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Early life high fat diet exposure maintains glucose tolerance and insulin sensitivity with a fatty liver and small brain size in the adult offspring

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Abstract

Diet during pregnancy has long lasting consequences on the offspring, warranting a study on the impact of early exposure to a high fat diet on the adult offspring. We hypothesized that a prenatal *n-6* enriched diet will have adverse metabolic outcomes on the adult offspring that may be reversed with a postnatal *n-3* enriched diet. To test this hypothesis, we examined the adult offspring from three groups: 1) *n-6* group: during gestation and lactation, dams consumed an *n-6* polyunsaturated fatty acid enriched diet, 2) *n-3* group: gestational *n-6* diet was followed by an *n-3* enriched diet during lactation, and 3) a control (CD) group that received standard diet throughout gestation and lactation. Offspring from all groups weaned to a control diet *ad libitum*. Beginning at postnatal day 2 ($p < 0.03$) and persisting at 360d in males ($p < 0.04$), an increase in hypothalamic AgRP expression occurred in the *n-6* and *n-3* groups, with an increase in food intake ($p = 0.01$), and the *n-3* group displaying lower body ($p < 0.03$) and brain ($p < 0.05$) weights. At 360d, the *n-6* and *n-3* groups remained glucose tolerant and insulin sensitive, with increased phosphorylated-AMP-activated protein kinase ($p < 0.05$). *n-6* group developed hepatic steatosis with reduced hepatic reflected as higher plasma *microRNA-122* ($p < 0.04$) that targets pAMPK. We conclude that early life exposure to *n-6* and *n-3* led to hypothalamic AgRP-related higher food intake, with *n-6* culminating in a fatty liver partially mitigated by postnatal *n-3*. While both diets preserved glucose tolerance and insulin sensitivity, postnatal *n-3* displayed detrimental effects on the brain.

Keywords

rats; high-fat diet; polyunsaturated fatty acids; brain; liver

1. Introduction

There is an obesity epidemic in the United States (US), which has extended into childhood. Between the ages of 12 to 19 years, 21% of US children are obese and 8% are severely

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obese [1]. This health crisis has led to an increased incidence of hypertension, dyslipidemia, cardiovascular disease, insulin resistance, and non-alcoholic fatty liver disease [2-7]. These diseases cause blindness, stroke, and renal and liver failure, all of which can be fatal and tax our health care economy [8,9]. Not surprisingly, 32% of women of childbearing age are obese. Once pregnant, obese women are at high risk for gestational hypertension and diabetes mellitus, both of which have adverse, long-lasting repercussions on the woman's health and that of the offspring extending into adulthood [10,11].

On this backdrop a considerable amount of attention has been given to the type of diet that pregnant women consume. There has been a large push to increase consumption of fat-reduced foods with the goal of reducing cholesterol concentrations, a major risk factor for cardiovascular accidents and myocardial infarctions [7,12]. However, these fat-reduced products have not abated the obesity epidemic and suggest that the replacement of fat with carbohydrates may negatively impact the adult phenotype of the offspring. For these reasons, it is important to determine the optimal diet during pregnancy and lactation [13]. The composition of fat, especially the *n*-6:*n*-3 polyunsaturated fatty acid ratio, is important for overall health. The recommended dietary *n*-6:*n*-3 fatty acid ratio is 2:1-4:1 for humans [14]. However, western diets typically exceed a ratio of 15:1 [15]. Excessive *n*-6 fatty acid intakes are associated with obesity, diabetes, and non-alcoholic fatty liver disease [16,17]. On the other hand, increased *n*-3 provisions are associated with improved glucose tolerance, insulin sensitivity and a decreased incidence of non-alcoholic fatty liver disease [18-21]. *n*-3 fatty acids also play a critical role in the development of the central nervous system and somatic growth [18,19].

Rodent investigations have shown that when maternal hyperglycemia due to maternal diabetes is encountered, the offspring develops glucose intolerance and insulin resistance [22-24]. More recently, a high fat diet consumed at least 2 months before conception and continued during gestation and lactation led to the adult offspring with fatty liver complicated by glucose intolerance and insulin resistance [25-27]. *n*-3 supplementation ameliorates this phenotype [28-30]. However, the birth weight of offspring exposed to a high fat diet was reduced as opposed to the expected increased birth weight [31,32], typically seen in humans. This observation raises the question whether the metabolic perturbations that manifest in the adult rodent offspring are because of fetal growth restriction or exposure to a high fat diet initiated prior to conception and continued during fetal life. Further, these studies support the premise that an abnormally high pre-conception body mass index signifying pre-existent obesity due to a high caloric diet has adverse effects on the fetus that may last into adult life. However in these investigations, it is difficult to separate the effects of pre-existing maternal obesity and its associated metabolic and inflammatory changes from the effect of maternal hypercaloric diet alone, on the ultimate phenotype of the adult offspring.

In our present study, we focused on diet content during pregnancy and lactation alone, rather than before conception, where a propensity towards pre-pregnancy obesity may confound the diet-induced observations. We hypothesized that an *n*-6 rich high fat diet during gestation initiated after conception and lasting through lactation in non-obese dams would lead to lesser obesity in the offspring, associated with milder glucose intolerance, insulin

resistance, and a fatty liver phenotype. Secondly, we hypothesized that an *n*-6 rich high fat diet during gestation followed by an *n*-3 rich high fat diet during lactation would mitigate the prenatal *n*-6 exposure induced changes, even if milder, by maintaining glucose tolerance and insulin sensitivity, and preventing fatty liver. To test these hypotheses, we used a prenatal and postnatal dietary modified rat model employing three groups, control with ad libitum access to regular chow diet (CD group), *n*-6 rich high fat diet during gestation and lactation (*n*-6 group), and *n*-6 rich high fat diet during gestation followed by an *n*-3 enriched high fat diet during lactation (*n*-3 group). All three groups had ad libitum access to water. To meet our objectives, we then compared insulin sensitivity and glucose tolerance, and the state of the liver in these three groups. Because polyunsaturated fatty acids play an important role in central nervous system development, we also assessed brain weight, certain neural markers and specific energy balance regulating hypothalamic neuropeptides.

2. Methods and Materials

2.1 Animals

2.1.1 Animal Care—Sprague-Dawley rats (Charles River Laboratories, Hollister, CA) were housed under 12h light/dark cycles at ~21-23°C and allowed ad libitum access to water [33]. The care and use of all animals in this study were approved by the Animal Research Committee of the University of California Los Angeles (protocol no. 1999-104-61A), and were in keeping with the guidelines issued by the National Institutes of Health.

2.1.2 Animal Model—The experimental design (Figure 1) consisted of creating two pregnant high fat exposed groups following conception towards preventing pre-pregnancy obesity in dams. Gestational day 2 through day 21, rats had ad libitum access to a control regular chow diet (CD; NIH-31 modified 7013 diet: Teklad Inc., Madison, WI) with the *n*-6:*n*-3 fatty acid ratio of 8.3:1 shown in table 1, or an *n*-6 enriched high fat diet (*n*-6; D12266B: Research Diet Inc., New Brunswick, NJ), the formulation of which is shown in table 1, resulting in a *n*-6:*n*-3 fatty acid ratio of 31.3:1 (table 1).

After birth, three groups of pups each maintained at 6 pups/litter, were created by subdividing the high fat *n*-6 enriched diet fed pregnant dams into either **1**) continuing to receive a high fat *n*-6 enriched diet (*n*-6) or **2**) replaced with a high fat *n*-3 enriched diet (*n*-3; D09101401: Research diets Inc., New Brunswick, NJ) with 10% menhaden oil (table 1), resulting in a *n*-6:*n*-3 fatty acid ratio of 0.9:1 (table 1), while **3**) the CD group continued to receive the regular chow diet, in all groups through the period of lactation (PN1-PN21). We opted to study these ratios of *n*-6:*n*-3 fatty acids (*n*-6 being ~4 times greater and *n*-3 being ~8 times lower than that found in the standard regular chow diet, which in turn was two to four times that recommended for humans) so we could determine the presence of phenotypic differences in the offspring between the groups both in early suckling and subsequent adult life. After weaning at 22d, males and females in all three groups were placed on a regular chow diet until the adult age of 360d (12 months).

2.2 Morphometric Measurements

Body weights, nose-rump, nose-tail lengths and organ weights were measured longitudinally through the life course of male and female offspring beginning from PN2 until 360d of age. At 360d of age, following euthanasia with intraperitoneal administration of phenobarbital (100 mg/kg) using a 27G 1/2 needle [34,35], organ weights were assessed.

2.3 Food and Water Intake

At adult ages of 30d, 90d, 180d and 360d of age, food intake was longitudinally assessed over a 24-hour period at each age by calculating the difference in the measured weight of food that was placed in the food container before and remaining after the 24-hour period (g-food/g-body weight). Similarly, water intake was also assessed over a 24-hour period by calculating the difference in the volume of water placed in the water container before and remaining after the 24-hour period (g-water/g-body weight). Evaporative losses were assessed by placing a water and food container attached to an empty cage with no animal in the same room, and these losses subtracted from the final amounts of food and water intake over a 24-hour period. All these measurements were performed first thing early in the morning soon after the maximal nocturnal ingestive phase as previously described [35,36].

2.4 Analysis of Brain Samples

2.4.1 Hypothalamic Neuropeptide mRNA Analysis—Hypothalamus from whole brain was removed under dissecting microscope (Olympus SZ40, Center Valley, PA) using established anatomical landmarks [37]. Total RNA from hypothalamus was extracted using RNAeasy lipid tissue kit (Qiagen, Valencia, CA). First strand cDNA was synthesized from 1 µg of DNase treated total RNA using Superscript II reverse transcriptase (Invitrogen Life Technologies, Carlsbad, CA), as previously described [35]. Quantitative real time PCR was performed as previously described [22,35,36]. Primers and Taqman probes for detection of specific genes in hypothalamus were designed using Primer Express Software (Applied Biosystems, Foster, CA) and are listed [36]. These designed forward and reverse primers generate corresponding DNA fragments after amplification. Taqman probes were synthesized and labeled with fluorescent dye, 6-carboxyfluorescein (FAM) on the 5'-end and N,N,N',N'-tetramethyl-6-carboxyrhodamine (TAMRA) on the 3'-end (Applied Biosystems, Foster CA). Taqman PCR was carried out using a StepOnePlus™ real-time PCR system (Applied Biosystems, Foster, CA). Real time PCR quantification was then performed using Taqman glyceraldehyde-3-phosphate dehydrogenase (GAPDH) or eukaryotic 18S rRNA (Applied Biosystems, Foster, CA) as internal controls. PCR amplifications were performed in triplicates. The amplification cycles consisted of 12 min at 95°C (hot start), followed by 40 cycles at 95°C for 30 sec (denaturation), 56°C for long isoform of the leptin receptor (ObRb) and melanocortin isoform 3 receptor (MC3-R); 58°C for cocaine and amphetamine regulated transcript (CART), neuropeptide Y (NPY) and melanocortin isoform 4 receptor (MC4-R); 60°C for AgRP and proopiomelanocortin (POMC) over 30 sec (annealing), and 72°C for 30 sec (extension) [35,36], using reagents from Applied Biosystems (Foster, CA). Relative quantification of PCR products were based on value differences between the target and GAPDH or 18S rRNA control using the comparative CT method, as previously described [22,35,36].

2.4.2 Brain Protein Analysis—Whole brains except hypothalamus were homogenized either in phosphate-buffered saline (PBS) containing protease inhibitors (20 µg/ml pepstatin A, 20 µg/ml leupeptin, 30 µg/ml aprotinin and 2 mM PMSF), 1% Nonidet P-40 and 5 mM EDTA or in cell lysis buffer (Cell Signaling Technology, Danvers, MA) as previously described [35,36]. Protein content was measured by the Bio-Rad Protein Assay (Bio-Rad, Hercules, CA). Homogenates (30 µg of protein) were subjected to sodium dodecyl sulfate polyacrylamide gel electrophoresis and the separated proteins transferred to nitrocellulose membrane filters (Bio-Rad, Hercules, CA). The blotted membranes were sequentially incubated in 3% bovine serum albumin or 5% nonfat dry milk and the primary antibody consisting of the rabbit anti postsynaptic density 95 (PSD95) and rabbit anti glucose transporter isoform 1 (Glut1, abcam, Cambridge, MA), rabbit anti Synaptophysin (SYP) (Millipore, Burlington, MA), rabbit anti glucose transporter isoform 3 (Glut3, a gift from Dr. Takata in Japan) and mouse anti-vinculin as an internal loading control (Sigma, St. Louis, MO). The proteins were visualized in Typhoon 9410 Phosphorimager (GE Healthcare Biosciences, Piscataway, NJ) by blotting with the enhanced chemiluminescence (ECL) plus detection kit (GE Healthcare BioSciences Corp., Piscataway, NJ) following horseradish peroxidase-labeled anti-rabbit IgG for anti PSD95, SYP, Glut1 and Glut3 or anti-mouse IgG for vinculin (GE Healthcare Biosciences Corp., Piscataway, NJ). Each protein was quantified by using Image Quant 5.2 software (GE Healthcare Biosciences, Piscataway, NJ), and normalized to vinculin.

2.5 Glucose (GTT) and Insulin (ITT) Tolerance Tests

360d old male and female offspring were lightly restrained in a plastic holder and allowed to acclimatize. Tail vein basal blood glucose was assessed after an overnight fast in the case of the GTT and non-fasting basal blood glucose in the case of the ITT [33]. The animals received a dose of glucose (1g/kg body weight) via the tail vein in the case of GTT or an intraperitoneal dose of insulin (0.75U/kg body weight) in the case of ITT [33]. Tail vein blood to assess glucose concentrations were drawn at various time points following either the glucose or the insulin administration to assess glucose tolerance and insulin sensitivity.

2.6. Analysis of Liver Samples

2.6.1. Liver Histology—360d old male animals were anesthetized with inhalational isoflurane. Liver tissues were taken and fixed in 4% paraformaldehyde in 0.1 M phosphate buffer, pH 7.4 and washed with PBS, infused in 20% sucrose in 0.1 M sodium phosphate buffer, pH 7.4, containing 0.02% sodium azide, and embedded in OCT compound as previously described [38]. Specimens were sectioned (10-µm thickness) with a cryostat Leica CM 1850 (Nussloch, Germany). Hematoxylin and Eosin stain and Oil red O stain were performed as previously described [39-41]. After nuclear stain with Hematoxylin (Vector Laboratories, Burlingame, USA), tissue samples were immersed briefly (30 sec) in 60% isopropanol and stained with Oil Red O (Sigma, St. Louis, Mo) for 15 min and briefly washed with 60% isopropanol followed by running tap water for 10 min and mounted. m thickness) washed with PBS and hydrated in distilled water. Separate sections were incubated in Picro Sirius Red solution (Abcam, Cambridge, MA) for 60 min following washing with PBS. The sections were next washed briefly with acetic acid solution, dehydrated through graded alcohol, and mounted. Images were visualized using a Nikon

Eclipse E-600 Microscope (Nikon, Melville, NY, USA) or a Leica DM1000 microscope equipped with the MC170 HD camera (Leica Microsystems; Heerbrugg, Switzerland), as previously described [35,42].

2.6.2. Hepatic Protein Analysis—As previously described [35,36], the blotted membranes from homogenates of liver (30 µg of protein) were sequentially incubated in the primary antibody consisting of the mouse anti-fatty acid synthetase (FAS, BD Biosciences, San Jose, CA), the rabbit anti-acetyl coenzyme-A carboxylase (ACC), 5 AMP-activated protein kinase (AMPK) and phosphorylated 5 AMP-activated protein kinase (pAMPK) (Cell Signaling Technology, Beverly, MA), and mouse anti-vinculin (internal loading control; Sigma, St. Louis, MO). The protein bands were visualized in Typhoon 9410 Phosphorimager (GE Healthcare Biosciences, Piscataway, NJ) by blotting with the enhanced chemiluminescence (ECL) plus detection kit (GE Healthcare BioSciences Corp., Piscataway, NJ) following horseradish peroxidase-labeled anti-rabbit IgG for anti ACC, AMPK, pAMPK, or anti-mouse IgG for anti-FAS and vinculin (GE Healthcare Biosciences Corp., Piscataway, NJ). Each protein band was quantified by using Image Quant 5.2 software (GE Healthcare Biosciences, Piscataway, NJ), and normalized to the vinculin protein bands.

2.6.3. Hepatic and Plasma miR-122 Analysis—Liver was collected and snap frozen in liquid nitrogen. Samples were stored at -80°C until the time of analysis. Tissue RNA was extracted using Zymo Direct-zol RNA miniprep kit (Irvine, CA) according to the manufacturer's instructions. miRNA specific cDNA was synthesized using TaqMan MicroRNA reverse transcription kit (Applied Biosystems, Foster City, CA). To analyze the relative expression of mature microRNA-122 (miR-122) in various samples, qPCR was performed using Taqman MicroRNA Assay kit specific for rno-miR-122 (Applied Biosystem assay, Foster City, CA). Expression of tissue miRNA was normalized using U6 RNA (Applied Biosystem assay, Foster City, CA) as an internal control as previously described [43].

For cell free miRNA, blood samples were collected and plasma was separated by centrifugation. Samples were stored at -80°C until further analysis. Plasma miRNA was isolated using miRNA easy Serum/Plasma kit from Qiagen according to the manufacturer's instructions (Hilden, Germany). miRNA specific cDNA synthesis and qPCR was done as previously described [43]. To analyze the concentration of cell free/secreted miR-122, synthetic Ce-miR-39 (Applied Biosystem assay, Foster City, CA) from *C. elegans* was used as a spike-in control during the isolation of miRNA from serum, as recommended by the manufacturer.

2.7. Statistical analyses

Sample size was determined to be $n=6/\text{group}$ by using the two-tailed unpaired t-test at 80% power to detect a difference (δ) of 1.4 - 2.0 with a significance level (α) of 0.05 (Graphpad Statmate version 2.0; San Diego, CA). All data was expressed as means \pm SEM. All experimental groups were compared following establishment of normality. Analysis of variance (ANOVA) was employed and F values determined. Once significance was established, inter-group differences were validated by the Fisher's paired least significance

difference test. Final *p* values were considered significant at < 0.05 . Statistical analyses were performed using StatView Version 5.0 (SAS institute Inc, Cary, NC) or GraphPad Prism version 7 (San Diego, CA) [35,44].

3. Results

3.1. Body Weights

Body weights for males (A) and females (B) are presented in table 2 for all three groups, CD, *n*-6 and *n*-3 spanning from PN2, PN21, 30d, 90d, 180d and 360d of age. For males and females, no inter-group differences were observed in body weights at PN2 and body weights for *n*-6 were comparable to CD at subsequent time points. However, at PN21, *n*-3 males and females weighed less than CD and *n*-6. This reduction in weight persisted until 30d of age. Thereafter, *n*-3 group was no different from the CD or *n*-6 groups at 90d, 180d and 360d (Table 2).

3.2. Morphometric measurements

As previously described, nose-rump, nose-tail lengths and organ weights at 360d of age for males and females are depicted in table 3 [33]. No differences were noted in either sex for length. When compared to CD, a significant reduction in brain weights was evident in both high fat groups in males, and in *n*-3 females only. When brain weights were expressed per average body weights (g-brain/g-body weight), brain weights of males in *n*-3 (brain:body weight ratio: $0.0026 \pm 2.49 \times 10^{-5}$) were significantly lower than those in *n*-6 ($0.0027 \pm 3.22 \times 10^{-5}$; $p=0.03$) with no significant difference when compared to CD ($0.0026 \pm 4.71 \times 10^{-5}$, $p=0.1$). Both *n*-3 ($0.0037 \pm 8.93 \times 10^{-5}$; $p=0.002$) and *n*-6 ($0.0039 \pm 5.97 \times 10^{-5}$; $p=0.02$) exposed female brain weights were significantly lower than that of CD ($0.0042 \pm 7.49 \times 10^{-5}$). While the male pancreas weighed less in *n*-6 versus the CD and *n*-3 groups, the female kidney weighed less in *n*-3 versus the *n*-6 group. There were no other weight differences observed in liver, brown adipose and white adipose tissues.

3.3. Food and Water Intake

Food and water intakes were examined longitudinally in males and females in the three experimental groups (Table 4 & 5). When compared to the CD group, an increase in food intake was seen in male *n*-3 and *n*-6 groups at 30d of age with no further change noted at later time points (Table 4). No differences were observed in females at the various time points. In contrast, at 30d, a significant reduction in water intake by the male *n*-3 group and the female *n*-6 group was evident when compared to the respective sex-matched CD group. In the male *n*-3 group, a reduction in water intake persisted at 90d of age when compared to CD and *n*-6 groups, while at 360d, this reduction was only evident when compared to the *n*-6 group (Table 5).

3.4. Glucose and Insulin Tolerance including Serum Profiles

At 360d of age, glucose tolerance and insulin sensitivity were assessed by glucose and insulin tolerance tests, respectively. In males, an enhancement of glucose tolerance with increased insulin sensitivity was evident in the *n*-3 group versus CD, while *n*-6 was no different from the CD group (Figure 2A, C). In contrast, no inter-group differences were

observed in glucose tolerance and insulin sensitivity in the females (Figure 2B, D). In 360d males, while total cholesterol decreased in the *n*-6 group when compared to CD, free fatty acids decreased in the *n*-3 group when compared to *n*-6 group (Table 6).

3.5. Hypothalamic Neuropeptides and Proteins in Brain

These sex-specific phenotypic observations noted mainly in males led to our subsequent brain and liver specific studies restricted to the male offspring alone. First, we investigated the hypothalamic expression of key neuropeptides and receptors that mediate energy balance in the male groups at two distinct time points, PN2 and 360d (Figure 3A). At PN2, higher expression of AgRP, ObRb and MC4-R was observed in the *n*-6 male group versus CD, with no change in NPY, MC3-R, POMC and CART. At 360d of age, the male offspring demonstrated a persistence in higher AgRP expression particularly in the *n*-3 group versus CD, with no other change in any of the other neuropeptides in both high fat male groups (Figure 3B). Since we noted a reduction in brain weights in both high fat groups in the 360d male offspring, we assessed certain synaptic markers along with glucose transporters in the adult male brain cortical regions. We noted no differences in PSD95 (post-synaptic marker) and SYP (axonal marker) expression. Further, while the blood-brain barrier and glial-cell specific glucose transporter isoform (Glut1) was not affected, there appeared to be a tendency towards lower values in the neuronal/synaptic Glut3 concentrations in the *n*-3 group when compared to CD and *n*-6 groups (Figure 3C).

3.5. Histology, Metabolic Proteins and miR-122 in Livers

Second and next, we examined hepatic histology (Figure 4) and proteins (FAS, ACC and AMPK) (Figure 5), and hepatic and plasma miR-122 (Figure 6) in the CD, *n*-6, *n*-3 male groups. We observed a fatty liver in the *n*-6 and *n*-3 groups. However, the amount of steatosis detected visually in the *n*-3 group was notably less than that seen in the *n*-6 group (Figure 4). In addition, fibrosis assessed as collagen (type I fibers) staining was noted within the liver parenchyma in only small collections more so in the *n*-6 group versus the *n*-3 group when compared to the CD group (Figure 4). Our exploration of the 360d hepatic fatty acid synthesizing enzymes revealed no differences in FAS or ACC proteins in either high fat group compared to the CD group (Figure 5A). In contrast, while no differences in total AMPK were seen, an increase in phosphorylated (p) AMPK was noted in both the *n*-6 and *n*-3 groups when compared to CD (Figure 5B). We next evaluated the impact on a microRNA that is predominantly expressed by hepatocytes that also targets pAMPK [45], namely *miR-122*, and observed lower expression in hepatic *miR-122* in *n*-6 and *n*-3 groups versus the CD group (Figure 6A). In contrast, *miR-122* concentrations were significantly higher in the plasma of the *n*-6 group when compared to CD (Figure 6B). However, in the *n*-3 group, concentrations in-between that seen in the *n*-6 and CD were observed (Figure 6B).

4. Discussion:

Our results reveal that *n*-6 diet during pregnancy beginning after conception and during lactation, and *n*-3 diet during lactation alone affect growth, brain size, metabolism, and the state of the liver in a sex-specific manner. Contrary to our hypothesis and expectation, both

male and female offspring exposed pre- and postnatally to *n*-6 diet during mother's pregnancy and lactation period are not overweight and are in fact (much more in males) glucose tolerant and insulin sensitive, thereby proving the null hypothesis in this case. However, despite being lean, these male rats have liver disease akin to non-alcoholic fatty liver disease. On the other hand, *n*-6 during pregnancy followed by *n*-3 during lactation lowered the offspring's body weight and conferred even further enhanced glucose tolerance and insulin sensitivity in males with some protection against later development of hepatic steatosis. Despite this relative protection, male *n*-6 and male and female *n*-3 had smaller brains compared to CD at 360d (12 months) of age.

We opted to introduce a high fat diet after conception, and not before conception, so that we could isolate the effect of maternal diet from maternal increase in pre-conception body mass and its associated effects on the offspring. Various investigations including our own in mice [46] have introduced a high fat diet prior to conception and continued this diet during pregnancy and lactation. This diet produces an overweight mother who gives birth to offspring who subsequently become obese and glucose intolerant. In contrast, our present study specifically questioned whether these metabolic changes were due to maternal high fat diet alone in the absence of maternal obesity. Further, we ensured a moderately high fat diet consisting of ~32 kcal% rather than the traditional ~60 kcal%, during pregnancy and lactation alone.

Prenatal exposure to *n*-6 enriched diet stimulated the male hypothalamic AgRP expression noted a day after birth. AgRP is orexigenic and increases the intake of fat enriched foods [26,47] as opposed to NPY which is also orexigenic, but targets carbohydrate enriched foods [26,48]. Along with the increase in AgRP, the expression of its associated receptor MC4-R (melanocortin isoform 4 receptor) also increased. Reduction in MC4-R due to mutations is associated with obesity [49,50], while an increase may signify a loss of weight. Similarly, mutations of *Obrb* (long form of the leptin receptor) cause mild obesity [51,52], while the increase observed in our study would mediate a lean phenotype. These hypothalamic changes that were examined only in the male were associated with a transient reduction in body weight, despite an increase in food intake in the male *n*-6 and *n*-3 groups. While the AgRP increase lasted until 360d in males, other hypothalamic changes reverted to normal without later body weight changes.

The high fat groups revealed glucose tolerance and insulin sensitivity, with the *n*-3 group revealing exaggerated glucose tolerance with heightened insulin sensitivity, particularly in the adult male offspring. This was accompanied by a decrease in free fatty acids in the male *n*-3 group when compared to CD. Additional confirmation was forthcoming when no changes in the liver fatty acid synthesizing enzymes were seen in males. An increase in hepatic pAMPK, a heterotrimeric serine-threonine kinase, further supports this lean phenotype, since pAMPK is a master regulator of cellular energy homeostasis by promoting catabolic pathways towards producing adenosine triphosphate. Thus, pAMPK promotes glycolysis, inhibits glycogen synthesis, enhances glucose transport and fatty acid oxidation. This change in hepatic pAMPK is in keeping with a high energy producing phenotype perhaps intended to match enhanced energy expenditure (not measured in our study).

Given these changes, we sought further confirmation, by examining the liver thoroughly. Other and our groups have demonstrated that *miR-122* is a liver-specific non-coding RNA post-translationally regulating gene expression [53]. Its function was noted to be in regulation of fatty acid and cholesterol metabolism, in replication of hepatitis C virus, as a tumor suppressor gene in hepatocellular carcinoma, and in regulation of iron homeostasis by modulating activators of the hormone hepcidin [54]. However, others have demonstrated that *miR-122* targets the 3'-UTR of the cation amino acid transporter isoform 1 (CAT-1), which provides bidirectional transport for the essential amino acids lysine and arginine. Recovery occurs when nutritional stressors disappear and human antigen R, an RNA-binding protein arrives in the cytoplasm from the nucleus and rescues CAT-1 from reductive regulation by *miR-122* [55]. In addition, pAMPK is specifically targeted in hepatocytes by *miR-122* suppressing its expression [45]. Given this background information, we considered *miR-122* to be an important indicator of hepatic cellular health. We found that the *n-6* and *n-3* groups had lower hepatic *miR-122* with simultaneous higher plasma (secreted by the hepatocytes) *miR-122* concentrations. Although the biological significance of these changes is not entirely uncovered by our present study suggesting a limitation, prior studies provide some clues. In non-alcoholic fatty liver disease, murine hepatic *miR-122* was down-regulated [56,57], while serum *miR-122* was up-regulated [58-60]. In other studies, upregulation of human serum *miR-122* predicted liver fibrosis [61]. In our present study, a high fat diet decreased liver specific *miR-122* concentrations and led to fatty liver changes with *n-6* being more pronounced than that seen with *n-3*. In addition, some amount of fibrosis was noted in the *n-6* group scattered within the parenchyma, that was notably absent in the *n-3* group. These histological changes are despite the offspring being lean, glucose tolerant, and insulin sensitive. Changes in *miR-122* may also signify changes in cholesterol biosynthesis and compromise to essential amino acid transport. In an independent study, we have shown that a western diet administered to pregnant mice led to an amino acid deficiency post-partum [62]. We speculate that a high fat diet exposure may have compromised the availability of essential amino acids causing a *miR-122* mediated reduction in amino acid transport as well. These concepts need future investigation, being a limitation of our present study. We did however observe higher hepatic pAMPK concentrations in the *n-6* group in the presence of lower hepatic *miR-122* expression, supporting prior evidence [45], that a reduction in *miR-122* may release pAMPK from its inhibitory regulation. In this study, introduction of an *n-3* diet postnatally led to a partial reversal of the adult phenotypic changes encountered in response to prenatal *n-6* dietary exposure, particularly in the male offspring.

Our findings are novel and attest to the fact that a high fat diet only during pregnancy and lactation can be protective of the offspring with respect to the metabolic phenotype, maintaining body weight and glucose-insulin homeostasis. Our present observation is in keeping with a previous report where the process of insulin resistance was averted in the male rat adult offspring under *n-3* supplementation during maternal pregnancy [18]. This lean glucose tolerant and insulin sensitive phenotype was noted by us despite a propensity towards higher hypothalamic AgRP induced food intake during early adult years. Thus, this higher food intake is more suggestive of an attempt at matching and supporting the presumed enhanced energy expenditure (indirect evidence being the elevated pAMPK). Further, our present study's high fat diet exposure during gestation and lactation, affording

protection to the offspring may also be related to reduced carbohydrate transfer from mother to fetus during a critical window of fetal (via placenta)/postnatal (via milk) development.

However, while both *n*-6 and *n*-3 diets during this window proved to be protective on the metabolic phenotype of the offspring, the detection of fatty liver and adverse brain findings were concerning. While our present study was not meant to fully examine brain development, the reduction in brain weight, while other organs for the most part remained unchanged, is concerning. Previous reports describing abnormal neuronal development and sensory function in offspring born to pregnant dams consuming *n*-3 further supports this concern [16,63].

While specific brain markers were not perturbed in our study, the subtle sign of a reduction of neuronal Glut3 concentrations, known for fueling neurotransmission [64-66], is concerning and requires future investigation. If this reduction in brain weight serves as a surrogate for inadequate neurogenesis, gliosis and/or myelination, there should be concern regarding the neurobehavior of these offspring. It is known that a pre-pregnancy and gestational high fat diet causes anxiolysis and other adverse changes in neurobehavior expressed by the offspring [67,68]. Further, investigations during embryonic day 21 revealed defects in peri-ventricular stem cell migration in the fetal brain [26] due to a high fat dietary exposure, supporting our current findings of aberrant brain growth in the offspring exposed prenatally and postnatally to a high fat diet. Thus, prior to recommending a high fat diet during gestation that is overlaid by postnatal *n*-6 or even *n*-3 (known to enhance infant visual and neural function) [69], one must weigh the potential metabolic (glucose tolerance and insulin sensitivity) benefits with the potential neurodevelopmental risks. Caution is recommended in ensuring that brain development and neurobehavior are optimized so that all offspring achieve their full potential.

We conclude that a high fat diet only during gestation and lactation (*n*-6 or *n*-3) has the propensity of affording metabolic protection to the adult offspring. This protection, results in lean, glucose tolerant and insulin sensitive male and female adult offspring. At the same time, the adult male offspring develops fatty liver. Non-alcoholic fatty liver disease is the most common chronic pediatric liver disorder [70], and predicted to become an indication for liver transplant, with increasing frequency. This same diet during gestation and lactation also detrimentally affected the brain weight of the adult offspring. It remains unclear currently how this change in brain weight impacts the neurological status, warranting future studies. However, like non-alcoholic fatty liver disease, the incidence of childhood neurodevelopmental disorders, including autism spectrum disorders, continues to grow [63]. Large epidemiological studies have made an association between maternal obesity and non-alcoholic fatty liver disease along with adverse neurological outcomes in the offspring [71]. In summary, we have shown that a high fat diet limited to *n*-6 in gestation and lactation, and *n*-3 in lactation only, produces a glucose tolerant and insulin sensitive offspring, but causes fatty liver and reduces brain weight.

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Abbreviations

ACC	acetyl coenzyme-A carboxylase
AgRP	agouti-related peptide
AMPK	5 AMP-activated protein kinase
ANOVA	analysis of variance
CART	cocaine and amphetamine regulated transcript
CAT-1	cation amino acid transporter isoform 1
CD	control regular chow diet
FAS	fatty acid synthetase
G	gestation
GAPDH	glyceraldehyde-3-phosphate dehydrogenase
Glut1	glucose transporter isoform 1
Glut3	glucose transporter isoform 3
GTT	glucose tolerance test
ITT	insulin tolerance test
MC3-R	melanocortin isoform 3 receptor
MC4-R	melanocortin isoform 4 receptor
miR-122	microRNA-122
<i>n</i>-3	dietary <i>n</i> -3 polyunsaturated fatty acids
<i>n</i>-6	dietary <i>n</i> -6 polyunsaturated fatty acids
NPY	neuropeptide Y
ObRb	long isoform of the leptin receptor
PBS	phosphate-buffered saline
POMC	proopiomelanocortin
PN	postnatal day
PSD95	postsynaptic density 95
pAMPK	phosphorylated 5 AMP-activated protein kinase
SYP	synaptophysin

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Highlights:

Gestational *n*-6 diet results in a glucose tolerant and insulin sensitive adult offspring.

Postnatal *n*-3 superimposition on gestational *n*-6 diet enhances insulin sensitivity in offspring.

Gestational *n*-6 diet enhances hypothalamic AgRP related food intake in the offspring.

Gestational *n*-6 diet results in adult onset fatty liver disease in male offspring.

Gestational *n*-6 with or without postnatal *n*-3 diet reduces brain weight in offspring.

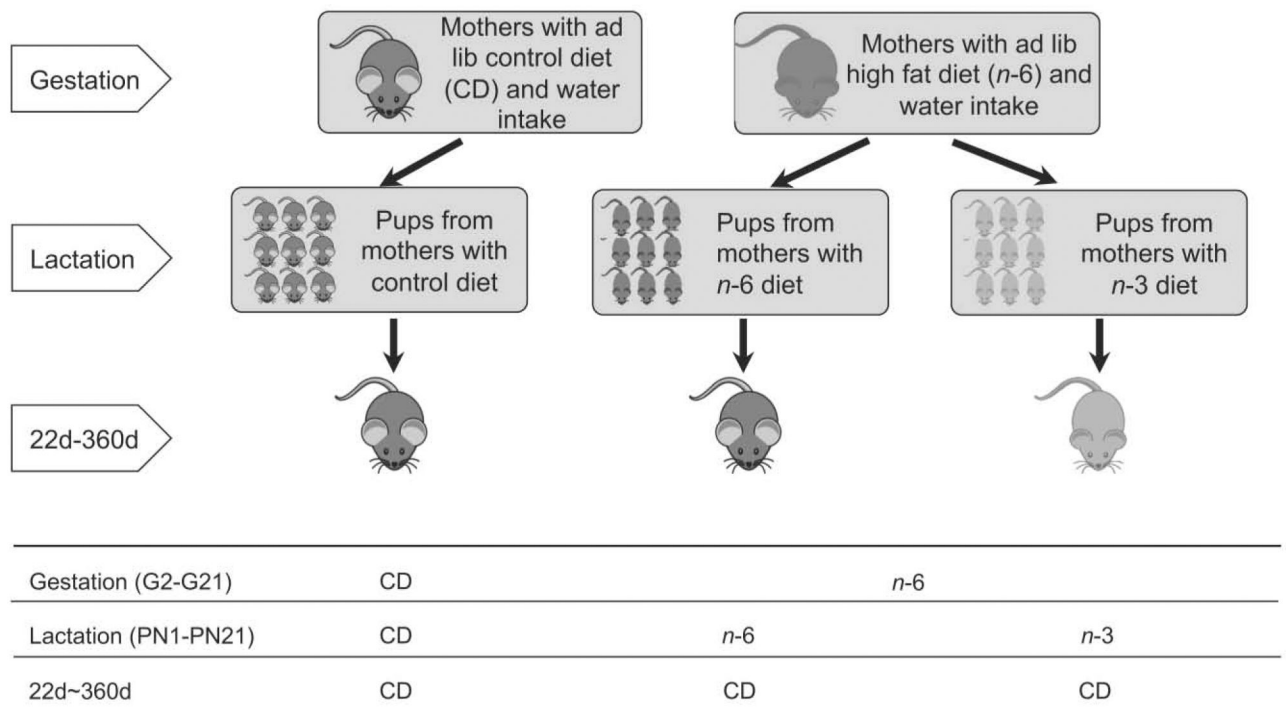


Figure 1: Scheme demonstrating the experimental design.

Three experimental groups were created: Two groups were created with ad lib access to either control regular chow diet (CD) or dietary *n*-6 polyunsaturated fatty acids (*n*-6) during gestation (G2-G21). The CD group continued on the same diet while the *n*-6 exposed group was sub-divided at birth into receiving *n*-6 or *n*-3 during lactation (PN1-21). After weaning, all three groups were given CD until day 360 of age.

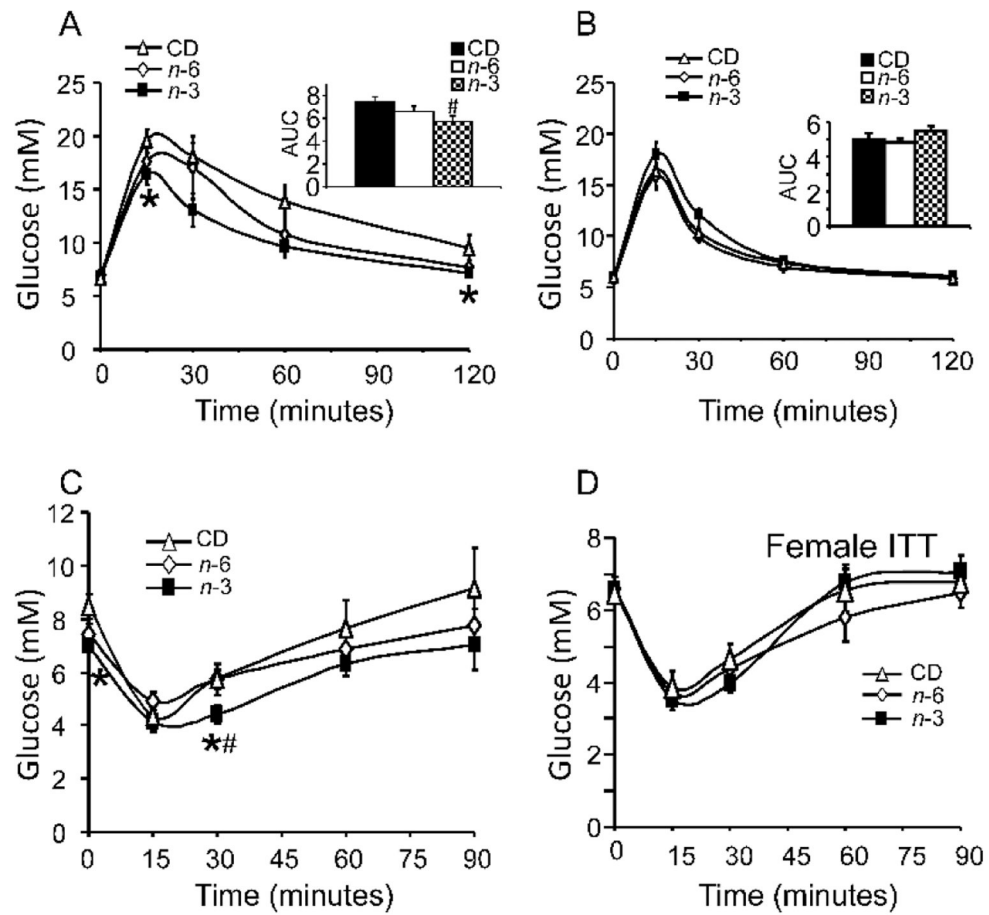


Figure 2: Glucose tolerance tests (A & B), and insulin tolerance tests (C & D) in CD, *n-3* and *n-6* groups.

A & B: GTTs in 360d male (A) and female (B) adult offspring in CD, *n-3* and *n-6*. N= 6~7 per each group. * $p < 0.05$ compared to CD. Inset: respective AUC calculations. # $p < 0.01$ compared to CD by one-way ANOVA and Fisher's PLSD test. C & D: ITTs in 360d male (C) and female adult offspring (D) in CD, *n-3* and *n-6*. N= 6 per each group. * $p < 0.04$ compared to CD. # $p < 0.04$ compared to *n-6* by one-way ANOVA and Fisher's PLSD test.

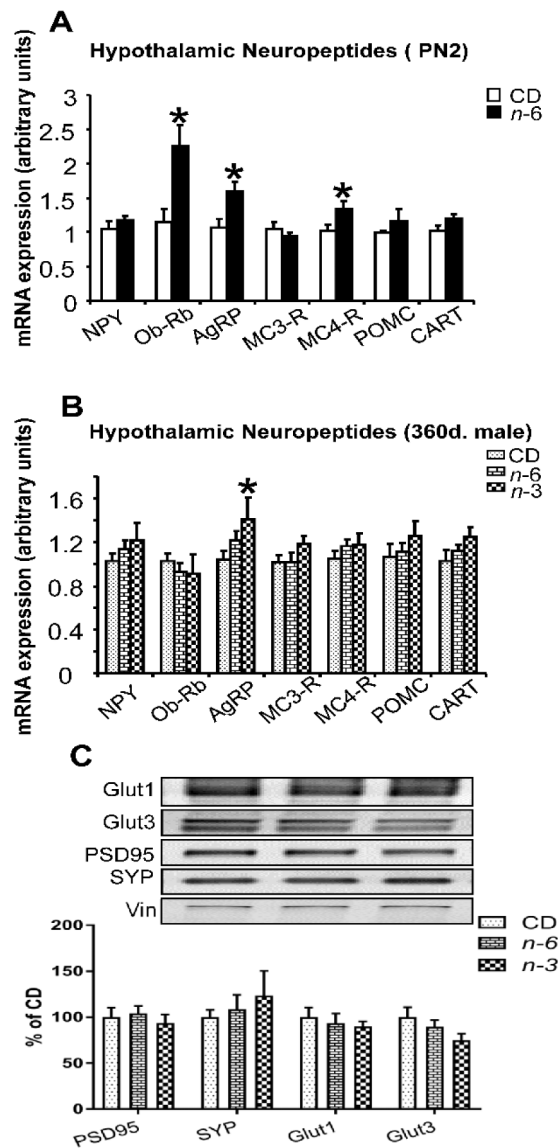


Figure 3: Hypothalamic neuropeptides (A & B) and proteins in brain (C).

A & B; Real-time quantitative RT-PCR analysis of male hypothalamic NPY, ObRb, AgRP, MC3-R, MC4-R, POMC and CART expression. CD and *n*-6 pups (male) at PN2 (A, *n*=6 per each group) or 360d (B, *n*=5~8 per each group). **p*<0.05 compared to CD. C; PSD95, SYP, Glut1 and Glut3 proteins assessed by Western blot analysis. There is no significant difference in the groups. *N*=5-8, each group. Unpaired t-test (A) or one-way ANOVA and Fisher's PLSD test (B & C).

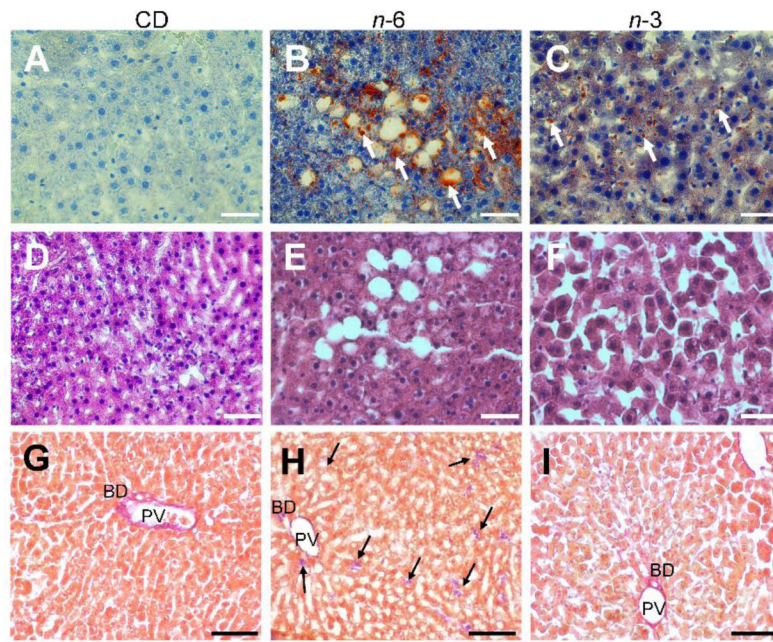


Figure 4: Liver Histology.

In 360d male offspring, morphology by Hematoxylin and Eosin stain (A-C), fat deposition by Oil red stain O (D-F) and collagen fiber (fibrosis) by Picro Sirius Red Stain (G-I) in liver sections of CD (A,D & G), *n*-6 (B,E & H) and *n*-3 (C,F & I) groups. *n*-6 group shows prominent fat deposits compared to *n*-3. In addition, *n*-6 group demonstrates inter-cellular parenchymal collagen deposits (arrows) more than that seen in the *n*-3 group. Collagen staining is present in the portal vein (PV) and bile ducts (BD) in all three groups. Scale bars, A-F: 50 μ m; G-I: 100 μ m

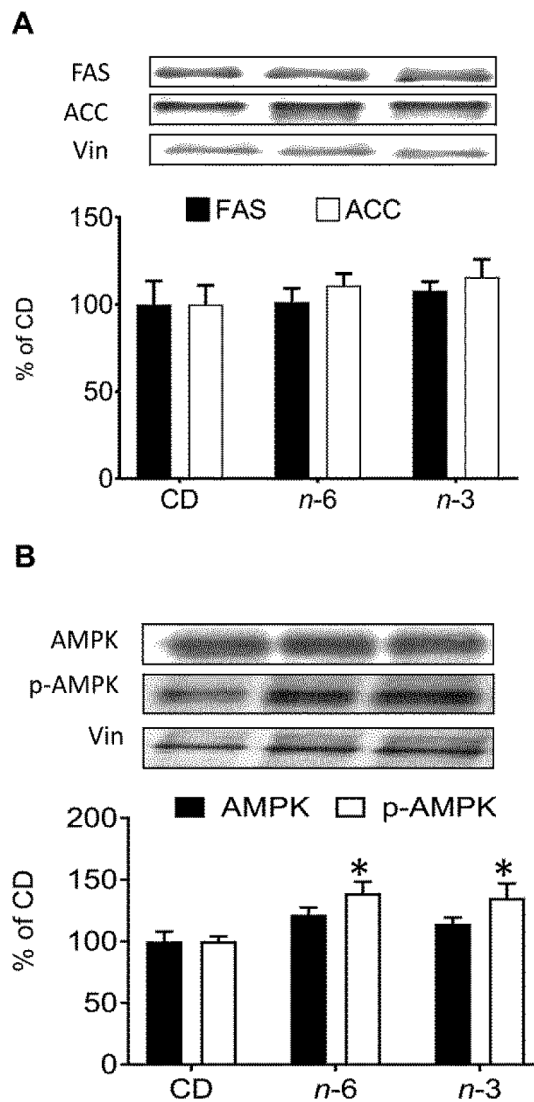


Figure 5: Hepatic fatty acid synthesizing metabolic proteins.

In male CD, *n-3* and *n-6* offspring at 360d, Top: Representative Western blots. Bottom: Densitometric quantification of corresponding protein concentrations. A, Fatty acid synthase (FAS) and Acetyl CoA carboxylase (ACC); B, AMPK (5' AMP-activated protein kinase) and phosphorylated AMPK (pAMPK). * $p < 0.05$ compared with CD, one-way ANOVA and Fisher's PLSD test. N=5-8, each group.

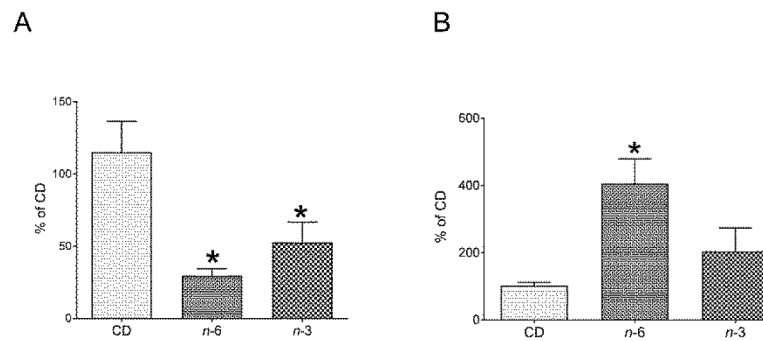


Figure 6: Hepatic and plasma *miR-122*.

In 360d male offspring, relative concentrations of hepatic (A) and plasma (B) *miR-122* concentrations determined by RT-qPCR. Hepatic *miR-122* concentrations were significantly lower in *n-6* (* $p=0.003$) and *n-3* (* $p=0.03$) when compared to CD group (A), as analyzed by one-way ANOVA and Fisher's PLSD test. Plasma concentrations of *miR-122* was significantly higher in *n-6* (* $p=0.04$) but not in *n-3* exposed male rat offspring when compared to the CD group, as analyzed by one-way ANOVA and Fisher's PLSD test. In the *n-3* group, a tendency towards higher values compared to the CD group, and a tendency towards lower values when compared to the *n-6* group is observed in plasma *miR-122* concentrations (B). $N=5-6$ in each group.

Table 1:

Fatty acid and ingredient composition employed in the study

Fatty acid (wt%)	Dietary treatments ^{a,b,c,d}		
	CD	n-6	n-3
4:0	0	0.9	0.3
6:0	0	0.5	0.2
8:0	0	0.3	0.1
10:0	0	0.7	0.3
12:0	0	0.8	0.3
14:0	0	2.8	5.9
14:1n-9	0	0.4	0.1
15:0	0	0	0.3
16:0	17.3	15.4	16.1
16:1n-9	0	0.6	7.0
16:2n-4	0	0	1.1
16:3n-9	0	0	1.1
16:4n-4	0	0	1.1
17:0	0	0	0.3
18:0	3.8	4.8	3.6
18:1n-9	25	25.6	16.3
18:2n-6	48.1	45.4	18.5
18:3n-3	5.8	1.5	1.6
18:4n-3	0	0	2.1
20:0	0	0.3	0.3
20:1	0	0	1.1
20:2	0	0	0.1
20:3n-6	0	0	0.3
20:4n-6	0	0	1.5
20:5n-3	0	0	9.9
21:5n-3	0	0	0.5
22:0	0	0	0.1

Fatty acid (wt%)	Dietary treatments <i>a,b,c,d</i>		
	CD	n-6	n-3
22:1	0	0	0.2
22:4n-6	0	0	0.1
22:5n-3	0	0	1.9
22:6n-3	0	0	7.1
24	0	0	0.4
24:1	0	0	0.1
SAT	21.1	26.4	27.7
MONO	25.0	26.7	24.9
PUFA	53.9	46.9	47.0
n-6 PUFA	48.1	45.4	20.4
n-3 PUFA	5.8	1.5	23.2
n-6:n-3	8.3	31.3	0.9

^aThe semipurified basal diet contains the following (g/kg): for n-6 and n-3, casein, 182; corn starch, 206.2; sucrose, 278.1; cellulose, 28.8; choline bitartrate, 1.9; salt mix, 38.4; vitamin mix, 10.5 (based on Research diet). For CD, corn starch, 193.52; oats, 100; wheat, 355; choline chloride, 60%; 1.48 NIH-31 vitamin mix, 3.5; NIH-31 mineral mix, 1.5 (based on Teklad diets).

^bMineral mix provides (mg/kg diet): for n-6 and n-3, CaHPO₄, 19200; MgO, 921.6; C₆H₅K₃O₇·H₂O, 8448; K₂SO₄, 1996.8; NaCl, 2841.6; CrK(SO₄)₂·12H₂O, 21.12; CU₂CO₃, 11.52; KIO₃, 0.38; C₆H₅FeO₇, 230.4; MnCO₃, 134.4; Na₂SeO₃, 0.38; ZnCO₃, 61.4. For CD, Fe, 174; Cu, 13.1; Zn, 205; Se, 0.4; Mn, 139; I, 1.4.

^cVitamin mix provides (mg/kg diet): for n-6 and n-3 diets, thiamine HCl, 7.7; niacin, 33; calcium pantothenate, 17.6; folic acid, 2.2; biotin, 0.22; cyanocobalamin (B12), 11; vitamin A palmitate (500,000 IU/g), 8.8; vitamin E acetate (500 IU/g), 110; vitamin D₃ (100,000 IU/g), 1.1. For CD; thiamine HCl, 66; riboflavin, 7; pyridoxine HCl, 9; calcium pantothenate, 36; folic acid, 70; biotin, 0.2; cyanocobalamin (B12), 1.54; vitamin A (25,560IU/Kg), vitamin E (a tocopherol acetate), 31, vitamin D₃ (cholecalciferol, 3620 IU/Kg).

^dDietary fat treatments include butter fat and corn oil for n-6 and n-3, soy oil for control diet (CD). Only n-3 diet includes 10% Menhaden oil. Total fat content is 155.6 g/kg for n-6 and n-3 diet, 62g/kg for CD.

^eSAT, total saturated fatty acids; MONO, total monounsaturated fatty acids; PUFA, total polyunsaturated fatty acids.

0 = not detected

Table 2:

Body Weight (g)

	PN 2			PN21			30d			90D			180D			360D		
	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3
Male	7.02 ± 0.1 4	7.25 ± 0.1 4	7.25 ± 0.1 4	58.58 ± 1.59	61.33 ± 0.76	54.00 ± 1.39 *, **	103.70 ± 2.17	109.25 ± 2.54	100.13 ± 3.70 **	572.50 ± 11.76	557.42 ± 17.46	570.75 ± 25.33	714.92 ± 17.42	696.33 ± 27.17	715.00 ± 37.58	867.50 ± 27.92	800.25 ± 37.04	815.86 ± 46.12
Female	6.57 ± 0.1 6	6.68 ± 0.1 1	6.68 ± 0.1 1	56.83 ± 2.35	58.75 ± 0.82	49.88 ± 0.64 *, **	95.40 ± 3.48	96.92 ± 2.46	85.00 ± 0.76 *, **	311.25 ± 12.38	310.92 ± 12.66	303.50 ± 10.28	377.58 ± 16.60	377.33 ± 23.86	377.50 ± 22.80	481.13 ± 25.68	498.42 ± 43.75	503.25 ± 31.60

Body weights of male (A) and female (B) offspring from three experimental groups: CD, n-6, and n-3 from PN2 to 360d are shown. In both males and females, data are shown as means ± SE (n=8-12)

* p<0.02 vs CD

** p<0.03 vs n-6 by one-way ANOVA and Fisher's PLSD test.

Table 3:

Organ weights and nose-tail or nose-rump length

	Male			Female		
	CD	n-6	n-3	CD	n-6	n-3
Nose-rump(cm)	28.56 ± 0.29	27.94 ± 0.27	28.00 ± 0.45	Nose-rump(cm)	24.00 ± 0.31	23.75 ± 0.17
Nose-tail(cm)	51.88 ± 0.58	51.88 ± 0.31	52.10 ± 0.95	Nose-tail(cm)	44.86 ± 0.20	44.42 ± 0.63
Brain(g)	2.30 ± 0.04	2.19 ± 0.03 *	2.12 ± 0.02 *	Brain(g)	1.93 ± 0.03	1.89 ± 0.05 *
Liver(g)	22.45 ± 1.04	22.36 ± 1.47	21.75 ± 2.18	Liver(g)	14.14 ± 1.87	13.18 ± 1.47
Kidney(g)	4.75 ± 0.25	4.46 ± 0.19	4.26 ± 0.38	Kidney(g)	3.14 ± 0.32	2.42 ± 0.10 **
Pancreas(g)	0.96 ± 0.05	1.54 ± 0.13 *	0.97 ± 0.12 **	Pancreas(g)	0.67 ± 0.08	0.63 ± 0.03
BAT(g)	0.75 ± 0.09	0.69 ± 0.05	0.68 ± 0.04	BAT(g)	0.56 ± 0.08	0.59 ± 0.05
WAT(g)	40.17 ± 2.38	33.54 ± 4.00	30.66 ± 2.30	WAT(g)	43.78 ± 7.32	37.08 ± 1.47

Data are shown as means ± SE (n=5-8)

* p<0.05 vs CD

** p<0.05 vs n-6 by one-way ANOVA and Fisher's PLSD test. BAT, brown adipose tissue; WAT, white adipose tissue.

Table 4:

Food Intake

	30d			90D			180D			360D		
	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3
Male	0.121 ± 0.009	0.149 ± 0.003*	0.148 ± 0.003*	0.049 ± 0.002	0.047 ± 0.001	0.045 ± 0.002	0.039 ± 0.002	0.037 ± 0.001	0.036 ± 0.002	0.032 ± 0.002	0.037 ± 0.002	0.031 ± 0.002#
Female	0.122 ± 0.006	0.132 ± 0.005	0.124 ± 0.007	0.054 ± 0.004	0.064 ± 0.006	0.06 ± 0.005	0.041 ± 0.001	0.042 ± 0.004	0.04 ± 0.006	0.041 ± 0.003	0.033 ± 0.003	0.035 ± 0.006

Food intake (g-food/g-body weight) in male and female offspring was measured at 30d, 90d, 180d and 360d. Data are shown as means ± SE (n=8-12)

* p<0.01 vs CD

p<0.06 vs n-6 by one-way ANOVA and Fisher's PLSD test.

Table 5:

Water Intake

	30d			90D			180D			360D		
	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3	CD	n-6	n-3
Male	0.380 ± 0.033	0.318 ± 0.020	0.276 ± 0.018*	0.063 ± 0.002	0.063 ± 0.002	0.049 ± 0.003*,**	0.051 ± 0.008	0.044 ± 0.003	0.042 ± 0.003	0.027 ± 0.002	0.032 ± 0.002	0.025 ± 0.002**
Female	0.391 ± . 038	0.290 ± 0.018*	0.324 ± 0.016	0.086 ± 0.017	0.069 ± 0.006	0.066 ± 0.003	0.062 ± 0.006	0.059 ± 0.014	0.055 ± 0.005	0.042 ± 0.003	0.034 ± 0.004	0.040 ± 0.001

Water intake (g-water/g-body weight) in male and female offspring was measured at 30d, 90d, 180d and 360d. Data are shown as means ± SE (n=8-12)

* p<0.02 vs CD

** p<0.03 vs n=6 by one-way ANOVA and Fisher's PLSD test.

Table 6:

Serum Lipid Profile

	CD	<i>n</i> -6	<i>n</i> -3
TG mg/dl	213.25±30.47	196.25 ±29.29	173.6 ±14.95
Tot chol. mg/dl	129 ± 11.5	89.3 ±4.59*	114.8 ±14.56
HDL mg/dl	79.75 ± 8.23	62.5 ±3.86	79.8 ±11.62
UC mg/dl	31.75 ± 3.28	24.375 ±1.34	30.4 ±4.98
FFA mg/dl	11.9 ± 0.76	13 ±0.68	9.4 ±0.67**

Plasma concentrations of various metabolites from 360-day-old offspring exposed to Chow Diet (CD), *n*-6 enriched high fat diet alone or with *n*-3 enriched high fat diet during the postnatal period (PN1-PN21). Female serum lipid profile was not measured. Data are shown as means ± SE (*n*=5-8)

* $p < 0.05$ vs CD

** $p < 0.004$ vs *n*-6 by one-way ANOVA and Fisher's PLSD test. *N*=5-8 for each group.