

Maternal obesity as a risk factor for brain development and mental health in the offspring

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Abstract

Maternal obesity plays a key role in the health trajectory of the offspring. Although research on this topic has largely focused on the potential of this condition to increase the risk for obesity in the offspring, it is becoming more and more evident that it can also significantly impact cognitive function and mental health. The mechanisms underlying these effects are starting to be elucidated and point to the placenta as a critical organ that may mediate changes in the response to stress, immune function and oxidative stress. Long-term effects of maternal obesity may rely upon epigenetic changes in selected genes that are involved in metabolic and trophic regulations of the brain. More recent evidence also indicates the gut microbiota as a potential mediator of these effects. Overall, understanding cause-effect relationships can allow the development of preventive measures that could rely upon dietary changes in the mother and the offspring. Addressing diets appears more feasible than developing new pharmacological targets and has the potential to affect the multiple interconnected physiological pathways engaged by these complex regulations, allowing prevention of both metabolic and mental disorders.

Keywords maternal obesity; pregnancy; fetal programming; mood disorders; placenta; oxidative stress.

Abbreviations

11 β -HSD: 11 β -hydroxysteroid dehydrogenase
AD: Alzheimer's disease
ADHD: attention deficit hyperactivity disorder
AHRR: aryl-hydrocarbon receptor repressor
ASD: autism spectrum disorder
BDNF: brain-derived neurotrophic factor
BMI: body mass index
CpG: 5'-cytosine-phosphate-guanine-3'
CVD: cardiovascular disease
DOHaD: developmental origin of health and disease
GDM: gestational diabetes mellitus
GC: glucocorticoid
GR: glucocorticoid receptors
GSH: glutathione
HPA: hypothalamic-pituitary-adrenal
HFD: high fat diet
KO: knock out
NAC: N-Acetyl-Cysteine
NAc: nucleus accumbens
OS: oxidative stress
PD: Parkinson's disease
PFC: prefrontal cortex
PBMC: peripheral blood mononuclear cells
PUFA: polyunsaturated fatty acid
RCT: randomized controlled trial
ROS: reactive oxygen species
T2D: type 2 diabetes
VTA: ventral tegmental area

Highlights

- Maternal obesity affects mental health in the offspring.
- Changes in redox signaling underlie both metabolic and cognitive effects of maternal obesity on the offspring.
- Dysbiosis and epigenetic changes could underlie the transgenerational transmission of the effects of maternal obesity.
- Quality of prenatal diet is a potentially modifiable target for reducing the risk of mental disorders in the offspring.

Introduction

Overweight and obesity are dramatically rising in low- and middle-income areas, particularly in urban settings (<https://www.who.int/end-childhood-obesity/publications/echo-report/en/>). The prevalence of obesity is increasing across all populations and age groups: although genetic factors may play a role in modulating vulnerability to weight gain and fat accumulation, they cannot explain the exponential increase in obesity we are currently witnessing (Congdon, 2019). Indeed, globalization and urbanization have gradually led to the so-called *nutrition transition* i.e. a reduction of physical activity associated to the increase in consumption of low-cost and easily accessible ultra-processed, energy-dense, nutrient-poor foods (<https://www.who.int/end-childhood-obesity/publications/echo-report/en/>). Such a spread of unhealthy lifestyles results in an energy unbalance that favors fat storage, eventually strengthening the ground for the settlement of an obesogenic environment (Townshend and Lake, 2017).

Maternal obesity affects 30% of pregnant women and excess weight gain occurs in 40% of gestations. The “Commission on Ending Childhood Obesity” (established by WHO) in its 2016 final report, has tackled the early life environment (including preconception and pregnancy) as a critical time for long-lasting and trans-generational effects of the metabolic derangement underlying obesity - and the associated comorbidities - as well as a window of opportunity to prevent them (<https://www.who.int/end-childhood-obesity/publications/echo-report/en/>). Although a great deal of research has focused on the mechanisms that can lead to offspring obesity, there is evidence for an effect of maternal obesity on cognition and mental health of the offspring (Buss et al., 2012; Casas et al., 2013; Hinkle et al., 2012; Hinkle et al., 2013; Rivera et al., 2015; Rodriguez, 2010). Indeed, exposure to maternal obesity or to an unhealthy maternal diet and metabolic diseases (diabetes and hypertension), can all increase the risk of for later-life cognitive disabilities and psychiatric disorders such as attention deficit hyperactivity disorder (ADHD), autism spectrum disorders (ASDs), anxiety, depression, schizophrenia and eating disorders in the offspring (Buss, et al., 2012; Dinan and Cryan, 2017; Hinkle, et al., 2012; Hinkle, et al., 2013; Rivera, et al., 2015; Yatsunenکو et al., 2012).

Obesity in pregnancy results in neuroendocrine, metabolic and immune/inflammatory changes which can affect fetal exposure to hormones and nutrients, disrupting the development of those neural pathways critical for the regulation of behavior and cognition. Inflammation during pregnancy may alter functional connectivity and may be associated with altered behavioral

regulation and reduced working memory performance in early childhood. Changes in microbiota composition as a result of maternal obesity may play a role in these effects. Indeed, the gut microbiota develops and stabilizes in early life phases (Yatsunenko, et al., 2012), which are crucial for the programming of tissues and organs, including the brain by e.g. modulation of the immune system (Dinan and Cryan, 2017;El Aidy et al., 2016) hormones, neurotransmitters and metabolites acting on brain physiology.

Postnatal nutrition (also targeting microbiota) may thus represent an important strategy for combating metabolic-associated cognitive impairment. In fact, the beneficial effects of diets, nutrients and foods (e.g. Mediterranean diets, omega-3) on mental health and cognitive performance are starting to be recognized (Bellisario et al., 2014;Bruce et al., 2002;Davari et al., 2013;Dimmitt et al., 2010;Dinan and Cryan, 2016;Kelly et al., 2016;Miliot et al., 2016;Mrizak et al., 2014;Patterson et al., 2016;Sherwin et al., 2016).

This review will focus on the contribution played by maternal obesity in shaping the metabolic and behavioral phenotype of the offspring by affecting the developmental programming of the fetus. Main emphasis will be given to changes in oxidative stress (OS) pathways and epigenetic modifications as possible mechanisms mediating the effect of maternal obesity and early exposure to high caloric diets (eg. westernized diet rich in fats and sugars, high fat diet - HFD). The most recent evidence coming from epidemiological and preclinical studies will be reviewed taking also into account the role played by maternal obesity in early programming of the offspring gut microbiota. Finally, the potentiality of strategies aimed at preventing/counteracting the negative effects of prenatal obesity will be considered, including antioxidants and probiotics/prebiotic administration during pregnancy and to the offspring.

Developmental Origin of Health and Disease (DOHaD). Evolutionary and epidemiological perspectives

Mammalian development unfolds as a gradual process, which is continuously adjusted to the needs and challenges posed by the environment (in this review we will specifically focus on the intrauterine development) within the constraints of the genetic asset of the individual. From an evolutionary perspective, the elevated plasticity characterizing the developmental program is perfectly suited at generating a plethora of best adapted phenotypes that allow the perpetuation of the species with regard to the eco-ethological niches they will colonize (Bateson, 2001; Bateson et al., 2004). While such plasticity helps the individual to adjust to changes in the environment, it can also provide a substrate for increased vulnerability later in life, thus resulting in a double edge sword. The overall outcome of the developmental program will depend heavily upon the interplay among the genetic background of the organism, the intrauterine milieu and the stability of the extra-uterine (external) environmental conditions with respect to those that primed fetal developmental trajectories (Bateson, et al., 2004).

Early life experiences thus become embedded biological traces in an animal's physiology, resulting either in increased vulnerability or to greater resilience for the onset of disease states. The seminal studies of Barker and colleagues have contributed to provide evidence for a strong association between environmental challenges during pregnancy, altered fetal growth and health outcomes later in life (Barker et al., 1993; Seckl, 1998) giving rise to the concept of the DOHaD. In fact, the observation that those regions in England showing the highest infant mortality, with regard to low birth weight, were also characterized by the highest rates of mortality from cardiovascular disease (CVD) led to the hypothesis that those babies who survived were at greater risk of CVD later in life (Barker, et al., 1993; Barker and Osmond, 1986; Barker et al., 1989). Most intriguingly, the relationship between reduced birth weight and the onset of diseases at adult age (e.g. hypertension or type 2 diabetes - T2D) was unrelated to lifestyle risk factors (Cirulli F. et al., 1994; Harris and Seckl, 2011; Leon et al., 1996; Osmond et al., 1993). In addition, studies on twins provide scarce evidence for a main role of genes in these associations (Baird et al., 2001; Bateson, et al., 2004) thus indicating that vulnerability to adult diseases might rely upon fetal life experiences as a result of the redirection of the developmental trajectory.

Although for historical reasons Barker's theory was focused on the effects of prenatal undernutrition (see for instance the studies on the "Dutch famine" eg. (Roseboom, 2019) and

references therein), nowadays it is rather maternal obesity and its sequelae for metabolic and mental health that poses serious public health concerns (Congdon, 2019; Jehn and Brewis, 2009). Indeed, recent studies clearly suggest that maternal obesity, both before becoming pregnant and throughout gestation, is associated with negative short- and long-term health outcomes. Among all, a general developmental delay in children and poorer physical and mental abilities during aging are most often observed (see for instance the studies carried out on the Helsinki Birth Cohort (Berry et al., 2018; Eriksson et al., 2014; LifeCycle Project-Maternal et al., 2019; Mina et al., 2017; Westberg et al., 2016)) with OS playing a key role as a mediator of both short and long-term effects (Berry, et al., 2018; Edlow, 2017). Higher maternal body mass index (BMI) has also been associated with shorter telomeres length as assessed in leukocytes in elderly women (Guzzardi et al., 2016) moreover, umbilical cord gene expression profiling has identified patterns consistent with neurodegeneration/premature brain aging in fetuses of obese compared with lean women (Edlow et al., 2016b) suggesting long-term sequelae of early metabolic stress. To this regard, Alzheimer's disease (AD), a neurodegenerative disease affecting old age, has been recently defined as 'Type-3-Diabetes' because of some common molecular and cellular mechanisms with type 1 diabetes and T2D and insulin resistance associated with memory impairment and cognitive decline in old subjects (Kandimalla et al., 2017) suggesting that an early obesogenic environment might set the stage for a shared vulnerability to both metabolic and mental health issues.

The obesogenic womb and its sequelae

While for the mother the negative effects of obesity are readily observed as they affect fertility (ability to become pregnant) and may lead to obstetric complications, the effects on the offspring may not be immediately evident at birth although they have the potential to have both short- and long-term consequences (Alfaradhi and Ozanne, 2011; Contu and Hawkes, 2017; Iozzo et al., 2014; Moussa et al., 2016; Tenenbaum-Gavish and Hod, 2013).

Most importantly, in addition to the effects on metabolic programming, long-term longitudinal and associative studies provide evidence for a direct association between prenatal metabolic stress and an increased risk to develop neuropsychiatric, mood disorders and cognitive disabilities (see (Howell and Powell, 2017) and references therein). It should be mentioned that being obese during childhood may, by itself, trigger the onset of behavioral and emotional issues leading in turn to stigmatization and poor socialization, all conditions that may reduce

educational attainment and trigger the onset or precipitation of psychiatric disorders (Miller et al., 2015;Pizzi and Vroman, 2013). However, there is now evidence that, independently from offspring obesity, maternal obesity is associated with neurobehavioral problems in the offspring, although proving cause-effect mechanisms is rather hard in humans, given the many confounds deriving from the many intervening (genetic and environmental) variables involved in these effects. Nonetheless, there are now quite a number of epidemiological studies indicating that compared to children born from mothers having normal weight, children of obese mothers have a greater chance to show behavioral problems or being diagnosed with a neurodevelopmental disorder, such as ADHD (Godfrey et al., 2017). Regarding cognitive deficits, studies on the UK Millennium cohort have provided evidence for a negative relationship between maternal BMI and children's general cognitive ability at 7 years (Basatemur et al., 2013). A large synthesis and meta-analyses of the literature has more recently evaluated the association between pre-pregnancy overweight or obesity (in relation to normal weight) and subsequent childhood neurodevelopmental outcomes (Sanchez et al., 2018). This meta-analysis supports previous preclinical and observational studies indicating that children born from mothers obese prior to and during pregnancy are at increased risk for neurodevelopmental disorders.

Notwithstanding the above-mentioned evidence, it is rather difficult to establish whether maternal obesity affects neurobehavioral development in the offspring directly. In addition, maternal intelligence quotient, socio-economic status, breastfeeding vs. formula, maternal mental health, maternal diet and other postnatal lifestyle influences may concur to effects of maternal obesity (Contu and Hawkes, 2017;Godfrey, et al., 2017;Soubry et al., 2013). These questions can be more easily asked using animal models which allow to distinguish the confounding variables present in human studies, which are mostly observational in nature. In animal models, maternal obesity is mainly modelled by feeding dams before and/or during gestation with high fat-diets (HFD). Overall results from these studies enlarge and strengthen clinical and epidemiological evidence, indicating that offspring of obese mothers are characterized by social impairments, anxiety and depressive-like symptoms, in addition to cognitive disability and hyperactivity (Sullivan et al., 2015). One main important question that has been asked through preclinical models is whether pre-pregnancy obesity and maternal obesity exert the same effects on the offspring. This is an important point that can redirect prevention policies: it is much easier to address weight gain problems during pregnancy than act upon an obesity condition that might be acquired much before gestation. A recent

preclinical study appears to support epidemiological evidence indicating that prenatal and pregnancy windows have independent programming effects on the offspring. Preconception exposure affects body composition and adiposity while gestation exposure affects metabolism and tissue immune cell phenotypes (Chang et al., 2019). Overall, the current evidence suggests that pre-pregnancy obesity, rather than weight gain during gestation, may be most harmful to the fetus.

The field is very complex as also shown by inconclusive evidence provided by a very recent meta-analysis addressing results coming from animal models (Menting et al., 2019). Indeed, while clear effects of maternal obesity were observed on locomotor activity and anxiety, both increased by the metabolic maternal challenge, no significant effect was revealed by the meta-analysis on learning and memory performance in the offspring (Menting, et al., 2019). Although animal models may have some limitations, they have also been fundamental for understanding that the effects of an obesogenic environment experienced during fetal life depend upon multiple and not-exclusive pathways, which may lead to long-term pathological outcomes.

Mechanisms underlying fetal programming by maternal obesity

In complex organisms there is a finely-tuned crosstalk between physiological and behavioral responses, ultimately leading to the avoidance or to the adaptation to challenges. While acute responses to short-term stressors are pivotal to restore homeostasis, the management of a chronic stress condition imposes a non-negligible burden to the organisms (“allostatic load”) that might affect growth, metabolism, reproduction, inflammatory/immune and neuroendocrine function (de Kloet et al., 2005;Maccari and Morley-Fletcher, 2007;McEwen, 1998;Seckl, 2004).

Pregnancy is a highly demanding task involving physiological adaptations leading to a shift in the homeostatic balance (Mannaerts et al., 2018). Such changes, which would result in a pathological state in non-pregnant women, are (usually) well tolerated in healthy (pregnant) subjects. Although obesity *per se* cannot be considered a pathological condition, it represents a major risk factor for the onset and/or precipitation of many metabolic and cardiovascular non-communicable disease in addition to mood disorders (Armani et al., 2017;Chaput et al., 2012;Wurtman and Wurtman, 2018). Thus, being pregnant and obese at the same time may turn out to be an extremely stressful condition leading to a maternal allostatic load that will be

affecting the fetus. This will engage multiple, not mutually exclusive, mechanisms (during sensitive developmental phases), affecting tissue organization and organs' physiology in the offspring (Harris and Seckl, 2011;Iozzo, et al., 2014). Among these mechanisms, hyperactivity of the HPA axis and the associated excessive glucocorticoids (GCs) secretion, OS and inflammation, as well as gut microbiota dysbiosis, (just to mention a few) may all contribute to mediate the effects of maternal obesity, ultimately leading to vulnerability to neurodevelopmental and psychiatric morbidity in the offspring which could be embedded, through epigenetic changes, in selected genes.

Maternal metabolic stress shaping brain development

There is abundant preclinical and clinical evidence to indicate that adverse socio-economic conditions, in addition to physical, emotional or sexual abuse, are well-established risk factors for mood disorders (Ehlert, 2013;McEwen, 2000;McLaughlin et al., 2015;Raikkonen and Pesonen, 2009;Roseboom, 2019;Roth and Sullivan, 2005). As the studies of Barker and colleagues (Barker, et al., 1993;Seckl, 1998) previously detailed in this review, have listed body weight at birth within the set of risk factors that can increase susceptibility for adult diseases, maternal obesity needs to be listed as another yet adverse condition capable to affect offspring's development. Although, counterintuitively, obesity may involve a form of malnourishment, as it entails "poor nutrition". Indeed, one has to consider that, even when food is relatively prevalent, it may lack vitamins, minerals or other fundamental nutrients.

In order to understand the mechanisms underlying the detrimental effects of maternal obesity on brain developmental trajectories, we need to consider that this condition is likely to rely upon the same mechanisms regulating responses to stress (Cirulli F., 2017;McEwen, 2000). Acute activation of this pathway represents a "positive stress response", allowing the organism to face an acute threat. Under circumstances of chronic or overwhelming adversity - and in the absence of social support – responses to stress may become maladaptive for the individual and lead to pathological states because of an inability to face the current threats (or "allostatic load") (McEwen et al., 2015). The signaling pathways involved in the elaboration of stressful and metabolic signals, which are deranged during early development, might then be significantly modified in their functioning when a "second hit" occurs during critical periods, such as adolescence, or at adulthood (Bock et al., 2014;Boersma et al., 2014;Cirulli F., Berry A, Alleva E., 2003a;Daskalakis et al., 2013;Ehlert, 2013;Entringer et al., 2015;Provencal and Binder, 2015). Thus, the effects of metabolic or psychological stressors experienced at later stages will

be more disruptive in those individuals who have been exposed to stressors early during development.

Comorbidity between metabolic and behavioral disorders suggests shared developmental pathways and common mediators. Our group, together with many others, have clearly identified neurotrophins, such as BDNF, and GCs as main effectors of brain plasticity and metabolic regulations in response to stressful events (Cirulli F., Berry A, Alleva E., 2003a; Cirulli F. and Alleva, 2009; Cirulli F. et al., 2003; Levine, 1957; McEwen, 2000; Meaney et al., 1989).

There is a plethora of data from clinical and preclinical studies suggesting that the type and quantity of micronutrients, high levels of leptin and insulin as well as elevations in inflammatory mediators, such as interleukins and tumor necrosis factor can play a fundamental role in affecting developmental trajectories of the fetus by crossing the placental barrier (Godfrey, et al., 2017). Central to the maternal-fetal regulation is indeed the placenta (Fig. 1). This temporary organ, the main mother-fetus interface, is able to convey stress signals to the fetus by modulating the entry of hormones, glucose, amino acids, vitamins and ions necessary for fetal growth and development (Bale et al., 2010; Cirulli F., 2017; Fowden et al., 2009). Since elevated levels of GCs can be detrimental for the fetus, a specific enzyme, the 11 β -hydroxysteroid dehydrogenase type 2 (11 β -HSD-2), acts as a shield, guaranteeing their rapid inactivation and allowing an optimal amount of these hormones to be transferred to the fetus for organs maturation. It is rather interesting that psychological stressors and metabolic/nutritional challenges can both affect the expression or the activity of the 11 β -HSD-2 enzyme, with high levels of GCs reaching the fetus, resulting in increased susceptibility for psychiatric disorders and metabolic complications linked to an increased risk of insulin resistance and T2D, in addition to CVD and metabolic disorders associated with obesity (Eriksson, et al., 2014; Mansur et al., 2015; Mina et al., 2015; Rivera, et al., 2015; Stout et al., 2015).

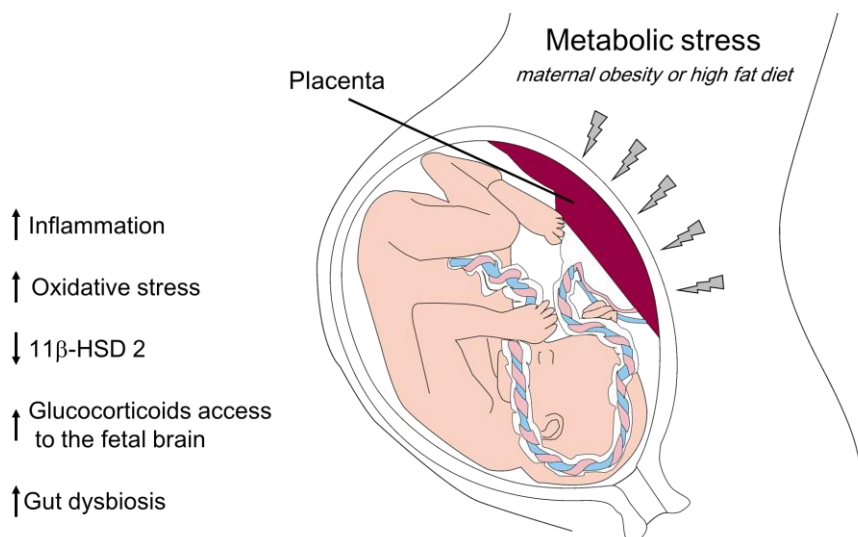


Fig. 1. Potential mechanisms underlying the effects of maternal obesity on fetal development. The placenta is the critical mother-fetus interface transducing metabolic stress into changes in the developing organism underlying the short- and long-term effects of maternal obesity (see text for a detailed description).

In recent studies we have clearly shown that HFD feeding during pregnancy (in mouse models), independently from maternal obesity, is a stressful challenge that is able to influence negatively the fetus, which is already exposed to excess GC as a result of maternal pregnancy status (Bellisario, et al., 2014; Bellisario et al., 2015; Eriksson, et al., 2014). HFD-feeding in female mice before conception and during pregnancy, in fact, increases levels of maternal stress hormones and is associated with both reduced 11 β -HSD-2 enzymatic activity and expression of the 11 β -HSD-1 gene in the placenta (Bellisario, et al., 2015). Exposure to a diet with high fat content has also been shown to disrupt maternal behavior at parturition with deleterious consequences on the offspring (Bellisario, et al., 2015). These dysregulations, observed in HFD-fed dams, might be caused by changes in neural activity in brain regions responsible for olfactory processing and social recognition, ultimately leading to inappropriate maternal behavior (Bellisario, et al., 2015). These effects are less apparent in animal models of reduced OS, especially in female subjects, suggesting that diet-induced metabolic dysfunction, neuroendocrine response and sex/gender all play a role in these regulations (see later in this review) (Bellisario, et al., 2014).

Worth noticing, these data also confirm clinical evidence showing that the presence of high levels of GC during gestation can affect the expression of GC-sensitive genes in the central nervous system, as well as in the periphery, with important effects on HPA axis function and

regulation (Mina, et al., 2017;Mina, et al., 2015). These effects are reminiscent of a solid literature in animal models indicating that prenatal maternal stress affects HPA axis activity and stress regulation through changes in GC receptors in limbic regions of the brain (Brunton and Russell, 2010;McGowan et al., 2008;Welberg and Seckl, 2001). We can thus hypothesize that, by affecting the amount of GC reaching the fetus, HFD during pregnancy could lead to similar short- and long-term effects already described for prenatal stress.

A further intriguing aspect is related to the modulating effect of the HPA axis on immune function and inflammation. The group of Sasaki (Sasaki et al., 2014) investigated the expression of several inflammatory genes linked to GC signaling (Sorrells et al., 2009) as a result of perinatal exposure to HFD (Bilbo and Tsang, 2010). Results suggest a dysregulation of pro- and anti-inflammatory genes linked to HFD exposure in utero with NF- κ B and IL-6 transcripts being increased in the hippocampus in HFD-exposed offspring. Moreover, increased expression of the cytokine IL-6 in the hypothalamus (De Souza et al., 2005) and cortex (White et al., 2009) is an established consequence of prolonged HFD exposure in utero.

Chronic systemic inflammation characterizes both maternal obesity and pregnancy. Obese pregnant women have higher levels of circulating pro-inflammatory cytokines while maternal BMI is directly correlated with (maternal) pro-inflammatory cytokine concentrations and the activation of pro-inflammatory placental pathways. Changes in fetal cytokine expression, fetal neuronal damage and changes in gene expression in the neonatal brain have been previously related to placental and intrauterine inflammation (Edlow, 2017). Inflammation during pregnancy might alter the connectivity of brain networks and be associated with altered neurobehavioral regulations and reduced memory performance in early childhood. Indeed, it has been recently reported that maternal IL-6 during pregnancy correlates with newborn brain connectivity and can predict future working memory in offspring (Rudolph et al., 2018). Thus, changes in inflammatory mediators could be important biomarkers to be used to estimate long-term effects of maternal obesity on brain function.

Role of the microbiota

Long-lasting effects on fetus and newborn resulting from exposure to maternal obesity during pregnancy could also be explained by an alteration in the composition of the gut microbiota during critical developmental windows (Cryan, 2016). Each individual owns a unique gut microbiota profile composed by different strains of bacteria. Changes in this balance may lead

to a condition of dysbiosis (Rinninella et al., 2019). Recent studies associate dysbiosis with different gastrointestinal and metabolic diseases such as diabetes and obesity but also with neurodegenerative and mental disorders, including AD and Parkinson's disease (PD), ASDs, schizophrenia and depression (Heiss and Olofsson, 2019). This suggests that the balance of the gut microbial composition may play a key role in the regulation of different developmental processes (De Palma et al., 2015;Dinan and Cryan, 2016;El Aidy, et al., 2016;Foster et al., 2016;Kelly, et al., 2016;Luczynski et al., 2016).

The early life phases are crucial for the development of the microbiota and to define its ultimate role in the programming of tissues and organs (Dimmitt, et al., 2010). A number of studies show that the gut microbiota influences brain function (Dinan and Cryan, 2016;El Aidy, et al., 2016;Foster, et al., 2016). Notwithstanding the fact that the infant gut microbiota has been worldwide considered to be determined after birth, recent evidences have shown how the whole intrauterine environment (placenta, amniotic fluid, meconium and umbilical cord blood) is actually colonized by specific bacterial species (Collado et al., 2016). These data robustly suggest the transmission of microorganisms from the mother to the fetus ahead of birth and this plays a pivotal role in the determination of the offspring microbiota (Aagaard et al., 2014;Collado, et al., 2016;Koleva et al., 2015;Satokari et al., 2009). Transfer of microbes and microbiome metabolites between mother and infant can also occur during delivery and lactation and can be affected by maternal health and metabolism, in addition to mode of delivery and use of antibiotics (Soderborg et al., 2016). Hence, dysbiosis and obesity during pregnancy can dramatically affect microbiota composition of the mother as well as microbial transmission and, in turn, microbial transmission to the fetus (Basu et al., 2011), maternal diet having one of the largest known impacts on these regulations.

The maternal gut microbiota differs in maternal obesity, particularly in the latter half of pregnancy, with overweight women being characterized by an increase in the Firmicutes phylum (*Staphylococcus*) as well as increases in some Proteobacteria (Soderborg, et al., 2016). Elevated levels of microbe-derived plasma endotoxin appear to be one potential mechanism by which maternal microbes in obesity could affect the developing fetus; this might increase translocation of bacteria-derived products across the intestinal mucosa, contributing to systemic and placental inflammation and insulin resistance (Basu, et al., 2011).

Adverse effects of maternal obesity or excess gestational weight gain have in fact been linked to inflammation in the placenta. Although the specific mechanisms through which the bacterial transmission is carried out are not currently well known, a new concept which is developing is

that the placenta microbiota may be the intermediary of the microbial passage from the mother to the baby and may affect fetal development (Pelzer et al., 2017). An important subject for future studies is the role of placenta microbiota and its interaction with other maternally-derived variables, including life style, diet, BMI and pregnancy complications, which could all alter placental microbiota (Pelzer, et al., 2017).

Role of Oxidative Stress

OS characterizes biological systems in aerobic conditions; it results from an unbalance between pro-oxidant and anti-oxidant molecules, with oxidants overriding the defense system of the organism. During normal healthy pregnancy, OS and its mediators - reactive oxygen species (ROS) - are increased within certain boundaries as the result of a physiological mild inflammatory state, to stimulate cell proliferation and proper fetal development (Dennery, 2007;Mannaerts, et al., 2018). However, if OS overrides safety levels, complications might arise both for the mother and the offspring (Edlow, 2017;Hracsko et al., 2008;Mannaerts, et al., 2018). OS is increased in many different pathological conditions including obesity, T2D, and metabolic syndrome and can be triggered by chronic consumption of HFD. There is clear evidence for disrupted oxidative signaling in psychiatric disorders. Alterations in multiple biomarkers of OS have been observed in ADHD, bipolar disorder, ASD, depression, and schizophrenia (Hovatta et al., 2010). Markers of oxidative damage to neurons have also been observed in post-mortem samples in several psychiatric diseases. It is of interest that mitochondrial disorders with a clear genetic origin are also associated with an elevated incidence of psychiatric disorders, especially mood disorders and psychosis. Increased ROS generation might lead to random damage to proteins, lipids and DNA and excessive OS during pregnancy might result in pathological conditions of the placenta, the embryo, and the fetus, also leading to epigenetic changes and altered gene expression in the fetus due to DNA damage (Del Rio et al., 2005;Mannaerts, et al., 2018). The placenta is a main source of OS and this organ, in obese women, is characterized by greater levels of OS markers when compared to that of lean women (Saben et al., 2014). Edlow and colleagues found an upregulation of genes related to OS response as a result a global gene expression profiling carried out in the amniotic fluid from obese women (Edlow et al., 2014). Intriguingly, among these, Apolipoprotein D, a protein that is highly expressed in the brain and is upregulated in psychiatric conditions and neurological disorders (Muffat and Walker, 2010;Sutcliffe and Thomas, 2002), was found to

be increased by nine-fold in fetuses of obese compared with lean women (Edlow, 2017;Edlow, et al., 2014).

A growing body of evidence suggest that OS may play a role in the etiology of mood disorders (Hovatta, et al., 2010). A number of studies indicate that exposure to chronic stress may perturb the overall body energy balance acting not only on the neuroendocrine system but also on mitochondrial remodeling, affecting OS balance in the body and in the brain (Picard and McEwen, 2014;Picard and McEwen, 2018;Picard et al., 2018). Recent work suggests that isoprostanes (biomarkers of lipid peroxidation) are selectively upregulated in adolescents who have experienced early childhood adversities, suggesting that dysregulation of stress-sensitive systems during early life stages can have persisting, and potentially deleterious, impact on brain structure-function development acting through OS-linked mechanisms (Horn et al., 2019). The mammalian brain is very sensitive to OS insults being characterized by high metabolic rate, poor antioxidant defenses and reduced capacity for cellular regeneration (Floyd and Carney, 1992). Thus, a prenatal insult such as an obesogenic womb, characterized by elevated levels of inflammation and OS, has the potential to affect dramatically the neurodevelopmental programming of the fetus, setting the stage for later life vulnerability to psychiatric disorders.

Preclinical data enlarge and corroborate this body of evidence and the p66Shc^{-/-} mouse model - that our group has thoroughly characterized - provides a striking example of how prenatal HFD feeding might impinge upon OS pathways to affect fetal programming and healthspan (Berry et al., 2012;Berry et al., 2007;Berry et al., 2010;Berry and Cirulli, 2013;Berry et al., 2008). p66Shc is a mammalian gene that regulates apoptosis by increasing intracellular OS and affects lipid metabolism by promoting fat storage (Berniakovich et al., 2008;Trinei et al., 2009). p66Shc is a protein acting specifically in the mitochondrion as a redox enzyme that generates H₂O₂ and whose function is in a cause-effect relationship with that of insulin/IGF1(Berniakovich, et al., 2008). Genetically modified mice lacking the p66Shc gene (p66Shc^{-/-}, knock out mice - KO) are resistant to OS insults and to HFD-induced obesity, resulting overall in a healthier and long-lived phenotype, showing greater brain and behavioral plasticity associated to increased central levels of the neurotrophin BDNF (Berry, et al., 2012;Berry, et al., 2007;Berry, et al., 2010;Berry and Cirulli, 2013;Berry, et al., 2008). Prenatal exposure to HFD in these KO mice leads to a gender-specific resilience to both stressful and metabolic challenges (Bellisario, et al., 2014). Administration of a HFD during peri-conceptional time and throughout pregnancy resulted, in fact, in reduced body weight at birth

and into a greater catch-up, particularly in males, while female offspring showed increased BMI as well as higher leptin levels. By contrast, p66Shc^{-/-} subjects were overall protected (Bellisario, et al., 2014), p66Shc^{-/-} females being characterized by an improved ability to cope with both metabolic and neuroendocrine stressors and showing enhanced glucose tolerance and insulin resistance. Moreover, while prenatal HFD led to a hyperactive HPA axis in wild type offspring, this was not observed in p66Shc^{-/-} mice (particularly in females) overall suggesting that sex-hormones might play an important part in the directing the effects of a metabolically stressful maternal challenge during fetal development (Bellisario, et al., 2014). Data on gender differences have been further extended by Edlow and colleagues (Edlow et al., 2016) who found that maternal obesity results specifically associated with sex-specific differences in fetal size and fetal brain gene expression signatures, males being the most vulnerable sex. In these studies, ROS metabolism was found to be affected by HFD (Edlow, et al., 2016) further strengthening the important role played by OS mediators in maternal obesity.

Translating these preclinical findings to humans, investigating a group of women enrolled from the Helsinki Birth Cohort, we found a long-term increase in p66Shc mRNA in peripheral blood mononuclear cells (PBMC) of old frail subjects born from obese mothers (Berry, et al., 2018). Moreover, when these women were stratified according to their BMI, PBMC p66Shc expression levels were reduced in subjects with a BMI \geq 30.4 kg/m² following physical exercise. This piece of data is particularly interesting for a number of reasons. First of all, it confirms a role for oxidative stress in human subjects in the signaling pathway linking maternal obesity to the health outcome in the offspring. Secondly it suggests that maternal obesity may play a role in the long-term programming of the mitochondrial function and metabolism (also through changes in the p66Shc expression) affecting the aging process. Moreover, it indicates that metabolic-challenging stimuli, such as physical exercise, may affect the expression of p66Shc, particularly in obese subjects, pointing to this molecule as a potential new target for therapeutic intervention studies (Berry, et al., 2018; Berry and Cirulli, 2013).

Long-lasting effects of maternal obesity - epigenetic signatures

Epigenetic modifications have been proposed as a key causal mechanism linking maternal adiposity and offspring outcome. Long-term effects of early nutritional experiences are potentially mediated by post-translational modifications of DNA, post-translational modification of histones and non-coding RNAs. One of the most interesting consequences of

these mechanisms is that they can account for transgenerational transmission of traits (Bock, et al., 2014;Daskalakis, et al., 2013;Provencal and Binder, 2015;Szyf, 2012;Szyf et al., 2005;Turecki et al., 2014).

To date, very little work has been performed to determine epigenetic changes in the brains of human offspring born to obese mothers. DNA methylation changes have been reported in cord blood and microRNA levels in amniotic fluid in human studies of maternal obesity, indicating a potential role for these epigenetic mechanisms in the long-term effects of maternal obesity on the offspring (Godfrey, et al., 2017). Cohort studies analyzing BMI extremes in these data sets found associations between maternal BMI and offspring DNA methylation at birth and at 3 years. However, more work is needed to study these regulations.

In candidate gene approach studies, one of the most important findings concerns the observation that aryl-hydrocarbon receptor repressor (AHRR) DNA methylation is 2.1% higher in offspring of obese vs. normal weight mothers; as robust links have been found between maternal smoking and offspring AHRR methylation (Reynolds et al., 2015), these observations suggest that methylation of these gene may be involved in the link between multiple adverse conditions, including maternal obesity, on offspring outcomes.

Among the classical epigenetic mechanisms, DNA methylation is an attractive target for investigation because levels of folic acid, a co-factor in the production of the methyl donor methionine, are decreased in the amniotic fluid of obese pregnant women (Contu and Hawkes, 2017;Mohd-Shukri et al., 2015). In clinical studies, maternal depressive symptoms during pregnancy have been found to correlate with increased DNA methylation of the GR (NR3C1) in male infants and to result in decreased BDNF IV DNA methylation in both sexes at 2 months of age (Braithwaite et al., 2015). BDNF plays also a critical role in the integration and optimization of behavioral and metabolic responses: by acting in the brain and periphery, it increases insulin sensitivity and parasympathetic tone (Cirulli F. and Alleva, 2009). Low levels of circulating BDNF characterize individuals with obesity and T2D, which implies a main role for this neurotrophin in obesity and metabolism (Mou et al., 2015). While methylation of the NRC31 has been related to prenatal early adversity in numerous studies (van der Knaap et al., 2015), these results indicate that genes involved both in stress responsivity and in feeding behavior are epigenetically regulated, supporting the notion of joint programming of these stress-activated pathways and allowing for more detailed studies of the mechanisms underlying comorbidity of mental and metabolic disorders (Rivera, et al., 2015).

Preclinical studies performed in animal models (Vucetic et al., 2010) have indicated decreased DNA methylation in the promoter regions of the mu-opioid receptor genes as well as globally in brain regions associated with reward such as the ventral tegmental area (VTA), PFC, and nucleus accumbens (NAc) of offspring born to obese mothers (Carlin et al., 2013). Supplementation of maternal HFD with methyl donors during gestation and lactation was able to restore some of these effects (Carlin, et al., 2013). Prenatal methyl supplementation has also been shown to reverse deficits in motivated behavior in offspring exposed to a HFD indicating that early life is a sensitive time during which dietary methyl donor supplementation can alter PFC-dependent cognitive behaviors (McKee et al., 2017). Differential expression of 37 microRNAs, rather than changes in DNA methylation (Edlow, 2017) have been reported in the brains of embryonic mice born to mothers fed a HFD vs. control diet.

Preventive strategies

As mentioned above, maternal obesity affects multiple interconnected physiological pathways engaged by complex regulations. Thus, addressing diets and dietary (particularly antioxidants) supplementation might provide a “broad-spectrum” promising a feasible strategy to prevent/counteract the disruptive effects of the obesogenic womb, especially considering the difficulty in the development of new target-specific pharmacological interventions. In this paragraph we will specifically focus on some of the most promising dietary intervention that include also oral supplementation with the antioxidant N-Acetyl-Cysteine (NAC).

NAC is the rate-limiting substrate in the biosynthesis of glutathione (GSH). It is a ROS scavenger and its clinical efficacy as well as safety have been recently documented in many pathological conditions (see Mokhtari et al., 2017 and references therein for a complete review). NAC is currently one of the most promising targets for neuropsychiatric disorders. Its most interesting feature is that its efficacy appears cross-diagnostic (acting on neurotransmitter systems such as glutamate, antioxidants as well as inflammatory pathways (Berk et al., 2013)), while being relatively safe. Determining precisely how antioxidants - in particular NAC -work is crucial both to understand the basic biology of mental disorders and in order to devise adjunctive therapies that can work on these pathways. Indeed, the apparent universality of NAC action is intriguing and implies that it may target downstream pathways underlying comorbidity between stress and metabolic responses. Although there is only preliminary data of the efficacy of NAC in many of psychiatric disorders, the field is expanding with many

additional trials that could be of interest also in the field of maternal obesity pathways (Berk, et al., 2013).

So far, some preclinical studies have focused on the effects of this molecule in the context of maternal obesity. NAC administration during pregnancy in mice might counteract apoptosis and ROS-related genotoxicity by increasing glutathione levels and decreasing mitochondrial membrane depolarization (Amin et al., 2008). Indeed, mice lacking the rate-limiting enzyme for GSH synthesis show a range of behavioral disorders and treatment with NAC reverses some of these deficits, restoring GSH levels (Berk, et al., 2013). NAC may also contribute to maintain oxidative balance through the action of the cysteine/cystine cycle (Elshorbagy et al., 2012). We have recently observed that the administration of NAC in animal models throughout pregnancy is able to buffer the effects of HFD on both the mother and the offspring. In our studies, offspring of mice supplemented with NAC during fetal life showed improved glucose tolerance, in addition to a reduced activation of the HPA axis, when exposed to stress (Berry et al., 2018). Worth noticing, results obtained through *in vivo* NAC supplementation parallel the protective effects observed in the p66Shc^{-/-} mice (a mouse model of reduced oxidative stress; see above) strengthening the link among metabolic stress experienced during fetal life, fetal programming and OS/mitochondrial pathways (Berry, et al., 2018; Berry, et al., 2007; Berry and Cirulli, 2013; Giorgio et al., 2012; Hovatta, et al., 2010).

In addition to NAC, plant-based-antioxidant-rich diets could protect against free radical production and oxidative damage, potentially preventing obesity and comorbidities. As an example, luteolin supplementation of obese, HFD-fed adult mice has been found to be associated with better cardiometabolic features and reduced inflammatory and oxidative stress markers (Gentile et al., 2018). Research on these diets could extend to the prenatal period in the future.

Poor prenatal diets have been often associated with poorer maternal and offspring mental health outcomes. Thus, the quality of maternal diet represents a potentially important and modifiable target for reducing the risk of mental disorders both in mothers and in the offspring (Lindsay et al., 2017). To this regard, longitudinal studies in clinical populations are warranted in order to provide further insights for the development and design of future and more effective intervention trials for improved maternal and child health outcomes (Lindsay, et al., 2017). So far, observational data in humans has pointed to polyunsaturated fatty acids (PUFAs), including omega-3 and omega-6 fatty acids, as possible candidate therapeutics in maternal obesity. In

particular, in human pregnancy studies data suggest that omega-3 PUFA may play a critical role not only on fetal neurodevelopment, but also for supporting positive maternal mood, decreasing stress, anxiety and depression in the pre- and postnatal periods (Lindsay, et al., 2017). Maternal omega-3 fatty acid deficiency has been associated with increased risk of offspring ASD and ADHD. Human pilot studies relying on the supplementation of obese pregnant women with omega-3 fatty acids demonstrated a reduction in maternal and placental inflammation (see (Lindsay, et al., 2017) for a review).

In preclinical studies it has been shown that prenatal exposure to dietary omega-3 fatty acids is able to stage key relationships between metabolic signals and BDNF methylation, creating an epigenetic memory and a reservoir of neuroplasticity that can protect the brain against later insults (Tyagi et al., 2015). In animal models, early life methyl donor supplementation has been used as a strategy to alter one carbon metabolism and DNA methylation in a sex-dependent manner (McKee, et al., 2017). In particular, supplementation of methyl donors in HFD fed dams has been able to attenuate weight gain and decrease fat preference (males) as well as to change gene expression and global hypomethylation in the brain of the offspring (McKee, et al., 2017). A methyl-balanced diet can also prevent the effects of prenatal stress on binge eating behavior (Schroeder et al., 2017). In addition, early micronutrient supplementation can protect from early life stress-induced cognitive impairments, thus reinforcing the notion of common pathways that are engaged by early nutrition and early life stress (Naninck et al., 2017).

We have previously argued that the negative effects of maternal obesity on mental disorders in the offspring may be mediated, at least in part, by gut dysbiosis. Changes in the microbiome and its metabolites offer the opportunity for non-invasive risk-screening and risk-reduction by tailored dietary formulations and diets, since earliest stages of life. Changing the composition of the gut microbiota through the use of probiotics and prebiotics could become a new strategy for reducing the risk of metabolic disorders in both the mother and offspring. In order to do this, it is necessary to identify clinically relevant markers and build early predictive models, which might help tailoring appropriate dietary interventions. Dietary interventions aimed at modifying gut microbiota development and composition in the first phases of life may promote cognitive and emotional development. The use of pre- and probiotics has been shown to rapidly modify the microbial community and reduce (at least temporarily) adiposity and chronic inflammation in animal models of obesity and in limited human studies. As far as clinical studies are concerned, there have only been a handful of randomized clinical trials (RCTs) that

examined the effects of probiotics administered during pregnancy with the aim of improving insulin sensitivity and reducing GDM diagnosis, with limited success. In one RCT either *Bifidobacterium lactis* alone or *B. lactis* plus *Lactobacillus rhamnosus* GG probiotic was administered to pregnant women 14 days before a scheduled Caesarean delivery (Baldassarre et al. 2018). Modulation of the gut microbiota by prebiotics, probiotics or by fecal microbiota transplantation has been used in the management of certain neurological disorders, including autism and depression, as well as their associated gastrointestinal symptoms, and increasing numbers of clinical trials suggest the beneficial effects of such treatments. Fermentable fiber influences gut microbiota composition and the production of short chain fatty acids; these, in turn, have been demonstrated to affect mood in animal models, as well as influence neurotransmitters such as serotonin although additional studies are needed to determine the role of different microbes in the physiology of the host and pathogenesis, as well as how the gut microbiota can be modulated for beneficial effects. Most of the interventions used mainly concern the prenatal phases, but they could be extended postnatally. It is rather difficult to discern the effects of prenatal maternal obesity to maternal obesity during lactation. Based upon an animal model, as an example, we were able to show that lactating mice fed a high fat diet during pregnancy (and not lactation) still showed inappropriate maternal behavior after parturition (Bellisario, et al., 2015). There is evidence from epidemiological studies that overweight and obese women are less likely to breastfeed than normal weight women (Amir and Donath, 2007).

We believe that it is especially important to stress here is that all the data above mentioned have to be taken with due caution given the relative paucity and high levels of heterogeneity of the studies present in the literature. In particular, it is imperative that further controlled trials be performed in order to determine the type of antioxidants vs pre- and probiotics and the most appropriate dose.

Based upon the above-mentioned considerations, and data gathered through RCTs, we can draw some hypothetical scenarios on the proper timing of an ideal intervention and their possible consequences. We expect that the negative effects of being exposed to maternal obesity might manifest themselves at different time points throughout life, also in relation to a second hit (being this a stressful event or another obesogenic stimulus), starting from the early developmental phases to adolescence, up until aging. We hypothesize different potential windows of intervention: the earlier we intervene the greater and long-lasting the benefit.

Ideally, the timing of the intervention should be set during brain development (before critical periods are closed) and, hopefully before the end of adolescence/puberty. Adolescence is a critical time for brain and behavioral development ultimately leading to the transition from childhood to adulthood. This peculiar time is characterized by a massive neuronal remodeling powerfully shaped by hormones, thus representing a very vulnerable time in life as well as a window of opportunity. With late interventions (middle-age or old age) we might still be able to act on residual neuronal plasticity, ameliorating the outcome of neurodegenerative disorders such as AD or PD (see Fig. 2).

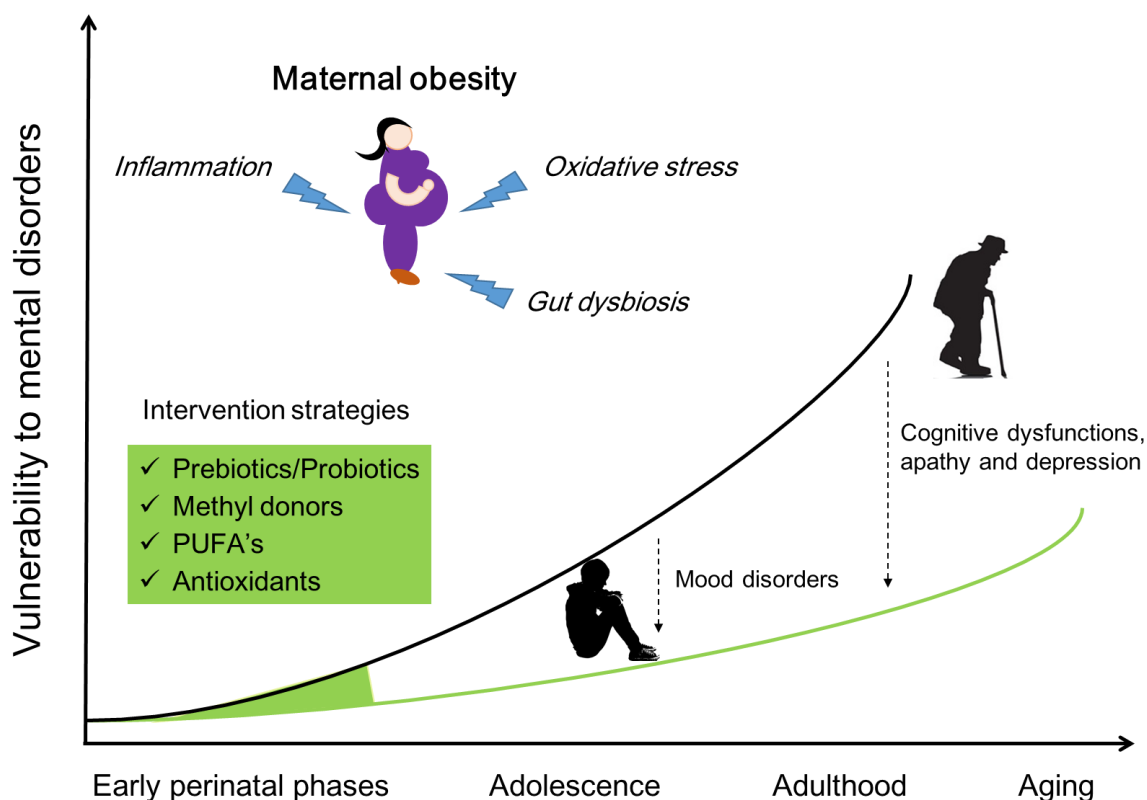


Fig. 2. Hypothetical scenarios on the proper timing of the interventions and their possible consequences. The ability of the organism to cope with the sequelae of the obesogenic womb (oxidative stress, gut dysbiosis and inflammation) decreases throughout life as a result of the physiological decrease in neuronal and behavioral plasticity. Different windows of intervention may be foreseen with earlier timing (perinatal phases until adolescence) leading to greater and longer-lasting benefits, later interventions (adulthood until aging) being able to act only on residual neuronal plasticity, ameliorating the outcome of age-related neurodegenerative disorders. The earlier the intervention (green area within the curves) the greater the potential to reduce disease risk later in life (dashed arrows). *Figure modified from Godfrey KM, Gluckman PD, Hanson MA. Developmental origins of metabolic disease: life course and intergenerational perspectives. Trends in endocrinology and metabolism: TEM. 2010;21(4):199-205.*

Conclusions

Although initial research linking prenatal development with major non-communicable disorders in later life focused on the effects of fetal undernutrition, nowadays it is rather maternal obesity and its sequelae for metabolic and mental health that poses serious public health concerns throughout the world (Congdon, 2019; Jehn and Brewis, 2009). Indeed, an ever-increasing body of epidemiological data - that are corroborated by preclinical evidence - suggest that exposure to an obesogenic womb might strongly impact offspring healthspan throughout life, setting the stage for an increased vulnerability to a number of pathological conditions including obesity, CVD, stroke, T2D and asthma (just to mention few), in addition to neuropsychiatric disorders. As recently pointed out by Godfrey and colleagues (Godfrey, et al., 2017), there is a need of thorough large-scale studies tackling, for example, the differential impact played by maternal diet and maternal obesity as well as maternal obesity during pregnancy vs obesity during the lactation period. There may be a mix of psychological, behavioral and cultural factors underlying such effects.

These studies should be performed including populations differing in terms of culture as well as ethnicity and collecting detailed data on the cognitive and behavioral phenotype of both parents and offspring; in addition, a comprehensive assessment of diet and of measures of adiposity should be performed (Godfrey, et al., 2017).

From an evolutionary perspective, foraging behavior in harsh environments, a condition opposed to maternal obesity and characterized by limited and intermittent food availability, has most likely contributed to shape our brain to deal more effectively with spatial navigation, decision-making, social relationships and even creativity (Berry and Cirulli, 2013; Mattson, 2019). Globalization and urbanization, by favoring sedentary lifestyle and the consumption of high-caloric foods, are gradually leading to a change in the energetic niche experienced both prenatally and early postnatally by humans with consequences that are hard to predict: the ultimate outcome of such a metabolic shift is now starting to show an impact on pre- as well as post-natal developmental trajectories with potential disruptive effects on cognitive processes and susceptibility to mental disorders. Epigenetic changes and dysbiosis linked to such processes have the potential to propagate these modifications to future generations.

While public health countermeasures that can rapidly break the vicious cycle of maternal and offspring obesity are mandatory, we are definitively in need of basic knowledge and appropriate longitudinal, controlled studies that can address the basic mechanisms underlying

such regulation, allowing for appropriate intervention but, more importantly, indicating appropriate biomarkers and effective preventive measures.

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Conflicts of Interest

The authors declare that there is no conflict of interest regarding the publication of this paper.

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