

# ГЕНЕТИКА ФОЛАТНОГО ЦИКЛУ, РІВЕНЬ ГОМОЦИСТЕЇНУ, ГОРМОНІВ ЩИТОПОДІБНОЇ ЗАЛОЗИ ТА ГІПОФІЗА У КРОВІ ДІТЕЙ ІЗ РАЙОНІВ, ЯКІ МЕЖУЮТЬ З ЧОРНОБИЛЬСЬКОЮ ЗОНОЮ ВІДЧУЖЕННЯ

## FOLATE CYCLE GENETICS, BLOOD HOMOCYSTEINE, THYROID AND PITUITARY HORMONES IN CHILDREN FROM DISTRICTS BORDERING THE CHERNOBYL EXCLUSION ZONE

<sup>1</sup>BANDAZHEVSKIY Yu.I.,  
<sup>2</sup>DUBOVA N.F.

<sup>1</sup>Ecology and Health  
Coordination and  
Analytical Centre,  
Ivankiv, Ukraine

<sup>2</sup>Shupyk National  
Healthcare University  
of Ukraine, Kyiv

The implementation of projects of the European Commission «Health and Ecological Programmes around the Chernobyl Exclusion Zone: Development, Training and Coordination of Health-Related Projects» and that of the Regional Council of Rhône-Alpes (France) enabled to examine the state of health of most children from districts bordering the Chernobyl exclusion zone. At the same time, hyperhomocysteinemia was first found in 73.2% of the examined children aged 12-17 years old from Ivankivskyi district and in 79.8% of the examined children

of the same age from Poliskyi district [1].

Hyperhomocysteinemia or elevation above the physiological level in a blood concentration of a sulfur-containing amino acid homocysteine signals that there is an unfavourable state of metabolism in a body. In adults, it may be combined with a number of severe conditions leading to disability and mortality [2-7]. Given the importance of this issue for medical science and practical health care, it is necessary to investigate the causes of this phenomenon. In particular, genetic changes in the folate

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<sup>1</sup>Бандажевський Ю.І., <sup>2</sup>Дубова Н.Ф.

<sup>1</sup>Координаційний аналітичний центр  
«Екологія і здоров'я», м. Іванків, Україна

<sup>2</sup>Національний університет охорони здоров'я  
України ім. П.Л. Шупика, м. Київ, Україна

**Метою роботи** було визначення взаємозв'язку між станом генетичного апарату фолатного циклу, змістом гомоцистеїну, гормонів щитоподібної залози та аденогіпофіза у крові дітей із Іванківського і Поліського районів Київської області, які межують з Чорнобильською зоною відчуження (ЧЗВ).

**Методи.** Імунохімічний, генетичний, статистичний.

**Результати.** У 158 дітей із Поліського району та 178 дітей із Іванківського визначався вміст у крові гомоцистеїну ( $H_{cy}$ ), тиреотропного гормону гіпофіза (ТТГ), вільного трийодтироніну ( $T_3$ ), вільного тироксину ( $T_4$ ), також оцінювався стан генетичної системи фолатного циклу (ФЦ). Середній вік обстежених склав ( $15,2 \pm 0,9$ ) років (95% ДІ 15,0-15,4 років). З урахуванням генотипів ФЦ було сформовано генетичні підгрупи, засновані на 100% представництві одного конкретного генотипу. Лабораторне обстеження проведено двічі: у квітні і грудні 2015 року (до і після пожеж у ЧЗВ). Встановлено, що у загальній групі дітей, а також у більшості аналізованих генетичних під-

груп рівень  $H_{cy}$  і гормонів щитоподібної залози (ЩЗ) у дітей із Іванківського району був достовірно вищим, ніж у дітей із Поліського. При цьому у загальній групі, а також у більшості генетичних підгруп дітей із Іванківського району між  $H_{cy}$  і ТТГ виявлено прямий кореляційний зв'язок, який був відсутнім у дітей із Поліського району. Рівень  $H_{cy}$  у крові дітей обох районів був достовірно вищим у підгрупі, що містить тільки гомозиготні варіанти алелі Т генетичного поліморфізму MTHFR:677 порівняно з підгрупами, що містять нейтральні алелі цього ж поліморфізму. При цьому були відсутні відмінності щодо ТТГ,  $T_3$  і  $T_4$ .

**Висновки.** Проведений аналіз дозволив встановити синхронну реакцію метаболічних циклів, що забезпечують обмін  $H_{cy}$  і гормонів ЩЗ в організмі підлітків незалежно від генотипів ФЦ. Кореляційний аналіз, а також результати статистичного аналізу свідчать про те, що підвищення рівня  $H_{cy}$  у крові дітей індукувало синтез ТТГ і  $T_3$ . Отримані результати свідчать про тісний зв'язок між гормоногенезом ЩЗ і обміном сірковмісних амінокислот метіоніну і  $H_{cy}$ . Найбільш імовірною причиною підвищення рівня  $H_{cy}$  і внаслідок цього гормонів гіпофізарно-тиреоїдної вісі у крові дітей із районів, що межують з ЧЗВ, є пожежі лісу, який містить довгоіснуючі радіоактивні елементи.

**Ключові слова:** гомоцистеїн, фолатний цикл, генетичні підгрупи, гормони щитоподібної залози і гіпофіза, підлітки, радіоактивно забруднені території.

cycle [8, 9] and thyroid diseases [10] are pointed out as causes of hyperhomocysteinemia. In this regard, it is important to identify a relationship between the state of folate cycle genetic apparatus and blood levels of homocysteine, thyroid and pituitary hormones in children living in districts bordering the Chernobyl exclusion zone. At the same time, one should take into account the effect of an external environmental factor associated with the Chernobyl exclusion zone on the body of children.

**The purpose of the current research was** to identify interrelations between the state of folate cycle genetic apparatus and blood levels of homocysteine, thyroid and adenohypophysis hormones in children from Ivankivskiy and Poliskiy districts, Kyiv region, bordering the Chernobyl exclusion zone.

**Material and methods.** The research was carried out in Poliskiy and Ivankivskiy districts, Kyiv region, whose territories are contaminated with radionuclides as a result of the accident at the Chernobyl nuclear power plant (a  $^{137}\text{Cs}$  soil contamination density is  $<2 \text{ Ci/sq.km}$  [11]). The age of the children at the time of the examination was  $(15.2 \pm 0.9)$  years old (95% CI 15.0-15.4 years old).

158 children from Poliskiy district and 178 children from Ivankivskiy district underwent laboratory examination. At the same time, we assessed blood levels of homocysteine ( $\text{H}_{\text{cy}}$ ), pituitary thyroid stimulating hormone (TSH), free triiodothyronine ( $\text{T}_3$ ), free thyroxine ( $\text{T}_4$ ) and the state of the genetic system of the folate cycle (FC).

The examined children had blood drawn from the ulnar vein after fasting in the morning to carry out the above studies. At the time of blood draws, all of them attended school. Blood from the children of Poliskiy district was drawn on 02.04.2015, from children of Ivankivskiy district on 18.12.2015.

The blood samples were analysed at a laboratory certified under quality standards with a financial support of the Regional Council of Rhône-Alpes (France) and were agreed with the parents.

TSH,  $\text{T}_3$  and  $\text{T}_4$  were determined using an electrochemiluminescent immunoassay (ECLIA) method. Analyser and test kit: Cobas 6000, Roche Diagnostics (Switzerland).

The blood  $\text{H}_{\text{cy}}$  concentration was measured using a chemiluminescent immunoassay (CLIA) method. Analyzer and test kit: Architect 1000 (ABBOT Diagnostics (USA)). The blood  $\text{H}_{\text{cy}}$  level in children of more than  $10 \mu\text{mol/L}$  was defined as hyperhomocysteinemia.

The following allelic variants were identified during genetic analysis of folate cycle: C677T and A1298C of the MTHFR gene (synthesis of the methylenetetrahydrofolate reductase enzyme), A2756G of the MTR gene (synthesis of the  $\text{B}_{12}$ -dependent methionine synthase enzyme) and A66G of the MTRR gene (synthesis of the methionine synthase reductase enzyme). A real-time PCR method was used. Analyser and test kit: DT-96 detecting thermocycler, DNA-Technology (Russia).

Taking into account the folate cycle genotypes, genetic subgroups were formed based on 100% representation of one specific genotype (table 1).

The statistical processing of the findings was performed using the IBM SPSS Statistics 22 software (USA). The arithmetic mean ( $M$ ), standard error of mean ( $m$ ), confidence interval for the mean value (95% CI), median ( $Me$ ), interquartile range

(IQR), minimum and maximum parameter values and percentiles were calculated for the variables analysed.

The distribution hypothesis was tested (a Kolmogorov-Smirnov test). All the studied parameters did not conform to the normal distribution law, thus, a non-parametric Mann-Whitney U test was used to compare values. The statistical significance of variables was assessed by determining a significance level for  $p$  with the help of the statistical software programme. The Student's  $t$ -test was used to compare relative values. The critical level of significance for the null hypothesis ( $p$ ) was set at 0.05. Associations between  $\text{H}_{\text{cy}}$ , TSH,  $\text{T}_3$  and  $\text{T}_4$  values were identified with the help of the Spearman's rank correlation coefficient ( $r_{\text{xy}}$ ). The strength of an association was assessed according to a typical scale: weak – 0 to 0.299; moderate – 0.3 to 0.699; strong – 0.7 to 1.0.

**Results and their discussion.** The level of  $\text{H}_{\text{cy}}$  and thyroid hormones ( $\text{T}_3$  and  $\text{T}_4$ ) was statistically significantly higher in children from Ivankivskiy district than in those from Poliskiy district in a total group of children (tables 2-4).

The  $\text{H}_{\text{cy}}$  level was statistically significantly higher in the blood of children from Ivankivskiy district comprising genetic subgroups 1, 2, 4, 5, 7, 8, 11, 12 than in that of children from Poliskiy district from similar subgroups (tables 2-4).

Table 1  
Subgroups of children with a specific folate cycle genotype

| Subgroup    | Genotype       | Number of children in subgroups |                      |
|-------------|----------------|---------------------------------|----------------------|
|             |                | Poliskiy district               | Ivankivskiy district |
| 1           | A/A MTR:2756   | 104                             | 106                  |
| 2           | A/G MTR:2756   | 45                              | 61                   |
| 3           | G/G MTR:2756   | 9                               | 11                   |
| 4           | A/A MTHFR:1298 | 82                              | 89                   |
| 5           | A/CMTHFR:1298  | 60                              | 80                   |
| 6           | C/C MTHFR:1298 | 16                              | 9                    |
| 7           | C/C MTHFR:677  | 79                              | 80                   |
| 8           | C/T MTHFR:677  | 60                              | 83                   |
| 9           | T/T MTHFR:677  | 19                              | 15                   |
| 10          | A/AMTRR:66     | 32                              | 27                   |
| 11          | A/G MTRR:66    | 72                              | 93                   |
| 12          | G/G MTRR:66    | 54                              | 58                   |
| Total group |                | 158                             | 178                  |

The level of thyroid hormones was higher in genetic subgroups 1, 5, 7, 8, 11, 12 of children from Ivankivskiy district than in similar subgroups of children from Poliskiy district. At the same time, this related to two hormones simultaneously ( $T_3$  and  $T_4$ ) in the subgroups 1, 5, 11, 12, and concerned only  $T_3$  in the subgroup 7 and only  $T_4$  in the subgroup 8.

Thus, a higher level of  $H_{cy}$  and thyroid hormones was simultaneously recorded in children from Ivankivskiy district compared to those from Poliskiy district in most of the analysed subgroups.

There were no differences between values of  $T_3$  and  $T_4$  with

the existing differences in the blood concentration of  $H_{cy}$  in the subgroups 2 and 4 of children from Ivankivskiy and Poliskiy districts (tables 2-4).

There were no differences both in relation to  $H_{cy}$  and both thyroid hormones between children from the above districts in the subgroups 3, 6, 9 and 10 (tables 2-4).

The TSH blood level did not differ statistically in the children from Poliskiy and Ivankivskiy districts in all subgroups except a subgroup 12 with the main G/G MTRR:66 genotype.

The  $H_{cy}$  blood level was statistically significantly higher in the subgroup 9 containing only

homozygous variants of the MTHFR:677 genetic polymorphism T allele compared to subgroups containing neutral C alleles of the same polymorphism – subgroups 7 and 8 both in the children from Poliskiy district and those from Ivankivskiy district (table 5). At the same time, there were no differences in relation to TSH,  $T_3$  and  $T_4$ .

A direct association was found between  $H_{cy}$  and TSH in a total group as well as in most genetic subgroups from Ivankivskiy district (table 6). A direct association was also recorded between  $H_{cy}$  and  $T_3$ , but in a smaller number of subgroups.

The  $H_{cy}$ -TSH and  $H_{cy}$ - $T_3$  associations were identified simultaneously in the subgroup 4 (table 6). There was no association between  $H_{cy}$  and  $T_4$  in the total group and in separate genetic subgroups.

There was no direct association between  $H_{cy}$  и TSH in the total group of children and all analysed subgroups of children from Poliskiy district. A direct association was observed between  $H_{cy}$  and  $T_3$  in the subgroups 1 and 12 (table 7). No association was found between  $H_{cy}$  and  $T_4$ .

The MTHFR:677 genetic polymorphism T allele was almost uniformly present in all subgroups of children from Poliskiy and Ivankivskiy districts formed on the basis of the MTR:A2756G, MTHFR:A1298C (A/A, A/C genotypes), MTRR:A66G genetic polymorphisms (table 8).

In this regard, the role of the above allele can not be considered as the main reason for a higher blood level of  $H_{cy}$  in children from Ivankivskiy district compared to those from Poliskiy district.

The findings show synergism in values of  $H_{cy}$ ,  $T_3$  and  $T_4$  in the blood of children from the groups of adolescents under study from the districts bordering the Chernobyl exclusion zone.

Higher levels of  $H_{cy}$  were found in most genetic subgroups of children from Ivankivskiy district compared to those from Poliskiy district. At the same time, more higher levels of thyroid hormones were recorded in 75.0% of cases.

No genetic subgroups were found in which there were differences between hormone levels

**Table 2**  
**Statistical characteristics of values of  $H_{cy}$ , TSH,  $T_3$  and  $T_4$  in genetic subgroups of children from Poliskiy district**

| Sub-group   | $H_{cy}$ , $\mu\text{mol/L}$ |             | TSH, mIU/mL |           | $T_3$ , pg/ml |           | $T_4$ , ng/dl |           |
|-------------|------------------------------|-------------|-------------|-----------|---------------|-----------|---------------|-----------|
|             | Me                           | IQR         | Me          | IQR       | Me            | IQR       | Me            | IQR       |
| 1           | 10.26                        | 8,39-13.56  | 1.82        | 1.39-2.43 | 4.04          | 3.79-4.59 | 1.19          | 1.09-1.29 |
| 2           | 9.37                         | 7.75-10.95  | 1.94        | 1.51-2.65 | 4.24          | 3.84-4.68 | 1.15          | 1.06-1.25 |
| 3           | 11.13                        | 10.17-12.32 | 2.10        | 1.19-2.41 | 4.37          | 3.53-5.63 | 1.04          | 0.98-1.32 |
| 4           | 10.14                        | 7.99-13.04  | 1.89        | 1.29-2.55 | 4.11          | 3.79-4.69 | 1.18          | 1.06-1.28 |
| 5           | 10.19                        | 9.01-13.19  | 1.82        | 1.47-2.37 | 4.02          | 3.71-4.57 | 1.17          | 1.08-1.28 |
| 6           | 10.31                        | 8.00 -12.69 | 1.93        | 1.62-3.22 | 4.18          | 3.92-4.64 | 1.20          | 1.10-1.35 |
| 7           | 9.44                         | 7.93-11.16  | 1.80        | 1.34-2.39 | 4.09          | 3.82-4.55 | 1.20          | 1.09-1.29 |
| 8           | 10.24                        | 8.42-13.18  | 1.82        | 1.42-2.55 | 4.02          | 3.75-4.68 | 1.15          | 1.04-1.25 |
| 9           | 14.47                        | 10.15-22.78 | 2.26        | 1.50-3.14 | 4.38          | 3.68-5.02 | 1.23          | 1.05-1.29 |
| 10          | 9.26                         | 8.03-13.13  | 1.92        | 1.67-2.36 | 4.20          | 3.85-4.69 | 1.21          | 1.12-1.34 |
| 11          | 10.08                        | 7.98-11.99  | 1.60        | 1.26-2.40 | 4.07          | 3.68-4.66 | 1.17          | 1.04-1.28 |
| 12          | 10.81                        | 9.13-14.18  | 2.27        | 1.43-2.82 | 4.08          | 3.84-4.53 | 1.17          | 1.09-1.28 |
| Total group | 10.17                        | 8.30 -13.10 | 1.86        | 1.41-2.48 | 4.10          | 3.79-4.62 | 1.18          | 1.07-1.28 |

**Table 3**  
**Statistical characteristics of values of  $H_{cy}$ , TSH,  $T_3$  and  $T_4$  in genetic subgroups of children from Ivankivskiy district**

| Sub-group   | $H_{cy}$ , $\mu\text{mol/L}$ |             | TSH, mIU/mL |           | $T_3$ , pg/ml |           | $T_4$ , ng/dl |           |
|-------------|------------------------------|-------------|-------------|-----------|---------------|-----------|---------------|-----------|
|             | Me                           | IQR         | Me          | IQR       | Me            | IQR       | Me            | IQR       |
| 1           | 11.93                        | 9.73-13.76  | 1.86        | 1.33-2.57 | 4.37          | 3.99-4.74 | 1.26          | 1.14-1.32 |
| 2           | 11.57                        | 10.23-13.33 | 1.72        | 1.30-2.22 | 4.41          | 4.00-4.81 | 1.21          | 1.09-1.30 |
| 3           | 9.46                         | 8.49-11.58  | 1.79        | 1.20-1.93 | 4.59          | 3.96-4.65 | 1.23          | 1.13-1.33 |
| 4           | 11.53                        | 9.64-14.22  | 1.85        | 1.29-2.43 | 4.35          | 3.88-4.75 | 1.22          | 1.12-1.31 |
| 5           | 11.91                        | 10.42-13.30 | 1.77        | 1.31-2.36 | 4.43          | 4.06-4.79 | 1.25          | 1.17-1.32 |
| 6           | 9.09                         | 7.91-12.56  | 1.93        | 1.45-2.73 | 4.38          | 4.05-4.91 | 1.13          | 1.00-1.19 |
| 7           | 11.36                        | 9.38-12.97  | 1.76        | 1.30-2.29 | 4.44          | 4.08-4.81 | 1.24          | 1.12-1.31 |
| 8           | 11.61                        | 9.86-13.31  | 1.83        | 1.30-2.22 | 4.33          | 3.99-4.66 | 1.23          | 1.14-1.30 |
| 9           | 16.59                        | 12.10-26.71 | 2.47        | 1.43-3.13 | 4.12          | 3.84-4.88 | 1.20          | 1.14-1.29 |
| 10          | 10.86                        | 9.19-13.23  | 1.79        | 1.39-2.77 | 4.41          | 3.96-4.60 | 1.25          | 1.14-1.33 |
| 11          | 11.58                        | 9.46-13.22  | 1.79        | 1.31-2.57 | 4.42          | 4.01-4.81 | 1.22          | 1.12-1.30 |
| 12          | 12.13                        | 10.74-14.49 | 1.83        | 1.27-2.24 | 4.33          | 3.99-4.77 | 1.27          | 1.15-1.31 |
| Total group | 11.62                        | 9.69 -13.39 | 1.80        | 1.30-2.42 | 4.40          | 3.99-4.75 | 1.23          | 1.14-1.31 |



**FOLATE CYCLE GENETICS, LEVEL OF HOMOCYSTEINE, THYROID AND PITUITARY HORMONES IN THE BLOOD AMONG THE CHILDREN FROM THE DISTRICTS BORDERING THE CHORNOBYL EXCLUSION ZONE**

**<sup>1</sup>Bandazhevskiy Yu.I., <sup>2</sup>Dubova N.F.**

<sup>1</sup>*Ecology and Health Coordination and Analytical Centre, Ivankiv, Ukraine*

<sup>2</sup>*Shupyk National Healthcare University of Ukraine, Kyiv*

**Objective:** The aim of the study was to identify interrelations between the state of folate cycle genetic apparatus and blood levels of homocysteine, thyroid and adenohypophysis hormones in the blood among the children from Ivankivskiy and Poliskiy districts, Kyiv region, bordering the Chornobyl exclusion zone (ChEZ).

**Methods:** Immunochemical, genetic, statistical ones.

**Results:** We measured blood levels of homocysteine ( $H_{cy}$ ), pituitary thyroid-stimulating hormone (TSH), free triiodothyronine ( $T_3$ ), free thyroxine ( $T_4$ ) in 158 children from Poliskiy district and 178 children from Ivankivskiy district and assessed the state of a genetic system of the folate cycle (FC). The average age of the examined was ( $15.2 \pm 0.9$ ) years old (95% CI 15.0–15.4 years old). Taking into account the FC genotypes, genetic subgroups were formed, they were based on 100% representation of one specific genotype. The laboratory examination was carried out twice: in April and December 2015 (before and after the fires in the ChEZ). It was found out that in the total group of the children, as well as in most of analyzed genetic subgroups, the level of  $H_{cy}$  and thyroid hor-

mones (TG) in the children from Ivankivskiy district was statistically significantly higher than in those from Poliskiy district. At the same time, a direct association between Hcy and TSH was observed in the total group, as well as in most of the genetic subgroups of children from Ivankivskiy district, and this association was absent in the children from Poliskiy district. The blood level of  $H_{cy}$  both in the children from Poliskiy district and in the children from Ivankivskiy district was statistically significantly higher in the subgroup containing only homozygous variants of the T allele of the MTHFR:677 genetic polymorphism in comparison with the subgroups containing neutral C alleles of the same polymorphism. At the same time, there were no differences for TSH,  $T_3$  and  $T_4$ .

**Conclusions:** The analysis enabled to establish a synchronous reaction of metabolic cycles ensuring the metabolism of  $H_{cy}$  and thyroid hormones in the adolescents, regardless of the FC genotypes. Correlation analysis, as well as the results of statistical analysis, indicate that an increase in the level of  $H_{cy}$  in the blood of children induced the synthesis of TSH and  $T_3$ . The results show a close relationship between thyroid hormone genesis and the metabolism of methionine sulfur-containing amino acids and  $H_{cy}$ . A forest fire, containing long-lived radioactive elements, is the most likely reason for the increase of  $H_{cy}$  level and, as a result, the hormones of the pituitary-thyroid axis in the blood of the children living in the districts, bordering the ChEZ.

**Keywords:** homocysteine, folate cycle, thyroid and pituitary hormones, adolescents, radiation contaminated areas.

and there were no differences between  $H_{cy}$  levels.

The high blood level of  $H_{cy}$  in children from Ivankivskiy district allowed to confirm synchronization of reactions of metabolism of  $H_{cy}$  and thyroid and pituitary hormones using a correlation analysis.

An increase in the  $H_{cy}$  level was found to induce the synthesis of TSH and  $T_3$  in the body of children living under conditions of radiation exposure associated with the Chornobyl nuclear power plant accident.

The absence of significant statistical differences in the levels of  $H_{cy}$ , TSH,  $T_3$  and  $T_4$  was recorded in the subgroup 9 where there are homozygous variants of the MTHFR:677 genetic polymorphism T risk allele which results in high blood Hcy levels regardless of environmental exposure.

**Results of statistical differences in variables of children from Poliskiy and Ivankivskiy districts**

Table 4

| Sub-group   | $H_{cy}$ , $\mu\text{mol/L}$ |                       | TSH, mIU/mL         |                       | $T_3$ , pg/ml       |                       | $T_4$ , ng/dl       |                       |
|-------------|------------------------------|-----------------------|---------------------|-----------------------|---------------------|-----------------------|---------------------|-----------------------|
|             | Mann-Whitney U test          | Significance level, p | Mann-Whitney U test | Significance level, p | Mann-Whitney U test | Significance level, p | Mann-Whitney U test | Significance level, p |
| 1           | 4384.000                     | 0.010                 | 5444.500            | 0.878                 | 4215.500            | 0.003                 | 4447.500            | 0.016                 |
| 2           | 721.500                      | 0.0001                | 1151.000            | 0.157                 | 1122.500            | 0.110                 | 1101.500            | 0.083                 |
| 3           | 35.000                       | 0.271                 | 39.000              | 0.425                 | 44.000              | 0.676                 | 38.000              | 0.382                 |
| 4           | 2659.500                     | 0.002                 | 3558.500            | 0.780                 | 3163.500            | 0.133                 | 3038.000            | 0.059                 |
| 5           | 1769.500                     | 0.008                 | 2215.000            | 0.436                 | 1659.500            | 0.002                 | 1709.000            | 0.004                 |
| 6           | 68.000                       | 0.821                 | 64.000              | 0.651                 | 58.000              | 0.428                 | 48.500              | 0.183                 |
| 7           | 2068.500                     | 0.0001                | 3013.500            | 0.614                 | 2183.500            | 0.001                 | 2857.500            | 0.297                 |
| 8           | 1917.000                     | 0.019                 | 2357.500            | 0.588                 | 2012.500            | 0.051                 | 1651.000            | 0.001                 |
| 9           | 134.000                      | 0.768                 | 141.500             | 0.972                 | 137.000             | 0.849                 | 137.000             | 0.849                 |
| 10          | 314.500                      | 0.074                 | 396.000             | 0.584                 | 431.000             | 0.988                 | 396.000             | 0.584                 |
| 11          | 2405.500                     | 0.002                 | 3042.500            | 0.315                 | 2486.500            | 0.005                 | 2661.500            | 0.024                 |
| 12          | 1201.500                     | 0.034                 | 1206.500            | 0.036                 | 1176.000            | 0.023                 | 1199.000            | 0.033                 |
| Total group | 10450.500                    | 0.0001                | 13319.000           | 0.403                 | 11115.000           | 0.001                 | 11461.500           | 0.003                 |

The results obtained make it possible to state about external environmental effects on the body of children living near the Chernobyl exclusion zone during the period between two blood draws.

The most probable environmental factor causing significant metabolic diseases in the body of most children living in settlements locat-

ed near the Chernobyl exclusion zone should be considered fires of forests, which have accumulated a huge amount of long-lived radioactive elements over the 30 post-Chernobyl years [12, 13].

A fire of forest and meadow vegetation occurred on an area of 10127 ha in the Chernobyl exclusion zone on April 26-29, 2015. Therefore the maximum

density of territory contamination in low-fire areas in some quarters of the Lubianskyi forestry was 1040 kBq/sq.m –  $^{137}\text{Cs}$ ; 368 kBq/sq.m –  $^{90}\text{Sr}$ ; 11.4 kBq/sq.m –  $^{238-240}\text{Pu}$  and 14.4 kBq/sq.m –  $^{241}\text{Am}$  [14].

Experimental research has established a negative effect of  $^{137}\text{Cs}$  and  $^{90}\text{Sr}$  radionuclides contained in oats consumed by laboratory animals on the metabolism of methionine, and thus its daughter product, homocysteine [15].

The burning of trees is also accompanied by the release into the environment of fine particles 0.1-2.5  $\mu\text{m}$  in size (PM<sub>2.5</sub>), including black carbon, capable of penetrating the respiratory tract, reaching the bronchioles and alveoli [16], causing metabolic disturbances in the cardiovascular system [17]. A positive association is known between black, as well as organic carbon and blood levels of  $\text{H}_{\text{cy}}$  in older men. At the same time, no associations were recorded between the level of  $\text{H}_{\text{cy}}$  and PM<sub>2.5</sub> [18].

The data obtained show a synchronous increase in the levels of  $\text{H}_{\text{cy}}$  and thyroid hormones in the body of children of most genetic subgroups after forest fires in the Chernobyl exclusion zone in 2015.

### Conclusions

Laboratory examination of children from districts bordering the Chernobyl exclusion zone in April and December 2015 enabled to establish a synchronous reaction of metabolic cycles supporting the metabolism of  $\text{H}_{\text{cy}}$  and thyroid hormones in a developing body, irrespective of folate cycle genotypes.

The conducted correlation analysis, as well as the results of statistical analysis, show that an increase in the level of  $\text{H}_{\text{cy}}$  in the blood of children induced the synthesis of TSH and  $\text{T}_3$ .

The absence of differences in levels of  $\text{H}_{\text{cy}}$ , TSH,  $\text{T}_3$  and  $\text{T}_4$  between similar subgroups of children from Ivankivskyi and Poliskyi districts that include 100% carriership of the T/T MTHFR:677 genotype is associated with the T risk allele causing high blood levels of  $\text{H}_{\text{cy}}$  regardless of environmental exposure.

The results obtained show a close relationship between thy-

**Table 5**  
**Statistically significant differences when comparing blood Hcy values in children of subgroups № 9<sup>1</sup> and № 7<sup>2</sup>, № 9<sup>1</sup> and № 8<sup>2</sup> from Poliskyi and Ivankivskyi districts**

| Variables                             | Poliskyi district |                 |                   |                 | Ivankivskyi district |                 |                   |                 |
|---------------------------------------|-------------------|-----------------|-------------------|-----------------|----------------------|-----------------|-------------------|-----------------|
|                                       | Subgroups 9 and 7 |                 | Subgroups 9 and 8 |                 | Subgroups 9 and 7    |                 | Subgroups 9 and 8 |                 |
|                                       | n                 | H <sub>cy</sub> | n                 | H <sub>cy</sub> | n                    | H <sub>cy</sub> | n                 | H <sub>cy</sub> |
| Average rank <sup>1</sup>             | 19                | 71.11           | 19                | 52.63           | 15                   | 68.27           | 15                | 66.77           |
| Average rank <sup>2</sup>             | 79                | 44.30           | 60                | 36.00           | 80                   | 44.20           | 83                | 46.38           |
| Mann-Whitney U test                   | 340.00            |                 | 330.00            |                 | 296.00               |                 | 363.50            |                 |
| Asymptotic significance (2-tailed), p | 0.0001            |                 | 0.006             |                 | 0.002                |                 | 0.011             |                 |

**Table 6**  
**Results of correlation analysis between values of H<sub>cy</sub> and T<sub>3</sub>, H<sub>cy</sub> and TSH in genetic subgroups of children of Ivankivskyi districts**

| Sub-group   | Genotype          | Correlation coefficient | Parameters                  |                           | Parameters                  |                |
|-------------|-------------------|-------------------------|-----------------------------|---------------------------|-----------------------------|----------------|
|             |                   |                         | H <sub>cy</sub> ,<br>μmol/L | T <sub>3</sub> ,<br>pg/ml | H <sub>cy</sub> ,<br>μmol/L | TSH,<br>mIU/mL |
| 2           | A/G MTR:2756      | Spearman's              | 0.129                       |                           | 0.278*                      |                |
|             |                   | p                       | 0.321                       |                           | 0.030                       |                |
|             |                   | N                       | 61                          |                           | 61                          |                |
| 4           | A/A<br>MTHFR:1298 | Spearman's              | 0.299**                     |                           | 0.288**                     |                |
|             |                   | p                       | 0.004                       |                           | 0.006                       |                |
|             |                   | N                       | 89                          |                           | 89                          |                |
| 6           | C/C<br>MTHFR:1298 | Spearman's              | 0.617                       |                           | 0.733*                      |                |
|             |                   | p                       | 0.077                       |                           | 0.025                       |                |
|             |                   | N                       | 9                           |                           | 9                           |                |
| 7           | C/C<br>MTHFR:677  | Spearman's              | 0.256*                      |                           | 0.023                       |                |
|             |                   | p                       | 0.022                       |                           | 0.838                       |                |
|             |                   | N                       | 80                          |                           | 80                          |                |
| 8           | C/T<br>MTHFR:677  | Spearman's              | 0.075                       |                           | 0.278*                      |                |
|             |                   | p                       | 0.498                       |                           | 0.011                       |                |
|             |                   | N                       | 83                          |                           | 83                          |                |
| 10          | A/A MTRR:66       | Spearman's              | 0.502**                     |                           | 0.108                       |                |
|             |                   | p                       | 0.008                       |                           | 0.592                       |                |
|             |                   | N                       | 27                          |                           | 27                          |                |
| 11          | A/G MTRR:66       | Spearman's              | 0.053                       |                           | 0.303**                     |                |
|             |                   | p                       | 0.612                       |                           | 0.003                       |                |
|             |                   | N                       | 93                          |                           | 93                          |                |
| Total group |                   | Spearman's              | 0.157*                      |                           | 0.206**                     |                |
|             |                   | p                       | 0.037                       |                           | 0.006                       |                |
|             |                   | N                       | 178                         |                           | 178                         |                |

Note: \* – correlation is significant at the <0.05 level (2-tailed).

\*\* – correlation is significant at the <0.01 level (2-tailed).

ГЕНЕТИКА ФОЛАТНОГО ЦИКЛА, УРОВЕНЬ ГОМОЦИСТЕИНА, ГОРМОНОВ ЩИТОВИДНОЙ ЖЕЛЕЗЫ И ГИПОФИЗА В КРОВИ ДЕТЕЙ ИЗ РАЙОНОВ, ГРАНИЧАЩИХ С ЧЕРНОБЫЛЬСКОЙ ЗОНОЙ ОТЧУЖДЕНИЯ  
**<sup>1</sup>Бандажевский Ю.И., <sup>2</sup>Дубовая Н.Ф.**

<sup>1</sup>Координационный аналитический центр «Экология и здоровье», г. Иванков, Украина

<sup>2</sup>Национальный университет охраны здоровья Украины им. П.Л. Шупика, г. Киев, Украина

**Цель работы:** определение взаимосвязи между состоянием генетического аппарата фолатного цикла, содержанием гомоцистеина, гормонов щитовидной железы и аденогипофиза в крови детей из Иванковского и Полесского районов Киевской области, граничащих с Чернобыльской зоной отчуждения (ЧЗО).

**Методы исследования.** Иммунохимический, генетический, статистический.

**Результаты.** У 158 детей из Полесского района и 178 детей из Иванковского определялось содержание в крови гомоцистеина ( $H_{cy}$ ), тиреотропного гормона гипофиза (ТТГ), свободного трийодтиронина ( $T_3$ ), свободного тироксина ( $T_4$ ), также оценивалось состояние генетической системы фолатного цикла (ФЦ). Средний возраст обследованных составил ( $15,2 \pm 0,9$ ) лет (95% ДИ 15,0-15,4 лет). С учетом генотипов ФЦ были сформированы генетические подгруппы, основанные на 100% представительстве одного конкретного генотипа. Лабораторное обследование проведено дважды: в апреле и декабре 2015 года (до и после пожаров в ЧЗО). Установлено, что в общей группе детей, а также большинстве анализируемых генетических подгрупп уровень  $H_{cy}$  и гормонов щитовидной железы

(ЩЖ) у детей из Иванковского района был достоверно выше, чем у детей из Полесского района. При этом в общей группе, а также в большинстве генетических подгрупп детей из Иванковского района между  $H_{cy}$  и ТТГ выявлена прямая корреляционная связь, отсутствующая у детей из Полесского района. Уровень  $H_{cy}$  в крови у детей из этих районов был достоверно выше в подгруппе, содержащей только гомозиготные варианты аллели Т генетического полиморфизма MTHFR:677, по сравнению с подгруппами, содержащими нейтральные аллели С этого же полиморфизма. При этом отсутствовали различия в отношении ТТГ,  $T_3$  и  $T_4$ .

**Выводы.** Проведенный анализ позволил установить синхронную реакцию метаболических циклов, обеспечивающих обмен  $H_{cy}$  и гормонов ЩЖ в организме подростков, независимо от генотипов ФЦ. Корреляционный анализ, а также результаты статистического анализа свидетельствуют о том, что повышение уровня  $H_{cy}$  в крови детей индуцировало синтез ТТГ и  $T_3$ . Полученные результаты говорят о тесной связи между гормоногенезом ЩЖ и обменом серосодержащих аминокислот метионина и гомоцистеина. Наиболее вероятной причиной повышения уровня  $H_{cy}$  и вследствие этого гормонов гипофизарно-тиреоидной оси в крови детей, проживающих в районах, граничащих с ЧЗО, стали пожары леса, содержащего долгоживущие радиоактивные элементы.

**Ключевые слова:** гомоцистеин, фолатный цикл, гормоны щитовидной железы и гипофиза, подростки, радиоактивно загрязненные территории.

roid gland hormonogenesis and metabolism of sulfur-containing amino acids – methionine and  $H_{cy}$ .

Fires of forests that contain long-lived radioactive elements are the most likely reason for an increase in the  $H_{cy}$  level and as a result pituitary-thyroid axis hormones in the blood of children living in districts bordering the Chornobyl exclusion zone.

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Table 7

#### Results of correlation analysis between values of $H_{cy}$ and $T_3$ , $H_{cy}$ and TSH in genetic subgroups of children of Poliskyi districts

| Subgroup    | Genotype     | Correlation coefficient | Parameters                   |                        | Parameters                   |                      |
|-------------|--------------|-------------------------|------------------------------|------------------------|------------------------------|----------------------|
|             |              |                         | $H_{cy}$ , $\mu\text{mol/L}$ | $T_3$ , $\text{pg/ml}$ | $H_{cy}$ , $\mu\text{mol/L}$ | TSH, $\text{mIU/mL}$ |
| 1           | A/A MTR:2756 | Spearman's              | 0.204*                       |                        | 0.010                        |                      |
|             |              | p                       | 0.038                        |                        | 0.917                        |                      |
|             |              | N                       | 104                          |                        | 104                          |                      |
| 12          | G/G MTRR:66  | Spearman's              | 0.424**                      |                        | 0.195                        |                      |
|             |              | p                       | 0.001                        |                        | 0.158                        |                      |
|             |              | N                       | 54                           |                        | 54                           |                      |
| Total group |              | Spearman's              | 0.130                        |                        | -0.002                       |                      |
|             |              | p                       | 0.102                        |                        | 0.984                        |                      |
|             |              | N                       | 158                          |                        | 158                          |                      |

Note: \* – correlation is significant at the  $<0.05$  level (2-tailed).

\*\* – correlation is significant at the  $<0.01$  level (2-tailed).

Table 8

#### Proportion of cases of carriership of MTHFR:C677T polymorphism T allele in genetic subgroups of children from Ivankivskiy and Poliskyi districts

| Sub-group   | Genotype       | Number of cases of carriership of MTHFR:C677T polymorphism T allele in subgroups of children |       |                   |       |
|-------------|----------------|--|-------|-------------------|-------|
|             |                | Ivankivskiy district   |       | Poliskyi district |       |
|             |                | Abs.   | %     | Abs.              | %     |
| 1           | A/A MTR:2756   | 61   | 57.55 | 50                | 48.08 |
| 2           | A/G MTR:2756   | 32   | 52.46 | 24                | 53.33 |
| 3           | G/G MTR:2756   | 5  | 45.45 | 5                 | 55.56 |
| 4           | A/A MTHFR:1298 | 60   | 67.42 | 54                | 65.85 |
| 5           | A/C MTHFR:1298 | 38   | 47.50 | 25                | 41.67 |
| 6           | C/C MTHFR:1298 | 0  | 0     | 0                 | 0     |
| 7           | C/C MTHFR:677  | 0  | 0     | 0                 | 0     |
| 8           | C/T MTHFR:677  | 83   | 100   | 60                | 100   |
| 9           | T/T MTHFR:677  | 15   | 100   | 19                | 100   |
| 10          | A/A MTRR:66    | 14   | 51.85 | 14                | 43.75 |
| 11          | A/G MTRR:66    | 53   | 56.99 | 37                | 51.39 |
| 12          | G/G MTRR:66    | 31   | 53.45 | 28                | 51.85 |
| Total group |                | 98   | 55.06 | 79                | 50.00 |

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