Full title: High-fat diet-induced obesity increases susceptibility to endogenous and ethyl

carbamate-induced somatic mutagenesis

Short title: High-fat diet and mutagenesis

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Abstract

Obesity is associated with increased risk for certain cancers. We conducted a set of experiments designed to investigate both the inherent mutagenicity of high-fat diet (HFD) and the effect of HFD on sensitivity to ethyl carbamate, a known mutagen, in a mouse model of HFD-induced obesity. Short term HFD exposure was found to induce both hepatic Cyp2e1 activity and ethenoadenosine, a marker of lipid peroxidation DNA damage. After one year, greater body mass in females was associated with higher frequency of Pig-a mutant erythrocytes as well as hepatic precancerous transformation, suggesting greater risk of tumor formation. HFD-induced obese female mice exhibited increased sensitivity to ethyl carbamate-induced mutagenesis compared to control mice. Together, these findings indicate that HFD exposure can induce both endogenously generated changes related to xenobiotic metabolism, mutagenesis, and pre-carcinogenesis, and sensitivity to exogenous mutagens, in a sex-specific manner.

Introduction

Obesity and overweight are highly prevalent in the United States and are associated with a number of conditions, including certain cancers. Of all new cancer cases diagnosed in 2015, 55% of those diagnosed in women and 24% of those diagnosed in men can be linked to obesity, including cancers of the breast, uterus, colon and liver[1]. In fact, incidence of obesity related cancers rose 7% between 2005 and 2014, while non-obesity related cancer declined by 13%, and the fastest growing obesity-related cancer type during that period was liver cancer, which rose by 29%[1,2]. Specifically among

women, rates of obesity-related cancers rose rapidly between 2001 and 2018, while

cancers due to other risk factors declined[3].

There are many factors underlying the progression from obesity to carcinogenesis,

including environmental exposures such as diet and mutagenic toxicants. In the context

of obesity and cancer risk, it is useful to consider diet as a type of environmental exposure

that leads to high internal levels of triglycerides and adipose tissue-secreted hormones

and proteins. Because obesity is often a lifelong condition, with childhood and adolescent

obesity predictably persisting into adulthood, this exposure can be chronic[4].

It has been known in the medical community that patients with obesity process

certain drugs at different rates than non-obese patients, due to altered levels of

cytochrome P4502E1, a promiscuous enzyme responsible for phase I metabolism of a

range of substrates, often to intermediate products with enhanced toxicity[5,6]. Despite

this, very few studies of diet-induced obesity in mouse models have measured enzymatic

activity of hepatic Cyp2e1, the majority favoring measurement of changes at the gene

expression level, [7–9]. CYP2E1/Cyp2e1 catalyzes both the genotoxic peroxidation of

endogenous lipids and synthesis of several toxic intermediates from environmental

mutagens, including ethyl carbamate[10,11]. Industrially, ethyl carbamate was used as

an intermediate in textile, pesticide and cosmetic manufacturing and is produced in food

and beverage fermentation[12]. While the risk to the general public is low, those that

regularly consume stone fruit brandy or fermented soy products may have a lifetime tumor

risk of 0.01%, 100-times the risk that is generally considered acceptable[13–15]. Ethyl

carbamate is largely desaturated by CYP2E1/Cyp2e1 to vinyl carbamate, and oxidized

into vinyl carbamate epoxide[10,16]. Vinyl carbamate epoxide is highly reactive with DNA,

forming stable, mutagenic etheno-DNA adducts[17]. These adducts are also formed in

the presence of excess lipids due to lipid peroxidation[18].

We have previously shown that high-fat diet (HFD) exposure in male C57BL/6J

mice leads to higher frequency of spontaneous somatic mutations using the Pig-a gene

mutation assay[19]. Here we present the results from three experiments including both

male and female mice. We hypothesize that the physiological state of HFD-induced

obesity, and resultant excess lipid accumulation provides activity-inducing substrate for

Cyp2e1, both raising endogenous levels of lipid peroxidation and metabolism of

exogenous compounds to genotoxic intermediate metabolites. First, we designed a short-

term HFD exposure experiment to determine if mice reared on HFD (60% calories from

fat) have higher hepatic Cyp2e1 activity and retain more ethenoadenosine adducts. We

then designed a long-term HFD exposure cohort to determine the effect of HFD over time

in both males and females on frequency of spontaneous somatic mutations using the Pig-

a gene mutation assay and markers of hepatic precancerous transformation. Given the

convergent mechanism of toxicity of HFD and ethyl carbamate we designed a third, two-

factorial experiment whereby mice with long-term HFD exposure were challenged with

ethyl carbamate and assessed for Pig-a gene mutation frequency. Together, this set of

experiments explores both the endogenous and exogenous factors underlying the

pathway to carcinogenicity in the context of diet-induced obesity.

Methods

General animal care, sample processing, and analysis

Mice were reared at the Tulane University Downtown Vivarium on a 12 hour

light/dark cycle. Research staff supplied C57BL/6J mice with HFD containing 60%

calories from fat from Research Diets, Inc. (D12492) or with a low-fat diet (LFD) containing

10% calories from fat (D12450J), *ad libitum*. The LFD formulation serves as the most

appropriate control condition to the HFD formulation because it is source-matched and

contains equivalent sucrose content. Water and veterinary care were provided by

vivarium staff. Organs were harvested and flash frozen in liquid nitrogen, serum or plasma

was isolated from whole blood, and all samples were stored at -80°C until further analysis

unless otherwise indicated. Prism software (GraphPad; San Diego, CA, USA) was used

to conduct statistical analysis and to generate figures. Unless otherwise indicated, column

charts present arithmetic mean ±standard error, and regression charts present individual

data points and resulting regression curve with a truncated x-axis. Statistical significance

was determined using α=0.05.

Short-term HFD experiment

Husbandry and sample collection

This experiment was conducted under Tulane IACUC protocol 4374. Mice were

maintained on HFD or LFD conditions starting at 8 weeks old, and the experiment was

terminated when mice reached 22 weeks old. Each dietary treatment group consisted of

three males and three females. Samples were collected following CO_2 overdose and

exsanguination.

Biomarker assays

Serum triglycerides and leptin content were analyzed via the Triglyceride

Quantification Assay Kit (Abcam, ab65336) and the Mouse/Rat Leptin Quantikine ELISA

Kit (R&D Systems, MOB00). To assess pre-mutagenic changes to DNA that might occur

in HFD mice, ethenoadenosine, a modification to the adenosine nucleoside that results

from lipid peroxidation, was measured in liver tissue using the OxiSelect Aldehyde-

Induced DNA Damage Kit (Ethenoadenosine Quantitation) (Cell BioLabs, STA-820).

Cyp2e1 activity assay

Cyp2e1 activity was measured in liver tissue by first isolating the rough

endoplasmic reticulum (RER) of liver cells and then incubating the isolated microsomal

fraction with *p*-nitrophenol, a Cyp2e1 substrate with colorimetric properties. Functional

RER isolation was achieved with a modified version of the Ca^{2+} precipitation protocol

outlined for Endoplasmic Reticulum Isolation Kit (Sigma-Aldrich, ER0100). The protocol

was followed as written, with the exception of the final resuspension, in which 50 mM

potassium phosphate with EDTA-free protease inhibitor cocktail tablets (Roche,

05892791001) replaced the supplied Isotonic Extraction Buffer. The conversion of *p*-

nitrophenol to *p*-nitrocatechol was quantified by colorimetric assessment at OD 535 nm

via spectrophotometry.

Statistical analysis

Differences in mean body mass, serum leptin, serum triglycerides, liver

ethenoadenosine, and liver Cyp2e1 activity were compared using the independent two-

way Student's t test within males and females separately. Body mass was also correlated

with liver ethenoadenosine and hepatic Cyp2e1 activity using simple linear regression

analysis and the slope of each curve was tested for significant deviation from zero using

an F test with the null hypothesis that the overall slope is zero. Correlation coefficient (R^2)

values were also reported to allow interpretation of the predictive value of body mass on

each outcome of interest. Sample size analyses were conducted using G*Power software

(Heinrich-Heine-Universität Düsseldorf; Dusseldorf, NRW, Germany)[19].

Long-term HFD experiment

Husbandry and sample collection

This set of experiments was conducted under the approval of Tulane University

IACUC protocol 546. Mice were maintained on HFD or LFD conditions from 8 weeks old

until 52 weeks old. Each dietary group consisted of 12 males and 12 females. Blood was

taken at three time points, when mice were 17, 33 and 52 weeks of age. This was

accomplished through repeated submandibular venipuncture under isoflurane

anesthetization[21]. Mouse livers were excised from 52 week old mice following final

submandibular venipuncture. Under isoflurane anesthetization, mice were exsanguinated

via cardiac puncture and half of the left hepatic lobe was excised.

Erythrocyte Pig-a mutant frequency analysis

152 Each blood sample was immediately transferred to a K_2EDTA-coated microtainers (BD,

153 365974) and kept on ice. Samples were packed into Whirl-Pak® bags and shipped to

154 Litron Laboratories (Rochester, NY). Pig-a mutant frequency analysis was carried out by

155 Litron research staff following *In Vivo* MutaFlow® kit protocols[19,22,23].

Hepatocyte proliferation index and Ki67 content analysis

A MicroFlow^PLUS Rat Liver (Prototype) kit was supplied *gratis* by Litron Laboratories

to Tulane University. Following excision, left liver lobes were placed into a tube of ice cold

HBSS with EDTA and 1% DMSO. Samples were transferred to 4°C, and within 6 hours,

each preserved liver was transferred into fresh buffer with EDTA and DMSO and stored

overnight. The following day, one gram of each sample was rinsed twice in Liver Rinse

Solution and then in Erythrocyte Clearing Solution. The liver samples were then digested,

and individual cells were isolated in collagenase. The outer cell membranes were lysed,

leaving nuclei intact, and SYTO 13 nucleic acid dye and an anti-Ki67-eFluor660 antibody

were applied. A MACSQuant Analyzer10 flow cytometer was used to acquire cellular

phenotype data and analysis was conducted using FlowLogic™ software (Inivai

Technologies; Mentone, VIC, Australia). Hepatocyte proliferation index (HPI) was

calculated using the following formula:

$$HPI = \frac{j + 2k + 3l}{j + k + l}, \text{where } j = 2\text{n events}, k = 4\text{n events}, l = 8\text{n events}$$

170 Statistical analysis

Body mass and mutant frequency were analyzed using mixed effects analysis of

variance (ANOVA) for repeated measures with Bonferroni's post hoc multiple

comparisons test performed at each time point between LFD and HFD subjects. Simple

linear regression analysis of the relationship between body mass and Pig-a mutant

frequency was also conducted at each time point, and the slope p value and R^2 were

reported as described for the short-term HFD experiment. HPI and Ki67 content was

compared between LFD and HFD groups using the two-tailed independent Student's t

test. Males and females were analyzed separately.

180 Ethyl carbamate challenge experiment

Husbandry and sample collection

This experiment was conducted under approved Tulane University IACUC protocol 36-105. 48 males and 48 females were reared on HFD or LFD from 8 weeks old until they reached 42 weeks old, when the experiment was terminated. Nine weeks prior to termination, at 33 weeks, mice were divided into groups of 6, and exposed to 0, 100, 200, or 400 mg/kg/day of ethyl carbamate (CAS # 51-79-6; Sigma, U2500, Lot # WXBC3505V) in deionized water via oral gavage for 3 consecutive days following the procedure outlined by Labash et al. [23]. At 42 weeks, to maximize yield and quality of blood samples, mice were anesthetized with no more than 5% isoflurane (v/v) and 100% oxygen saturation at a 1 L/min flow rate. The sternal cavity was opened for full visualization of the heart, and blood was collected from the left ventricle with a heparin-coated 22G needle and 1 mL syringe. The blood was then transferred to a microtainer containing EDTA (BD, 365974) to prevent coagulation. Immediately after collection, 60 μL of each sample was transferred to a tube containing an anticoagulant solution provided by Litron Laboratories and kept on ice for further processing following the Mouse MutaFlow^{PLUS} Kit manual (Litron Laboratories, v170925). Exsanguinated tissues were excised and flash frozen in liquid nitrogen and stored at -80°C.

Pig-a mutant frequency analysis

Pig-a mutant frequency analysis was conducted at Tulane University using a modified MutaFlow method. In summary, white blood cells and platelets were removed with Lympholyte®-Mammal (CedarLane, CL5110) and remaining reticulocytes, erythrocytes, and platelets were labeled with Anti-CD24-PE (Miltenyi, 130-110-688) and Anti-CD61-PE (Miltenyi, 130-102-628) antibodies and PE-conjugated magnetic

microbeads (Miltenyi, 130-048-801). A small portion of each sample was retained for cell

density determination and the remainder was enriched for mutant cells via

immunomagnetic separation on LS columns (Miltenyi, 130-042-401) within a magnetic

field. SYTO 13 nucleic acid dye (Invitrogen, S7575) was added to discriminate mature

erythrocytes from immature reticulocytes and remaining lymphocytes. Total mature

erythrocyte and *Pig-a* mutant erythrocyte counts were enumerated via the Miltenyi

MACSQuant Analyzer 10 flow cytometer. *Pig-a* mutant erythrocytes were classified as

cells exhibiting absence of the PE fluorescence signal and intracellular SYTO 13,

indicating a lack of both CD24 surface markers and nucleic acids. Mutant frequency was

calculated as the number of mutant erythrocytes per 10^6 total mature erythrocytes.

Hepatic Cyp2e1 activity

Hepatic Cyp2e1 activity was quantified at described above for the short-term HFD

experiment.

Statistical analysis

Pig-a mutant frequency was analyzed using two-way ANOVA with diet and ethyl

carbamate dose as the two factors, with Bonferroni's post hoc comparison test between

LFD and HFD groups within each dose group. A dose-response curve was plotted to

compare the effect of ethyl carbamate dose on Pig-a mutant frequency between LFD and

HFD groups, with slope comparison and R^2 calculation as well as an F test to compare

dose-response curve slopes between HFD and LFD groups. Cyp2e1 activity was

analyzed using a two-tailed independent Student's t test and simple linear regression, as

described for the short-term experiment. Males and females were analyzed separately.

Results

Short-term HFD experiment

Males gain more weight than females and exhibit biomarkers of diet-induced obesity after short term HFD treatment

All mice survived the duration of the short-term HFD experiment. When mice reached 22 weeks of age, the experiment was terminated. HFD males were significantly heavier (t_4=3.85, p=0.018) than LFD males and HFD females were also heavier than LFD females, though this difference was not statistically significant (t_4=2.48, p=0.068) (Fig 1A). HFD males exhibited significantly elevated serum leptin (t_4=4.745, p=0.009) and elevated serum triglycerides, (t_4=2.756, p=0.051) (Fig 1B-C). Serum leptin and triglycerides were not significantly elevated in HFD females (t_4=2.23, p=0.09; t_4=1.453, p=0.22) (Fig 1B-C).

Fig 1. Effects of diet on body mass, serum leptin, and serum triglycerides. Body mass (A), total serum triglycerides (B), and serum leptin (C) of female (left) and male (right) mice at collection (22 weeks). * p<0.05, **p<0.01, two-tailed independent Student's t test conducted within males and females separately, between LFD and HFD groups (n=3).

Short term HFD exposure induces DNA damage linked to lipid peroxidation and increased activity of xenobiotic metabolizing enzyme Cyp2e1

13 weeks of HFD treatment led to elevated levels of the adducted nucleotide

ethenoadenosine, a marker of lipid peroxide-induced DNA damage. Female and males

exhibited similarly elevated levels of liver etheno adenosine, though this difference was

significant in females (t_4=3.897, p=0.0176) and not significant in males (t_4=1.609,

p=0.1829) (Fig 2A). To further evaluate the effect of HFD-induced weight gain on lipid

peroxidation, linear regression analysis was conducted to determine the correlation

between body mass and liver ethenoadenosine. Among females, regression analysis

suggested a correlation between body mass and liver ethenoadenosine though not

significantly (R^2=0.5557, p=0.0889; Fig 2B, Table 1). Among males, body mass was

significantly positively correlated with liver ethenoadenosine content with an R^2 value of

0.77 (p=0.0216; Fig 2C, Table 1).

Fig 2. Effect of diet on liver ethenoadenosine. (A) Mean liver ethenoadenosine in

females and males, LFD compared to HFD. (B) Correlation of body mass and liver

ethenoadenosine in female and (C) males by simple linear regression. *p<0.05 per two-

tailed independent Student's t test conducted within males and females separately,

between LFD and HFD groups (n=3).

Table 1. Simple linear regression correlating body mass and liver ethenoadenosine

in female and male mice following short term HFD exposure.

Sex	Equation	R^2	Deviation from zero p value
Female	y = 1.970x-20.68	0.5557	0.0889
Male	y = 2.998x-87.49	0.7700	**0.0216**

Bold type indicates the slope deviates significantly from zero (α=0.05).

To evaluate the potential for HFD-induced sensitivity to xenobiotic toxification,

enzymatic activity of Cyp2e1 was measured in liver. Hepatic Cyp2e1 activity was elevated

in males (t_4=3.076, p=0.0371), but not significantly elevated in females (t_4=1.851,

p=0.1378; Fig 3). Simple linear regression suggested a correlation between body mass

and hepatic Cyp2e1 activity, but the association was not statistically significant in males

(R^2=0.5805, p=0.0783) or females (R^2=0.6416, p=0.0555) (Table 2). This pilot study was

limited by a small sample size, but post hoc power analysis indicated a sample size of

n=11 would yield significant differences in male ethenoadenosine, given the observed

effect size of 1.3. A sample size of n=8 per group would yield significant elevation in

female Cyp2e1, given the observed effect size of 1. Together, these results suggest an

increase in Cyp2e1-mediated synthesis of genotoxic lipid metabolites due to short-term

HFD exposure alone.

Fig 3. Effects of diet on liver Cyp2e1 activity. (A) Mean hepatic Cyp2e1 activity,

reported as fold difference between LFD and HFD groups, within males and females

separately. (B) Correlation of body mass and Cyp2e1 activity in female and (C) males by

simple linear regression. *p<0.05 per two-tailed independent Student's t test conducted

within males and females separately, between LFD and HFD groups (n=3).

Table 2. Simple linear regression correlating body mass and liver Cyp2e1 activity

in male and female mice following short term HFD exposure.

Sex	Equation	R^2	Deviation from zero p value
Female	y=0.07249x-0.5783	0.6416	0.0555
Male	y=0.09217x-2.023	0.5805	0.0783

Long-term HFD experiment

Higher rates of somatic mutations and precancerous markers develop over time with long term HFD in females.

To assess the mutagenicity of long-term HFD exposure, the erythrocyte Pig-a mutation assay was conducted with sequential blood sampling at 17, 33 and 52 weeks old. By week 52, one LFD female was euthanized due to injury, and two HFD males died due to undetermined causes. Both HFD males and HFD females were significantly heavier than their LFD counterparts at each sampling timepoint. HFD females were 6.98 ($t_{17.11}$=6.334, p<0.0001), 19.99 ($t_{15.12}$=8.281, p<0.0001), and 32.10 ($t_{14.92}$=13.27, p<0.0001) grams heavier than LFD females at 17, 33, and 52 weeks, respectively (Fig 4A). HFD male were 10.62 ($t_{14.24}$=8.151, p<0.0001), 14.83 ($t_{11.34}$=11.18, p<0.0001), and 15.90 ($t_{11.07}$=9.494, p<0.0001) grams heavier than LFD males at 17, 33, and 52 weeks, respectively (Fig 4B).

Fig 4. Effect of diet on body mass at each sampling timepoint. Male (A) and female (B) body mass in LFD and HFD groups at 17, 33 and 52 weeks of age. **** p<0.0001 per mixed effects analysis (for repeated measures) with Bonferroni's multiple comparisons test, between LFD and HFD groups at each timepoint, conducted in males and females independently (n=10-12).

One extreme outlier LFD female was excluded from all Pig-a mutant frequency analyses due to very high frequency of Pig-a mutant erythrocytes detected at 33 and 52 weeks (>4 standard deviations). Both raw and fold-difference erythrocyte mutant

frequency values were reported to normalize interexperiment variability. Despite a trend

of an approximately 1.5-2-fold elevation, there was no significant difference in mutant

frequency at any sampling timepoint between LFD and HFD females [24,25]. Because of

the variability in body mass observed in response to HFD, simple linear regression was

performed to assess the association between body mass and mutant frequency at each

sampling time point. Simple linear regression for the association between body mass and

erythrocyte mutant frequency per million total erythrocytes indicated the curve slope was

not significantly different from zero at 17 weeks (Fig 5C, Table 3). By week 33, there was

significant positive association between body mass and mutant frequency, with an R^2 of

0.2004 and significantly non-zero slope (p=0.0322). By week 52, there was a stronger

association, with an R^2 of 0.2518 and a significant slope (p=0.0286) (Fig 5D-E), indicating

higher body mass was associated with higher incidence of somatic mutations, and this

association grew stronger with age in females. Among males, while there was a similar

trend towards higher mutant frequency among the HFD group, there was no significant

differences between the LFD and HFD groups at any time point (Fig 6A-B), and there was

no significant correlation between body mass and mutant frequency at any sampling

timepoint (Fig 6C-E, Table 4).

Figure 5. Effects of diet and body mass on spontaneous mutation frequency in erythrocytes at 17, 33 and 52 weeks of age in females. (A) Mutant frequency per million erythrocytes, and (B) Mutant frequency fold-difference between LFD and HFD groups at each time point, per mixed effects analysis (for repeated measures) with Bonferroni's multiple comparisons test, between LFD and HFD group sat each timepoint (n=10-12). (C) Correlation between body mass and erythrocyte mutant frequency per

million cells at 17 weeks. (D) Correlation between body mass and erythrocyte mutant frequency per million cells at 33 weeks. (E) Correlation between body mass and erythrocyte mutant frequency per million cells at 52 weeks.

Table 3. Simple linear regression correlating body mass and Pig-a mutant frequency in females after 17, 33 and 52 weeks of dietary exposure.

Week	Equation	R^2	Deviation from zero p value
17	y=0.01042x+0.07179	0.01873	0.5763
33	y=0.008555x-0.01226	0.2004	**0.0322**
52	y=0.007024x+0.06099	0.2518	**0.0286**

Bold type indicates the slope deviates significantly from zero (α=0.05).

Fig 6. Effects of diet and body mass on spontaneous mutation frequency in erythrocytes at 17, 33 and 52 weeks of age in males. (A) Mutant frequency per million erythrocytes, and (B) Mutant frequency fold-difference between LFD and HFD groups at each time point, per mixed effects analysis (for repeated measures) with Bonferroni's multiple comparisons test, between LFD and HFD group sat each timepoint (n=10-12). (C) Correlation between body mass and erythrocyte mutant frequency per million cells at 17 weeks in males. (D) Correlation between body mass and erythrocyte mutant frequency per million cells at 33 weeks in males. (E) Correlation between body mass and erythrocyte mutant frequency per million cells at 52 weeks in males.

Table 4. Simple linear regression correlating body mass and Pig-a mutant frequency in males after 17, 33 and 52 weeks of dietary exposure.

Age (weeks)	Equation	R^2	Deviation from zero p value
17	y=0.01836x-0.4354	0.1339	0.0939
33	y=0.05517x-1.474	0.1504	0.1118
52	y=0.0112x-0.2016	0.02489	0.4946

To assess the impact of long term HFD exposure on development of precancerous hepatic biomarkers, hepatocyte nuclei were assessed for proliferative capacity by calculating the HPI and the nuclear Ki67 percent positivity. The HPI was significantly higher in HFD females compared to LFD females at 52 weeks old (t_6=8.171, p=0.0002), while there was no difference between HFD and LFD males (t_6=1.886, p=0.1082) (Fig 7A). Similarly, the percentage of hepatic nuclei that were Ki67+ was greater in HFD females than in LFD females (t_6=4.204, p=0.0057), with no difference seen between HFD and LFD males (t_6=0.1165, p=0.9110) (Fig 7B). These findings indicate HFD is a risk factor for cellular and molecular markers that are suggestive of precancerous transformation in females.

Fig 7. Effects of diet on precancerous hepatic markers at 52 weeks of age. (A) HPI in females (left) and males (right), (B) Percentage of hepatocyte nuclei that contained the Ki67 nuclear marker in females (left and males (right). ** p<0.01, *** p<0.001 per two-tailed independent Student's t test between LFD and HFD groups, conducted separately in males and females (n=3-5).

Ethyl carbamate challenge experiment

HFD increases sensitivity to ethyl carbamate-induced somatic mutagenesis in females.

To determine the effect of HFD on sensitivity to a known mutagen, mice were reared on HFD or LFD starting at 7 weeks of age, exposed to ethyl carbamate at 33 weeks, and the experiment was terminated following a 9 week mutation manifestation period. Dietary conditions were maintained for the entirety of the experiment, and 93 of 96 total mice survived until termination at 42 weeks old. One female HFD mouse was sacrificed at 17 weeks due to excessive pruritus with unhealed dermal lesions and lack of weight gain. One male HFD mouse was sacrificed at 22 weeks due to sudden rapid weight loss, several large obstructive growths were observed in the colon during necropsy. One female LFD mouse was cannibalized at 29 weeks.

Final body mass did not differ between ethyl carbamate dose groups, and all HFD groups were significantly heavier than their corresponding LFD controls (S1 Fig). Similarly to the long-term HFD only experiment, the proportional difference between HFD and LFD female body weight was greater than the body mass difference between LFD and HFD males. HFD exposure led to elevated liver triglycerides and elevated plasma leptin (S2 Fig) as well as AST and ALT in both males and females, indicating physiological obesity and liver damage due to HFD exposure (S3 Fig).

To determine the effect of longer term HFD exposure alone on the potential for generation of toxic lipid metabolites, hepatic Cyp2e1 activity was also assessed at 42 weeks in samples that had not been exposed to ethyl carbamate (vehicle controls). Hepatic activity of Cyp2e1, the enzyme that metabolizes ethyl carbamate to its mutagenic metabolite, vinyl carbamate, was elevated approximately 2.5-fold in females (t_9=2.872,

p=0.0184; Fig 8). There was no significant difference in hepatic Cyp2e1 activity in males

due to die (t_9=1.337, p=0.2141; Fig 8). Higher body mass was significantly associated

with higher hepatic Cyp2e1 activity in vehicle control females (R^2 =0.5083, p<0.05), but

not males (R^2=0.08934, p=0.4015) (Table 5). Because of the elevated endogenous

Cyp2e1 activity, females with long-term HFD exposure are predisposed to generation of

genotoxic lipid peroxides and vinyl carbamate epoxides.

Fig 8. Effects of diet on hepatic Cyp2e1 activity in 0 mg/kg/day ethyl carbamate

dose group. (A) Mean hepatic Cyp2e1 activity, reported as fold difference between LFD

and HFD groups, within males and females separately (n=5-6). (B) Correlation of body

mass and Cyp2e1 activity in female and (C) males by simple linear regression. * p<0.05

per Student's t test.

Table 5. Simple linear regression correlating body mass and liver Cyp2e1 activity

in male and female mice following 52 weeks of dietary exposure.

Sex	Equation	R^2	Deviation from zero p value
Female	y=0.07241x-0.9335	0.5083	**0.0138**
Male	y=0.01470x+0.4762	0.08934	0.4015

Bold type indicates the slope deviates significantly from zero (α=0.05).

HFD exposure in females led to increased susceptibility to somatic mutations

induced by exposure to 400 mg/kg/day ethyl carbamate (t_{38}=4.67, p=0.0001; Fig 9A). The

difference was on average 3.8-fold higher in the HFD group than the LFD group, within

those who received 400 mg/kg/day ethyl carbamate (t_{38}=4.67, p=0.0001; Fig 9B). To

determine if HFD females were more sensitive than LFD females to the same ethyl

carbamate dose regimen, dose response curves were plotted with regression analysis

between ethyl carbamate dose and Pig-a mutant frequency response. Simple linear

regression indicated the dose-response slope does not significantly differ from zero in the

LFD group (p=0.3872) and there was no correlation between ethyl carbamate dose and

mutant frequency response (R^2=0.0376; Fig 9C, Table 6). The HFD regression curve is

highly significantly different from zero (p<0.0001) and there is a strong correlation

between ethyl carbamate dose and mutant frequency (R^2 =0.5975; Fig 9C, Table 6). The

slope of the HFD curve is significantly higher than the LFD curve, indicating HFD

significantly increases the susceptibility of female mice to ethyl carbamate induced

somatic mutagenesis (p=0.0003; Table 6).

Fig 9. Effects of diet and body mass on mutation frequency in erythrocytes of

female mice exposed to 0, 100, 200, and 400 mg/kg/day ethyl carbamate. (A) Mutant

frequency per million erythrocytes in each ethyl carbamate dose group. (B) Mutant

frequency fold-difference between LFD and HFD groups in each ethyl carbamate dose

group. (C) Dose-response curves for ethyl carbamate dose and erythrocyte mutant

frequency with simple linear regression analysis, points represent mean ±standard error.

*** p<0.001 per two-way ANOVA with Tukey's multiple comparison test between LFD and

HFD groups (n=4-5).

Table 6. Simple linear regression dose-response analysis correlating ethyl

carbamate dose and Pig-a mutant frequency response in females.

Diet	Equation	R^2	Deviation from zero p value	Slope comparison p value
LFD	y=0.0006029x + 0.1626	0.4032	0.3872	**0.0003**
HFD	y=0.005166x - 0.02898	0.5924	**<0.0001**	

Bold type indicates the slope or slope difference deviates significantly from zero (α=0.05).

Among males there was no significant difference between the response of the HFD and

LFD groups to any level of ethyl carbamate treatment (Fig 10A). Though there was a

trend suggesting HFD males exhibit an approximately 2-fold increase in Pig-a mutant

frequency, regardless of ethyl carbamate treatment, these differences were not

statistically significant (Fig 10B). Linear regression analysis of the correlation between

ethyl carbamate dose and mutant frequency with slope comparison between the HFD

and LFD groups suggest a positive trend, although the slopes were not significant and

the R^2 values were weak (LFD: p=0.1032, R^2=0.1486; HFD: p=0.0951, R^2=0.1471; Fig

10C). There was ultimately no difference between the slopes for the LFD and HFD curves,

suggesting both dietary groups are equally sensitive to ethyl carbamate-induced

mutagenesis (p=0.5654, Table 7).

Fig 10. Effects of diet and body mass on mutation frequency in erythrocytes of

male mice exposed to 0, 100, 200, and 400 mg/kg/day ethyl carbamate. (A) Mutant

frequency per million erythrocytes in each ethyl carbamate dose group. (B) Mutant

frequency fold-difference between LFD and HFD groups in each ethyl carbamate dose

group. (C) Dose-response curves for ethyl carbamate and erythrocyte mutant frequency

with simple linear regression analysis, points represent mean ±standard error.

Table 7: Simple linear regression dose response analysis between ethyl carbamate dose and Pig-a mutant frequency response in males.

Diet	Equation	R^2	Deviation from zero p value	Slope comparison p value
LFD	y=0.0002403x+0.0.1553	0.1486	0.1032	
HFD	y=0.003947x+0.5569	0.1471	0.0951	0.5654

Discussion

The purpose of this set of experiments was to examine elements of the pathway that link diet-induced obesity to a physiological state primed for carcinogenesis from three angles. First we wanted to determine the early effects of short-term HFD exposure on two key outcomes linked to lipid peroxidation, ethenoadenosine adducts in genomic DNA and hepatic Cyp2e1 activity. Cyp2e1 catalyzes the reduction of lipids to reactive peroxides. This phenomenon is well studied in the context of alcoholic liver disease, methionine-choline deficient diet-induced NASH, and human NAFLD, but few studies have linked early Cyp2e1 induction and ethenoadenosine production where short term HFD was the only exposure in a controlled mouse model. Because of the small sample size used in this experiment, and the variability of body weight seen as a result of HFD, linear regression analysis was also performed to determine the relationship between body mass and the lipid peroxidation endpoints of interest. At the early time point, males and females responded somewhat differently to HFD. Males had significantly higher body mass, serum triglycerides, and serum leptin after 13 weeks of HFD, and females followed similar trends but were less responsive (Fig 1A-C). HFD females did have significantly higher liver ethenoadenosine levels than LFD females (Fig 2A-B), indicating even small, non-

significant increases in body mass and Cyp2e1 activity (Fig 3A-B) are linked to

accumulation of mutagenic DNA damage. Males exhibited similar trends. While HFD

males did not have significantly heightened levels of ethenoadenosine, male body mass

correlated significantly with liver ethenoadenosine. (Fig 2A and 2C, Table 1). Similarly,

though only males had significantly elevated Cyp2e1 activity, both males and females

exhibited a nearly significant correlation between Cyp2e1 activity and body mass (Fig 3A

and 3C, Table 2) . Together, these experiments were instrumental in suggesting that even

short-term HFD exposure alone leads to a hepatic environment primed for mutagenesis,

prompting the next set of experiments.

A year-long cohort was used to determine the effects of long-term HFD exposure

on somatic mutagenesis and pre-cancerous hepatic transformation in both male and

female mice. This was the first time this type of experiment was conducted including both

sexes. Surprisingly, though the short-term experiment indicated females are initially less

responsive, females in this long-term experiment were more responsive to HFD as they

aged than males, as indicated by the fact that HFD females were twice as heavy as LFD

females, while HFD males were less than 50% heavier than LFD males (Fig 4). This was

largely due to the fact that even males on the LFD become obese over time while females

do not, a phenomenon well-documented in the C57BL/6J strain[26]. Because of this

tendency of males to gain weight over time, regardless of diet, we postulate that both LFD

and HFD males represent some degree of biological overweight/obesity.

In females, linear regression analysis demonstrates that the association between

body mass and Pig-a mutant frequency strengthens with age and continued HFD

exposure (Fig 5C-D, Table 3), suggesting that the length of the duration of obesity over

the lifespan is linked to degree of cancer risk. While somatic mutagenesis is a key

component of the pathway from obesity to carcinogenesis, increased cell proliferation rate

is also often used as a biomarker to predict the likelihood of future cancer development

and progression[27,28]. Here we demonstrated that hepatocytes isolated from 52 week

old HFD females were more likely to be in an active proliferation state than age-matched

LFD controls, indicated by HPI and the percentage of hepatocytes with Ki67+ nuclei (Fig

7A-B). Both LFD and HFD males exhibited proliferation biomarker levels similar to those

found in HFD females. This result is consistent with our other findings that LFD and HFD

males exhibit similar characteristics at baseline.

The final set of experiments was designed to test the hypothesis that HFD-induced

obesity increases sensitivity to ethyl carbamate, a known Cyp2e1 substrate and

hepatotoxicant. Hepatic Cyp2e1 activity analysis was first repeated in the vehicle treated

group of this cohort to determine the effect of longer-term (until 42 weeks old) HFD. In

contrast to the short-term HFD cohort, females exhibited approximately 2.3-fold higher

Cyp2e1 activity due to HFD, while males had relatively similar activity regardless of diet

(Fig 8). HFD females developed, on average 3.8-times more somatic mutations in

response to high-dose ethyl carbamate than LFD females (Fig 9B). In fact, obesity

appears to potentiate the mutagenicity of ethyl carbamate in females, as indicated by the

significance of the ethyl carbamate-mutant frequency dose-response slope among HFD,

but not LFD, females (Fig 9C, Table 6). Among HFD males, mutagenicity of ethyl

carbamate appears to saturate between 100 and 400 mg/kg/day, with a peak at 200

mg/kg/day (Fig 10A-C). HFD males tended to exhibit higher mutant frequencies than LFD

males in each ethyl carbamate dose group, however the dose-response curves for LFD

and HFD males shared a common slope (Fig 10C, Table 7). These results suggests that

males on both diets were similarly sensitive to ethyl carbamate, further strengthening our

postulation that C57BL/6J male on the HFD and LFD formulation used in these studies

do not differ sufficiently. We attribute the overall lack of association between diet and

mutant frequency with the underlying weight gain observed in LFD males, and this

supports our assertion that studies that only include male C57Bl6/J mice are not sufficient

to explore the interactive effects of high-fat diet-induced obesity and environmental

toxicants.

Interpretation of sex differences observed in these three sets of experiments is

complicated by the different trends in weight gain between males and females. That is,

males on the LFD gained weight consistently throughout the duration of the study, while

females on the LFD maintain proportionally lower body masses. Increased weight gain

by LFD males maintained for longer periods of time may mask differences that might be

evident a study that included in truly lean (non-overweight, non-obese) males as a control

group. It is likely that increased metabolism of ethyl carbamate to mutagenic vinyl

carbamate epoxide by Cyp2e1 is at least partially responsible for the increased *Pig-a*

mutant frequency observed in HFD females. This observation, in conjunction with the lack

of obesity-related Cyp2e1 activity elevation and response to ethyl carbamate due to HFD,

further emphasizes that C57Bl6/J males are not the most suitable model for long-term

studies of diet-induced obesity with toxicant exposure. Additional calorie restriction or

reduced dietary fat may be required to generate a male equivalent of the female non-

obese phenotype within this model system.

Conclusions

The findings presented here emphasize the importance of studying obesity as a

mutagen and a carcinogen and the necessity of including both sexes in diet-induced

obesity research. The current paradigm regarding body weight and chemical safety is that

excess body weight is protective, due to the larger volume of distribution. This is a

dangerous assumption for chemicals that are metabolized by CYP2E1, if not other

CYP450s as well. If the purpose of chemical risk assessment is to protect the most

vulnerable groups, obesity must be factored in. In our female model, obesity appears to

act to both directly induce *de novo* mutations and hepatic proliferation, and to increase

sensitivity to xenobiotic-induced mutagenesis. We demonstrated that obesity is likely

involved in both the initiation (mutagenesis) and promotion (proliferation) stages of cancer

progression, however, further studies are needed to investigate the differences in male

and female response to HFD exposure. The limited scale of our experiment rendered

this research group unable to follow tumor formation, but it will be important to conduct

studies in the context of diet-induced HCC and other cancers as well.

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