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Sleep Loss Activates Cellular Markers of Inflammation: Sex Differences

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Abstract

Sleep disturbance is associated with inflammation and related disorders including cardiovascular disease, arthritis, and diabetes mellitus. Given sex differences in the prevalence of inflammatory disorders with stronger associations in females, this study was undertaken to test the effects of sleep loss on cellular mechanisms that contribute to proinflammatory cytokine activity. In 26 healthy adults (11 females; 15 males), monocyte intracellular proinflammatory cytokine production was repeatedly assessed at 08:00, 12:00, 16:00, 20:00, and 23:00 h during a baseline period and after partial sleep deprivation (awake from 11 PM to 3 AM). In the morning after a night of sleep loss, monocyte production of interleukin 6 and tumor necrosis factor- α differentially changed between the two sexes. Whereas both females and males showed a marked increase in the lipopolysaccharide (LPS) - stimulated production of IL-6 and TNF- α in the morning immediately after PSD, production of these cytokines during the early- and late evening was increased in the females as compared to decreases in the males. Sleep loss induces a functional alteration of monocyte proinflammatory cytokine responses with females showing greater cellular immune activation as compared to changes in males. These results have implications for understanding the role of sleep disturbance in the differential risk profile for inflammatory disorders between the sexes.

Introduction

Epidemiologic data indicate that sleep disturbance and short sleep duration adversely impact human physical health (Gangwisch et al., 2007) and mortality risk (Dew et al., 2003; Gangwisch et al., 2008; Shankar et al., 2008). Although the biological basis for these links between sleep and health are not known, risk of a wide spectrum of medical conditions including cardiovascular disease, arthritis, diabetes, certain cancers, obesity, and functional decline is associated with activation of cellular signals that initiate expression of inflammatory cytokines (Karin, 2005; Volpato et al., 2001).

Sleep loss has been found to have consequences on inflammatory mechanisms. Experimental sleep deprivation induces increases in circulating levels of inflammatory markers such as interleukin-6 (IL-6), tumor necrosis factor- α (TNF- α), and C-reactive protein (CRP) (Irwin et al., 2004; Meier-Ewert et al., 2004; Shearer et al., 2001). In addition, early night partial sleep deprivation (PSD) activates cellular expression of IL-6 and TNF- α , with effects on up-stream

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sources of cellular inflammatory cytokine expression (Irwin et al., 2006). Moreover, such up-regulation of inflammatory gene expression following such sleep loss (Irwin et al., 2006) is due to activation of nuclear factor (NF)- κ B transcription control (Irwin et al., 2008) that plays a key role in controlling cellular expression of pro-inflammatory genes (Pascual and Glass, 2006). Activation of NF- κ B following sleep loss, however, occurs primarily in females, but not in males (Irwin et al., 2008).

There are well-known sex differences in the prevalence of inflammatory disorders, with females being between 2-9 times as likely as males to develop autoimmune disorders including arthritis and diabetes (Whitacre, 2001). In addition, whereas cardiovascular disease is thought to have an inflammatory basis and is more prevalent in males than females, recent epidemiologic data indicate that subjective symptoms of disturbed sleep and short sleep duration were associated with a greater risk of cardiovascular disease in females than in males, even after control for relevant confounders such as body mass index and physical activity (Cappuccio et al., 2007; Newman et al., 2000). Moreover, Suarez (Suarez, 2008) recently found that poor sleep quality and self-reported prolonged sleep latency were associated with increases of CRP and IL-6 in females but not males. Hence, in this study we sought to determine sex differences in the effects of experimental sleep loss on the inflammatory response by measuring the production of proinflammatory cytokines by monocytes following ligation of the Toll-like receptor 4 (TLR4) with lipopolysaccharide (LPS). TLRs mediate innate immune responses to common pathogens (Cook et al., 2004), and aberrant increases of TLR activity has been linked to inflammatory diseases such as rheumatoid arthritis (Andreaskos et al., 2004), Crohn's disease (Andreaskos et al., 2004), and heart failure (Satoh et al., 2005).

Methods

Twenty six subjects (eleven women and fifteen men) who were medically healthy, as determined by medical history, physical examination, and laboratory testing, were recruited between October 2006 and June 2008. All subjects fulfilled criteria for Never Mentally Ill as determined by Structured Clinical Interview for Diagnostic and Statistical Manual - IV (SCID). Females and males did not differ in mean \pm SD age (37.2 ± 9.6 vs. 36.3 ± 10.4 years; $t = 0.2$, $P = 0.82$), education level (15.9 ± 1.8 vs. 15.3 ± 1.8 ; $t = 0.9$, $P = 0.37$) mean body mass index [BMI] (23.8 ± 3.8 vs. 24.9 ± 3.7 ; $t = 0.7$, $P = 0.47$), or ethnicity (nonwhite: 36.4% vs. 20.0%; $\chi^2 = 0.86$, $P = 0.35$) 8=26.9 [SD=3.6]; 35.7% were non-white). Without restricting assessments to a single point in the menstrual cycle, females and males did not differ in mean levels of progesterone (1.2 ± 0.6 ng/ml vs. 1.0 ± 0.6 ; $t = 1.1$, $P = 0.30$) or estradiol (67.1 ± 62.3 ng/ml vs. 37.5 ± 9.7 ng/ml; $t = 1.9$, $P = 0.08$). Past or current Axis I DSM-IV psychiatric disorder, use of psychotropic medication, regular use of nonsteroidal anti-inflammatory medications, and tobacco smoking were exclusionary. Subjects regularly slept between 22:30 and 7:30 h as confirmed by 2-week sleep diaries. The study was approved by the UCLA Institutional Review Board.

Subjects participated in a PSD protocol that was conducted on the UCLA General Clinical Research Center (GCRC) as previously described (Irwin et al., 2006). The PSD night immediately followed the baseline night and all subjects remained on the GCRC during the day with behavioral monitoring to prevent any napping behavior. Acquisition of blood samples occurred via an indwelling venous forearm catheter at 08:00, 12:00, 16:00, 20:00, and 23:00 h during baseline and after PSD. Hence, a total of 5 measures were obtained prior to sleep deprivation (i.e., baseline) with 5 additional measures obtained after PSD. Blood samples were held at room temperature between collection and assay, as we have found up to a 10 h- interval does not impact assay results; baseline and PSD samples were treated similarly in terms of hold times.

Monocyte intracellular production of IL-6 and TNF- α was assessed by flow cytometry using peridinin chlorophyll protein—labeled CD14 monoclonal antibody, allophycocyanin-labeled anti-TNF- α monoclonal antibody, and phycoerythrin-labeled IL-6 antibody. (Collado-Hidalgo et al., 2006). Heparin-treated blood (1 mL) was mixed with 100 pg/mL LPS (Sigma, St. Louis, MO) and 10 Ag/mL brefeldin A (Sigma) and incubated for 4 hours at 37°C in a platform mixer followed by an overnight incubation at 4°C. RBCs were lysed in fluorescence-activated cell sorting lysing solution (BD Biosciences), remaining cells were permeabilized in fluorescence-activated cell sorting permeabilizing buffer (BD Biosciences), and fluorescence conjugated antibodies were added for 30 minutes at room temperature in the dark. Cells were then washed and resuspended in 1% paraformaldehyde for assay on a Coulter Elite flow cytometer using the Coulter Elite software. Forward scatter and side scatter were used to gate on monocytes and granulocytes. About 12,000 CD14+ events were counted to determine the percentage of cytokine-secreting monocytes, with quadrant coordinates set based on unstimulated monocytes cells. To determine percentage of stimulated cells expressing TNF- α , IL-6, or co-expressing TNF- α and IL-6, unstimulated cytokine-positive event percentages were subtracted from stimulated percentages to obtain net stimulated cytokine positive event percentages. We have previously reported that PSD does not alter absolute numbers of monocytes (Irwin et al. 1996).

Statistical Analyses

Data were analyzed using SAS version 9.13 for Windows. To determine the effects of PSD on monocyte intracellular proinflammatory cytokine expression in females and males, repeated measures mixed model ANOVA was performed using a 2 (sex: female, male) \times 2 (condition: baseline, PSD) \times 5 (time: 08:00, 12:00, 16:00, 20:00, 23:00 h) design, covarying for age, education, ethnicity, BMI, progesterone and estradiol. Analyses were conducted for monocytes expressing TNF- α , IL-6, or co-expressing TNF- α and IL-6, as well as the aggregate.

Results

Partial night sleep deprivation induced differential changes in the capacity of monocytes to express TNF- α and IL-6 as compared to the baseline condition, with a condition \times time interaction for the cells expressing TNF- α , IL-6, co-expressing TNF- α and IL-6, and the aggregate measure (Figure 1) ($F(4,183) = 3.9, P < 0.01$; $F(4,183) = 25.8, P < 0.001$; $F(4,183) = 8.2, P < 0.001$; $F(4,183) = 5.8, P < 0.001$). Notably, both females and males showed a marked increase in the stimulated production of IL-6 and TNF- α in the morning immediately after PSD, consistent with our prior findings (Irwin et al., 2006). The fixed effect for time was significant for the cells expressing TNF- α , IL-6, co-expressing TNF- α and IL-6, and the aggregate measure ($F(4,184) = 4.3, P < 0.01$; $F(4,184) = 18.7, P < 0.001$; $F(4,184) = 21.1, P < 0.001$; $F(4,184) = 11.5, P < 0.001$). In addition, there were condition effects for cells expressing TNF- α , IL-6, co-expressing TNF- α and IL-6 ($F(4,184) = 8.2, P < 0.01$; $F(4,184) = 15.7, P < 0.001$; $F(4,184) = 6.8, P < 0.01$) but not for the aggregate measure ($F(1,185) = 0.8, P = 0.37$).

There were sex differences in the capacity of monocytes to express proinflammatory cytokines following PSD, with significant condition \times sex interactions for cells expressing TNF- α , co-expressing TNF- α and IL-6, and the aggregate measure (Figure 1) ($F(1,184) = 3.8, P < 0.06$; $F(1,184) = 14.4, P < 0.001$; $F(1,184) = 13.7, P > 0.001$) but not for cells expressing IL-6 ($F(1,184) = 0.3, P = 0.84$). Additional analyses examined differential responses at individual timepoints and found condition \times sex interactions for cells co-expressing TNF- α and IL-6 and the aggregate measure at two timepoints: 20:00 ($F(1,19.0) = 9.4, P < 0.01$; $F(1,19.0) = 6.0, P < 0.05$) and 23:00 ($F(1,19.0) = 4.7, P < 0.05$; $F(1,19.9) = 7.4, P < 0.05$) in which females showed increases in the co- and aggregate expression of TNF- α and IL-6 as compared to

decreases in men. Figure 2 displays the results for the aggregate measure of TNF- α and IL-6 expression across the individual timepoints. Similar responses between the sexes were found at 8:00, 12:00, and 16:00 (all $P > 0.1$) and for cells expressing TNF- α or IL-6 at all timepoints (all $P > 0.1$). No main effect for sex found for cells expressing TNF- α , IL-6, co-expressing TNF- α and IL-6, and the aggregate (all $P > 0.1$).

Discussion

Here we present evidence that acute sleep loss induces a differential alteration in a functional cellular innate immune response. In the morning after a night of sleep loss, LPS ligation of TLR4 or CD14 triggered significantly greater production of IL-6 and TNF- α in peripheral blood monocyte populations relative to morning levels following uninterrupted sleep. Whereas both females and males showed a similar increase in morning levels of IL-6 and TNF- α , the two sexes differed in cellular immune activation during the subsequent daytime period. As compared to responses in males, females showed increases in the cellular expression of these proinflammatory cytokines during the early- and late evening hours. Consistent with these observations, Suarez (2008) found that poor sleep quality and prolonged sleep latency were associated with increases of circulating levels of CRP and IL-6 in females, but not in males. Moreover, our prior findings demonstrate sex differences in nuclear translocation of NF- κ B following sleep loss (Irwin et al., 2006), in which females show elevations of NF- κ B in the morning following sleep loss. Although prior work did not evaluate nocturnal levels of NF- κ B following PSD, it is possible that this transcription factor is similarly activated in both sexes during the night, which in turn contributes to similar increases in morning levels of IL-6 and TNF- α . To account for sex differences in the daytime expression of IL-6 and TNF- α , we hypothesize that such activation of NF- κ B persists into the morning in females, but not in males, and drives subsequent elevations of proinflammatory cytokines during the latter part of the day in females. Together, these data further demonstrate that sleep loss enhances inflammatory biology in females as compared to changes in males, with implications for understanding the increased risk profile for inflammatory disorders in females.

Transient changes in monocyte production of proinflammatory cytokines, even when extended across the course of the day, might not translate into increased risk for disease. However, it is known that aberrant increases of TLR activity are found in association with rheumatoid arthritis (Andreaskos et al., 2004), Crohn's disease (Andreaskos et al., 2004), and heart failure (Sato et al., 2005). Furthermore, increases in monocyte responses to LPS stimulation correlate with increases in circulating inflammatory markers (Collado-Hidalgo et al., 2006), and small elevations in circulating inflammatory mediators, for example, are associated with the syndrome of insulin resistance and type II diabetes mellitus, independent of adiposity (Festa et al., 2000).

PSD induces marked increases in cardiovascular responses (Irwin and Ziegler, 2005) as well as sympathoadrenal activity upon awakening (Irwin et al., 1999; Irwin and Ziegler, 2005). In turn, adrenergic output is known to facilitate *in vivo* release of inflammatory mediators into circulating blood (Friedman and Irwin, 1997; Johnson et al., 2005). Although catecholamines *in vitro* are reported to suppress proinflammatory cytokine production (Cole et al., 1998; Wahle et al., 2005), physiological concentrations of norepinephrine are reported to be sufficient to result in a significant dose dependent increase of NF- κ B-binding activity *in vitro* (Bierhaus et al., 2003), which might drive increases in the monocyte expression of proinflammatory cytokines. To our knowledge, no prior study has examined whether sex differences alter adrenergic outflow following sleep loss, although in women, but not men, sympathovagal balance as indexed by heart rate variability has been found to be negatively associated with monocyte IL-6 expression (O'Connor M et al., 2007). Moreover in women, but not in men, vagal tone is positively associated with production of this cytokine (O'Connor M et al.,

2007). In turn, circadian variation in sympathetic output might contribute to the circadian variation in stimulated monocyte production of proinflammatory cytokines, which we have previously demonstrated (O'Connor et al., 2007; Irwin et al., 2006). Finally, it is possible that differences in reproductive hormones including changes in levels of such hormones across the menstrual phase contributed to these differences, although these findings were demonstrated even after levels of progesterone and estradiol were statistically adjusted.

Loss of sleep during only part of the night is one of the most common complaints of persons who experience environmental or psychological stress (Akerstedt et al., 1990; McDermott et al., 1997). In this study, PSD was used as an experimental model of difficulties falling asleep. Our results show that a modest amount of sleep loss differentially activates cellular markers of inflammation in females as compared to responses in males. Given these data, further investigations are needed to define the effects of sleep loss on inflammatory mechanisms in females and males, with implications for understanding increased risk of chronic inflammatory disorders in females.

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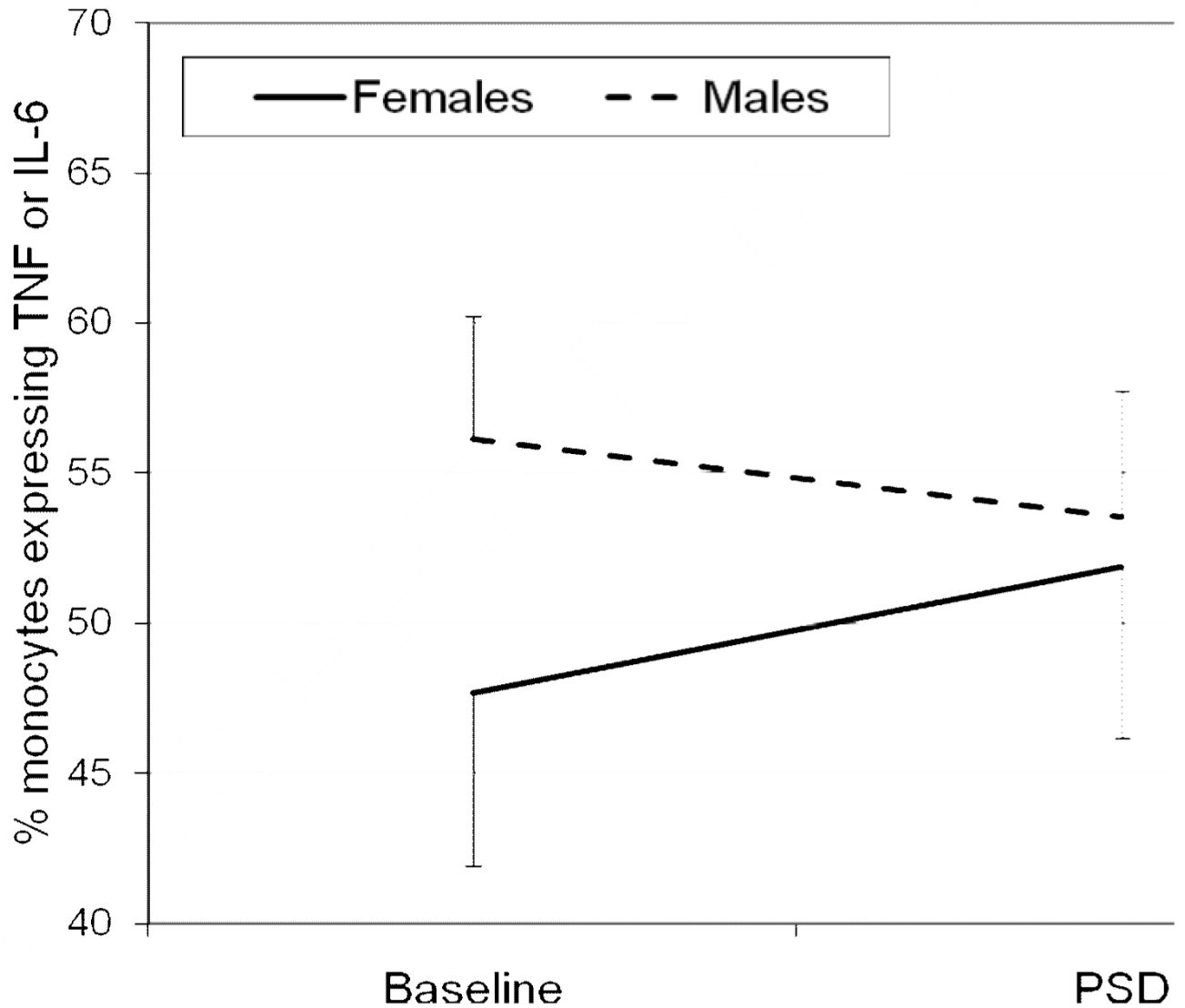


Figure 1. Sex differences in aggregate expression of interleukin (IL) 6 and tumor necrosis factor- α (TNF) in lipopolysaccharide-stimulated CD14+ cells between baseline and partial sleep deprivation PSD conditions. A significant condition \times sex interaction in the aggregate expression of TNF- α and IL-6 was found in which females showed relative increases in aggregate expression of TNF- α and IL-6 as compared to decreases in men ($F(1,184) = 13.7$, $P > 0.001$). Data are represented as mean \pm SEM.

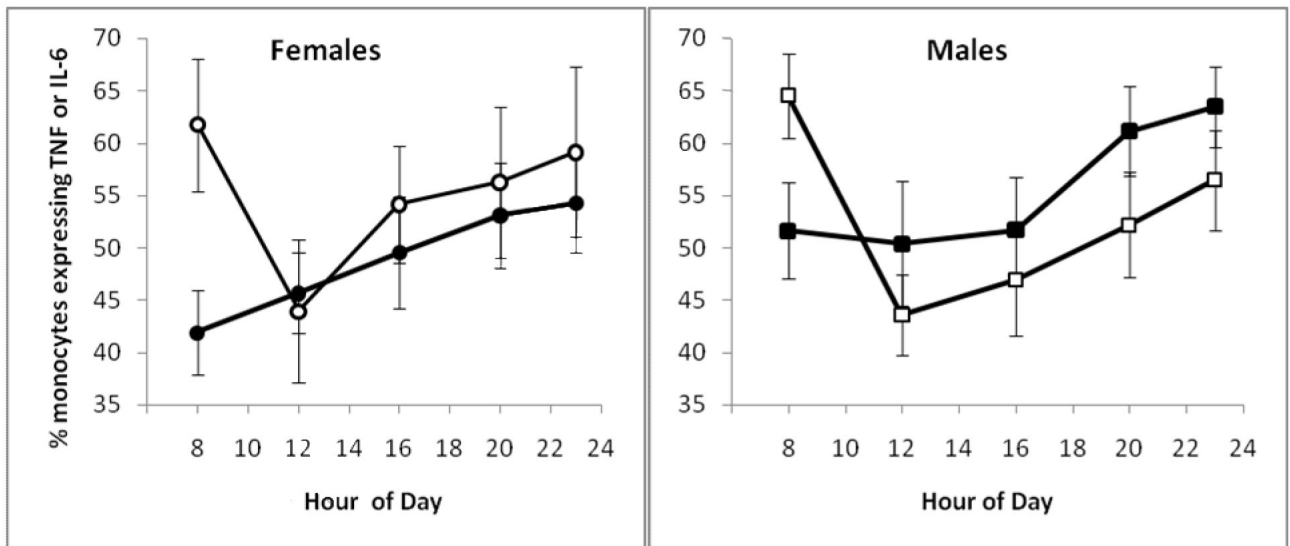


Figure 2.

Sex differences in aggregate expression of interleukin (IL) 6 and tumor necrosis factor- α (TNF) in lipopolysaccharide-stimulated CD14+ cells between baseline (●—●) and partial sleep deprivation (○—○)PSD conditions. Significant condition \times sex interactions in the aggregate expression of TNF- α and IL-6 was found at two timepoints: 20:00 ($F(1,19.0) = 6.0, P < 0.05$) and 23:00 ($F(1,19.9) = 7.4, P < 0.05$) in which females showed increases in aggregate expression of TNF- α and IL-6 as compared to decreases in men. Data are represented as mean \pm SEM.