Diurnal Variation in Secretory Immunoglobulin A Concentration

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The aim of this study was to investigate the effect of time of day on secretory immunoglobulin A (sIgA). Six males participated in this study. Saliva samples were collected for each subject at 06:00, 06:30, 07:00, 07:30, 11:30, 18:00, 22:30 and 06:00 next morning on both routine diet day and prescribed diet day. sIgA concentration was determined by an enzyme linked immunosorbent assay, and total saliva protein concentration was determined by BCA protein kit. There was no significant diurnal variation for saliva flow rate in each day. sIgA concentration and secretion rate were high in the morning and low in the afternoon and evening on both days, and the levels of sIgA concentration at 18:00 and 22:30 were significantly lower than that at 06:00 next morning on the prescribed diet day. Both sIgA concentration and secretion rate had no significant difference between routine diet day and prescribed diet day. The levels of total saliva protein concentration and secretion rate at 18:00 were higher in routine diet day than in prescribed diet day. These results suggested that sIgA concentration showed diurnal variation on prescribed diet day.

Keyword: secretory immunoglobulin A, diurnal variation, protein

INTRODUCTION

Immunoglobulin A (IgA) is the predominant immunoglobulin contained in secretions of the mucosal immune system. Secretory IgA (sIgA) is found in saliva, intestinal secretions, bronchoalveolar lavage fluids, urine, tears, and other mucosal fluids (Brandtzaeg, 1985; Underdown et al., 1994). sIgA is a dimer, containing a polypeptide (J chain) that links two IgA molecules as well as a polypeptide secretory component (SC) responsible for sIgA resistance to proteolysis in an enzyme-rich environment. The molecular weight of sIgA is approximately 2.5 times greater than that of serum IgA (Underdown et al., 1994). Compared with other antibodies that bathe the mucosal surfaces, sIgA has a relatively
short half-life (3–6 days) and a high synthesis rate [66 mg/kg/day] (Schaffer et al., 1991).

The secretory immune system of mucosal tissues such as the upper respiratory tract is considered the first barrier to colonization by pathogenic microorganisms causing upper respiratory tract infection (URTI). SlgA is a major effector of host defense against microorganisms causing URTI. IgA inhibits attachment and replication of pathogenic microorganisms, preventing their entry into the body. The level of IgA in mucosal fluids is more closely correlated with resistance to URTI than serum antibodies (Mackinnon et al., 1994). A decrease in the concentration of slgA was considered as a possible causal factor in the increased susceptibility of athletes to URTI. SlgA has been shown to decrease during a 7-month training program in elite swimmers and correlate inversely with the number of URTI (Gleeson et al., 1999). After acute prolonged exercise, slgA has also been shown to decrease (McDowell et al., 1992; Mackinnon et al., 1993; Steerenberg et al., 1997).

However, virtually all of physiological and psychological variables show evidence of large rhythmic changes (24-hour rhythmicity), and many physiological responses to exercise are influenced by the time of day effect (Hill et al., 1992; Shephard et al., 1997). SlgA concentration has also been shown to exhibit diurnal variation with a significant decrease during the morning hours reaching a plateau by noon, thereafter, slgA concentration remained stable (Walsh et al., 2002). So, studies show a decrease in slgA after prolonged exercise performed in the morning cannot make clear conclusions about whether the decrease in slgA was a result of the exercise per se or due to the passage of time (Walsh et al., 2002). We are interested in studying the effects of exercise on slgA, and we wished to establish the effects of prolonged exercise on slgA responses independent of any diurnal variation in slgA. So as the first step, we investigated the effect of time of day on slgA concentration.

MATERIALS AND METHODS

Subjects and experimental protocol

The subjects were six healthy males who were studying in Fukuoka University (22–42 years old). The aim of the study and the procedures involved were explained to the subjects before their written consent was obtained.

Six subjects were randomly divided into two groups. Saliva samples were collected for each of them on two days, without any exercise. At the first day, one group had a routine diet and another group had a prescribed diet (1950 kcal/d, protein, fat and carbohydrate: 16.6%, 21.5%, 61.9%, respectively). Then, the two groups were exchanged at the third day by having the prescribed diet and routine diet, respectively. When the subjects performed the prescribed diet test, they were required to eat food at the same time and to drink water only. The subjects arrived at the laboratory after an overnight fast on the testing morning. They were allowed to drink water with the exception of the 10-min period before each saliva samples was obtained.

Saliva collection and assay

Each subject was asked to swallow in order to empty the mouth, then, unstimulated whole saliva was collected over a 4 min period into the tubes at 06:00, 06:30, 07:00, 07:30, 11:30, 18:00, 22:30 and 06:00 next morning (n–06:00) on each test. Salivettes were cen-
trifuged at 4700×g for 10 min at 4°C. Saliva mass was recorded to the nearest milligram and samples were stored at -30°C until analysis.

Secretory IgA was determined by sandwich ELISA. Each well of microtiter plates (NUNC–IMMUNO PLATE, Nunc A/S, Denmark) was coated at 37°C for 1 h with 100μl per well of an α chain specific goat anti-human IgA (Biosource, Camarillo, CA) diluted 1:1000 with coating buffer (0.05 M carbonate–bicarbonate buffer, pH 9.6). The wells were washed three times with PBS (0.01 M phosphate buffer containing 0.15 M NaCl, pH 7.2) containing 0.05% Tween–20 (Nacalai Tesque, INC. Kyoto, Japan). Unspecific binding was blocked by incubating wells with 300μl of blocking reagent, 10% Block Ace (BA), for 1 h at 37°C or overnight at 4°C. Plates were washed again as previously described. 50μl of standards or saliva samples diluted with 10% BA were added to the wells and incubated for 1 h at 37°C. Unbound IgA was removed by washing three times. 100μl α chain specific goat anti-human IgA conjugated with horseradish peroxidase (Biosource, Camarillo, CA) were added at dilution 1:1000 in 10% BA and incubated for 1 h at 37°C. After washing, 100μl of substrate buffer (0.006% H2O2–0.1 M citrate buffer (pH 4.0): dH2O: 2, 2’-Azino-bis (3-ethylbenzothiazoline-6-sulfonic Acid) Diammonium Salt (ABTS) = 5: 4.5: 0.5) were added, and plates were incubated for several minutes at room temperature. The color reaction was stopped by the addition of 100μl 1.5% oxalic acid. The plates were read on a microplate reader (MPRA4, TOSHO CORPORATION, Tokyo, Japan) in dual wavelength mode (415 and 492 nm). Control of IgA was included on each plate. To avoid interassay variability, all samples from one athlete were assayed on the same microtiter plate.

The absolute concentration (μg/ml) of IgA was determined by regression analysis. The relationship between known concentrations of IgA (standards) and absorbance was used to interpolate IgA concentration in the samples. The saliva flow rate in ml/min was calculated from measurement of the saliva divided by the collection time. Saliva was weighed to the nearest mg and saliva density was assumed to be 1.00 mg/ml. The secretion rate of IgA (μg/min) was calculated by multiplying the absolute concentration by saliva flow rate (ml/min).

Saliva samples were analysed for total saliva protein (mg/ml) using bovine serum albumin (BSA) as standard (BCA Protein Assay Reagent Kit, PIERCE, Rockford, IL). Saliva protein secretion rate (mg/min) was calculated by multiplying saliva flow rate (ml/min) by saliva total protein concentration (mg/ml).

Statistical
All values are shown as means ± SD. Statistical evaluation of the changes in the concentration of saliva constituents and saliva flow rate was carried out using repeated-measure analysis of variance (ANOVA). Post-hoc analysis was performed using the Turkey test where appropriate. The accepted level of significance was p<0.05.

RESULTS

Diurnal variation in saliva flow rate

Saliva flow rates were ranged from 0.13 to 0.52 ml/min on routine diet day, and 0.15 to 0.63 ml/min on prescribed diet day at the first saliva collect (06:00). The flow rate
increased to the top at 07:30 before the breakfast, and then decreased until the next morning on both days (Fig. 1). But there was no significant diurnal variation for saliva flow rate on each day, and no significant difference between routine diet day and prescribed diet day.

Fig. 1. Diurnal variation in saliva flow rate on routine diet day and prescribed diet day. Values are mean ± SE (n=6). ▲; routine diet; □, prescribed diet; n=06:00, 06:00 at next morning.

Fig. 2. Diurnal variation in sIgA concentration (A) and secretion rate (B) on routine diet day and prescribed diet day. Values are mean ± SE (n=6). ▲; routine diet; □, prescribed diet; n=06:00, 06:00 next morning. Values containing an asterisk mark are significantly different from the values in n=06:00 on prescribed diet day at p<0.01.
Diurnal variation in sIgA

On routine diet day, sIgA concentrations were ranged from 77.29 to 201.63 μg/ml, and were high in the morning and low in the afternoon and evening. The lowest level of sIgA concentration was shown at 22:30. On prescribed diet day, sIgA concentrations at 18:00 and 22:30 were significantly lower than that at next morning (n=06:00) (p<0.01) (Fig. 2A). sIgA secretion rate was also found to be high in the morning and low in the after-

![Graph A: Protein concentration in routine and prescribed diet day](image)

![Graph B: Protein secretion rate in routine and prescribed diet day](image)

![Graph C: sIgA to protein ratio in routine and prescribed diet day](image)

Fig. 3. Diurnal variation in total saliva protein concentration (A), protein secretion rate (B) and sIgA to protein ratio (C) on routine diet day and prescribed diet day. Values are mean ± SE (n=6). ▲: routine diet; □: prescribed diet; n=06:00, 06:00 at next morning. Values containing an asterisk mark are significantly different from the values in prescribed diet day at p<0.05.
noon and evening in each day. The highest levels of slgA secretion rate were shown at 07:30 and 11:30 on routine diet day and prescribed diet day, respectively (Fig. 2B). However, there was no significant diurnal variation for slgA secretion rate on both days. Both slgA concentration and secretion rate did not show significant difference between routine diet day and prescribed diet day.

**Diurnal variation in total saliva protein**

There was no significant diurnal variation for total saliva protein concentration, protein secretion rate and slgA to protein ratio on both days. However, the levels of protein concentration and secretion rate at 18:00 were found to be higher in routine diet day than in prescribed diet day \( (p<0.05) \) (Fig. 3A, 3B), and slgA to protein ratio also showed a peak at 07:00 on each day (Fig. 3C).

**DISCUSSION**

The findings of the present study agreement with the study that showed slgA concentration was high in the morning and low in the evening (Dimitriou et al., 2002). Previous studies that determined diurnal variation in slgA are very limited. In the study of Dimitriou et al. (2002), they also found that slgA secretion rate and saliva flow rate were higher in the evening than in the morning in contrast with slgA concentration. However, Miletic et al. (1996) found that slgA secretion rate were significantly lower in the elderly than in the young, and the level of slgA secretion rate also lower at afternoon than in morning in young person. In addition, Walsh et al. (2002) reported that a significant decrease in slgA concentration in the morning reaching a plateau by 12:30, and a significant decrease in slgA secretion rate at 10:30 reaching a plateau by 12:30.

Athletes appear to be at high risk of developing URTI during intense training and after major competition (Makinnon et al., 1994). A decrease in the concentration of slgA was considered as a possible causal factor in the increased susceptibility of athletes to URTI (Gleeson et al., 1999). After acute prolonged exercise, slgA has been shown to decrease (McDowell et al., 1992), increase (Schouten et al., 1988), or remain unchanged (Walsh et al., 1999). Studies on slgA in response to exercise appear equivocal. However, many physiological responses to exercise are influenced by the time of day effect (Hill et al., 1992; Shophard et al., 1997), and the effects of time of day on the immunoeendocrinological response to exercise are not clear (Trine et al., 1995). Taken together our results and previous studies suggest that experimenters wishing to determine the effects of exercise (or other factors) on the slgA response should control for the diurnal variation in slgA concentration (Walsh et al., 2002).

In addition, some authors express slgA as slgA relative to total saliva protein ratio. This accounted for changes in saliva volume that may have occurred due to drying of the oral surfaces as a result of exercise, and reflected IgA output relative to other proteins (Mackinnon et al., 1993). However, There was no significant diurnal variation for slgA to protein ratio in this study. If slgA is the major component of salivary protein to increase when subjects relax, expressing changes in slgA as a secretion rate seems more appropriate than expressing slgA in relation to changes in total protein concentration (Reid et
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al., 2001). Bishop et al. (2000) reported that carbohydrate and fluid intake influence the slgA and saliva flow rate response to prolonged submaximal exercise. Another study suggested that the exercise–induced change in slgA level was not affected by protein supplements (Krzywickowski et al., 2001). To investigate whether food intake affects slgA level, we also determined the diurnal variation in slgA with the prescribed diet to compare the routine diet. However, there was no significant difference on slgA concentration between prescribed diet and routine diet.

Secretory IgA is the predominant salivary immunoglobulin. The primary function of slgA is opsonization of foreign invaders at the port of entrance to the body (respiratory, gastrointestinal, vaginal, and anal mucosae) and blockade of their infectivity. This is especially important in the case of antiviral immunity in which binding of slgA abolishes the ability of viruses to penetrate into and productively infect cells (Schaffer et al., 1991). Use of slgA as an immune marker has numerous advantages over measurement of T and B cell in the blood. At first, compared to the blood tests, saliva collection do not have any risk for the subjects under study. Saliva is easily retrieved and can be collected as many times daily as is needed for a study. Even small children can be easily tested. Second, slgA synthesis rate is high and half-life is short. Thus any change in slgA concentration can be observed immediately and directly correlated with the causative agent or procedure. In addition, slgA synthesis is T cell dependent and changes in synthesis can be correlated directly to T and B cell activation. At last, there is cooperation between distant compartments in the mucosal immune system, and therefore, the findings in saliva may be generalized to the entire mucosal system (Miletic et al., 1996). In conclusion, these data suggested that slgA concentration showed diurnal variation on prescribed diet day.

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