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Depressive Symptoms, *n-6:n-3* Fatty Acids, and Inflammation in Older Adults

Janice K. Kiecolt-Glaser, Ph.D., Martha A. Belury, Ph.D., Kyle Porter, M.A.S., David Beversdorf, M.D., Stanley Lemeshow, Ph.D., and Ronald Glaser, Ph.D.

From the Departments of Psychiatry (J.K-G), Human Nutrition (M. A. B.), Ohio State University Center for Biostatistics (K. P., S. L.), Neurology (D. B.), Ohio State University School of Public Health (S.L.), Molecular Virology, Immunology, and Medical Genetics (R.G.), The Ohio State Institute for Behavioral Medicine Research (J.K-G, S. L., R.G.); Ohio State University

Abstract

Objective—Depression and stress promote proinflammatory cytokine production. Dietary intakes of omega-3 (*n-3*) and omega-6 (*n-6*) polyunsaturated fatty acids (PUFAs) also influence inflammation; high *n-6:n-3* ratios enhance proinflammatory cytokine production, while *n-3* has anti-inflammatory properties. This study addressed how interactions between PUFA levels and depressive symptoms were related to proinflammatory cytokine synthesis.

Methods—Blood samples from 43 older adults (mean age = 66.67, SD=10.09) provided data on PUFAs and TNF- α , IL-6, and sIL-6r. Depressive symptoms were assessed by the CES-D.

Results—Depressive symptoms and *n-6:n-3* ratios worked together to enhance proinflammatory cytokines beyond the contribution provided by either variable alone, with substantial variance explained by their interaction: 13% for IL-6 and 31% for TNF- α , whereas full models accounted for 18% and 40%, respectively. Although predicted cytokine levels were fairly consistent across *n-6:n-3* ratios with low depressive symptoms, higher *n-6:n-3* ratios were associated with progressively elevated TNF- α and IL-6 levels as depressive symptoms increased. Higher levels of sIL-6r were associated with higher *n-6:n-3* ratios. Six individuals who met criteria for major depressive disorder had higher *n-6:n-3* ratios and TNF- α , IL-6, and sIL-6r levels than those who did not meet criteria; excluding these 6 individuals reduced the variance explained by the depressive symptoms and *n-6:n-3* ratio interaction.

Conclusions—Diets with high *n-6:n-3* PUFA ratios may enhance risk for both depression and inflammatory diseases.

Keywords

depression; proinflammatory cytokines; omega-3; psychoneuroimmunology

Depression is the most common psychiatric illness, and both major depression and subthreshold depressive symptoms carry substantial health risks (1,2). A number of well-controlled prospective studies have linked depressive symptoms with coronary heart disease (CHD), the leading cause of death in the United States (3); moreover, the nearly 60-fold variance in the annual prevalence of major depression across countries is very similar to the pattern for cardiovascular disease, which shows a strong comorbidity for depression (4,5).

Epidemiological studies have demonstrated significant inverse relationships between annual fish consumption and prevalence of major depression (4). Fish oil is the prime source for two key omega-3 (*n*-3) polyunsaturated fatty acids (PUFAs), eicosapentaenoic acid (EPA) and docosahexaenoic acid (DHA). Several laboratories have provided evidence that depressed patients have, on average, lower plasma levels of *n*-3 PUFAs than nondepressed controls; furthermore, there are relationships within these populations between severity of depressive symptoms and lower plasma levels of the *n*-3 PUFAs (3–6). What is more, all but one of four randomized controlled trials reported significant improvement in the treatment of depression following *n*-3 PUFA supplementation compared to nonsupplemented controls (6). Significant relationships between lower *n*-3 PUFA plasma levels and greater negative mood have also been documented in nonpsychiatric populations (7). Omega-6 (*n*-6) PUFAs are also implicated in depression, with higher *n*-6:*n*-3 ratios observed in depressed patients compared to nondepressed controls (3–6).

Arachidonic acid (AA) derived *n*-6 eicosanoids (primarily from refined vegetable oils such as corn, sunflower, and safflower) increase the production of proinflammatory cytokines, operating as precursors of the proinflammatory eicosanoids of the prostaglandin (PG)₂-series (5). In contrast, the *n*-3 PUFAs, found most abundantly in fish, fish oil, walnuts, wheat germ, and flaxseed can curb the production of AA-derived eicosanoids (5,8). Thus it is not surprising that both higher levels of *n*-3 PUFAs as well as lower *n*-6:*n*-3 ratios are associated with lower proinflammatory cytokine production (9).

The fatty acid composition of the modern Western diet has changed dramatically during the last century, and these changes are thought to be related to increases in inflammatory-related diseases, including depression and cardiovascular disease (6,10). For example, the early hunter-gatherer diet had an *n*-6:*n*-3 ratio of 2:1 to 3:1 (11). However, during the last century, the typical Western diet underwent fundamental alterations with the enormous growth in refined vegetable oil use, a central *n*-6 source that replaced *n*-3 PUFAs from fish, wild game, nuts, seeds, and green leafy vegetables (5,11); in the contemporary North American diet, the *n*-6:*n*-3 ratio is 15:1 to 17:1 (12,13). In fact, the heightened *n*-6:*n*-3 ratio in the Western diet since 1913 has been suggested as a central stimulus for the sharply increased incidence of major depression (4,5).

The *n*-6:*n*-3 ratio is related to serotonergic as well as catecholaminergic neurotransmission (14), providing mechanistic pathways that help explain the relationships with negative mood. Alterations in serotonin (5-HT) receptor numbers and function provoked by changes in PUFAs have been used to link fatty acids with contemporary theories of depression (5,15). Moreover, immune activation may interact with 5-HT functioning to promote depression (5), because PUFAs influence the synthesis of proinflammatory cytokines.

Furthermore, both syndromal depression and depressive symptoms can enhance production of proinflammatory cytokines, including interleukins 1 and 6 (IL-1 and IL-6) and tumor necrosis factor alpha (TNF- α) (2,16–21). Brief laboratory stressors can also provoke transient increases in proinflammatory cytokines (1,22,23). Moreover, stress and depression may effectively “prime” inflammatory responses, promoting larger cytokine increases in reaction to subsequent stressors and/or pathogens (19,24–27). Thus, chronic or longer-term stress and depression can elevate both acute and chronic proinflammatory cytokine production (16,20,23,28). Importantly, these are bi-directional relationships; cytokines have substantial effects on the central nervous system (CNS), producing and enhancing negative moods, as well as physical symptoms such as lethargy and fatigue (18). Indeed, there is evidence that cytokines play a role in the neuroendocrine and behavioral features of depressive disorders (18).

In a provocative study relevant to the present investigation, students who had higher $n\text{-}6:n\text{-}3$ ratios (above the mean) before exams demonstrated greater TNF- α production by lipopolysaccharide (LPS) and mitogen-stimulated peripheral blood leukocytes (PBLs) during exams than those with lower ratios (29). These data suggest that the $n\text{-}6:n\text{-}3$ ratio may influence the proinflammatory response to stressors; because TNF- α and IL-6 are produced by a variety of types of cells, serum cytokine levels may better reflect the overall inflammatory profile than stimulated PBLs or *ex vivo* production (8). Indeed, the aging and depression literatures have focused on proinflammatory cytokine levels assessed in serum (2,30).

Accordingly, we were interested in how depressive symptoms and the $n\text{-}6:n\text{-}3$ ratio would be related to serum TNF- α , IL-6, and the IL-6 soluble receptor (sIL-6r). We studied an older population because proinflammatory cytokine production is elevated following menopause or andropause even in the absence of infection, trauma, or stress (31). Both higher $n\text{-}6:n\text{-}3$ ratios and higher levels of depressive symptoms have been associated with greater production of proinflammatory cytokines; however, in accord with the student examination study (29), we expected that higher $n\text{-}6:n\text{-}3$ ratios and higher levels of depressive symptoms would predict higher levels of TNF- α , IL-6, and sIL-6r than either alone.

Methods

Subjects

The subjects were part of a larger project on stress and health in older adults. Subjects were recruited via notices placed in community and university newspapers, senior citizen centers, a collaborating neurologist, as well as through the Alzheimer's Disease Association; thus, the 43 participants included 18 spouses who were currently providing at least 5 hours of week of care for a spouse with Alzheimer's disease or another progressive dementia, and 25 noncaregivers who were demographically indistinguishable from caregivers but had no similar responsibilities. Inclusion of dementia spousal caregivers in this study was desirable because caregivers have, on average, higher levels of syndromal depression as well as depressive symptoms than noncaregivers (28). Subjects with immunologically-related health problems such as cancer or recent surgeries were excluded during recruitment, as well as those taking any medications with broad immunological consequences. These data were collected between September, 2004 and September, 2005.

For this study we used all individuals in the current cohort who were not taking statins, steroids, or medications for diabetes. Of the 23 subjects excluded for these medications, 10 were caregivers and 13 were noncaregivers; 9 of the 28 caregivers who were part of the larger cohort met current criteria for MDD, compared to 4/38 noncaregivers, $\chi^2(1) = 4.32$, $P=.03$. However, when we eliminated subjects based on our medication exclusions, we ended up with 3 caregivers and 3 noncaregivers and a nonsignificant group difference in terms of current MDD, $\chi^2(1) = 0.19$, $P=.66$. Thus, our medication exclusions substantially reduced the number of strained caregivers, and may also have attenuated group-related differences in cytokines (16,28).

The average age of the 18 men and 25 women in this sample was 66.67 (SD=10.09; the range was 40–86 with one subject under 50, 12 who were 50–59, 13 who were 60–69, 12 who were 70–79, and 5 who were 80 or older); the median education was partial college; three were African American; all were married. The Ohio State University Biomedical Research Review Committee approved the project; all subjects gave written informed consent prior to participation.

Assessment of Mental Health, Depressive Symptoms, and Health-Related Behaviors

The Diagnostic Interview for Genetic Studies (DIGS) (32), developed for the differential diagnosis of major mood and psychotic disorders, provided data on both current and lifetime disorders using DSM-IV criteria. Interviews were administered by well-trained graduate psychology or nursing students who were supervised by a psychologist with extensive experience with structured interviews.

The Center for Epidemiological Studies Depression Scale (CES-D) assessed the severity of depressive symptoms (33,34). Studies have shown acceptable test-retest reliability and excellent construct validity (34). Widely used, the CES-D has distinguished depressed from non-depressed participants in community and clinical samples (34).

We collected health-related data to assess the possibility that relationships among depressive symptoms, inflammatory markers, and PUFAs might simply reflect the contribution of other variables. Health questions from the Older Adults Resources Survey (OARS) (35) assessed underlying diseases. Several studies have found excellent agreement between self-reports and hospital or physician records for specific conditions of interest, including myocardial infarction, stroke, and diabetes (36,37). Assessment of health-related behaviors included body mass index (BMI), smoking, medications, and alcohol intake (38). Two questions assessed exercise (39).

The Pittsburgh Sleep Quality Index (PSQI) (40), a self-rated questionnaire, provides data on sleep quality and disturbances over a one-month interval. The PSQI has good diagnostic sensitivity and specificity in distinguishing good and poor sleepers (40). We also asked about the amount of sleep the prior night.

Immunological Assays

All blood samples were drawn between 8 AM and 10 AM to control for diurnal variation. TNF- α , IL-6, and sIL-6r levels were assayed using Quantikine High Sensitivity Immunoassay kits (R&D), per kit instructions, as described elsewhere (19,28). Serum samples were frozen 45 min - 1 hr after collection. Serum samples were run undiluted in duplicate.

Fatty Acid Analyses

Lipids were extracted from plasma using chloroform: methanol (2:1, v/v) with 0.2 vol. 0.88% KCl (41). Fatty acid methyl esters of the fractions were prepared by incubating the fractions with tetramethylguanidine at 100°C (42) and analyzed by gas chromatography (Hewlett Packard) using a 30-m Omegawax 320 (Supelco-Sigma) capillary column. The helium flow rate was 30 ml/min and oven temperature ramped beginning at 175°C and held for 4 min then increased to 220°C at a rate of 3°C/min as described previously (43). Retention times were compared to authentic standards for fatty acid methyl esters (Supelco-Sigma, St. Louis, MO and Matreya, Inc, Pleasant Gap, PA). Blood samples were not drawn in a fasting state.

Statistical Analysis

Linear regression models were fit to each cytokine, with the *n*-6:*n*-3 ratio and the CES-D score as the independent variables. The interaction between the *n*-6:*n*-3 ratio and CES-D was also included in each model. Several sociodemographic variables, including age and gender, as well as health behavior and medication usage variables were assessed as potential confounders or effect modifiers. A variable was considered a confounder if inclusion in the model resulted in a change of more than 15% in one of the model coefficients. A variable was considered an effect modifier if there was a significant interaction between that variable

and a variable in the primary model. To correct for skewness in the dependent variables, base 10 log-transformed data were used. Statistical significance was evaluated using 2-sided tests at the .05 level of significance.

Fatty acid and cytokine levels were compared for subjects with and without current major depression using analysis of variance (ANOVA) when the data were approximately normally distributed with homogeneous variance. Wilcoxon's rank sum test was used when the data were not normally distributed.

Results

Table 1 shows the percentages of the key fatty acids. Two subjects with extreme *n-6:n-3* ratios of 49.35 and 168.18 (> 9 SDs from the mean) were excluded from analyses using PUFA data as well as from the group means shown in the table. AA/EPA and AA/DHA ratios allow comparisons with recent reports linking these ratios with depression (3,7).

Table 2 provides data on the depressive symptoms and cytokines. The correlation between the *n-6:n-3* ratio and CES-D scores was $r(39) = .24$, $P = .12$.

Table 3 shows the significant interaction between the *n-6:n-3* ratio and depressive symptoms (CES-D) in the model for IL-6 ($F_{1,37} = 5.76$, $P = .02$) as well as in the model for TNF- α , ($F_{1,37} = 19.55$, $P < .001$). For both the IL-6 and TNF- α models the relationship between cytokine levels and the *n-6:n-3* ratio was modified by depressive symptoms. Figures 1a and 1b show that the cytokine levels were relatively consistent at lower depressive symptom levels, while at higher levels there was a marked increase in cytokine levels as the *n-6:n-3* ratio increased. The interactions explained substantial amounts of the variance beyond that explained by the individual contributions of depressive symptoms and the *n-6:n-3* ratio. The full model explained 18% of the total variance in IL-6, an increase of 13% over the model without the interaction. The full model explained 40% of the total variance in TNF- α , an increase of 31% over the model without the interaction.

To assess the estimation of cytokine levels by different fatty acid measures, we compared the results obtained using the *n-6:n-3* ratio – the relative balance of *n-6* and *n-3* – with models that used either *n-6* or *n-3* individually. Each of these models contained one of either the *n-6:n-3* ratio, *n-6* alone, or *n-3* alone as one independent variable; models also contained CES-D as the second independent variable and the interaction between the two independent variables. For IL-6, the model using the *n-6:n-3* ratio accounted for 18% of the total variance, while the model with *n-3* alone accounted for 12% of the total variance, and the model with *n-6* alone accounted for less than 1%. For TNF- α , the *n-6:n-3* ratio model accounted for 40%, the *n-3* model 27%, and the *n-6* model 3%. Thus, in estimating IL-6 and TNF- α levels, the *n-6:n-3* ratio explained more of the total variation than either *n-3* or *n-6* alone.

Another multiple regression model was fit with sIL-6r as the dependent variable. The *n-6:n-3* ratio and CES-D were the independent variables. The *n-6:n-3* ratio was significantly associated with sIL-6r ($F_{1,38} = 4.17$, $P = .05$). The interaction between the *n-6:n-3* ratio and CES-D was not significant and the interaction term was dropped from the model. Model coefficients were 4.46 (SE = .11) for the intercept, .016 (.008) for the *n-6:n-3* ratio, and -.004 (.005) for CES-D.

Major Depression

Six participants met criteria for current major depression; 5 of these were taking antidepressants. The *n-6:n-3* ratio was significantly higher in those with current major

depression than among those without ($F_{1,39} = 10.72$, $P = .002$; 17.92 ± 3.25 vs. 13.49 ± 3.04), while $n-3$ ($P = .12$) and $n-6$ ($P = .17$) differences were nonsignificant.

Participants with current major depression also had significantly higher cytokine production than those who were not clinically depressed. TNF- α was higher in the former than the latter (Wilcoxon $P = .05$; 3.16 ± 3.63 vs. 1.49 ± 0.84), as were IL-6 ($F_{1,41} = 4.20$, $P = .05$; 2.86 ± 3.55 vs. 1.22 ± 0.89), and sIL-6r ($F_{1,41} = 4.05$, $P = .05$; 63.93 ± 28.81 vs. 45.74 ± 17.35 ng/mL). Cytokine means and standard deviations are reported on the raw scale, while log-transformed values were used in the analyses.

Exclusion of all participants who met criteria for major depression from the analysis predictably affected the results, reducing the variance explained by the interaction for IL-6 to a nonsignificant increment; however, for TNF- α the full model explained 26% of the variance, with the interaction accounting for 10% of that variance even in this reduced sample.

As described earlier, the relationship between depressive symptoms and cytokines is bidirectional, and thus we calculated how much variance in depressive symptoms was explained by the measured cytokine levels together with $n-6:n-3$. For IL-6, the interaction was not significant at the 0.05 level ($P=0.109$). However, the model with the interaction did explain substantially more of the variance in depressive symptoms than the model without the interaction, increasing from 6% to 13%. The interaction between TNF- α and $n-6:n-3$ was significant ($P=0.046$) in predicting depressive symptoms. The model with the interaction explained 18% of the variance in depressive symptoms, compared to 8% explained by the model without the interaction.

Depressive Symptoms, Cytokines, Fatty Acids, Sociodemographic Data, and Health Behaviors

Sociodemographic and health behavior variables that were significantly correlated with cytokine levels, fatty acid levels, or depressive symptoms were evaluated as potential confounders or effect modifiers by adding these variables to the models for IL-6, TNF- α , and sIL-6r. None of age, level of education, BMI, hours of sleep the prior night, or exercise (hours of vigorous physical activity per week) were significantly correlated with cytokine levels, fatty acid levels, or depressive symptoms, all $P_s > .18$. Caregivers and controls did not differ significantly on depressive symptoms, $P=.24$, the $n-6:n-3$ ratio, $P=.74$, IL-6r, $P=.82$, IL-6, $P=.51$, or TNF- α , $P=.25$. Two subjects were smokers. Typical weekly alcohol intake was negatively associated with IL-6, $r_s = -.47$, $P=.002$; the median number of drinks reported was 1, and only 2 subjects reported more than 7 alcoholic drinks in an average week, (12 and 14 drinks per week, respectively). In addition, sleep over the last month as measured by the PSQI (full scale) was significantly correlated with depressive symptoms, $r(41) = .51$, $P = .001$. Including the PSQI and alcohol intake in the models for IL-6, TNF- α , and sIL-6r did not change the significant coefficients by more than 15% for any of the models. Thus, neither of these variables confounded the association between cytokine levels, the $n-6:n-3$ ratio, and depressive symptoms.

Medication use and chronic health problems were also considered as potential confounders. The most common medications taken by subjects were angiotensin (10 subjects), antidepressants (9), NSAIDs (9), calcium channel blockers (8), anticoagulants (6), thyroid supplements (6), osteoporosis medications (6), antacids (5), diuretics (5), beta blockers (4), and estrogen supplements (4). Variables for the use of each of these medications were added individually to the models for IL-6, TNF- α , and sIL-6r. A variable indicating subjects not taking any medications (11 subjects) was also considered to evaluate the effect of chronic health problems. The CES-D by $n-6:n-3$ ratio interaction coefficient did not change by more

than 15% for any of the medications in the IL-6 or TNF- α models. In the sIL-6r models, antidepressant use had a slight confounding effect on the association between sIL-6r and the n -6: n -3 ratio, decreasing the coefficient for the n -6: n -3 ratio by 16%. However, there was no significant interaction between antidepressant use and the n -6: n -3 ratio, so antidepressant use was not an effect modifier and the relationship between sIL-6r and the n -6: n -3 ratio was consistent among those taking and those not taking antidepressants.

Potential confounders for IL-6, TNF- α , and sIL-6r models were assessed to determine if the significant relationships among cytokines, fatty acids, and depressive symptoms might be a function of differences in sociodemographic variables, health behaviors, or medication use. We found no evidence that the significant interactions between the n -6: n -3 ratio and CES-D in the models for IL-6 and TNF- α or the significant association between the n -6: n -3 ratio and sIL-6r were simply a function of chronic health problems, medications, or health habits.

Discussion

Higher levels of depressive symptoms as well as higher n -6: n -3 ratios worked together to markedly enhance proinflammatory cytokines beyond the contribution provided by either variable alone. Indeed, the amount of variance explained by the interactions alone was substantial, 13% for IL-6 and 31% for TNF- α ; the full models accounted for 18% and 40%, respectively. These data support and extend the results reported for students taking examinations (29): we found the same pattern they reported for TNF- α with both IL-6 and TNF- α . We also found that individuals who met criteria for syndromal depression had significantly higher n -6: n -3 ratios as well as higher TNF- α , IL-6, and sIL-6r levels than those who did not meet criteria. Moreover, because we were able to show the relationships in serum cytokine levels rather than with levels of cytokines produced by stimulated PBLs, the data provide an important bridge with the aging and depression literatures. Thus, a diet with a high ratio of n -6 to n -3 ratio may enhance risk for both depression as well as inflammatory-related diseases.

Indeed, proinflammatory cytokines influence the onset and course of a spectrum of conditions associated with aging, including CHD, osteoporosis, arthritis, type 2 diabetes, about 15% of cancers, Alzheimer's disease, and periodontal disease (31,44,45). In fact, more globally, chronic inflammation has been suggested as one key biological mechanism that may fuel declines in physical function leading to frailty, disability, and, ultimately, death (30,46,47).

The fact that key inflammatory pathways are influenced by both stress and diet provides one obvious mechanism for our findings. Transcription factor nuclear factor kappa B (NF- κ B) activation upregulates proinflammatory cytokine production (48–50). Psychological stress promotes NF- κ B activation (49,50), providing a mechanism for translating psychological stress into mononuclear cell activation (49). For example, patients with major depression demonstrated significantly greater stress-induced plasma IL-6 levels and mononuclear cell NF- κ B activation than non-depressed controls, and the magnitude of the increase following a laboratory stressor was correlated with depressive symptoms (51).

In contrast, two key n -3 PUFAs, EPA and DHA, can substantially decrease lipopolysaccharide-induced TNF- α expression by blocking NF- κ B activation (48). Indeed, recent work suggests that even modest supplementation with n -3 PUFAs reduces plasma norepinephrine, an important link to stress responses via NF- κ B (52,53).

In contrast to IL-6 and TNF- α , sIL-6r was significantly related to the n -6: n -3 ratio, but not to depressive symptoms, or to the interaction between the two. However, patients who met criteria for major depression had higher sIL-6r levels than those who did not meet criteria.

Mechanistically, sIL-6R forms a ligand-receptor complex, and thus can regulate the impact of IL-6 on inflammation (54). In effect, the sIL-6R expands the cell types that can respond to IL-6, and thus it is not surprising that higher sIL-6R levels have been associated with inflammatory disorders in addition to depression, including arthritis, asthma, and inflammatory bowel disease (54).

Proinflammatory cytokine secretion increases with age (31), and stress and depression can further enhance age-related production (16–19,23,28). In this context it is interesting to note that older subjects show larger immunological changes in *n*-3 supplementation studies, and also respond with greater increases in plasma EPA and DHA and larger decreases in AA than younger participants (55). The ties between age-associated diseases and proinflammatory cytokine production (31) highlight the potential importance of these changes.

Analyses of fatty acids in plasma reflect the past few weeks or months of dietary intake; longitudinal studies suggest that reliability is good over periods from six months to two to three years (56,57), and thus it is not surprising that PUFA levels do not appear to change significantly in response to stressors (29). We did not have food frequency questionnaire data which would have been a useful adjunct to evaluate factors such as differences in dietary intake that could be related to depressive symptoms. However, although food frequency questionnaires are well validated and widely used, fatty acid exposure depends not only on intake, but also absorption and metabolism (9), and thus plasma PUFA levels provide a key endpoint. Our *n*-6:*n*-3 ratio of 14:1 closely approximates the contemporary North American diet; other studies have reported *n*-6:*n*-3 ratios of 15:1 to 17:1 (12,13).

Some of the strongest data linking PUFAs and serum proinflammatory cytokine levels comes from a large cross-sectional study in which researchers assessed serum fatty acids, as well as key inflammatory markers (9). Higher levels of *n*-3 PUFAs were associated with lower IL-6 and TNF- α (9). Importantly, in both cases there was an evident dose-response relationship between the *n*-3 PUFAs and the cytokines (9).

This study is limited by its cross-sectional nature and small sample size; indeed, when individuals who met criteria for major depression were eliminated from analyses, the consequent restricted ranges for both depressive symptoms and *n*-6:*n*-3 ratios attenuated the variance explained for IL-6 and TNF- α (although the TNF- α interaction remained significant). We do not see the attenuated variance as a reflection of the unique contributions of clinical depression, because an exam stress study clearly demonstrated that the joint contributions of stressors and higher *n*-6:*n*-3 ratios enhance proinflammatory cytokine production independent of syndromal depression (29).

Indeed, although Maes et al. (29) excluded any student who had a current or past Axis I psychiatric disorder, and any student who reported a major negative life event in the prior year, they found substantially enhanced TNF- α production during exams in students with higher *n*-6:*n*-3 ratios (> 400%) compared to those with lower ratios (~50%), despite their small sample, *n*=27. Importantly, their “high” group’s *n*-6:*n*-3 ratio was 13.38 (SD=3.29), compared to our sample’s *n*-6:*n*-3 mean of 14.13 (SD=3.42), while their “low” group had a mean of 6.15 (SD=1.20); only 4 of our subjects would have fallen into their low group. Thus, together their data and ours suggest that the joint contributions of higher *n*-6:*n*-3 ratios and depressive symptoms or stress can markedly enhance proinflammatory cytokine production; a randomized controlled supplementation trial would provide the optimal cause-and-effect demonstration.

Epidemiological studies have clearly linked elevated proinflammatory cytokines with depressive symptoms (2), as well as elevated *n*-6:*n*-3 ratios (9). However, other researchers

have not always found depression-cytokine relationships or have reported mixed results for different cytokines (18). Our data suggest that a lower $n-6:n-3$ ratio offers some protection, particularly as depressive symptoms increase; accordingly, it is possible that differences in participants' dietary $n-3$ intakes may have contributed to the variability in depression-cytokine relationships observed among other studies (18).

Furthermore, increased inflammatory activity has been associated with treatment-resistant depression in several studies (18), and preliminary evidence suggests that $n-3$ supplementation is beneficial for treatment-resistant depression (58,59). If the combination of a high $n-6:n-3$ ratio concomitant with high depressive symptoms promotes inflammation, then our data suggest that one obvious potential benefit of $n-3$ supplementation would be decrements in cytokine production; as noted earlier, cytokines have substantial CNS effects, including the production and enhancement of negative moods, and thus supplementation may be beneficial because it interrupts a maladaptive feedback loop (18).

In summary, our findings highlight ways in which diet may enhance or inhibit depression-related inflammation among older adults. These behavior-dietary-immune interactions have important implications for both mental and physical health.

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Acronyms

AA	arachidonic acid
BMI	body mass index
CES-D	Center for Epidemiological Studies Depression Scale
DHA	docosahexanoic acid
DIGS	Diagnostic Interview for Genetic Studies
EPA	eicosapentaenoic acid
IL-6	interleukin 6
sIL-6r	IL-6 soluble receptor
$n-3$	omega-3
$n-6$	omega-6
NF- κ B	nuclear factor kappa B
PSQI	Pittsburgh Sleep Quality Index
PUFA	polyunsaturated fatty acid
TNF- α	tumor necrosis factor alpha

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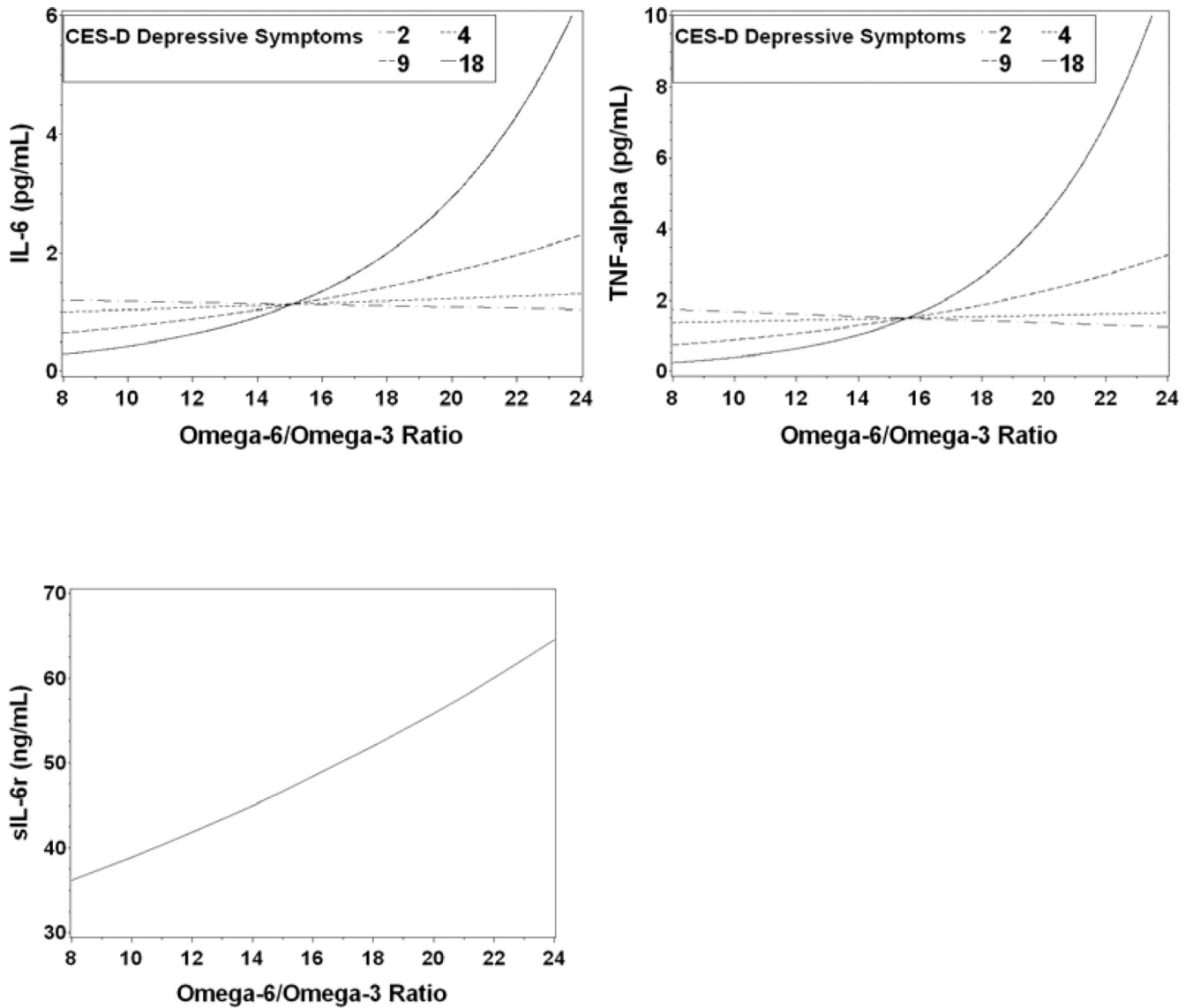


Figure 1.

a-c. IL-6 (1a) and TNF- α (1b) plots showing the predicted cytokine response by n -6: n -3 ratio separately for different CES-D depressive symptom levels. Although the analyses used continuous variables, predicted cytokine values are plotted for the midpoint of each depressive symptom quartile (2,4,9,18) to illustrate the effect of depression on the cytokine n -6: n -3 ratio relationship. The predicted values for IL-6 and TNF- α are non-linear because the model was on the log scale and then the predicted values were transformed back to the original scale.

In the IL-6sr model (1c) the n -6: n -3 ratio main effect was significant, and the ratio did not differ significantly by depression. Therefore only one response curve is on the plot, at the mean depression level of 7.4. Responses at other depression levels would yield parallel curves.

Table 1

Relative fatty acid composition of plasma phospholipids

Fatty Acid	Means (% fatty acid)	SD
<i>n</i>-3 Series		
Alpha Linolenic 18:3 <i>n</i> -3	0.64	0.22
Stearidonic 18:4 <i>n</i> -3	0.00	0.00
Eicosatetraenoic 20:4 <i>n</i> -3	0.33	0.16
Eicosapentaenoic 20:5 <i>n</i> -3	0.73	0.57
Docosapentaenoic (<i>n</i> -3) 22:5 <i>n</i> -3	0.19	0.13
Docosahexaenoic 22:6 <i>n</i> -3	0.82	0.25
Total EPA and DHA	1.55	0.72
<i>n</i> -3 Total	2.71	0.84
<i>n</i>-6 Series		
Linoleic 18:2 <i>n</i> -6	25.7	5.01
Gamma-linoleic 18:3 <i>n</i> -6	0.02	0.03
Dihomogammalinolenic 20:3 <i>n</i> -6	1.08	2.72
Arachidonic 20:4 <i>n</i> -6	10.21	5.82
Docosadienoic 22:2 <i>n</i> -6	0.28	0.16
<i>n</i> -6 total	38.53	5.98
Ratios		
<i>n</i> -6: <i>n</i> -3 ratio	14.13	3.42
AA/EPA ratio	19.14	27.41
AA/DHA ratio	12.27	10.38

Table 2

Means and SDs for Depressive Symptoms and Cytokines

Measure	Means	SD
CES-D (depressive symptoms)	7.41	5.86
IL-6, pg/ml (raw)	1.48	1.61
TNF- α , pg/ml (raw)	1.76	1.61
sIL-6r, pg/ml (raw)	48643.90	20364.04

Table 3

Parameter estimates and standard errors for IL-6 and TNF- α *

Cytokine	Regression Parameter Estimates (Standard Errors)			Interaction p-value
	Intercept	<i>n-6:n-3</i> Ratio	Depressive Symptoms	
IL-6 (log ₁₀)	.275 (.267)	-.015 (.019)	-.083 (.034)	.005 (.002)
TNF- α (log ₁₀)	.535 (.187)	-.023 (.013)	-.110 (.024)	.007 (.002)

*The parameter estimates can be interpreted in terms of the slope of the regression of the cytokine on the *n-6:n-3* ratio dependent on the level of depressive symptoms. For example, the slope for the regression of log-transformed IL-6 on *n-6:n-3* ratio is (-).015 + .005-depressive symptoms. Therefore the slope is zero when depressive symptoms are equal to 3.0 and increases by .005 for each unit increase in depressive symptoms.