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Original Article Impact of partial sleep deprivation on immune markers

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ABSTRACT

Background: Sleep quality is considered to be an important predictor of immunity. Lack of sleep therefore may reduce immunity, thereby increasing the susceptibility to respiratory pathogens. A previous study showed that reduced sleep duration was associated with an increased likelihood of the common cold. It is important to understand the role of sleep in altering immune responses to understand how sleep deprivation leads to an increased susceptibility to the common cold or other respiratory infections. *Objective:* We sought to examine the impact of partial sleep deprivation on various immune markers.

Patients and methods: Fifty-two healthy volunteers were partially sleep deprived for one night. We took blood samples before the sleep deprivation, immediately after, and 4 and 7 days after sleep deprivation. We measured various immune markers and used a generalized estimating equation (GEE) to examine the differences in the repeated measures.

Results: CD4, CD8, CD14, and CD16 all showed significant time-dependent changes, but CD3 did not. The most striking time-dependent change was observed for the mitogen proliferation assay and for HLA-DR. There was a significant decrease in the mitogen proliferation values and HLA-DR immediately after the sleep deprivation experiment, which started to rise again on day 4 and normalized by day 7.

Conclusions: The transiently impaired mitogen proliferation, the decreased HLA-DR, the upregulated CD14, and the variations in CD4 and CD8 that we observed in temporal relationship with partial sleep deprivation could be one possible explanation for the increased susceptibility to respiratory infections reported after reduced sleep duration.

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1. Introduction

Sleep is commonly considered a restorative process with supportive influences on immune functions [1]. A modest amount of sleep loss alters molecular processes that drive cellular immune activation and induce inflammatory cytokines and inflammatory markers [2,3]. Experimental studies have demonstrated that sleep deprivation results in poorer immune function (i.e., reduced natural killer cell activity, suppressed IL-2 production) [2,4,5], as well as increased circulating levels of inflammatory markers (i.e., IL-6, tumor necrosis factor α [TNF- α], C-reactive protein) [4,6,7]. Sleep loss induces a functional alteration of the monocyte proinflammatory cytokine response [2,4]. Sleep deprivation also has been found to attenuate antibody response to immunizations [8].

Because sleep quality is considered to be an important predictor of immunity, in turn it also may increase one's susceptibility to the

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common cold [9]. A graded average sleep duration was associated with an increased likelihood of a cold [9]. The common cold is the most frequent illness managed in general practice [10], with an enormous economic impact [11,12].

It is important to understand the role of sleep in altering immune responses to understand how sleep deprivation leads to an increased susceptibility to the common cold or other respiratory infections. In our study, we sought to examine the impact of partial sleep deprivation on various immune markers.

2. Methods

2.1. Participants

Healthy volunteers between the age of 21 and 65 years not involved in shift work or night duties were recruited by local advertisement from the general population in Singapore. Equal numbers of men and women were selected. Specific exclusion criteria included sleep disorders, use of sleeping tablets within the past 8 weeks, long-haul flight within the past 8 weeks, any current



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illness (i.e., current respiratory infections, current or history of psychiatric disease), alcohol abuse, autoimmune diseases or any diseases associated with immunosuppression, immunosuppressive therapy (i.e., steroids, inhaled steroids), and any endocrine disorders. Written informed consent was provided by all subjects. The study was approved by the institutional review board of Singapore.

2.2. Test site and conditions

The overnight partial sleep deprivation experiment started at 9:00 pm and lasted for 10 h, until 7:00 am. All subjects stayed in one room without beds and remained seated throughout most of the night. Subjects were given a light meal and snacks and drinks. Sleeping tablets and alcohol- and caffeine-containing beverages were not permitted. The study coordinator stayed in the room for continuous supervision, and the subjects played card games or talked for most of the night. If subjects did fall asleep, the study coordinator recorded the time and disrupted their sleep after a maximum of 2 h. A total sleeping time of a maximum of 3 h was allowed. The total sleeping time was recorded for every subject. Subjects recorded their subjective stress and fatigue ratings on a Likert scale (values 1-4 with 4 being most stressed or tired, respectively) at baseline, day 1, and day 7. Venous samples were taken before and after the night with sleep deprivation and again on days 4 and 7. To exclude circadian variation, blood samples were taken at the same time of the day at 11:00 am.

2.3. Laboratory analyses

Venous samples were analyzed for complete blood cell count, immunologic markers, HLA-DR expression, and the mitogen proliferation assay. White blood cells were separated from erythrocytes using the hemolysis method [13]. Flow cytometry analysis was performed using the staining protocol, as previously described [14]. CD3, CD4, CD8, CD14, and CD16 lymphocyte blood cell counts were measured. In addition, we examined the expression of HLA-DR on monocytes [15].

2.3.1. Mitogen assay

Lymphocyte proliferation in response to the nonspecific mitogen phytohemagglutinin was determined by ³H-thymidine incorporation. Peripheral blood mononuclear cells were isolated by the Ficoll-Paque (specific gravity, 1.077) gradient density method. The peripheral blood sample was centrifuged at 600 g at 18 °C for 20 min. The peripheral blood mononuclear cell layer was collected and washed with phosphate buffer saline solution containing 0.4% sodium citrate to remove platelets. The cells were resuspended in HyClone RPMI 1640 media supplemented with 10% fetal calf serum (PAA Laboratories), 100 U/mL penicillin, and 100 µg/mL streptomycin (Gibco) at a final concentration of 1 × 10⁶/mL. The mitogen phytohemagglutinin (Sigma–Aldrich, Singapore) was added at a final concentration of 2 µg/mL for 48 h

Marker	Description
CD3	Lineage marker: T cells
CD4	Lineage marker: helper T cells
CD8	Lineage marker: cytotoxic T cells
CD14	Lineage marker: monocytes
CD16	Lineage marker: neutrophils
HLA-DR	Major histocompatibility complex class II molecule
Mitogen proliferation assay	Functional evaluation of the immune system

to stimulate lymphocyte proliferation and levels of proliferation quantitated by a ³H-thymidine incorporation assay. Table 1 summarizes the markers and their main functions.

2.4. Statistical analysis

We used the generalized estimating equation (GEE) to examine the difference in parameter estimates at days 1, 4, and 7 with baseline. The advantage of GEE models is that they allow for the correlation within each subject across the follow-up periods [16]. Because each subject was measured several times in the course of our study, such a correlation should not be ignored; therefore, GEE is the most appropriate statistical tool. Because the outcome is continuous, we specified a Gaussian distribution for the family, along with an identity link for the model. We used the exchangeable correlation matrix in the model. Data analysis was performed in Stata V10.2 (StataCorp, College Station, TX, USA) and level of significance was set at 5%.

3. Results

Fifty-two subjects (26 men and 26 women) with a mean age of 31 years (range, 21–65 y) were included in our study. The median sleeping time was 2 h (range, 0–3 h). The mean stress ratings on the Likert scale at baseline and days 1 and 7 were 2.12, 2.98, and 2.19. The mean ratings for fatigue at baseline and on days 1 and 7 were: 2.53, 3.82, and 2.57. Study subjects therefore showed the

Table 2

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Generalized estimating equations showing the differences on days 1, 4, and 7 after a night of partial sleep deprivation compared to the day before sleep deprivation (baseline).

Parameter	Mean Difference	95% Confidence interval		P value
CD3				
Day 1	2.45	-0.84	5 73	144
Day 4	-1.18	-4.46	2.11	.482
Day 7	2.03	-1.25	5.32	.225
Before (mean value)	65.96			
CD4				
Day 1	67.05	14 04	120.05	013
Day 4	-77.96	-130.97	-24 96	004
Day 7	-19.56	-72.57	33.44	.469
Before (mean value)	678.55			
CD8				
Day 1	51 55	16.90	86 19	004
Day 4	_41.67	-76.32	_7.03	018
Day 7	-16.48	-51.13	18 16	351
Before (mean value)	337.14	51115	10110	1001
CD14				
Day 1	1 14	0.39	1 90	003
Day 4	-1.15	-1.91	-0.40	.003
Day 7	0.12	-0.64	0.10	765
Before (mean value)	4.65	0.01	0.07	.705
CD16				
Day 1	-4 69	_9.57	0.20	06
Day 4	7.05	3.03	12 79	.00
Day 7	0.71	-4 17	5 59	775
Before (mean value)	76.59		0100	
ער א וע				
Day 1	_7 34	-10.07	-4 62	< 001
Day 4	-4.85	-7.57	-2.12	< 001
Day 7	0.40	-2.32	3 13	771
Before (mean value)	27.20	2.52	5.15	.,,1
Mitogen				
Day 1	_39996.85	_48281 01	_31711.80	< 001
Day 4	-15403.24		_7118.19	< 001
Day 7	-4072.50	-12357 55	4212.56	335
Before (mean value)	82173.13	12337.33	1212.50	



Fig. 1. Lymphocyte proliferation with stimulation from phytohemagglutinin on days 1, 4, and 7 after a partially sleep-deprived night in comparison to the baseline (day 0).

highest extent of fatigue and stress immediately after the partially sleep-deprived night.

The results of the complete blood cell count for all subjects were within reference range at baseline, and there were no significant changes in the absolute values and proportion of leucocytes, lymphocytes, and eosinophils immediately after the partial sleep deprivation or after 4 or 7 days (data not shown). Partial sleep deprivation over one night also did not result in any changes in the T-cell marker CD3. However, there were statistically significant changes for CD4 and CD8, which showed an increase immediately after the experiment but then decreased by day 4 and returned to baseline levels by day 7 (Table 2). CD14 increased on the day after the partially sleep-deprived night and then significantly decreased below the baseline, whereas CD16 had changes in the opposite direction (Table 2).

The most striking time dependent change was observed for the mitogen proliferation assay and for HLA-DR. There was a significant decrease in the mitogen proliferation values immediately after the sleep-deprivation experiment, which started to rise again on day 4 and normalized by day 7 (Table 2; Fig. 1). HLA-DR significantly decreased on days 1 and 4 after the sleep-deprived night and normalized to baseline values by day 7 (Table 2).

4. Discussion

We studied the effect of a single night with partial sleep deprivation (sleep duration <3 h) on various immune markers and on functional leucocyte proliferation assays. Because partial sleep deprivation in daily life is more common than total sleep deprivation, we decided to only measure the effect of partial sleep deprivation. All measured markers showed statistically significant changes, except for CD3. CD4 and CD8 both immediately increased after the sleep deprivation then decreased below the baseline on day 4 and normalized to baseline values by day 7. Variations in healthy volunteers over time in CD4 counts have been described [17]; however, the initial activation of both CD4 and CD8 on day 1 and subsequent suppression of both markers by day 4 were statistically different to the baseline and both normalized after 7 days to baseline values. Therefore, these changes are unlikely to be a normal variation but appear to be the result of a time-dependent transient effect of sleep deprivation on early CD4/CD8 activation, followed by short-lived suppression. The clinical importance of these findings is unclear. Because the absolute values of CD4 did not fall below clinically relevant values of 500, the depression of CD4 on day 4 therefore does not present true severe immune suppression but only transient mild immune suppression.

The protein CD14 is a membrane-bound protein on the surface of mononuclear cells which recognizes bacterial lipopolysaccharide [18]. In an animal model, CD14 was shown to be critical to initiate neutrophil recruitment into the airways and was found to be an essential mediator of lipopolysaccharide-induced airway disease [19]. The upregulated CD14 that we found immediately after a sleep-deprived night may act as a marker for disease activity.

CD16 can be found on the surface of natural killer cells and neutrophil leukocytes [20]. The phenotypic changes that we saw after sleep deprivation possibly provide a cellular basis for an increase in natural killer cell activity, which is observed in the common cold [21].

Furthermore, we found significant changes in HLA-DR and the mitogen proliferation assay, with the most predominant change on the first day after sleep deprivation, which started to increase again by day 4 and normalized by day 7. HLA-DR is an antigen presentation molecule present on antigen-presenting cells such as monocytes and B cells. An increased level of HLA-DR expression is normally associated with activation in antigen-presenting cells in response to proinflammatory signals, potentiating the cell's ability to induce stronger T-cell proliferation and respond to maturation signals [15]. Likewise a decrease in HLA-DR expression is associated with anti-inflammatory or immunosuppressive signals, which can potentially lead to a decreased ability of monocytes to induce T-cell proliferation. Low levels of HLA-DR expression on peripheral blood monocytes have been demonstrated to correlate with the risk for infection in surgery, trauma, and adult transplant patients [22]. The decrease in HLA-DR expression after partial sleep deprivation may be a potential signal of an increased susceptibility to infection.

The lymphocyte proliferative response can be viewed as a functional evaluation of the immune system [23]. The lymphocyte proliferation assay showed a clear linear time-dependent response in our GEE models. There was a marked decrease following the sleep deprivation that rose on day 4 and normalized by day 7 (similar levels compared to baseline).

We are unsure of the clinical implications of this transient decrease for HLA-DR and mitogen proliferation. However, both indicate a certain degree of immunosuppression and decreased ability to induce T-cell proliferation. Impaired immune function in healthy individuals possibly represents a suboptimal level of immune functioning that should not be compared with depressed immune status, which may occur in individuals who are immunosuppressed.

Our study showed an inherent variability in the parameters studied. Possible solutions include increasing the sample size to even out the noise or adding studies to determine the intraclass coefficient. However, the method that we chose was the GEE, which is a robust statistical approach used to longitudinally analyze how subjects are measured at different points in time. The advantage of GEE models is that they allow for the correlation within each subject across the follow-up periods [16], and the comparisons therefore are in line with the baseline of each subject, which should reduce the possibility of spurious results.

In summary, although our study does not answer the question of whether or not these changes result in clinically relevant outcomes, we do postulate that the transiently impaired mitogen proliferation, the decreased HLA-DR, the upregulated CD14, and the variations in CD4 and CD8 that we observed in temporal relationship with partial sleep deprivation could be one possible explanation for the increased susceptibility to respiratory infections reported after reduced sleep duration. Further studies are needed to investigate if the transient immune impairment observed in our study indeed translates to a higher risk for acquiring respiratory infections.

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Conflict of interest

The ICMJE Uniform Disclosure Form for Potential Conflicts of Interest associated with this article can be viewed by clicking on the following link: http://dx.doi.org/10.1016/j.sleep.2013.07.001.

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