THE MOTHER AND THE YOUNG CHILD: WHO PROGRAMS WHOM

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ABSTRACT

Intrauterine life is regulated by an epigenetic programme which adapts fetal life to the current and presumed conditions. The fetal programming determined by maternal nutritional disorders is at the origin of some metabolic and cardiovascular diseases of childhood and adult life. The maternal programming guides the child’s immunity, circadian rhythm, and stress response. The maternal and paternal imprinted genes exert a contradictory action upon fetal development (the war of the sexes). In turn, the fetus programs the maternal physiology through placental hormones and the cell transfer in the maternal organism (feto-maternal microchimerism).

Keywords: maternal fetal programming, metabolism, immunity, circadian rhythm, stress, parental conflict, feto-maternal programming

INTRODUCTION

In the model centered on the genetic concept, parental influence on the development of the offsprings was subordinated to a predefined genetic program (1). The accumulation of clinical and experimental studies has highlighted that in fact genes express themselves differently according to the environment. One of the most shocking medical hyposthesis at the end of the 20th century was the connection between the prematurity/low birth weight and the chronic pathology of the adult age (2-4). Considering that in terms of life cycle the most important stage of natural selection takes place prenatally, numerous studies have proven that there is a connection between nutrition, the mother’s life conditions and the intrauterine and subsequent development of the child, these being at the origin of some somatic and psychic disorders. In other words, the intrauterine life is part of a rigorous developing program in which no detail is insignificant. All of these adaptations are made possible by changes in gene expression without changes in DNA sequences, that is by epigenetic mechanisms. The epigenetic inheritance regards the transmission of some epigenetic markers to the offsprings (1), the phenotype being the result of the interference of genetic factors and those originated in the environment (epigenetic).

TABLE 1. Critical periods in prenatal development

<table>
<thead>
<tr>
<th>Organs/systems</th>
<th>Maximum sensitivity</th>
<th>Development until</th>
</tr>
</thead>
<tbody>
<tr>
<td>CNS</td>
<td>4-8 weeks</td>
<td>Post-natal – adult</td>
</tr>
<tr>
<td>Heart</td>
<td>5-9 weeks</td>
<td>12 weeks</td>
</tr>
<tr>
<td>Limbs</td>
<td>6-10 weeks</td>
<td>12 weeks</td>
</tr>
<tr>
<td>Eyes</td>
<td>6-10 weeks</td>
<td>Term</td>
</tr>
<tr>
<td>Auditory system</td>
<td>6-11 weeks</td>
<td>13 weeks</td>
</tr>
<tr>
<td>Pancreas (insulin secretion)</td>
<td>7-? weeks</td>
<td>Term</td>
</tr>
<tr>
<td>Kidneys</td>
<td>8-32 (34) weeks</td>
<td>Term</td>
</tr>
</tbody>
</table>

In the light of the collected data, fetal programming is considered to be at the origin of the events that take place during conception, intrauterine life, childhood, and adulthood, being connected by the epigenetic background. Thus, certain diseases which occurred in the first part of life become predictors of adult pathology. Some studies which became “classic” followed the over time evolution of some subjects exposed to some nutritional restrictions during intrauterine life. During the famine in the Netherlands (1944-1945), during the Second World War, the medium food intake ranged between 300 and 600 calories per day. The children born during this time manifested after 50 years a structure of pathology which was linked to the period of damaged pregnancy (7).
TABLE 2. A synthesis of types of pathology in adults aged over 50, according to the period of maternal nutritional deprivation in the Netherlands, 1944-1945 (7)

<table>
<thead>
<tr>
<th>First trimester</th>
<th>Second trimester</th>
<th>Third trimester</th>
</tr>
</thead>
<tbody>
<tr>
<td>Glucose intolerance</td>
<td>Glucose intolerance</td>
<td>Glucose intolerance</td>
</tr>
<tr>
<td>Cardiovascular disease</td>
<td>Pulmonary disease</td>
<td>Intolerance</td>
</tr>
<tr>
<td>Arterial hypertension (AH)</td>
<td>Renal disease</td>
<td>Cardiovascular disease</td>
</tr>
<tr>
<td>Dyslipidemia</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Obesity</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Emotional disorders</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

The trend of birth weight increase linked to the high rate of overweight or obese women was noticed in various countries during the last decades. Maternal chronic hyperglycemia is responsible for the accumulation of an abundant adipose tissue in newborns. Paradoxically, the consequences of fetal macrosomia (birth weight of over 4,000-4,500 g) are similar to those of children with low birth weight in terms of the occurrence of diabetes mellitus (DM) during adolescence or adulthood (8). A study compared the weight of children born before the bariatric surgery for obesity (macrosomia of 34.8%) with that of a group of women after the same type of surgery (macrosomia of 7.7%) (9). The data suggest the intergenerational transmission of the decrease in glucose tolerance and of obesity of nongenetic origin. Numerous clinical and experimental data support the epigenetic involvement in the appearance of obesity, of NIDDM, AH, and other diseases.

MATERNAL PROGRAMMING OF THE CHILD’S IMMUNITY

Except for its nutritive role, maternal milk plays other major biological roles: the formation and modulation of the immunity, the live cell transfer through maternal milk, the microflora formation of the newborn and infant. The capacity of maternal milk to develop and modulate neonatal and later immunity is based on the transmission of an enormous quantity of information expressed through different molecules (cytokines, interleukins, defensins, hormones, growth factors, immunoglobulins etc.), cells and microorganisms. The process of transmission of the maternal antibodies to the neonatal naïve immune system is initiated transplacentally under the action of fetal Fc receptors. Immunoglobulins of the type IgG, IgM and IgA have been identified in colostrum and mature milk. The persistence of maternal antibodies in the nursling’s circulation continues until the age of 9 months and is dependent on the beginning of their active production (10). Abundantly represented in the maternal milk, IgA plays an important role; as secretary IgA, it connects to the mucus layer of the epithelial intestinal cells, forming a barrier against the aggressive pathogens, before the contact with the epithelial cells (11). As a response to its own intestinal antigens, the maternal immune system produces IgA antibodies; the immune information is transferred to the mammary plasmocytes, which, in turn, take over the production of the same type of antibodies (the entomammary system). This results in the bacterial intestinal colonization of the newborn with a commensal flora similar to that of the maternal intestine and the limitation of pathogen proliferation. The effect of transferring maternal antibodies to the newborns and infants manifests only for a relatively short period but it leads to the irreversible programming of the immune repertoire, suggesting that, in some cases, the maternal antibodies may influence the immune function along many generations (12). In the maternal milk there are live bacteria which come from the mammary ducts and areola, but they are partially intrinsic components of the milk (13). Observations that maternal milk is not sterile, even when it is aseptically collected, support the idea that this is a natural bacterial colonizing source of the neonatal intestine (12). The origin of the bacteria identified in milk is found in the maternal intestine; the observation is backed up by the finding that identical bacterial DNA fragments can be identified in the maternal stool, blood, milk, and in the nurslings’ stools (10). The transfer of the viable bacteria or their genetic material is ensured by maternal mononuclear cells both during the pregnancy and after (15). The population of the gastrointestinal tract begins before birth as the fetus ingests amniotic fluid which contains microbial germs (14). Even if the composition of the microbiota changes with age, the microbial flora acquired from the mother at birth and during the natural feeding will influence the development of intestinal microbial environment in the long run. The immune programming of the intestine through lactation is also assumed by the transfer of live and active leukocytes, among which most of them are macrophage and polymorphonuclear. The population of lymphocytes transferred to the nursling consists of Ly T motile and interactive, involved in the active transfer of immunity (16,17); among these, Foxp3 (Treg) play an important role in constituting the immune tolerogenic response. The primary bacterial colonization of the intestine is the child’s first immunity lesson.

THE PROGRAMMING OF THE CIRCADIAN RHYTHM OF THE INFANTS THROUGH NATURAL FEEDING

Human physiology and behaviour are driven by the circadian rhythm governed by the light-dark cy-
circles and the energy metabolism, under the regulation of CNS (18,19). The formation of the circadian rhythm in newborns and infants sets the rhythm for the wake/sleep periods and feeding demands. The composition of the maternal milk varies during the day. The concentration of glucocorticoids is three times higher in the morning milk than in the samples collected in the afternoon or in the evening, stimulating the activity, while the melatonin level is close to the limit of detection during the day, but during the night it can grow up to 280 +/- 34 pmol/l (melatonin being known as "the sleep hormone") (20,21). This way the natural feeding works through chronosignals, helping the baby to differentiate between day and night periods in the first months of life.

THE BEHAVIORAL PROGRAMMING OF THE BABY THROUGH TACTILE STIMULATION

It is known that during the first years of life the baby displays a fascinating behaviour, which strengthens the emotional connection with the mother (father or close people). This complex system of interpersonal relationship is genetically programmed, its purpose is to ensure protection and it is common among all mammal babies. Known as attachment theory (22), it objectifies the young child’s need for protection, calmness and the provision of the food intake. The attachment behaviour strongly manifests during alarm situations accompanied by emotional stress.

THE PROGRAMMING OF STRESS RESPONSE

Under experimental conditions (in rodents) it was found that the careful nursing of the babies through the maternal tactile contact is responsible for a moderate stress response (as compared to the babies deprived of maternal care) in the next stages of life (23,24). Stress response involves a series of changes of the hypothalamic-pituitary-adrenal axis (HPA) expressed by the increase of ACTH, of cortisol, of catecholamines etc. In children, too, it was found that there is a connection between physical contact, the amount of care and the range of stress response, proven by the fact that the careful maternal nourishing diminishes the intensity of reactivity to stressful factors of the environment. In these situations, at the molecular level, the increase of GR (glucocorticoid receptor) gene in the hippocampus has been found, followed by the decrease in the stress response of the HPA axis, change that is epigenetically transmitted to the later stages of age, too (24) (25). Oberlander et al (26) have looked into the relation between the maternal depressed/anxious behaviour in the third trimester of pregnancy and the increase of the degree of methylation of the cellular receptor gene for the glucocorticoids (GRG) followed by the increase of cortisol levels at the age of 3 months according to saliva testing. In the case of naturally fed 5-month-old nurslings, the decrease of methylation degree of GR gene and of cortisol reactivity was found, that is the natural feeding contributes to the physiological regulation of behaviour (27). The infant’s crying (especially over extended periods) reflects the baby’s stress and discomfort and represents a part of the emergency system in the infant-parent communication (28,29).

Due to the fact that glucocorticoids freely diffuse through the cell membrane it is important that there is another regulatory mechanism which would limit their access from the mother into the fetal circulation, avoiding the excessive gene expression. Under physiological conditions there is a glucocorticoid gradient of 100-1000 times higher in maternal circulation compared to the fetal circulation. The deficiencies in maternal diet produce the decrease in activity of the placental enzyme 11 β-hydroxysteroid-dehydrogenase (11β HSD2). Under standard conditions this transforms the active glucocorticoids into forms without physiological activity, thus regulating the access of maternal corticosterone to the fetus. If the placental barrier is not fully functional, the excessive maternal glucocorticoid transit stimulates the HPA fetal axis, with later somatic and behaviour changes (anxiety, stress response, AH) (44).

THE PARENTAL CONFLICT IN FETAL PROGRAMMING

The fetal development is controlled by the mother functional unit, placenta and fetus. Apparently paradoxical, the development of the fetus is determined by the contradictory interaction between maternal and paternal genes in the conflict named the genome war. The antagonistic effects are determined by the imprinted genes (which come from only one parent) which originate in the ovule or sperm. In this case,
some maternal genes can be expressed and the paternal ones completely inhibited or vice versa, some of the paternal genes can be active at the expense of the maternal ones.

**WAR OF THE SEXES**

The mother “is interested” in her survival, the conservation of the substrates and the diminishing of fetal weight, she transmits genes which limit fetal growth. The father, “interested in” increasing the fetal growth, fetal survival, higher birth weight, transmits paternal genes such as IGF-2. The factors IGF-1 and IGF-2 stimulate growth, acting upon the IGF-1r receptor. The second type of receptor, whose gene is of exclusive maternal provenance, IGF-2r, opposes growth by trapping and degradating of IGF-2 (31,32).

**IS THERE A CHILD-MOTHER PROGRAMMING?**

The first opinions on the complex connection between mother and fetus date from 1993, when Haig (34) expressed the opinion that fetal genes increase the nutrient transfer towards the fetal metabolism, while the maternal ones limit the excess of their transit. Although this opinion may be considered excessive, one should take into consideration the effects of vascular and endocrine changes produced by the fetus on the maternal organism.

The physiological pregnancy is accompanied by insulin resistance as a physiological adaption of the mother to provide an adequate carbohydrate intake for the fetus in rapid growth (35). The hormones produced by the placenta regulate the intake, the absorption, the use and the transfer of maternal nutrients to the fetus (36), at the same time, being at the origin of insulin resistance and, possibly, of gestational diabetes. The bidirectional exchange of live cells between mother and fetus has been highlighted in different studies, but the functions of fetal cells in the maternal tissues are not clearly defined (38-40). Some of the cells of the fetus transit the placenta even from the first weeks of the first trimester; feto-maternal cell transfer probably takes place in all of pregnancies (39).
**FIGURE 2.** The interaction between IGF-1, IGF-2 and their receptors in the regulation of fetal growth in the experimental model in rodents: IGF-1 and IGF-2 stimulate the intrauterine growth, acting upon the receptor IGF-1r. The second type of receptor, IGF-2r, opposes growth by degradation of IGF-2 (33)

**TABLE 3.** Mother-child relation during the pregnancy and after. The theory of cooperation and conflict (the feto-maternal negotiation) can clarify the role of the microchimerism of the fetal cells in the mother’s health (38,41)

<table>
<thead>
<tr>
<th>The presence of fetal cells</th>
<th>Baby’s interest</th>
<th>Mother’s interest</th>
</tr>
</thead>
<tbody>
<tr>
<td>Brain</td>
<td>The manipulation of the parental neuroendocrine system (oxytocin, prolactin) for nutrition and strengthening the affection towards the baby</td>
<td>The presence of fetal paternal** DNA cells in the brain many years after birth of a son; integration in the neuronal circuits. Role in differentiation of neurons</td>
</tr>
<tr>
<td>Thyroid</td>
<td>The increase of heat production</td>
<td>The regulation of the heat production at a level which permits the optimization of the use of resources at present and for the future offsprings</td>
</tr>
<tr>
<td>Breast</td>
<td>The fetal cells which migrated at the breast level increase the production of milk through stimulation factors or the differentiation in mammary cells</td>
<td>The modulation of the milk production; granting the resources at present and for the offsprings</td>
</tr>
<tr>
<td>Immune system</td>
<td>The induction of maternal tolerance Avoiding detection and destruction by the maternal immune system</td>
<td>The tolerance of semi-allogeneic fetal material. The elimination of fetal cells which emphasize the resource transfer at a level above the optimal one</td>
</tr>
</tbody>
</table>

**Identified by the presence of the Y chromosome**
FETO-MATERNAL MICROCHIMERISM

Feto-maternal microchimerism (the presence of a small number of cells originating in a subject genetically different from the host) is characterised by the preservation of a number of fetal cells in the maternal tissues for a long time (decades) (40). Their transfer may start shortly after the implantation (38). The fetal cells circulate in the maternal body, they can colonise different tissues and can be found in different maternal tissues and organs (e.g. blood, bone marrow, brain, liver, tegument) (36). Their role was interpreted either as being in the child’s interest, or in the interest of the mother.

Microchimerism involves all types of fetal cells and trophoblasts, but especially fetal stem cells (42). The presence of some stem-like cells in different cellular lines indicates their capacity to transform into adult hematopoietic cells in all the lines (42), in endothelial cells, neurons, smooth muscle cells and cardiomyocytes (41). Their migration and their role in healing the lesions produced by the caesarean section by producing type I and III collagen and TGF-β (43) have been proven. On the other hand, the microchimeric fetal cells, by their escaping from the maternal immune system, were involved in the pathogenesis of some cancers and some autoimmune diseases (e.g. SLE, scleroderma) (38,41).

CONCLUSION

Despite all the contradictory aspects, it is obvious that the exchange of hormones, immunity, cells represents a bidirectional mother-child path of communication, which continues in other ways the entire life.

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