Effects on prolactin secretion and binding to dopaminergic receptors in sleep-deprived lupus-prone mice

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Sleep disturbances have far-reaching effects on the neuroendocrine and immune systems and may be linked to disease manifestation. Sleep deprivation can accelerate the onset of lupus in NZB/NZWF₁ mice, an animal model of severe systemic lupus erythematosus. High prolactin (PRL) concentrations are involved in the pathogenesis of systemic lupus erythematosus in human beings, as well as in NZB/NZWF₁ mice. We hypothesized that PRL could be involved in the earlier onset of the disease in sleep-deprived NZB/NZWF₁ mice. We also investigated its binding to dopaminergic receptors, since PRL secretion is mainly controlled by dopamine. Female NZB/NZWF₁ mice aged 9 weeks were deprived of sleep using the multiple platform method. Blood samples were taken for the determination of PRL concentrations and quantitative receptor autoradiography was used to map binding of the tritiated dopaminergic receptor ligands [3H]-SCH23390, [3H]-raclopride and [3H]-WIN35,428 to D₁ and D₂ dopaminergic receptors and dopamine transporter sites throughout the brain, respectively. Sleep deprivation induced a significant decrease in plasma PRL secretion (2.58 ± 0.95 ng/mL) compared with the control group (25.25 ± 9.18 ng/mL). The binding to D₁ and D₂ binding sites was not significantly affected by sleep deprivation; however, dopamine transporter binding was significantly increased in subdivisions of the caudate-putamen - posterior (16.52 \pm 0.5 vs 14.44 \pm 0.6), dorsolateral (18.84 \pm 0.7 vs 15.97 ± 0.7) and ventrolateral (24.99 ± 0.5 vs 22.54 ± 0.7 µCi/g), in the sleep-deprived mice when compared to the control group. These results suggest that PRL is not the main mechanism involved in the earlier onset of the disease observed in sleepdeprived NZB/NZWF₁ mice and the reduction of PRL concentrations after sleep deprivation may be mediated by modifications in the dopamine transporter sites of the caudate-putamen.

Key words: Systemic lupus erythematosus; NZB/NZWF₁ mice; Prolactin; Stress; Autoradiography; Dopamine transporter

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Introduction

Systemic lupus erythematosus (SLE) is a prototype autoimmune rheumatic disease, which presents many immunological abnormalities, such as B cell hyperactivity, antinuclear antibodies (ANA), and immune complex deposition that can lead to arthritis, skin rash, and glomerulone-phritis. The strongest risk factor for the development of SLE is gender, since it tends to develop or is exacerbated during pregnancy and the postpartum period (1). The

autoimmune-prone NZB/NZWF₁ mouse is an excellent model for SLE. Like humans with SLE, these mice display a pathognomonic ANA response that includes anti-double-stranded DNA, and they spontaneously develop fatal glomerulonephritis. Also, similar to humans, the disease is most frequent in female mice (2).

Accumulating evidence suggests that prolactin (PRL) is involved in the pathogenesis of SLE. PRL plays a significant role in the regulation of the humoral and cellular immune responses in physiological as well as pathological states,

300 B.D. Palma et al.

such as autoimmune diseases (3). Increased serum PRL levels have been reported in lupus patients of both genders, and have been associated with accelerated disease expression in lupus-prone mice (4,5). The primary signal for PRL secretion is under tonic inhibitory control by hypothalamic dopamine (for a review, see Ref. 6). Since elevated secretion of PRL is common in SLE, we hypothesized that this endocrine imbalance is a consequence of impaired dopaminergic regulation in the central nervous system. The results of some studies are consistent with the hypothesis that lupus-like disease compromises dopaminergic neurotransmission in the central nervous system (7,8).

SLE patients display a variety of neurologic manifestations, which may include sleep disturbances (9). According to Valencia-Flores and colleagues (10), these patients are sleepier during the day by virtue of sleep fragmentation, with more arousals and sleep stage transitions. In addition, the disease is exacerbated by sleep disruption. Recent data from our laboratory indicated that the NZB/ NZWF₁ mice subjected to sleep deprivation (SD) show an earlier onset of the disease reflected by increased numbers of ANA (11). There is evidence to support the view that sleep disturbances lead to hormonal, neurochemical and immunological alterations that may be linked to disease manifestation (12-16). For instance, we recently demonstrated that there was an increase in circulating levels of corticosterone in NZB/NZWF₁ animals as the disease progressed, and this effect was more evident in sleep-deprived mice (17).

In view of these considerations, the aim of the present study was to examine the impact of SD on the pattern of PRL secretion as well as the regulation of dopaminergic receptors in multiple brain regions in NZB/NZWF $_1$ mice. We were particularly interested in the hypothesis that SD may lead to hormonal and neurochemical changes since these factors are associated with lupus onset. To this end, we employed a well-established procedure for producing SD and examined its effects on PRL secretion. This was complemented by quantitative autoradiographic analyses of [3 H]-SCH23390, [3 H]-raclopride and [3 H]-WIN35,428 binding in order to detect possible changes in D $_1$ and D $_2$ receptors and in dopamine transporter (DAT) sites, respectively, throughout the brain of sleep-deprived mice after SD.

Material and Methods

Animals

New Zealand black (NZB, females) and New Zealand white (NZW, males) mice were obtained from Universidade of São Paulo (São Paulo, SP) and were mated in our Research Laboratory to produce NZB/NZWF₁ hybrids.

After weaning, NZB/NZWF $_1$ mice were housed in groups of 6 in plastic cages filled with hardwood bedding, receiving water and rodent chow *ad libitum*. The animals were kept in a room with controlled lighting (12-h light/dark cycle) and temperature (24 \pm 2°C). Due to the fact that murine lupus shows a preponderance in females, only this gender was used in the present study. All procedures were approved by the Ethics Committee of UNIFESP (CEP #1163/01) and carried out in accordance with the rules and regulations on animal care of the National Institutes of Health (http://www.nih.gov/).

Sleep deprivation

Female NZB/NZWF₁ mice aged 10 weeks (a period when they were considered to be healthy) were subjected to SD using the platform method. The method of SD used was an adaptation of the multiple platform method, originally developed for rats (18). The technique is based on the muscle atonia that accompanies paradoxical sleep (19). Briefly, 12 narrow circular platforms (3 cm in diameter) were placed inside a tiled tank (41 x 34 x 17 cm) filled with water to within 1 cm below the upper border of the platform. Groups of 6 mice were placed on the platforms in each tank, an arrangement that allowed them to move inside the tank, jumping from one platform to the other. In this procedure, the animals are aroused from sleep when the loss of muscle tone leads them to fall off the platform. This method produces a consistent amount of sleep reduction in mice (20).

Mice were randomly assigned to two groups containing 12 mice each: control mice remained in their home-cages in the SD room and sleep deprivation (SD) mice were deprived of sleep for two periods of 96 h each separated by an interval of three days. During the interval, mice were placed back in their home-cage. Throughout the study both groups had free access to food and water. This experimental protocol was carried out in an attempt to simulate a chronic condition of SD (similar to that which is observed in chronic inflammatory disease) (11).

Effect of SD on PRL secretion in NZB/NZWF₁ mice (Experiment 1)

Immediately after the end of SD, the animals (N = 12) were rapidly decapitated and trunk blood was collected into tubes containing EDTA.

Hormone determination. Immediately after sampling, blood was centrifuged at 2500 rpm at 4°C for 10 min and plasma was separated and stored at -80°C. Radioimmunoassay for mouse PRL was performed by the National Hormone and Peptide Program (USA). The detection limit of the assay was 1.0 ng/mL and intra-assay variation was 7%.

Assessment of D₁ and D₂ receptors and DAT binding after SD in NZB/NZWF₁ mice (Experiment 2)

Immediately after the end of SD and decapitation of the animals in Experiment 1, the brain was rapidly removed, frozen over dry ice, and stored at -80°C.

Autoradiography procedures. Coronal cryostat sections (20 µm) were cut at -18°C from the olfactory bulbs to the substantia nigra, mounted onto lysine-coated slides and then stored at -80°C. Binding assays for D₁ and D₂ receptors and DAT followed the procedures of Nobrega et al. (21) and Wilson et al. (22). Briefly, to remove the endogenous ligands, the slide-mounted sections were pre-incubated at room temperature in 50 mM Tris buffer, pH 7.4, for 30 min for D₁ binding; in 50 mM Tris buffer, pH 7.4, for 15 min for D₂ binding, and in 25 mM Tris buffer, pH 7.7, for 20 min for DAT. To label D₁ sites, sections were incubated with 2 nM [³H]-SCH23390 (Perkin Elmer, USA; 85 Ci/mmol) for 90 min at 37°C. For D₂ receptors, the sections were incubated with 2 nM [3H]-raclopride (Perkin Elmer; 87 Ci/mmol) for 120 min at room temperature. For DAT sites, the sections were incubated with [3H]-WIN35,428 (Perkin Elmer; 85.6 Ci/mmol) for 120 min at room temperature.

Non-specific binding was defined as binding in the presence of 2 μ M butaclamol (Sigma, USA), 10 μ M sulpiride (Sigma) or 30 μ M cocaine (Sigma), for D₁, D₂ and DAT sites, respectively. Slides were then rinsed in cold buffer, followed by cold distilled water, then air-dried and exposed to Kodak Biomax (Scientific Imaging Film) for 4 weeks for [³H]-SCH23390 and [³H]-WIN35,428, or 5 weeks for [³H]-raclopride in the presence of calibrated standards. Densitometric analyses were performed using an M2 MCID system (Imaging Research, Canada) on coded films. Anatomical regions were defined according to the atlas of Franklin and Paxinos (23) and analyzed without knowledge of the group membership of the animals.

Statistical analysis

In both experiments, data were analyzed by the Student *t*-test. Data are reported as means \pm SEM. A P value ≤ 0.05 was considered to be statistically significant.

Results

Effect of SD on plasma PRL secretion in NZB/NZWF₁ mice

Figure 1 shows that PRL concentrations were ten times lower (2.58 ng/mL) in sleep-deprived mice compared to control mice (25.25 ng/mL; P < 0.02).

Assessments of D₁ and D₂ receptors and DAT binding after SD in NZB/NZWF₁ mice

[3H]-SCH23390 and [3H]-raclopride binding to D₁ and

 D_2 receptors. Binding to D_1 and D_2 receptors did not differ statistically among sleep-deprived mice and the control group in any of the brain regions analyzed (Tables 1 and 2).

[³H]-WIN35,428 binding to DAT sites. As shown in Table 3, [³H]-WIN35,428 binding was significantly increased in three subdivisions of the caudate-putamen in the sleep-deprived mice when compared to the control group. Figure 2 illustrates increased binding in three subdivisions of the caudate-putamen (posterior, dorsolateral and ventrolateral).

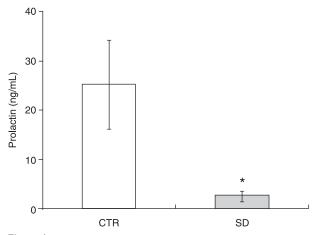


Figure 1. Plasma prolactin concentrations in sleep-deprived (SD) and control (CTR) NZB/NZWF₁ mice. Data are reported as means \pm SEM. *P < 0.02 compared to control (Student *t*-test).

Table 1. Sleep deprivation has no effect on [³H]-SCH23390 binding to D1 receptors.

	Control (N = 8)	Sleep deprivation (N = 10)
Caudate putamen		
Anterior	29.58 ± 0.98	29.41 ± 0.65
Posterior	24.30 ± 0.48	22.37 ± 0.74
Dorsomedial	26.42 ± 1.02	26.70 ± 0.79
Dorsolateral	26.92 ± 0.94	27.40 ± 0.71
Ventrolateral	28.49 ± 1.11	29.32 ± 0.80
Nucleus accumbens	23.72 ± 1.05	25.61 ± 0.49
Core	26.57 ± 0.92	27.53 ± 0.62
Shell	22.77 ± 1.12	23.54 ± 0.69
Olfactory tubercle	26.52 ± 1.09	28.78 ± 0.77
Substantia nigra	15.16 ± 0.56	15.58 ± 0.39
Reticular part	15.80 ± 0.68	16.50 ± 0.35
Compact part	13.00 ± 0.40	13.26 ± 0.43
Lateral part	6.82 ± 0.32	8.00 ± 0.65
Ventral tegmental area	2.47 ± 0.25	2.71 ± 0.15
Nucleus of the ansa lenticularis	9.44 ± 0.27	8.77 ± 0.32
Globus pallidus	3.20 ± 0.17	2.86 ± 0.17
Amygdala	12.10 ± 0.32	12.55 ± 0.58

Data are reported as means ± SEM (pmol/g tissue). There were no statistical differences between sleep-deprived and control mice (Student *t*-test).

302 B.D. Palma et al.

Table 2. Sleep deprivation has no effect on $[^3H]$ -raclopride binding to D_2 receptors.

	Control (N = 8)	Sleep deprivation (N = 10)
Caudate putamen		
Anterior	8.30 ± 0.15	8.79 ± 0.27
Posterior	9.46 ± 0.19	9.32 ± 0.22
Dorsomedial	8.52 ± 0.19	8.46 ± 0.26
Dorsolateral	11.53 ± 0.19	11.48 ± 0.38
Ventrolateral	11.53 ± 0.19	12.75 ± 0.46
Nucleus accumbens	5.57 ± 0.19	5.79 ± 0.21
Core	5.68 ± 0.37	6.18 ± 0.20
Shell	5.21 ± 0.32	5.34 ± 0.30
Olfactory tubercle	6.14 ± 0.16	6.25 ± 0.16
Substantia nigra	3.41 ± 0.12	3.13 ± 0.17

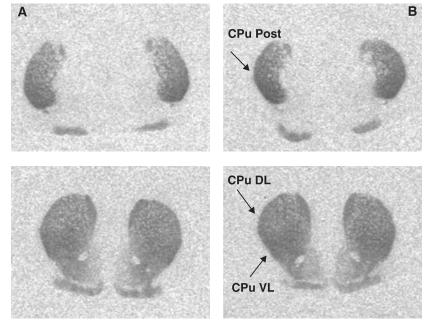
Data are reported as means \pm SEM (pmol/g tissue). There were no statistical differences between sleep-deprived and control mice (Student t-test).

Table 3. [³H]-WIN35,428 binding to dopamine transporter.

	Control	Sleep deprivation (N = 10)
	(N = 8)	
Caudate putamen		
Anterior	15.05 ± 0.53	16.12 ± 0.69
Posterior	14.44 ± 0.65	16.52 ± 0.51*
Dorsomedial	15.84 ± 0.88	16.40 ± 0.67
Dorsolateral	15.97 ± 0.72	18.84 ± 0.73*
Ventrolateral	22.54 ± 0.71	24.99 ± 0.54*
Nucleus accumbens		
Core	14.68 ± 0.60	15.15 ± 0.67
Shell	7.25 ± 0.26	7.30 ± 0.29
Olfactory tubercle	12.36 ± 0.48	13.06 ± 0.39
Substantia nigra		
Compact part	7.90 ± 0.30	7.69 ± 0.40
Reticular part	3.09 ± 0.15	2.76 ± 0.14
Ventral tegmental area	7.95 ± 0.09	7.92 ± 0.41

Data are reported as means \pm SEM (pmol/g tissue). *P < 0.05 compared to control (Student *t*-test).

Figure 2. Illustration of increased dopamine transporter binding in the caudate-putamen nucleus after sleep deprivation. *A*, Control; *B*, sleep deprivation. CPu Post = posterior caudate-putamen; CPu DL = dorsolateral caudate-putamen; CPu VL = ventrolateral caudate-putamen.



Discussion

There is increasing evidence that PRL can exacerbate SLE, particularly, in experimental models (5). Previous data from our laboratory indicated that the NZB/NZWF₁ mice subjected to SD had an earlier onset of the disease as reflected by an increased number of ANA (11). In the present study, we had hypothesized that the SD could increase PRL secretion, and this would be involved in an earlier onset of the disease. However, we observed a

significant reduction in plasma PRL concentrations in sleep-deprived mice.

To our knowledge, the present study provides the first quantification of PRL in NZB/NZWF₁ mice subjected to SD. Previous studies had already reported the effects of SD on PRL secretion. Everson and Crowley (24) reported decreased PRL concentrations in sleep-deprived rats, while Andersen et al. (25) found that the rats subjected to 96 h of SD had higher PRL concentrations than controls. Factors that may contribute to these inconsistencies may be the

use of different animal species and gender or different methods and durations of SD.

There are different possible explanations for the decreased PRL secretion observed after SD in the current study. First, sleep itself is involved in PRL secretion. PRL concentrations are elevated during sleep, even if sleep is delayed (26). Moreover, short periods of SD and sleep fragmentation in humans are associated with lower nocturnal PRL levels in comparison to normal sleep (27). Second, several lines of evidence support the notion that SD is a stressful stimulus (28,29). Thus, SD can be considered a type of biological stress given that sleep is essential to life and to health. It is well known that stress in a number of forms induces PRL secretion (for a review, see Ref. 6). Although PRL reliably increases in response to acute stress, PRL responses to chronic stress become inhibited over continuous exposure (30). The reduction in PRL in this condition is presumably due to an increase in dopamine release at the level of the median eminence (31). Our results agree with the notion that SD, as a type of chronic stress, leads to a reduction in PRL secretion.

Another point that needs to be considered is the fact that PRL secretion is inhibited by an increase of dopaminergic activity (for a review, see Ref. 6). Since this is the main mechanism of PRL secretion (32), we speculated whether altered dopaminergic activity could be involved. Indeed, previous SD studies have reported dopaminergic alterations, including augmented responses to dopaminergic agonists (33,34) and upregulation of brain D₂ dopamine receptors (35). With respect to quantitative receptor autoradiography, we observed that D₁ and D₂ binding sites were not significantly affected by SD in any brain region analyzed in NZB/NZWF₁ mice. However, we observed that DAT binding was significantly increased in subdivisions of the caudate-putamen in sleep-deprived mice. The increase in DAT binding observed in the present study could reflect increased extracellular dopamine concentrations and, as such, could be related to the observed decrease of PRL levels. However, it is important to mention that the regulation of DAT and other neurotransmitter transporters is still not completely elucidated. A review of the literature suggests that DAT play a physiological role in the regulation of dopamine release and consequently of PRL secretion. Females lacking DAT show an impaired ability to nurse their young (36). Moreover, administration of DAT blockers has been shown to inhibit PRL secretion and to disrupt the estrous cycle of the rat (37).

The main regulation of PRL secretion is mediated by three populations of hypothalamic neuroendocrine dopaminergic neurons (tuberoinfundibular, tuberohypophyseal and periventricular-hypophyseal) (6). However, some studies have also linked PRL function to striatal dopaminergic activity. Thus, Malven (37) demonstrated an inhibition of PRL release in conscious sheep following stimulation of nucleus accumbens and caudate-putamen. Also, consistent with this framework is a finding by Harlan et al. (38) that immunoreactive PRL cells are localized in many regions of the rat brain, including the caudate-putamen. In a recent study, Pi et al. (39) detected the expression of the long form of PRL receptor mRNA in the caudate-putamen. It is thus conceivable that the augmented DAT binding in the caudate-putamen observed in our study could be involved in the reduction of PRL secretion after SD.

In summary, the present results add further evidence that SD induces physiological alterations. Our data suggest that PRL is not involved in the early onset of disease in sleep-deprived NZB/NZWF $_1$ mice, and the observed reduction in PRL levels after SD may be mediated by modifications of the DAT sites in the caudate-putamen.

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304 B.D. Palma et al.

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