Inflammation and emotion regulation: Findings from the MIDUS II study

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evident in various psychiatric groups, linking inflammatory biomarkers with expressive suppression (Khan et al., 2020; Osphina et al., 2021; Powers et al., 2016). However, previous studies linking inflammation and ER have been limited by focusing on specific populations (e.g., bereaved spouses, psychiatric groups) and/or measurement of a narrow range of cytokines and abbreviated ER assessments. The present study builds on prior literature and more broadly investigates whether individual differences in levels of inflammatory cytokines are associated with expressive suppression and cognitive reappraisal in a national sample of middle-aged adults. In particular, we hypothesized that expressive suppression would be associated with higher inflammation, while cognitive reappraisal would correlate with lower inflammation.

2. Methods

We analyzed data from The Midlife in the United States (MIDUS) study, a longitudinal assessment of a national U.S. sample of adults developed to investigate how psychological, behavioral, and social factors relate to and influence mental and physical health (Brim et al., 2004). The first wave of the study (MIDUS I) collected sociodemographic and psychosocial data from 7108 English-speaking Americans, aged 25–74 years, living in the contiguous 48 states with access to at least one telephone (recruited by random digit dialing). Of the original MIDUS I participants, 4963 individuals were re-contacted and successfully completed the second wave (MIDUS II) of the study, which included a phone interview and completion of self-reports. As an additional assessment of MIDUS II, a subset of participants (n = 1255) completed the Biomarkers project between 2002 and 2006 to investigate biological mechanisms underlying behavioral and psychosocial factors, which included a detailed medical history and collection of biological samples (i.e., blood, urine, and saliva). A subset of participants (n = 331) who completed the Biomarker project also completed the Neuroscience Project between 2004 and 2009, which included assessment of the neural circuitry involved in affective style, regulation and reactivity via self-reports, EMG, EEG, and eyeblink startle magnitude at rest and in response to the presentation of affective imagery (Ryff and Davidson, 2010; Ryff et al., 2021).

2.1. Participants

Of the 331 participants who completed the MIDUS II Biomarker and Neuroscience projects, inflammatory biomarkers and the full ER measure were obtained from 117 participants; the following analyses focus on this subsample of participants. All procedures were approved by the Institutional Review Boards at the University of Wisconsin, the University of California Los Angeles, Pennsylvania State University, and Georgetown University; all participants provided informed consent.

2.2. Measures

Inflammatory biomarkers assessed in the Biomarker Project included: CRP, Interleukin Adhesion Molecule-1 (ICAM-1), IL-6, fibrinogen, and E-selectin. Additional inflammatory biomarker data were extracted using stored MIDUS II samples and later added to the MIDUS Refresher Biomarker protocol, which included interleukin-8 (IL-8), interleukin-10 (IL-10) and TNF-α, as well as a re-assay of CRP samples falling below the assay range. Assay procedures for all inflammatory markers have been described previously (Dienberg Love et al., 2010) and are available at the MIDUS website. In short, fasting blood samples were collected from each participant before breakfast on their second day of their hospital stay. Samples were subsequently stored in a −85°C freezer until assay, where cytokines were assayed in the MIDUS Biocore Laboratory (University of Wisconsin, Madison, WI), and ICAM-1, E-selectin, fibrinogen and CRP assays were performed at the Laboratory for Clinical Biochemistry Research (University of Vermont, Burlington, VT). IL-8, IL-10 and TNF-α were assayed via immunoelectrochemiluminescence, which included using a V-plex Custom Human Cytokine Kit (catalog #K151A0H-2) manufactured by Meso Scale Diagnostics, Rockville, MD. IL-6 was measured using the Quantikine High Sensitivity ELISA kit #HS600B manufactured by R&D Systems, Minneapolis, MN. ICAM-1 was assayed by an ELISA assay produced by Parameter Human sICAM-1 Immunoassay, R&D Systems, Minneapolis, MN. E-selectin was measured using a high-sensitivity ELISA assay (Parameter Human s-Selectin Immunoassay, R&D Systems, Minneapolis, MN). Fibrinogen was evaluated using a BNII nephelometer (N Antiserum to Human Fibrinogen; Dade Behring Inc. Deerfield, IL). CRP was analyzed using the BNII nephelometer from Dade Behring utilizing a particle-enhanced immunonephelometric assay. Finally, samples falling below the assay range for CRP were re-assayed by immunoelectrochemiluminescence using a high-sensitivity assay kit (Meso Scale Diagnostics #K1515TG).

The full 10-item version of the Emotion Regulation Questionnaire (ERQ) was utilized to assess use of specific ER strategies (Gross and John, 2003). The cognitive reappraisal and expressive suppression scales consist of six and four items, respectively. Participants were instructed to indicate using a 7-point scale (from 1 strongly disagree to 7 strongly agree) to what extent they agree with each item, with higher scores reflecting a higher endorsement of that strategy use. The ERQ has demonstrated adequate reliability, with alpha of 0.79 for cognitive reappraisal and 0.73 for expressive suppression, and good test-retest reliability (0.69 for both scales) (Gross and John, 2003).

Participant characteristics known to influence ER and inflammation were obtained during the biomarker collection and MIDUS II assessments, including sociodemographic factors (age and sex), total number of medications ingested (which includes prescription, over the counter, and alternative medications), body mass index (BMI), and depressive symptomatology total score indexed using the Center for Epidemiological Studies Depression Inventory (CESD; Radloff, 1997) (Appleton et al., 2013; Ellis et al., 2019).

2.3. Statistical analyses

Participant characteristics, including demographics, ER and inflammatory biomarker levels were computed for the sample. As inflammatory markers were skewed, we applied log-transformation to achieve normality, and removed any additional outliers via visual inspection. Next, we calculated Pearson’s full and partial correlations between ER and inflammatory biomarkers, controlling for: age, sex, BMI, total number of prescribed medications, and total depression score. In order to evaluate whether inflammatory biomarkers predicted use of ER strategies, we conducted separate hierarchical linear regression models for expressive suppression and cognitive reappraisal with demographic and clinical covariates in block 1, and inflammatory biomarkers in block 2. All analyses were performed using SPSS version 24 (IBM Corp., Armonk, NY, USA). For all analyses, alpha was set at 0.05 and tests were two-tailed.

3. Results

In the study sample (N = 117), participants were aged 37–84 years (M = 53.84, SD = 9.99), and 52.1% were female. Only 78 participants disclosed racial and educational status, of which 96.15% self-reported as white, and 50% earned a bachelor’s degree or higher. Follow-up analyses between responders and non-responders for racial and educational status revealed no difference between any ER strategies or inflammatory biomarkers. Full participant characteristics are presented in Table 1. Full and partial Pearson’s correlations between ER strategies and inflammatory biomarkers are presented in Table 2. Results for full correlations demonstrated significant negative relationships between expressive suppression and IL-10, TNF-α, and ICAM-1, with greater levels of IL-10, TNF-α, and ICAM-1 linked to lower expressive suppression. Results for partial correlations yielded similar results, with


4. Discussion

Results from the present study expand upon previous literature investigating the link between inflammation and ER strategies in healthy adults. The most critical findings show that expressive suppression use is correlated with immune biomarkers, particularly IL-10, TNF-α, and ICAM-1, with increased inflammation relating to lower use of suppression.

These findings stand in contrast to previous reports suggesting a positive relationship between expressive suppression use and levels of CRP, IL-2, IL-6, and TNF-α (Appleton et al., 2013; Ellis et al., 2019; Lopez et al., 2020). Theoretically, one recent biobehavioral model (Renna, 2021) proposes that ER mediates the influence of negative affective experience on inflammation in a bi-directional manner, with continuous use of negative emotions increasing immune system dysregulation and promoting a cascade of maladaptive ER strategies (including suppression use) (Mennin et al., 2005), all of which subsequently increase risk for poor long-term health outcomes. Thus, our contradictory results invite speculation on the specific mechanistic properties of the investigated inflammatory biomarkers.

IL-10 is conceptualized as a major anti-inflammatory agent, and plays an important role in preventing inflammatory and autoimmune pathologies (Saraiva and O’garra, 2010). Therefore, lower levels of the anti-inflammatory IL-10 may promote subsequent increased levels of pro-inflammatory agents, which has previously been observed in association with higher expressive suppression use. Indeed, one prior study investigating the relationships between inflammation and emotional intelligence (including domains of emotion awareness and ER) discovered a positive relationship, with higher IL-10 levels linked to better emotional intelligence (Jung et al., 2019). IL-10 is often induced by inflammatory stimuli and has been observed to play a central role in limiting inflammatory response via multiple pathways (Saraiva and O’garra, 2010); as such, additional research is necessary to investigate the mechanistic effects of IL-10 on specific neuroanatomical regions implicated in ER use.

The other two significant biomarkers under investigation (TNF-α and ICAM-1), however, are considered pro-inflammatory (Rubel et al., 2001; Van de Stolpe and Van der Saag, 1996). Of these, TNF-α is considered the master immune regulator as its activation impacts related inflammatory responses such as vasodilation, leukocyte adhesion to epithelium through expression of adhesion molecules (such as ICAM-1), and blood coagulation (of which fibrinogen is an essential contributor) (Zelová and Hosek, 2013). Therefore, the significant and trend-level negative relationships of ICAM-1 and fibrinogen with expressive suppression (respectively) may be due to the primary significant negative relationship evident between TNF-α and expressive suppression. Prior studies assessing expressive suppression use and TNF-α have been limited and inconsistent, with one study demonstrating no association in a healthy sample (Jung et al., 2019), and another study showing higher TNF-α relating to greater expressive suppression use in bereaved spouses (Lopez et al., 2020). Still, these contradictory results conflict with our current results, suggesting that more research using similar demographic samples is necessary to more accurately assess the relationship between TNF-α and expressive suppression. Mechanistically, initial CNS research demonstrated neurodegenerative effects of increased TNF-α in the brain via neuronal apoptosis, leading to subsequent impairments in cognitive functioning and eventual pathogenesis of neurodegenerative diseases in various populations as a result of activation of TNF receptor 1 (McAfoose and Baune, 2009; Kogan et al., 2018, 2019). Interestingly, TNF-α has also recently shown to exert neuroprotective effects, with increased synaptic plasticity via stimulation of TNF receptor 2 (Fischer & Maier, 2015). One model of cytokine function argues for consideration of the specific level of TNF-α as well as the sample population under investigation (McAfoose and Baune, 2009), as prior studies indicate that optimal levels of TNF-α in immunologically non-challenged individuals is essential for normal cognitive functioning.

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### Table 1: Participant characteristics – MIDUS subsample.

| Sociodemographic characteristics | M (SD) | N (%) |
|----------------------------------|-------|-------|
| Age                              | 53.84 (9.99) | 117 |
| Sex: female (% female)           | 61 (52.1%) | 78 |
| Race: white (% white)            | 75 (96.1%) | 115 |
| Education: bachelor’s degree or higher (%) | 39 (50.0%) | 41 (35.0%) |
| Current depression ratings       | 8.09 (6.32) | 117 |
| Total # of prescription medications per person | 4.16 (3.84) | 117 |
| BMI                              | 30.27 (6.21) | 117 |

| Health Indicators | M (SD) | N (%) |
|-------------------|-------|-------|
| Age               | 53.84 (9.99) | 117 |
| Sex: female (% female) | 61 (52.1%) | 78 |
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| BMI                              | 30.27 (6.21) | 117 |

### Emotion Regulation

| Emotion Regulation | M (SD) | N (%) |
|--------------------|-------|-------|
| Expressive Suppression | 3.45 (1.24) | 117 |
| Cognitive Reappraisal | 4.97 (0.90) | 117 |

### Inflammation Levels

| Inflammation Levels | M (SD) | N (%) |
|---------------------|-------|-------|
| IL-6 (pg/mL)        | 1.13 (1.17) | 117 |
| IL-8 (pg/mL)        | 15.58 (8.77) | 117 |
| IL-10 (pg/mL)       | 0.29 (0.25) | 117 |
| TNF-α (pg/mL)       | 1.92 (0.53) | 117 |
| CRP (ng/mL)         | 2.60 (2.63) | 117 |
| E-selectin (ng/mL)  | 62.03 (27.96) | 117 |
| ICAM-1 (ng/mL)      | 214.36 (81.44) | 117 |
| Fibrinogen (mg/dL)  | 357.21 (71.03) | 117 |

Note: N = 117. BMI: Body Mass Index; IL: interleukin; TNF-α: tumor necrosis factor-α; CRP: C-reactive protein; ICAM-1: Intercellular Adhesion Molecule-1.

As indicated in Table 3, for the hierarchical regression model predicting expressive suppression, no significant covariates were found in block 1. However, significant covariates in block 2 included age (β = 0.25, p = 0.04), sex (β = -0.20, p = 0.05), and a trend for depressive symptomatology (β = 0.16, p = 0.09). Significant inflammatory predictors included TNF-α (β = -0.24, p = 0.03) and fibrinogen at trend-level (β = -0.22, p = 0.07). The model including inflammation markers was significant, accounting for 23% of the variance (F13, 113) = 2.32, p = 0.01. Results for the hierarchical regression model predicting cognitive reappraisal yielded no significant predictors, neither covariates nor inflammatory biomarkers.

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expressive suppression negatively correlating with TNF-α and ICAM-1, and at trend level for IL-10. Neither full nor partial correlations yielded significant associations between inflammatory biomarkers and cognitive reappraisal.

As indicated in Table 3, for the hierarchical regression model predicting expressive suppression, no significant covariates were found in block 1. However, significant covariates in block 2 included age (β = 0.25, p = 0.04), sex (β = -0.20, p = 0.05), and a trend for depressive symptomatology (β = 0.16, p = 0.09). Significant inflammatory predictors included TNF-α (β = -0.24, p = 0.03) and fibrinogen at trend-level (β = -0.22, p = 0.07). The model including inflammation markers was significant, accounting for 23% of the variance (F13, 113) = 2.32, p = 0.01. Results for the hierarchical regression model predicting cognitive reappraisal yielded no significant predictors, neither covariates nor inflammatory biomarkers.

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Note: N = 117. BMI: Body Mass Index; IL: interleukin; TNF-α: tumor necrosis factor-α; CRP: C-reactive protein; ICAM-1: Intercellular Adhesion Molecule-1.

a N = 78.

b N = 115.
c N = 116.
Future research should investigate potential optimal levels of TNF-α in both immunologically impaired and unimpaired populations in relation to cognition, including adaptive ER use.

Our results showed no relationships between cognitive reappraisal use and inflammatory biomarker levels. Prior studies assessing this link have yielded inconsistent results. While greater cognitive reappraisal use has been associated with lower inflammatory biomarker levels (Appleton et al., 2013; Brown et al., 2020), other studies have shown no relationship (Ellis et al., 2019; Khan et al., 2020; Lopez et al., 2020). Possible explanations for this inconsistency may relate to sample size and characteristics, the manner in which inflammatory biomarkers are extracted (peripheral blood vs. nasal shedding), the specific inflammatory markers assessed, and lack of relevant covariates in statistical models. Future studies should aim to address these discrepancies.

Our results should be considered in light of a number of limitations in the present study. First, given the use of a small sample of convenience, our results should be considered exploratory. As this was a cross-sectional study, causality cannot be inferred. Given our and prior results of significant demographic covariates (e.g., age and sex), larger samples must be employed to assess the impact of mediators on ER, as well as assessing a broader range of inflammatory markers. Related to demographics, only 78 individuals of our subsample (N = 117) elected to report their race and educational status; therefore, the actual demographic breakdown of race and education remains unclear. While this explains why we did not include these variables as covariates in our correlation and regression analyses, future studies must account for these factors, given that previous literature suggests differential inflammation levels as a function of race (Stepanikova et al., 2017) and level of education (Friedman and Herd, 2010). There is also the possibility of reporting bias when using self-report assessments, such that participants’ perceptions on how they generally regulate their emotions differs on their actual daily ER use. ER was not recorded on the same day as their blood draw for inflammatory assessment; future research should collect repeated assessments of ER during daily life and emotion-eliciting situations to examine whether specific ER-strategy use is linked to concurrent or subsequent changes in inflammation. Additionally, our sample skewed older, with 18 out of 44 women indicating

(Saune et al., 2008). Overall, these observations may partly explain our current results, suggesting that an ideal level of activation of a particular TNF-α sub-type may correlate with optimal cognitive performance.
menopause as their current menstrual status. As menopause has been characterized by an increase in pro-inflammatory biomarkers attributed to estrogen deprivation (Gameiro et al., 2010), menstrual status must be considered in subsequent studies. Finally, given the current study’s exploratory nature, corrections were not made for performing multiple statistical tests.

In summary, our results offer further evidence for a relationship between inflammation levels and ER, specifically between TNF-α and expressive suppression ER-strategy use. Understanding the underlying etiological mechanisms of adaptive ER is crucial for promoting improved physical and mental well-being. For example, chronic experience of negative emotions has been linked to a range of diseases such as cancer, infectious and autoimmune diseases, whose onset and course may be impacted by the immune system (Kiecolt-Glaser et al., 2002).

Relatedly, maladaptive regulation of emotions, particularly persistent use of expressive suppression has been linked to a spectrum of disorders such as cardiovascular disease, chronic pain, arthritis and cancer (Panagopoulou et al., 2002), with one possible explanation for this relationship pointing towards altered immune response (Ellis et al., 2019). Indeed, the recent field of psychoneuroimmunology aims to discover the pathways through which negative emotionality and maladaptive ER influences physical health via altered immune functioning (Renna, 2021). Failure to adequately regulate ones’ emotions perpetuates negative affective experience, which prolongs physiological activation and heightened immune levels, thereby increasing the risk of long-term health problems. Therefore, future research aiming to improve well-being may develop pharmaceutical and/or psychotherapeutic interventions that may dually target both enhanced immune response as well as effective ER.

Declaration of competing interest
All authors report no conflict of interests.

Data availability
Data will be made available on request.

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Appendix A. List of Abbreviations

| Abbreviation | Definition |
|--------------|------------|
| BMI          | Body Mass Index |
| CESD         | Center for Epidemiological Studies Depression Inventory |
| CNS          | Central Nervous System |
| CRP          | C-reactive protein |
| ER           | Emotion Regulation |
| ERQ          | Emotion Regulation Questionnaire |
| ICAM-1       | Intercellular Adhesion Molecule-1 |
| IL-2         | Interleukin-2 |
| IL-6         | Interleukin-6 |
| IL-8         | Interleukin-8 |
| IL-10        | Interleukin-10 |
| IFN-γ        | Interferon-γ |
| MIDUS        | Midlife in the United States |
| TNF-α        | Tumor Necrosis Factor-α |

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