Changes in MOTS-c Level in the Blood of Pregnant Women with Metabolic Disorders

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Abstract: MOTS-c peptide is a member of the group of mitochondria-derived peptides (MDP). It is a product of the open reading frame in the 12S RNA gene. Due to its features and functions in the body, this peptide is classified as a hormone. The first publications indicated that this hormone improves insulin sensitivity and lowers body weight in obese animals. This suggests that it may be an important peptide in maintaining the body’s energy homeostasis. The aim of our work was to investigate the potential role of MOTS-c peptide during pregnancy, which is a condition prone to metabolic disorders. The research covered healthy, obese women and women with thyroid disorders. The obtained results indicated an increase in the concentration of MOTS-c in the blood of mothers and newborns in the obese group as compared to the healthy control group and a corresponding decrease in the concentration of this peptide in mothers and newborns in the group with hypothyroidism compared to the obese group. Moreover, we also observed a strong positive correlation between the concentration of MOTS-c in maternal blood and in umbilical cord blood. In summary, the MOTS-c peptide shows changes in blood concentration in various physiological states and may, in the future, become an important tool in the fight against metabolic diseases such as obesity or type 2 diabetes.

Keywords: MOTS-c; obesity; mothers’ blood; umbilical cord blood

1. Introduction

MOTS-c is one of the most intriguing peptides and belongs to the mitochondria-derived peptides (MDP) family. It was first discovered in 2015 by Lee et al. [1]. The term “MOTS-c” is an acronym defined as the mitochondrial open-reading-frame of the twelve S rRNA type-c. MOTS-c is the second member of the MDP family and was discovered almost 20 years ago.
The first member of the MDP family, humanin, was discovered in 2001 and 2003 by three independent teams using different methods [2–4]. The unique feature of MOTS-c and humanin peptides is their localization within the sequence coding for mitochondrial 12S RNA. Moreover, based on its systemic actions in organisms, the MOTS-c peptide is classified as a hormone [1] and can serve as a transcription factor [5]. MOTS-c is a product of an open reading frame that is a 51 bp segment, and the mature peptide has 16 aa. Since its discovery, a number of reports describing the influence of MOTS-c on the metabolism have been published. The significant role of MOTS-c was first documented in the works by Lee et al. [1,6,7]. The authors showed that MOTS-c is a crucial element in energy homeostasis and metabolic processes. It was shown to improve diet-induced obesity, decrease muscle insulin resistance, stimulate glucose uptake, and enhance pentose and purine synthesis [1,6]. Moreover, it was shown that the MOTS-c protein level is regulated by androgens (DHT—dihydrotestosterone) in combination with insulin in gravid uterine and placental ferroptosis in PCOS-like rats [8]. Improvement of insulin sensitivity through MOTS-c has been shown to be closely related to the adiponectin pathway [9]. In addition, it has been shown that in pre-diabetic and diabetic individuals, blood levels of MOTS-c decrease significantly [10]. Moreover, MOTS-c can extend the lifespan of cells and possibly the whole organism [11,12]. It is interesting to note that MOTS-c is closely related to physical activity and exercise [13,14].

The importance of MOTS-c in metabolism is a compelling reason to analyze its role in pregnancy, which is associated with an increased risk of metabolic abnormalities [15–17]. Therefore, this study aimed to investigate the potential changes in the levels of MOTS-c in healthy pregnant women and women with metabolic disorders such as obesity and hypothyroidism, and to document the metabolic state of these women by blood biochemical analysis. The second aim of the study was to explore the possible production of MOTS-c in newborns and measure its levels in cord blood shortly after birth.

2. Materials and Methods

2.1. Ethics

The study was conducted according to the principles stated in the Declaration of Helsinki. The protocol was approved by the Clinical Research Ethics Committee of Poznan University of Medical Sciences (approval number 997/18).

2.2. Material

All subjects in this study were Caucasian. The test material used was maternal peripheral blood collected on the day of childbirth (MB) and umbilical cord blood (CB). The parameters recorded for mothers included in the study were as follows: age (years), height (cm), body weight (kg), body mass index (BMI, kg/m²), and body weight just after birth. Obesity was defined as a BMI value of >30 according to the WHO standards [18]. The parameters measured for newborns were as follows: body weight (kg), head circumference (cm), chest circumference (cm), abdominal circumference (cm), thigh circumference (cm), and arm circumference (cm). The data on the characteristics of mothers and newborns are presented in Tables 1 and 2 [19].

Table 1. Characteristics of mothers.

| PARAMETER            | Non-Obese (n-18) | Obese (n-22) | Hypothyroidism (n-17) |
|----------------------|------------------|--------------|-----------------------|
| **BEFORE PREGNANCY** |                  |              |                       |
| Women's age (years)  | 30.09 ± 1.300    | 32.62 ± 1.243| 29.75 ± 1.448         |
| Height (cm)          | 168.3 ± 1.571    | 167.8 ± 1.245| 166.8 ± 1.422         |
| Body weight (kg)     | 63.88 ± 1.204    | 90.08 ± 3.536** | 67.25 ± 3.112## |
| BMI (kg/m²)          | 22.54 ± 0.194    | 32.29 ± 1.107** | 24.32 ± 1.282## |
| **ON THE DAY OF BIRTH** |              |              |                       |
| Body weight (kg)     | 79.58 ± 1.903    | 98.32 ± 3.116** | 80.58 ± 2.861## |

Values are presented as mean ± S.E.M. Statistically significant differences are marked for *p < 0.01 (**). The # and ## symbols were used by analogy but in comparison to the obese group.
2.3. Metabolic and Hormonal Profile in Blood Serum

Metabolic parameters were determined in the blood samples using commercially available colorimetric tests. The levels of glucose, cholesterol (total, high-density lipoprotein, and low-density lipoprotein), and triglycerides (TGs) were measured using kits from Pointe Scientific according to the manufacturer’s instructions. The sample and reagent were analyzed using 96-well microplates after their volume was adjusted using the micro method. Absorbances were read on a Synergy 2 apparatus (Biotek, Winooski, VT, USA). Hormone concentrations were determined by enzyme-linked immunosorbent assay (ELISA) or radioimmunooassay (RIA) kits—serum MOTS-c levels were determined using ELISA kits (test range 4.985–600 pg/mL, sensitivity: 2.659 pg/mL, cat. no: 201-11-3562, Sunred, Shanghai, China), while other hormones were measured by the following tests: insulin—Human Insulin-Specific RIA (test range: 2–200 µU/mL; cat. no: HI-14K, Merck Millipore, Burlington, MA, USA), adiponectin—Adiponectin ELISA (test range: 0.6–31,000 µg/L; cat. no: E09, Mediagnost, Reutlingen, Germany), and leptin—Multi-Species Leptin RIA (test range: 1–50 ng/mL; cat. no: XL-85K, Merck Millipore, Burlington, MA, USA). For RIA tests, gamma radiation was quantified using the Wallac Wizard 1470 Gamma Counter (Perkin Elmer, Waltham, MA, USA). ELISA measurements were carried out on a Synergy 2 microplate reader (Biotek, Winooski, VT, USA) [20,21].

2.4. Statistical Analysis

Statistical analysis was performed using unpaired Student’s t-test and Mann–Whitney test. All data were normally distributed except for the hypothyroidism groups, where the Mann–Whitney test was used. Statistical significance was assumed at \( p < 0.05 \) and \( p < 0.01 \) and denoted by * and **, respectively, in comparison to the control group. The # and ## symbols were used by analogy but in comparison to the obese group. On the column graph, the letters are indicators of statistical differences and \( p < 0.05 \) and \( p < 0.01 \) are denoted by lowercase and uppercase letters, respectively, in comparison to the columns indicated by the letters. Correlations between the levels of MOTS-c in maternal blood and umbilical cord blood were analyzed using Pearson’s correlation model and linear regression. All statistical analyses were conducted in GraphPad Prism 6.0 software (GraphPad Software, San Diego, CA, USA).

3. Results

3.1. Body Weight and General Metabolic Profile

The body weight recorded before pregnancy and BMI unequivocally indicated obesity in one group of analyzed women. On the other hand, female subjects with hypothyroidism showed a lower body weight as well as BMI in comparison to obese subjects. A similar trend of body weight and BMI was also observed in the analyzed women on the day of birth (Table 1). Several parameters measured in the newborns did not show any statistically significant differences (Table 2). A detailed description of the metabolic state of the women included in the study is presented in Table 3. Parameters such as TG and leptin concentrations were found to be significantly elevated, while the adiponectin level was diminished in the serum of the obese women group. These observations support the notion

| PARAMETER                  | Non-Obese       | Obese           | Hypothyroidism |
|----------------------------|------------------|-----------------|----------------|
| Child gender (M/F)         | 9/9              | 10/12           | 10/7           |
| Body weight (kg)           | 3.555 ± 0.146    | 3.544 ± 0.177   | 3.447 ± 0.090  |
| Head circumference (cm)    | 36.11 ± 0.5536   | 36.3 ± 0.5287   | 36.06 ± 0.3147 |
| Chest circumference (cm)   | 34.11 ± 0.5710   | 34.25 ± 0.5519  | 34.06 ± 0.3028 |
| Abdominal circumference (cm)| 33.33 ± 0.8205   | 33.35 ± 0.5679  | 34.65 ± 0.3202 |
| Thigh circumference (cm)   | 12.39 ± 0.2574   | 12.5 ± 0.2565   | 12.24 ± 0.2016 |
| Arm circumference (cm)     | 10.17 ± 0.2021   | 10.22 ± 0.2365  | 10 ± 0.1485    |

Values are presented as mean ± S.E.M.
that this group of participants had obesity. Similar results were observed in women with hypothyroidism: a decrease in the levels of T3, fT3, and T4 and an increase in the level of leptin (Table 3).

Table 3. Metabolic characteristics of the study subjects.

| Parameter               | Non-Obese | Obese | Hypothyroidism |
|-------------------------|-----------|-------|----------------|
|                         | MB CB     | MB CB | MB CB          |
| Glucose (mg/dL)         | 96.74 ± 4.39 | 88.70 ± 4.06 | 103.7 ± 2.62 | 97.32 ± 3.67 | 97.93 ± 2.72 | 90.01 ± 3.97 |
| NEFA (mmol/L)           | 0.651 ± 0.055 | 0.408 ± 0.042 | 0.893 ± 0.094 | 0.418 ± 0.045 | 0.749 ± 0.081 | 0.542 ± 0.070 |
| Cholesterol (mg/dL)     | 197.7 ± 9.46 | 185.1 ± 23.80 | 244.7 ± 8.79 | 172.6 ± 7.15 | 216.9 ± 11.66 | 206.4 ± 16.22 |
| Triglycerides (mg/dL)   | 233.9 ± 24.23 | 135.4 ± 20.40 | 339.3 ± 28.74 | 121.7 ± 22.65 | 267.1 ± 28.43 | 155.5 ± 24.43 |
| Adiponectin (µg/mL)     | 9.95 ± 0.72 | 35.06 ± 2.62 | 7.84 ± 0.60 | 35.22 ± 2.93 | 8.36 ± 0.87 | 32.04 ± 2.75 |
| NEFA (mmol/L)           | 0.651 ± 0.055 | 0.408 ± 0.042 | 0.893 ± 0.094 | 0.418 ± 0.045 | 0.749 ± 0.081 | 0.542 ± 0.070 |
| Leptin (ng/mL)          | 8.09 ± 0.45 | 7.59 ± 0.59 | 9.78 ± 0.28 | 6.84 ± 0.71 | 9.43 ± 0.22 | 6.55 ± 0.87 |
| Insulin (ng/mL)         | 11.42 ± 1.23 | 3.53 ± 0.69 | 16.71 ± 2.76 | 9.25 ± 4.33 | 10.88 ± 1.56 | 3.98 ± 1.03 |
| TSH (mIU/L)             | 1.62 ± 0.18 | 8.95 ± 1.59 | 2.13 ± 0.47 | 9.75 ± 1.87 | 1.57 ± 0.17 | 11.85 ± 2.52 |
| T3 (ng/mL)              | 1.48 ± 0.07 | 0.64 ± 0.07 | 1.44 ± 0.06 | 0.75 ± 0.15 | 1.15 ± 0.09 | 0.67 ± 0.05 |
| fT3 (pg/mL)             | 2.451 ± 0.14 | 1.211 ± 0.13 | 2.235 ± 0.15 | 1.268 ± 0.20 | 1.917 ± 0.15 | 1.235 ± 0.21 |
| T4 (µg/dL)              | 8.76 ± 0.43 | 8.68 ± 0.26 | 8.58 ± 0.31 | 8.10 ± 0.28 | 8.22 ± 0.41 | 8.74 ± 0.25 |
| FT4 (ng/mL)             | 1.13 ± 0.03 | 1.07 ± 0.03 | 1.12 ± 0.03 | 1.15 ± 0.02 | 1.05 ± 0.02 | 1.10 ± 0.02 |

Values are presented as mean ± S.E.M. Statistically significant differences are marked for \( p < 0.05 \) (*) and \( p < 0.01 \) (**) vs. non-obese or \( p < 0.05 \) (#) vs. obese. MB—mother’s blood; CB—cord blood.

3.2. MOTS-c Concentration Changes

No significant differences were found among the MOTS-c concentrations in the maternal blood serum and cord blood serum in the healthy group (Figure 1). Similarly, no differences were noticed in the obese group or in subjects with hypothyroidism. Interesting differences were noted between the MOTS-c levels in the maternal blood serum of the healthy women and obese women, and a significant increase in this level was found in obese women (\( p < 0.048 \), Figure 1). Similarly, an increased concentration of MOTS-c was observed in the cord blood in the obese group (\( p < 0.045 \), Figure 1).

A distinct decrease in MOTS-c levels was observed in maternal blood in the subjects with hypothyroidism in comparison to obese women (\( p < 0.003 \), Figure 1). Similarly, a decrease in MOTS-c concentration was observed in the cord blood in subjects with hypothyroidism compared to the obese group (\( p < 0.039 \), Figure 1).

![Figure 1. Changes in MOTS-c concentration in maternal and cord blood in non-obese and obese mothers. Values are presented as mean ± standard error of the mean (S.E.M.). Statistically significant differences are marked for \( p < 0.05 \) (*) and \( p < 0.01 \) (**) MB—mother’s blood; CB—cord blood.](image-url)
The correlation analysis of the MOTS-c levels determined in all groups showed a strong positive correlation between the hormone levels in maternal blood and cord blood ($r = 0.8812$, $p < 0.001$, Figure 2).

![Figure 2](image.png)

**Figure 2.** Correlation between MOTS-c level in mother’s blood and umbilical cord blood. Values for $r$ and $p$ are indicated in the graph. Solid and dashed lines show the means and 95% confidence intervals, respectively, following linear regression analysis. The $r$-value indicates correlation, and the $p$-value indicates the significance of the correlation.

### 4. Discussion

The obtained results shed some light on the role of MOTS-c in an organism. Until now, this peptide has received relatively little attention, and thus each piece of information could be important in the future. We provide novel findings derived directly from human physiology that add to our understanding of MOTS-c.

First, the biometric body features and biochemical characteristics of the participants were analyzed. The obtained data confirmed the physiological state of the healthy, obese, and hypothyroidism women included in the study. Similarly, biometric analysis was performed for newborns, which showed no significant differences. Furthermore, the metabolic and hormone profiles of all three groups of pregnant women with different metabolic states were explored. Similar parameters were measured in cord blood collected immediately after birth. The main part of the research analyzed the concentration of MOTS-c in maternal and cord blood and the correlation between the levels of MOTS-c in maternal blood and cord blood, which is a component of the newborn circulation.

The potential role of MOTS-c peptide in the regulation of metabolism and energy homeostasis seems to be of importance [1,22,23]. As stated above, MOTS-c is a product of the mitochondrial open reading frame of the 12S RNA gene and thus potentially present in all cells of an organism. The 12S gene is constantly transcribed, and the further formation of MOTS-c results from the subsequent translation mechanisms. This feature can be important for the metabolism of energy substrates and maintaining the energy balance in each cell. These assumptions seem to be true because MOTS-c has been reported to influence the metabolism of both carbohydrates and lipids [1,24,25]. Hence, we decided to investigate the levels of MOTS-c in pregnant women as pregnancy can be considered an important factor modifying the energy balance [26,27].

In our study, we did not observe any changes between maternal blood and cord blood in any of the analyzed groups. Moreover, we found a clear correlation between the MOTS-c levels in maternal blood and cord blood. This may suggest two facts: first, MOTS-c is
expressed in the mother’s tissues and can easily cross the placental barrier; second, MOTS-c is also expressed in the placental tissue and in the fetus but cannot cross the placental barrier. Both assumptions can be true, but further analysis is needed to confirm one of them. Unfortunately, we did not have access to additional tissues (e.g., placental tissue) to at least partially verify these hypotheses, which is the main limitation of our work. The next step is to determine the expression of MOTS-c in the placenta as well as its immunofluorescent localization in the cells of this organ. This is important because MOTS-c has also been shown to regulate the metabolism of human placenta-derived mesenchymal cells [28].

Second, our results showed a clear increase in MOTS-c in obese mothers’ blood in comparison to healthy control women. This observation confirms that MOTS-c is an important factor regulating metabolism, possibly also as a factor in preventing damage to pancreatic islets [29]. An increase in the level of MOTS-c implies that the organism attempts to maintain high glucose utilization and limit the growth of adipose tissue. The increased concentration of MOTS-c in cord blood in the healthy and obese groups supports this assumption and allows it to be extended to the growing fetus as well. Interestingly, a similar increase was not observed in children, men, or non-pregnant women. This is probably due to a significant physiological difference in the state of pregnancy compared to other organisms [23,30].

Third, a decrease in the concentration of MOTS-c was observed in maternal blood and cord blood in subjects with hypothyroidism in comparison to the obese group. Hypothyroidism is generally characterized by a lower metabolism rate and possibly partially by a decrease in the MOTS-c level in the blood. We suppose that lower levels of MOTS-c can be one of the causes and an effect of hypothyroidism, which leads to a lower metabolism rate. In this context, MOTS-c can be a potential activator of metabolism, which could alleviate some of the symptoms of hypothyroidism.

5. Conclusions

In summary, MOTS-c, as a promising factor that plays a crucial role in the energy balance, should be intensively investigated. Understanding the role of this peptide could help to partially explain some of the intricacies of human metabolism and perhaps develop novel therapeutic methods [31].

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Data Availability Statement: The data presented in this study are available on reasonable request from the corresponding author.

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