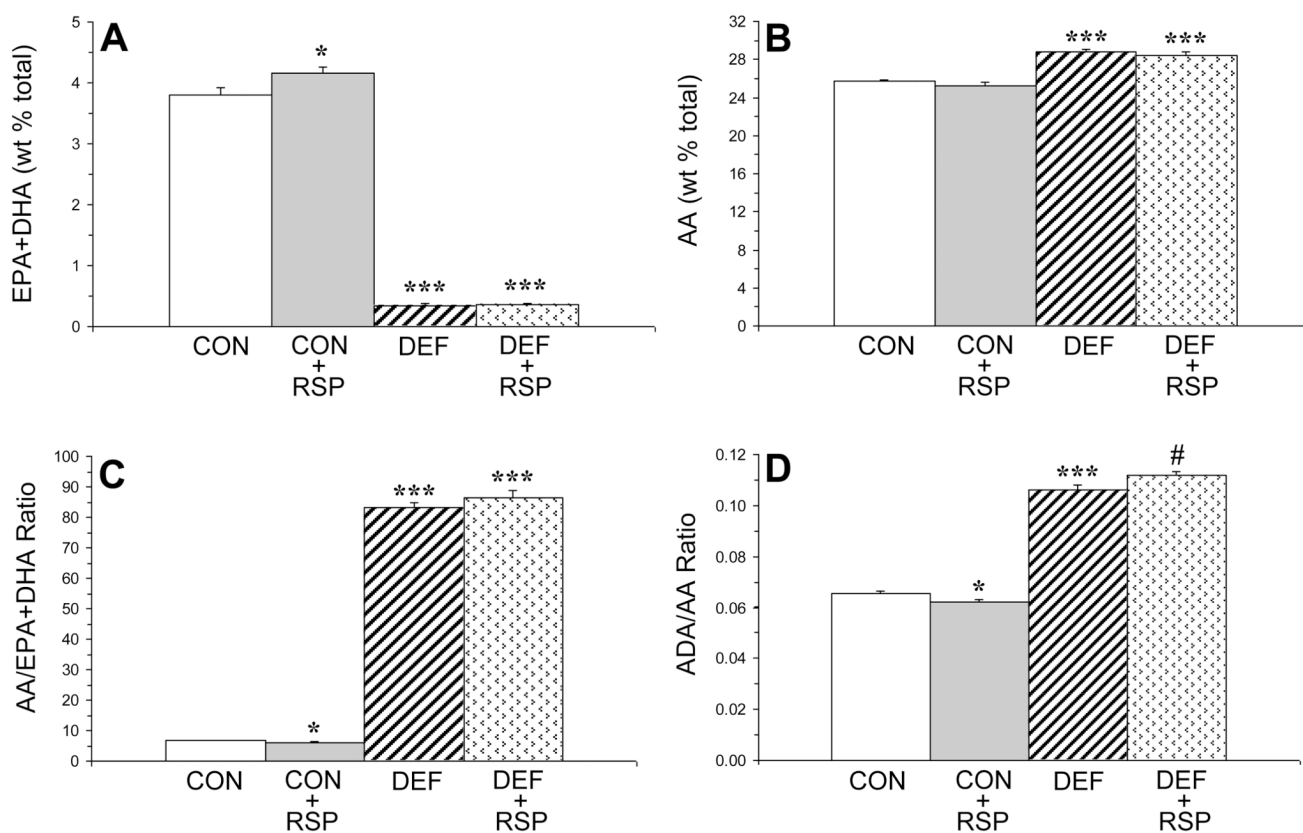
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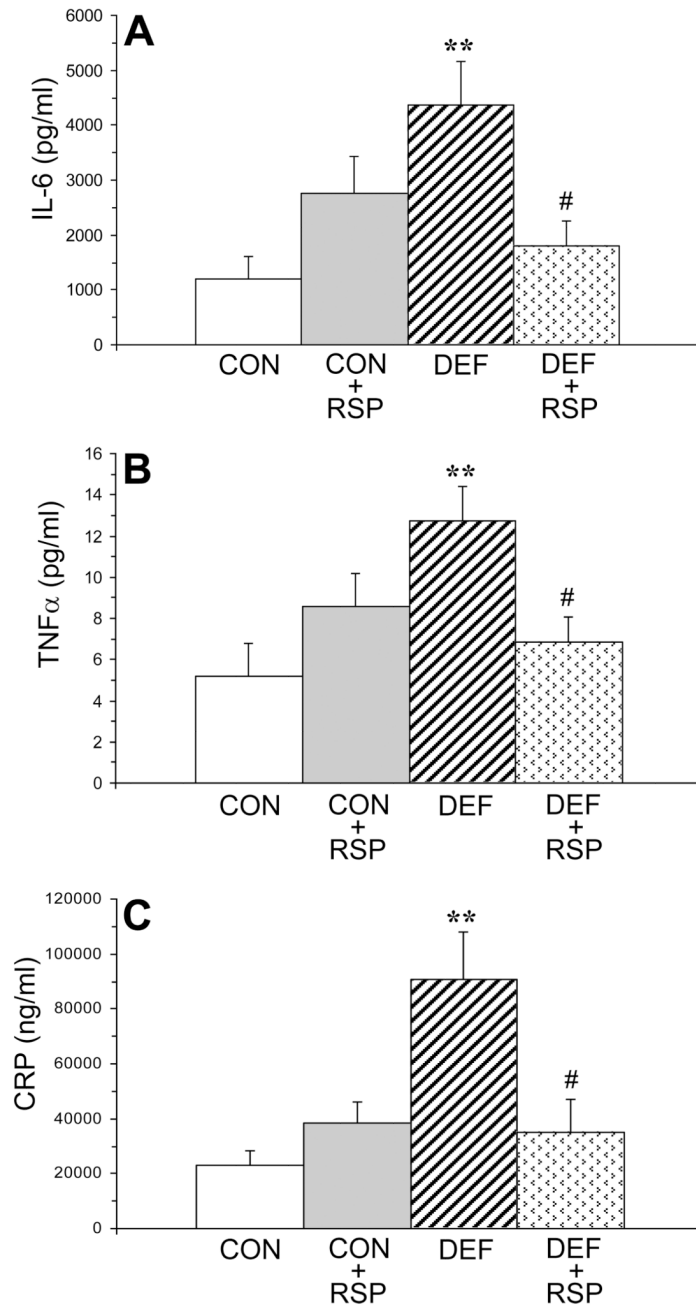
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**Figure 1.**

Erythrocyte composition (wt % total fatty acid composition) of combined long-chain *n*-3 fatty acids, eicosapentaenoic acid (EPA, 20:5*n*-3)+docosahexaenoic acid (DHA, 22:6*n*-3) ('omega-3 index')(A) and arachidonic acid (AA, 20:4*n*-6)(B), and the AA/EPA+DHA (C) and the adrenic acid (ADA, 22:4*n*-6)/AA (D) ratios in control (CON, *n*=10), control +risperidone (CON+RSP, *n*=10), *n*-3-deficient (DEF, *n*=10), and *n*-3-deficient+risperidone (DEF+RSP, *n*=10) rats. Values are group means  $\pm$  S.E.M. \* $P \leq 0.05$ , \*\*\* $P \leq 0.0001$  vs. controls, # $P \leq 0.05$  vs. DEF rats.





**Figure 2.**

Plasma concentrations of interleukin-6 (IL-6, pg/mL)(A), tumor necrosis factor-alpha (TNF $\alpha$ , pg/ml)(B), and C-reactive protein (CRP, ng/ml)(C) in control (CON,  $n=10$ ), control +risperidone (CON+RSP,  $n=10$ ),  $n$ -3-deficient (DEF,  $n=10$ ), and  $n$ -3-deficient+risperidone (DEF+RSP,  $n=10$ ) rats. Values are group means  $\pm$  S.E.M. \*\* $P \leq 0.01$  vs. controls, # $P \leq 0.05$  vs. DEF rats.