Diet-dependence of metabolic perturbations mediated by the endocrine disruptor tolylfluanid

Shane M Regnier1,2,3, Andrew G Kirkley3,4, Daniel Ruiz1,3, Wakanene Kamau3, Qian Wu5, Kurunthachalam Kannan5 and Robert M Sargis6

1Committee on Molecular Metabolism and Nutrition, Chicago, Illinois, USA
2Pritzker School of Medicine, Chicago, Illinois, USA
3University of Chicago, Chicago, Illinois, USA
4Committee on Molecular Pathogenesis and Molecular Medicine, Chicago, Illinois, USA
5Wadsworth Center, New York Department of Health, Albany, New York, USA
6Division of Endocrinology, Diabetes, and Metabolism, Department of Medicine, University of Illinois at Chicago, Chicago, Illinois, USA

Abstract

Emerging evidence implicates environmental endocrine-disrupting chemicals (EDCs) in the pathogenesis of metabolic diseases such as obesity and diabetes; however, the interactions between EDCs and traditional risk factors in disease pathogenesis remain incompletely characterized. The present study interrogates the interaction of the EDC tolylfluanid (TF) and traditional dietary stressors in the promotion of metabolic dysfunction. Eight-week-old male C57BL/6 mice were fed a high-fat, high-sucrose diet (HFHSD) or a high-sucrose diet (HSD), with or without TF supplementation at 100 μg/g, for 12 weeks. Food intake, body weight and visceral adiposity were quantified. Glucose homeostasis was interrogated by intraperitoneal glucose and insulin tolerance tests at 9 and 10 weeks of exposure, respectively. After 12 weeks of dietary exposure, metabolic cage analyses were performed to interrogate nutrient handling and energy expenditure. In the background of an HFHSD, TF promoted glucose intolerance; however, weight gain and insulin sensitivity were unchanged, and visceral adiposity was reduced. In the background of an HSD, TF increased visceral adiposity; however, glucose tolerance and insulin sensitivity were unchanged, while weight gain was reduced. Thus, these analyses reveal that the metabolic perturbations induced by dietary exposure to TF, including the directionality of alterations in body weight gain, visceral adiposity and glucose homeostasis, are influenced by dietary macronutrient composition, suggesting that populations may exhibit distinct metabolic risks based on their unique dietary characteristics.

Key Words
- tolylfluanid
- endocrine disruptor
- diet
- adiposity
- glucose tolerance

Introduction

Global metabolic health has deteriorated dramatically over the last several decades with the emergence of the obesity and diabetes epidemics. Nearly 69% of the U.S. population is now overweight, with 35% currently obese (1) and approximately 50% projected to be obese by 2030 (2). Obesity is a major risk factor for the development of cardiovascular disease, obstructive sleep apnea and nonalcoholic fatty liver disease (3, 4), and the societal expansion in fat mass is thought to underlie much of the increasing prevalence of diabetes (5). Worldwide, 415 million individuals are estimated to have diabetes, with this number projected to reach a staggering
Recently, exposures to endocrine-disrupting chemicals (EDCs) have been implicated as novel metabolic disease risk factors. Indeed, synthetic chemical production correlates with the increasing prevalence of overweight and obesity (9) and diabetes (10). One compound of interest is the fungicide and antifouling agent tolylfluanid (TF). Utilized in agricultural regions outside of the U.S., TF is one of the most commonly detected fungicides in Europe where it has also been identified in groundwater (11, 12, 13, 14). Human exposure likely occurs through ingestion of contaminated food and water as well as occupationally (15, 16). However, current human exposure levels have not been tabulated. While not approved for use in the U.S., TF is permitted on imported foodstuffs, including apples, hops, grapes and tomatoes (17).

Previously, we have shown that TF augments adipogenesis in vitro (18) and promotes insulin resistance in rodent and human adipose tissues exposed ex vivo (19). These effects are postulated to result from activation of adipocytic glucocorticoid receptor (GR) signaling (18, 20). In recent studies, male C57BL/6 mice exposed to TF via a standard laboratory chow diet for 12 weeks exhibited increased adiposity, glucose intolerance and insulin resistance with concomitant impairments in adipose insulin sensitivity (21). While this study highlighted a potential role for TF in metabolic deterioration, the impact of this novel metabolic disruptor in the context of dietary changes which are characteristic of modernity is unknown. To this end, the present study expands upon prior work to elucidate the diet-dependent impact of TF on energy metabolism.

Methods

Animal husbandry, tissue processing, diets and EDC exposure

Eight-week-old male C57BL/6 mice were obtained from Jackson Laboratories and housed in pairs at 22.2 ± 1°C cycle. Mice were fed *ad libitum* with one of the following diets: a high-fat, high-sucrose diet (HFD), with fat content primarily derived from milk fat similar to that previously shown to disrupt metabolic homeostasis in mice (22), or a low-fat, high-carbohydrate diet enriched in sucrose (HSD) (Harlan Laboratories, Madison, WI, USA) (Table 1). Animals exposed to TF received an identical diet supplemented with 100 µg/g TF at the time of manufacturing to recapitulate regulatory policy that defines tolerance limits as residue levels per unit weight and to address controversies in TF regulations (23). Animals were weighed biweekly, and food pellets were weighed and replaced weekly. Animals were treated humanely in accordance with protocols approved by the Institutional Animal Care and Use Committee at the University of Chicago. At study termination, animals were euthanized by isoflurane inhalation followed by exsanguination via cardiac puncture followed by adipose depot dissection and quantitation. Adipose samples for histology were fixed in 10% formalin for 24 h before paraffin embedding and staining with hematoxylin and eosin.

Intraperitoneal glucose tolerance test (IP-GTT) and serum insulin levels

After nine weeks of exposure, mice were fasted for 6 h beginning at 07:30 h. Fasting blood glucose levels were measured using a Freestyle Lite glucometer (Abbott Laboratories). Mice then received an IP injection of dextrose (2 g/kg body weight) with blood glucose concentrations measured serially for 120 min. Fasting blood samples (20–25 µL) for serum insulin measurement were obtained using heparinized tubes, placed immediately on ice, centrifuged at 1500 g for 15 min at 4°C and concentrations were determined using a commercial insulin ELISA kit as per the manufacturer’s instructions (Millipore).

### Table 1 Macronutrient composition of murine diets.

<table>
<thead>
<tr>
<th>Diet (code, energy density)</th>
<th>High-fat, high-sucrose (HFHSD)* (TD.120438, 4.7 kcal/g)</th>
<th>High-sucrose (HSD)** (TD.140437, 3.8 kcal/g)</th>
</tr>
</thead>
<tbody>
<tr>
<td>Macronutrient</td>
<td>% kcal from</td>
<td>% kcal from</td>
</tr>
<tr>
<td>Protein</td>
<td>17.3</td>
<td>14.8</td>
</tr>
<tr>
<td>Carbohydrate</td>
<td>47.6</td>
<td>47.6</td>
</tr>
<tr>
<td>Fat</td>
<td>23.2</td>
<td>44.6</td>
</tr>
<tr>
<td>Saturated</td>
<td>14.0</td>
<td>27.2</td>
</tr>
<tr>
<td>Monounsaturated</td>
<td>7.0</td>
<td>13.6</td>
</tr>
<tr>
<td>Polyunsaturated</td>
<td>2.0</td>
<td>3.9</td>
</tr>
</tbody>
</table>

*Sucrose 340.0 g/kg; anhydrous milk fat 210 g/kg; **sucrose 703.0 g/kg; anhydrous milk fat 20.0 g/kg.
Intraperitoneal insulin tolerance test (IP-ITT)

After 10 weeks of exposure, mice were fasted for 3h beginning at 09:00h. Fasting blood glucose levels were measured as above. Mice then received an IP injection of Humalog insulin (0.5 U/kg body weight; Eli Lilly) with blood glucose concentrations measured serially for 90 min.

Metabolic cage analysis

After 12 weeks on the experimental diets, indirect calorimetric measurements were performed using the LabMaster System (TSE Systems) as previously described (21). The respiratory exchange ratio (RER) was measured as the ratio of O₂ inhalation to CO₂ production, averaged over a 12-h period. Metabolic flexibility was defined as the absolute change in RER between successive light and dark cycles. Energy expenditure, activity and food consumption were also quantified.

Measurement of dietary TF content

Chemicals and standards

Anhydrous sodium sulfate, sodium chloride and acetonitrile were purchased from Mallinckrodt (St Louis, MO, USA). TF (99.6%) was purchased from Fluka (St Louis, MO, USA).

Sample preparation

The extraction of TF followed the QuEChERS method (24), with slight modifications. Briefly, 0.5 g of diet pellet was transferred into a 15-mL polypropylene tube and homogenized in 10 mL of acetonitrile by an Ultra-Turrax T25 homogenizer. After the addition of 1 g of sodium sulfate and 0.25 g of sodium chloride, the samples were shaken in an orbital shaker for 40 min. The samples were centrifuged at 5000 g for 5 min. One milliliter of the supernatant was transferred into an amber autosampler vial for instrumental analysis.

Instrumental analysis

Instrumental analysis of TF was performed on an Agilent 1100 high-performance liquid chromatography system (Agilent) coupled with an API 2000 tandem mass spectrometer (AB SCIEX, Framingham, MA, USA) (HPLC–MS/MS). Five microliters of the sample extract was injected into a BetaSil C18 column (5 µm; 150 mm × 2.1 mm; Thermo Scientific). The HPLC mobile phase started with 20% A (100% acetonitrile) and 80% B (0.2% formic acid in water) for 2 min, increased to 95% A at 10 min followed by a 4-min hold and decreased to 20% A at 15 min followed by a 5-min hold reverting to original conditions. The tandem mass spectrometry was operated in the positive ionization mode with the ion spray voltage of 5500 V. The mass transition for TF was 347 > 137. The dwell time was 400 ms. Collision energy was 32 V for TF. All other parameters were optimized by infusing the standards in the tune mode.

Quality assurance and quality control

Procedural blanks were analyzed through the entire analytical procedure with 10 mL of acetonitrile for every five samples. Duplicate samples were analyzed for every five samples. Matrix spike tests were carried out for each sample type by spiking a known amount of the target compounds at three different concentrations in triplicates. The concentrations in the extracts designed for the matrix spike tests were 200, 1000 and 5000 ng/mL for TF. The respective mean recovery of TF through the analytical procedure was 85% for the HFHSD and 88% for the HSD. Matrix-matched calibration curves for each of the diets were used to calculate the concentrations of TF in samples. The limit of quantification was 1 µg/g for TF.

Statistical analyses

Adiposity was defined as adipose tissue weight divided by total body weight. Time-dependent variables were assessed by two-way ANOVA with multiple comparisons and Holm–Sidak post hoc test. Glucose and insulin tolerance were measured as the AUC of glucose over time by the trapezoidal method. For all endpoint analyses, individual groups were compared by F-testing to determine differences in variance; for F<0.05, t-tests were performed with Welch’s correction, whereas for F>0.05, standard Student’s t-tests were performed. All analyses were performed using GraphPad Prism version 6.0. A value of P<0.05 was considered statistically significant.

Results

Diet-dependent effects on caloric intake and TF exposure

Caloric excess drives metabolic dysfunction; however, the macronutrient content of diets may influence the
ultimate metabolic phenotype, including the impact of EDCs on energy homeostasis. The impact of TF on metabolic outcomes was analyzed within the context of two diets differing in macronutrient composition (Table 1). In neither the HFHSD (Fig. 1A and B) nor the HSD (Fig. 1C and D) did TF alter weekly or total caloric intake. TF was added to the dietary mix at the time of manufacturing at a concentration of 100 μg/g (0.2879 mmol/kg); final concentrations of TF were measured by HPLC–MS/MS and found to be 64.2 ± 7.59 mg/kg (0.185 ± 0.022 mmol/kg) in the HFHSD and 62.0 ± 0.40 mg/kg (0.179 ± 0.001 mmol/kg) in the HSD, which were not significantly different between diets (P > 0.05). Based on the quantity of food consumed and the final concentration of TF in each diet, weekly and total TF exposures were not significantly different between HFHSD- and HSD-fed mice (Fig. 1E and F). Taken together, these data suggest that TF does not modulate total food consumption.

Incorporating TF into an HFHSD impairs glucose tolerance and reduces visceral adiposity

Because consumption of a Western diet is associated with obesity (7), studies were undertaken to interrogate the potential interaction between TF and macronutrient contents of the diet in promoting metabolic dysfunction. In mice fed an HFHSD, the addition of TF did not modulate weight accrual (Fig. 2A and B). Despite the lack of body weight differences, after 12 weeks of feeding, mice on an HFHSD with TF were found to have a 14.7% reduction in visceral adiposity relative to the HFHSD controls (Fig. 2C). Histologic analysis of perigonadal/epididymal adipose tissue identified a significant increase in the number of adipocytes per high-power field (hpf) in the TF-treated group (Fig. 2D). Despite this reduction in adiposity, mice fed HFHSD with TF displayed impaired glucose intolerance relative to HFHSD controls (Fig. 2E), with a 20.1% increase in...
the glucose AUC (Fig. 2F). At this time, fasting insulin levels were not significantly different between groups (39.35 ± 3.79 µU/mL for HFHSD vs 40.16 ± 4.88 µU/mL for HFHSD + TF (P > 0.05)). Furthermore, despite these differences in glucose tolerance, systemic insulin sensitivity was not altered by TF exposure in an HFHSD (Fig. 2G and H). These findings suggest that consumption of an HFHSD supplemented with TF decreases adiposity and glucose tolerance, without altering body weight gain or insulin sensitivity.

In prior studies using a standard laboratory chow diet, TF altered fuel utilization preference and metabolic flexibility while disrupting diurnal rhythms of energy expenditure and food intake (21). To expand upon these findings, mice were phenotyped in metabolic cages following 12 weeks of exposure to TF in the setting of an HFHSD. In the background of an HFHSD, no significant differences were identified in TF-treated mice when assessing the RER, metabolic flexibility, absolute and corrected energy expenditures, activity or food consumption (Fig. 3A–F). Thus, in the setting of an HFHSD, TF exposure does not alter fuel preference, energy expenditure or metabolic circadian rhythms.

Incorporating TF into an HSD increases adiposity, reduces weight gain and modulates fuel utilization preference and energy expenditure

High dietary sugar consumption is a potent metabolic stressor with the capacity to disrupt insulin-glucose homeostasis and promote obesity development (25). Mice fed an HSD with TF exhibited reduced weight gain relative to HSD controls, ultimately resulting in a 23.3% decrease in fractional weight gain after 12 weeks (Fig. 4A and B). Interestingly, although the addition of TF to an HSD led to reduced weight gain, the same mice were found to have a 12.3% increase in visceral adiposity, with specific increases in perigonadal/epididymal, perirenal and mesenteric adipose depots (Fig. 4C). No significant difference in the number of adipocytes per hpf was identified with TF exposure (Fig. 4D). When the same mice were metabolically challenged via IP-GTT and IP-ITT, no differences in glucose tolerance (Fig. 4E and F) or insulin sensitivity (Fig. 4G and H) were identified. In addition, fasting insulin levels at the time of IP-GTT were not significantly different (8.21 ± 0.64 µU/mL for HSD vs 10.09 ± 1.27 µU/mL for HSD + TF (P > 0.05)). Thus, consumption of an HSD supplemented with TF promotes
Diet-dependent metabolic effects of tolylfluanid

Figure 3
Metabolic cage analyses on a high-fat, high-sucrose diet. Eight-week-old male C57BL/6 mice were provided a high-fat, high-sucrose diet (HFHSD) with or without tolylfluanid (TF) added at the time of manufacturing at a concentration of 100 μg/g for 12 weeks. After 12 weeks’ exposure, mice were singly housed in metabolic cages and continued on their respective diets. The respiratory exchange ratio (RER) was calculated by the ratio of O₂ inhalation to CO₂ production, with higher values suggesting an increased preference for carbohydrate as fuel relative to lipid. There were no differences in circadian fuel preference (Panel A) or in metabolic flexibility (Panel B). There were no differences in absolute or body mass-adjusted energy expenditure (Panels C and D, respectively). There were no differences in activity (Panel E) or food consumption (Panel F). Data are presented as means ± s.e.m. N = 4 per group.

Figure 4
Metabolic impact of TF exposure in the context of a high-sucrose diet. Eight-week-old male C57BL/6 mice were provided a high-sucrose diet (HSD) with or without tolylfluanid (TF) added at the time of manufacturing at a concentration of 100 μg/g for 12 weeks. Body weight was measured twice weekly (Panel A) with fractional body weight gain calculated after 12 weeks, with a significant reduction in weight gain in the TF-exposed group (Panel B). At the end of 12 weeks, mice were euthanized and visceral adipose depots (perigonadal/epididymal, perirenal, mesenteric) harvested, weighed and sectioned for histological analysis. Adiposity was calculated as adipose weight normalized to body weight. Exposure to TF led to an increase in adiposity for all visceral depots (Panel C), without a difference in the number of adipocytes per high-power field (Panel D). After 9 weeks’ exposure, glucose tolerance testing (Panel E) did not reveal significant differences between groups (Panel F). After 10 weeks’ exposure, insulin tolerance testing was performed (Panel G), without significant differences identified between groups (Panel H). Data are presented as means ± s.e.m. N = 23–24 for body weight and adiposity measures, and 20–21 for glucose and insulin tolerance testing. *P < 0.05; **P < 0.01; ***P < 0.001.
increased visceral adiposity with a concurrent reduction in weight gain; however, despite this increase in adiposity, insulin–glucose homeostasis was preserved.

When HSD-fed mice were analyzed in metabolic cages, exposure to TF increased RER during dark cycles, suggesting enhanced utilization of carbohydrate relative to fat as an energy source (Fig. 5A). This change occurred without impacting metabolic flexibility, likely because RER was also non-significantly higher during the light cycle as well (Fig. 5B). Exposure to TF also led to a significant increase in absolute energy expenditure during the light cycle (Fig. 5C). When corrected for total body weight, TF treatment increased energy expenditure during both light and dark cycles (Fig. 5D). TF supplementation of the HSD did not affect activity (Fig. 5E). Food consumption during the dark cycle was increased in the TF-exposed group (Fig. 5F). Of note, daily food intake in the HSD group, but not the TF-exposed group on HSD, was significantly reduced during the metabolic cage analysis relative to the calculated daily food consumption during the other studies discussed herein (data not shown; $P<0.05$). Taken together, these findings suggest that in the setting of an HSD, TF exposure modulates fuel utilization, energy expenditure and food consumption.

**Discussion**

Human metabolic health has deteriorated dramatically in recent decades, concurrent with alterations in diet composition and increasing exposure to environmental pollutants. We have postulated that the interactions among various metabolic stressors underlie an individual’s disintegration of metabolic integrity (26). Previously, the phenylsulfamide fungicide and booster biocide TF, when consumed in the background of a standard laboratory chow diet (Teklad 2018), promoted increased body weight gain, visceral adiposity, glucose intolerance and insulin resistance (21). In the present study, the interaction between dietary stressors and EDC exposure in promoting metabolic dysfunction was examined utilizing the same EDC, animal model and route of exposure. When coupled
with an HFHSD, TF promoted glucose intolerance; however, adiposity was reduced, while weight gain and insulin sensitivity remained unchanged (Fig. 2). In contrast, addition of TF to an HSD led to increased adiposity; however, weight gain was reduced, while glucose tolerance and insulin sensitivity were unaffected (Fig. 4). Importantly, weekly and total TF exposures were equivalent between the HFHSD and the HSD (Fig. 1). Thus, these findings suggest that the macronutrient composition of the diet through which TF exposure occurs alters TF-dependent metabolic outcomes.

While EDCs have emerged as contributors to metabolic disease pathogenesis, it is well recognized that calorically dense diets are potent contributors to the observed increase in obesity and related metabolic diseases. In animal models, mice fed a high-fat diet are known to develop obesity, insulin resistance and glucose intolerance (27). In addition to high levels of saturated and trans fats, Western diets are characterized by high quantities of sugar, particularly sucrose and high fructose corn syrup, principally from added sweeteners (28). Importantly, consumption of sucrose as a caloric sweetener in the U.S. increased 43% between 1950 and 2000, with per capita consumption now over 65 pounds annually (28). Interestingly, when fed a sucrose-supplemented diet, pregnant mice developed increased adiposity and dyslipidemia (29). In another study, when given a diet high in both sucrose and fat, male mice developed obesity and metabolic dysfunction in only 4 weeks (22). While the impact of high-fat diets on the metabolic effects of EDCs has begun to be investigated (30, 31), the impact of high sugar diets has not received much attention. Based on the data presented herein, further work in this area is needed as EDC effects on metabolic outcomes may be nutrient-dependent.

Metabolic cage analyses allow for precise examination of movement, nutrient consumption and energy expenditure. In prior studies, consumption of a standard chow diet with TF for 12 weeks promoted a state of relative metabolic inflexibility and an increased reliance on lipids as an energy source during the dark cycle (21). This metabolic disruption is directionally similar to that observed in mice fed a high-fat diet (32), and in obese humans (33). In the present study, mice fed an HSD supplemented with TF exhibited enhanced carbohydrate utilization during the dark cycle relative to HSD controls (Fig. 5A). Of note, in the HSD group, daily food consumption was significantly reduced during the metabolic cage analysis in which mice were singly housed, relative to the calculated daily food consumption under standard pair-housing for the other studies discussed herein; this finding was not observed for the group fed TF-enriched HSD nor either HFHSD group. This finding may be secondary to the effects of cumulative exposure to TF over 12 weeks in the setting of an HSD or to diet-specific TF-mediated changes in response to social isolation. This reduced caloric intake by HSD-fed controls may partially explain the relative increase in food consumption of TF-treated mice consuming an HSD during the metabolic cage analyses (Fig. 5F), since TF did not alter caloric intake over the duration of the 12-week exposure study (Fig. 1C and D). Furthermore, the increase in carbohydrate-rich food consumption by TF-treated HSD mice may contribute to the observed carbohydrate utilization preference (Fig. 5A). Alternatively, this enhanced carbohydrate utilization with TF may reflect TF-mediated shifts in nutrient handling, as a prior study showed that TF reduced expression of fatty acid oxidation genes (21). Interestingly, the increased energy expenditure observed in TF-treated HSD-fed mice (Fig. 5C and D) may contribute to the reduction in weight gain identified through 12 weeks of TF exposure (Fig. 4A and B).

The findings presented herein provide insights into the interplay between environmental contaminants and diet composition, which may have important effects on metabolic health disparities. Black and Hispanic adults and children have increased rates of overweight and obesity relative to Caucasians (1, 34). While many factors contribute to this, minorities and the socioeconomic disadvantaged are more likely to reside in food deserts, regions defined by limited access to fresh foods, forcing increased reliance on prepackaged and fast foods high in the same macronutrients utilized in the present study (35). Unsurprisingly, individuals in the same demographic groups have a higher prevalence of obesity, diabetes and cardiovascular disease (36). In addition to these dietary stressors, low-income and minority communities also suffer a greater burden of pollutant exposure (37). For example, urinary concentrations of the common EDC bisphenol A are inversely correlated with income (38), while demographic analyses demonstrated a disproportionately higher percentage of minority individuals living nearest to high-risk chemical exposure areas (39). Consequently, understanding the interplay between diet and EDC exposure will be vital for addressing the contribution of disproportionate environmental exposures to obesity and diabetes disparities (37).

While the present findings suggest that TF differentially modulates energy handling based upon dietary context,
this study has several limitations. Although post-processing concentrations of TF were equivalent between the HFHSD and the HSD, tissue concentrations were not assessed, precluding direct comparison of TF-dependent effects between diets. Because TF is lipophilic (log octanol-to-water partition coefficient ($K_{ow}$) of 3.9 (47)), differential gastrointestinal absorption related to dietary fat content is a possibility, as shown for diethylhexyl phthalate (40). Furthermore, although consumption of TF was equivalent across diets, caloric intake was not (Fig. 1), thus precluding direct comparison between diets. Despite these limitations, the observed differences in the directionality of TF effects on metabolic outcome measures suggest that diet macronutrient content modulates TF-mediated alterations on weight gain, adiposity and glucose tolerance. These findings provide justification for future investigations using a pair-feeding model matching both exposure and caloric intake while ensuring comparable internal TF levels. In addition, future studies may expand the scope of dietary content to include other common components of the Western diet, such as fructose and other animal fats. Finally, future studies hope to decipher the molecular mechanisms by which TF interacts with dietary macronutrients to alter energy homeostasis. Given the importance of GR signaling in regulating energy metabolism and the capacity of TF to modulate signaling through this pathway (18, 20, 41, 42), efforts to understand the impact of TF on GR signaling in the context of dietary manipulation will be particularly important for linking disparate metabolic phenotypes to underlying molecular mechanisms. This is particularly important in the context of studies suggesting that the impact of TF on metabolic physiology may not be universal (43).

The present study extends prior findings demonstrating TF-induced metabolic disruption by identifying novel interactions between TF and diet macronutrient content in the disruption of key metabolic parameters, including weight gain, adiposity and glucose tolerance. Evidence that diet composition modulates the metabolic consequences of pesticide residues suggests that Environmental Protection Agency tolerance limits may need modification based on the dietary vehicle through which exposure occurs. Moreover, these findings highlight the important role of diet in mediating outcomes of EDC exposure, a crucial consideration for understanding the complexity of factors responsible for the pathogenesis of obesity and diabetes.

Declaration of interest
The authors declare that they have no conflicts pertinent to this manuscript.

Funding
Work supported by the National Institutes of Health: K08-ES019176 and R21-ES021354 to R M S, T32-HD07009 to S M R and D R, and a Pilot and Feasibility Grant from the NIH-funded University of Chicago Diabetes Research and Training Center (P60-DK020595 to R M S) and the ChicAg Center for Health and Environment (CACHET, P30-ES027792 to R M S). This work was also supported by the American Diabetes Association (1-17-JDF-033 to R M S).

Acknowledgements
The authors would like to thank Nicole Massad for her contributions.

References
14 Reemtsma T, Alder L & Banasiak U. Emerging pesticide metabolites in groundwater and surface water as determined by the application of


18 Sargs RM, Johnson DN, Choudhury RA & Brady MJ. Environmental endocrine disruptors promote adipogenesis in the 3T3-L1 cell line through glucocorticoid receptor activation. Obesity 2010 18 1283–1288. (https://doi.org/10.1038/oby.2009.419)


20 Neel BA, Brady MJ & Sargs RM. The endocrine disrupting chemical tolyfluanid alters adipocyte metabolism via glucocorticoid receptor activation. Molecular Endocrinology 2013 27 394–406. (https://doi.org/10.1210/me.2012-1270)


23 European Food Safety Authority. Reasoned opinion on the review of the existing maximum residue levels (MRLs) for tolyfluanid according to Article 12 of Regulation (EC) No 396/2005. European Food Safety Authority Journal 2013 11 3300. (https://doi.org/10.2903/j.efsa.2013.3300)


25 DiNiccolantonio JJ & Berger A. Added sugars drive nutrient and energy deficit in obesity: a new paradigm. Open Heart 2016 3 e000469. (https://doi.org/10.1136/openhrt-2016-000469)


29 Toop CR, Mulhuisser BS, O’Dea K & Gentili S. Consumption of sucrose, but not high fructose corn syrup, leads to increased adiposity and dyslipidaemia in the pregnant and lactating rat. Journal of Developmental Origins of Health and Disease 2015 6 38–46. (https://doi.org/10.1017/S2040174414000610)


Received in final form 18 November 2017
Accepted 28 November 2017
Accepted Preprint published online 29 November 2017

License.