



Dietary fat quality impacts metabolic impairments of type 2 diabetes risk differently in male and female CD-1[®] mice

Allison L. Unger¹, Thomas L. Jetton² and Jana Kraft^{1,2*}

¹Department of Animal and Veterinary Sciences, The University of Vermont, Burlington, VT 05405, USA

²Department of Medicine, Division of Endocrinology, Metabolism and Diabetes, The University of Vermont, Colchester, VT 05446, USA

(Submitted 19 May 2021 – Final revision received 30 August 2021 – Accepted 28 September 2021 – First published online 4 October 2021)

Abstract

Metabolic impairments associated with type 2 diabetes, including insulin resistance and loss of glycaemic control, disproportionately impact the elderly. Lifestyle interventions, such as manipulation of dietary fat quality (i.e. fatty acid (FA) composition), have been shown to favourably modulate metabolic health. Yet, whether or not chronic consumption of beneficial FAs can protect against metabolic derangements and disease risk during ageing is not well defined. We sought to evaluate whether long-term dietary supplementation of fish-, dairy- or echium-derived FAs to the average FA profile in a U.S. American diet may offset metabolic impairments in males and females during ageing. One-month-old CD-1[®] mice were fed isoenergetic, high-fat (40%) diets with the fat content composed of either 100% control fat blend (CO) or 70% CO with 30% fish oil, dairy fat or echium oil for 13 months. Every 3 months, parameters of glucose homeostasis were evaluated via glucose and insulin tolerance tests. Glucose tolerance improved in males consuming a diet supplemented with fish oil or echium oil as ageing progressed, but not in females. Yet, females were more metabolically protected than males regardless of age. Additionally, Spearman correlations were performed between indices of glucose homeostasis and previously reported measurements of diet-derived FA content in tissues and colonic bacterial composition, which also revealed sex-specific associations. This study provides evidence that long-term dietary fat quality influences risk factors of metabolic diseases during ageing in a sex-dependent manner; thus, sex is a critical factor to be considered in future dietary strategies to mitigate type 2 diabetes risk.

Key words: Ageing: Dairy fat: Echium oil: Fish oil: Glucose tolerance: Insulin sensitivity: Sex differences

The USA Census Bureau has reported a staggering increase in the elderly population worldwide (i.e. 65 years and older). As of 2015, there are approximately 617 million aged individuals, representing 9% of the total population^(1–3). From a healthcare perspective, a significant concern is the heightened susceptibility of the elderly to disease. In particular, insulin resistance, weight gain, and progressive pancreatic β -cell dysfunction are concomitant with the ageing process^(4–7), and thus ageing is considered an important risk factor for metabolic diseases such as type 2 diabetes (T2D)^(8,9). With T2D prevalence doubled in the elderly (20% from 9%) compared with the general population⁽¹⁰⁾ and the rising life expectancy of the older population⁽¹⁾, T2D in the elderly is a fast-growing public health issue.

Lifestyle intervention is considered to be the most effective strategy to prevent metabolic diseases, even compared with traditional medication regimens^(11–15). For example, physical activity, weight loss, and restriction of total energy or macronutrients consumed (e.g. carbohydrates) are commonly employed to

combat metabolic abnormalities⁽¹⁶⁾. Recently, manipulation of dietary fat composition (i.e. fatty acid (FA) profile) to mitigate metabolic impairments has also become of considerable scientific interest. The benefits of supplementing the diet with cold-water fatty fish or its derived oils are well-established, underscoring the favourable physiologic effects of long-chain *n*-3 FAs (i.e. EPA, 20:5 and DHA, 22:6)^(17–19). Furthermore, epidemiological studies indicate that consumption of full-fat dairy products, with a uniquely complex dietary FA composition (> 400 FAs⁽²⁰⁾; short-, odd- and branched-chain FAs, 18:1 and 18:2 *trans* isomers)⁽²¹⁾, can reduce metabolic derangements and disease risk^(22–25). However, detailed clinical evidence and mechanistic data validating this relationship are not available. Additionally, the impact of consumption of plant-derived *n*-3 and *n*-6 FAs on T2D susceptibility is not clear^(26–29). Echium oil is a promising source of pro-health FAs^(27,30) because of its unusually high content of α -linolenic acid (18:3 *n*-3), stearidonic acid (18:4 *n*-3), and γ -linolenic acid (18:3 *n*-6)⁽³¹⁾, an *n*-6 FA with purported

Abbreviations: BO, butter oil; EO, echium oil; FA, fatty acid; FO, fish oil; GTT, glucose tolerance test; ITT, insulin tolerance test; T2D, type 2 diabetes.

* **Corresponding author:** Jana Kraft, email jana.kraft@uvm.edu



anti-inflammatory properties⁽³²⁾. Importantly, whether or not chronic consumption of beneficial FAs can protect against metabolic abnormalities and disease development during ageing is unclear.

We sought to determine whether long-term effects of diets differing in fat quality influence metabolic derangements associated with T2D risk in male and female CD-1[®] mice. We hypothesised that dietary supplementation with fish-, dairy- and echium oil would attenuate metabolic impairments induced by consumption of a fat with a FA composition representative of the average U.S. American diet. Our specific objectives were to (i) measure glucose tolerance and insulin sensitivity every 3 months of experimental feeding, (ii) assess the effects of dietary fat quality, sex, and age on glucose tolerance and insulin sensitivity and (iii) investigate potential diet-induced biological mechanisms influencing glucose tolerance and insulin sensitivity in CD-1[®] mice.

Experimental methods

Animals and experimental design

The outbred CD-1[®] mouse stock (IGS #022) was chosen as a pre-clinical model to assess the metabolic effects of a long-term nutritional intervention throughout different life stages because of its genetic heterogeneity⁽³³⁾ (representative of a human population) and documented longevity⁽³⁴⁾. After weaning (3 weeks of age), male and female CD-1[®] mice ($n = 81$, $n = 40-41$ per sex) were shipped from Charles River (Raleigh, NC, USA) to The University of Vermont (Colchester, VT, USA). Upon arrival, mice were immediately randomised (no specific randomisation technique used) into same-sex pairs and one group of three male cage mates. Cages were kept in ventilated racks (Thorens Caging Systems, Hazelton, PA, USA) on a 12 h light/dark cycle at 23-6°C with 64 % humidity. Wooden blocks and igloo shelters were provided as enrichment.

For 1 week, while mice adjusted to their new housing conditions, mice were fed a standard laboratory pelleted chow (26 % protein, 60 % carbohydrate, and 14 % fat; LabDiet, St. Louis, MO, USA). At 4 weeks of age, mice ($n = 10-11$ per sex/diet group) were switched over to one of four isoenergetic high-fat (40 % of total energy) diets. Investigators were not blinded to the group allocation for any stage of the experiment but were blinded during analysis. Mice consumed their designated diets until the conclusion of the study (14 months of age). Throughout the study, mice had *ad libitum* access to feed and water.

All experimental procedures with animals were approved by and performed in accordance with The University of Vermont Institutional Animal Care and Use Committee (protocol #16-007). Animal protocols also adhered to the ARRIVE Guidelines for Reporting Animal Research (The ARRIVE Essential 10 and The Recommended Set)⁽³⁵⁾.

Experimental diets

Experimental diets were formulated to be equivalent to one another with respect to macronutrients (protein, carbohydrate, and fat as 17, 43, and 40 % energy, respectively) and differed

only in FA composition (Table 1). Each experimental diet was designed to supply 40 % of daily kcal as fat, 5 % greater than what is generally recommended by USA dietary guidelines⁽³⁶⁾, yet still feasible for human consumption. The fat content in the control diet (CO) was composed of an in-house designed USA fat blend, formulated to mimic the FA composition of the average U.S. American diet⁽³⁷⁾. The fat content of the other three experimental diets contained 70 % of the USA fat blend and 30 % of a supplementary fat source, either (i) dairy fat derived from butter oil (BO diet), (ii) echium oil from the seed of *Echium plantagineum* (EO diet) or (iii) fish (menhaden) oil (FO diet). Supplementation of the BO-, EO- and FO-diet with 30 % of the respective fat source intentionally sought to represent a dietary pattern whereby an average U.S. American substitutes fat derived from dairy, echium or fish in place of a portion of their typical fat source. In-house preparation and FA analysis of the experimental fat blends⁽³⁸⁾ and a complete composition and ingredient list of the

Table 1. Composition (g/kg) of experimental high-fat diets

	CO	FO	BO	EO
Diet component (g/kg diet)				
Protein	200	200	200	200
Carbohydrate	500	500	500	500
Fat	210	210	210	210
Basal diet mix*	790	790	790	790
USA fat blend†	210	147	147	147
Fish oil supplement‡	–	63	–	–
Dairy fat supplement	–	–	63	–
Echium oil supplement	–	–	–	63
FA classes (g/kg diet)§				
∑ SFAs	70	66	86	61
∑ MUFAs	81	67	76	73
∑ PUFAs	44	48	34	75
∑ Dairy-derived FAs	3	6	9	2
∑ Short- and medium-chain FAs¶	1	1	3	1
∑ BCFAs**	0	1	1	0
18:1 n-1	0	0	1	0
∑ Echium-derived FAs††	4	5	3	34
18:3 c6,c9,c12 (n-6)	0	0	0	6
18:3 c9,c12,c15 (n-3)	4	3	3	22
18:4 c6,c9,c12,c15 (n-3)	0	1	0	7
∑ Fish-derived FAs‡‡	0	14	0	0
20:5 c5,c8,c11,c14,c17 (n-3)	0	7	0	0
22:5 c7,c10,c13,c16,c19 (n-3)	0	1	0	0
22:6 c4,c7,c10,c13,c16,c19 (n-3)	0	5	0	0

BCFAs, branched-chain fatty acids. CO, 100 % USA fat blend. FO, 70 % CO supplemented with 30 % fish oil. BO, 70 % CO supplemented with 30 % dairy (butter) fat. EO, 70 % CO supplemented with 30 % echium oil. FA, fatty acid.

* Formulated by Research Diets Inc.

† USA fat blend consisted of lard, walnut oil, high-oleic sunflower oil, coconut oil and palm oil in a ratio of 18:8:3:6:2:8:1:8:1:0.

‡ Fish oil supplement was derived from menhaden and supplied by Research Diets Inc.

§ Analysis of FAs was performed via gas-liquid chromatography as described in Unger *et al.*⁽³¹⁾. Data presented as 0 g/kg of diet signify <0.5 g/kg.

|| Dairy-derived FAs include the sum of 4:0–11:0, 13:0 *iso*, 13:0 *aiso*, 14:0 *iso*, 15:0, 15:0 *iso*, 15:0 *aiso*, 16:0 *iso*, 16:1 *iso*, 17:0, 17:0 *iso*, 17:0 *aiso*, 18:0 *iso*, 18:1 *trans* isomers, 18:2 *trans* isomers and 18:2 conjugated isomers.

¶ Short- and medium-chain FAs include the sum of dairy-derived FAs 4:0–11:0.

** BCFAs include the sum of 13:0 *iso*, 13:0 *aiso*, 14:0 *iso*, 15:0 *iso*, 15:0 *aiso*, 16:0 *iso*, 17:0 *iso*, 17:0 *aiso* and 18:0 *iso*.

†† Echium-derived FAs include the sum of 18:3 c6,c9,c12 (n-6), 18:3 c9,c12,c15 (n-3) and 18:4 c6,c9,c12,c15 (n-3).

‡‡ Fish-derived FAs include the sum of 20:5 c5,c8,c11,c14,c17 (n-3), 22:5 c7,c10,c13,c16,c19 (n-3), and 22:6 c4,c7,c10,c13,c16,c19 (n-3).



experimental pelleted diets⁽³¹⁾ (formulated by Research Diets, Inc. (Brunswick, NJ, USA)) have been previously described.

Parameters of health and glucose homeostasis

Each mouse was considered an experimental unit. For the duration of the study, feed intake and body weight were measured to assess growth and health. Feed intake was evaluated weekly on a cage basis. Body weight was measured every week for the first 3 months and subsequently every month. If an animal was observed to rapidly lose a significant amount of weight (10–20% of body weight within one month), animals were euthanised.

Every month during the intervention, whole blood was collected via tail-nick for determination of fed (09.00) glucose and insulin concentrations. In addition, mice were subjected to intraperitoneal glucose tolerance tests (GTTs) and insulin tolerance tests (ITTs) every 3 months to evaluate glucose homeostasis and insulin sensitivity. These procedures have been previously described in detail in Unger *et al.*⁽³⁹⁾. Briefly, after a 6 h fast, GTTs (120 min in duration) began with an intraperitoneal injection of sterile 2 g/kg glucose (Sigma-Aldrich, St. Louis, MO, USA). The Zeitgeber time was 3 (i.e. 10.00, as lights on in our facility is 07.00) for both glucose and ITTs; thus, food removal and cage changes commenced at 04.00 (Zeitgeber time of 21) for a 6 h fast. From this procedure, the following measurements were assessed: fasted glucose and insulin, homeostatic model of insulin resistance (HOMA-IR), GTT area under the curve of glucose (AUC), and final (120 min) blood glucose measurement of GTT. Similarly, fasted mice were challenged during ITTs (60 min in duration) with intraperitoneal administration of insulin (Humulin® R (U-100); 0.75 U/kg; Eli Lilly and Company, Indianapolis, IN, USA). From this procedure, the following measurements were assessed: ITT AUC, final (60 min) blood glucose measurement of ITT, and percent change in blood glucose during ITT.

To minimise potential confounding of results, such as treatment order, during procedures and measurements, two to three cages of each sex and treatment group were assessed together. Procedures and measurements were then staggered over the course of approximately 1 week. In this manner, procedure day and procedure time of day were controlled for each treatment group and sex. For example, fasting of mice on a cage basis as preparation for GTTs and ITTs was staggered every 20 min beginning at 04.00, with glucose or insulin administration, respectively, staggered accordingly starting at 10.00 to achieve a 6-h fast.

Colonic bacterial composition

The effects of the experimental diets on the colonic bacterial composition (relative abundance and diversity indices) in this cohort of mice, assessed at 10.5 and 13.5 months of age, have been established previously⁽³⁸⁾. To examine the relationship between dietary fat quality, metabolic health, and colonic bacteria, we performed correlation tests between parameters of glucose homeostasis at 12 months of experimental feeding with published data of colonic bacterial composition measured before harvest (13.5 months of age).

Tissue collection and analysis

We also sought to correlate dietary FA tissue content with metabolic responses to the experimental diets. Previous measurements of diet-derived FAs (percent of total FA methyl esters) into body tissues collected at harvest (14 months of age) have been reported in Unger *et al.*⁽³¹⁾.

Measurement of circulating estradiol

Circulating estradiol (E2) concentrations were measured in a subset of female mice at sexual maturity (5 months of age) and at harvest (14 months of age). Plasma E2 was measured via a commercial murine E2 ELISA kit (Calbiotech Inc., El Cajon, CA, USA) following the protocol provided by the manufacturer.

Statistical analysis

All data, except for plasma E2 concentrations, were analysed via linear mixed model with an unstructured covariance in IBM SPSS Statistics for Macintosh, Version 25.0 (MIXED function; IBM Corp., Armonk, NY, USA). Diet, sex, and time (age) were included as fixed effects, while body weight was included as a covariate. Normality in the distribution of residuals in the model was evaluated via Q-Q plots, and data that were not normally distributed were transformed. The following dependent variables were transformed as follows: feed efficiency, body weight, body weight gain, ITT AUC via square root; fed glucose and insulin, fasted glucose and insulin, GTT AUC, HOMA-IR, and final blood glucose measurement of GTT and ITT via log. If significance for a main effect or an interaction between two main effects on a dependent variable was present, unadjusted pairwise differences (COMPARE function) were examined. An unadjusted post hoc test was specifically chosen due to the large number of interactions in the model. Spearman correlations were performed in R (version 3.4.2). For all data, significance was determined as $P < 0.05$, and a trend was determined as $P = 0.05–0.10$. Non-transformed data are stated in all tables and figures. Heat maps for correlations were generated in R (version 3.4.2), and all other graphic visualisations of the data were created with GraphPad Prism version 7.00 (GraphPad Software, La Jolla, CA, USA). Plasma E2 concentrations at 5 and 14 months of age in females were compared via a two-sample t test in JMP® (Version 15, SAS Institute Inc., Cary, NC, USA).

Sample size determinations were performed using an Excel-based power calculator⁽⁴⁰⁾ and considering 10% attrition. The minimum number of mice required for these studies was based on our previous study in chronically diet-induced obese mice⁽⁴¹⁾, whereby frank insulin resistance was initially detectable by ITT, yielding an effect size of ~ 0.5 . This was based on a two-sided t test (P of 0.05 at 80% power).

During statistical analysis, outcome measurements were not available, and thus not included, for a given animal if premature mortality occurred. For example, when analysing parameters of glucose homeostasis derived from GTTs and ITTs, there were five animal mortalities due to euthanasia secondary to severe weight loss ($n = 1$, BO-fed male), fatal fight wounds ($n = 1$, FO-fed male), coronary blood clot ($n = 1$, CO-fed male), atrial

thrombosis ($n = 1$, EO-fed female), and an unknown cause ($n = 1$, CO-fed female). Data points for parameters of glucose homeostasis were also excluded if the animal failed to respond appropriately to the glucose or insulin challenge (e.g. no change in blood glucose or severe hypoglycaemia, respectively). We attributed an occasional lack of response to technical issues with intraperitoneal injection. Exclusion was made without consideration of assigned treatment or sex of the animal.

Results

Feed intake, weight gain and feed efficiency

Diet had no effect on feed intake, animal weight, weight gain, or feed efficiency of mice (Table 2). As expected, animal weight and weight gain were 60 and 110 % greater in males compared with females, respectively, which was also reflected in a 100 % greater feed efficiency in males ($P < 0.001$; Table 3). Animal weight and weight gain increased over time ($P < 0.001$), with a weight gain of 13.3 g by 3 months of experimental feeding and 26.3 g by 12 months of experimental feeding (Table 4). Accordingly, feed efficiency steadily decreased as mice aged ($P < 0.001$; Table 4). An interaction between the main effects of diet and sex on feed intake ($P = 0.004$) and weight gain ($P = 0.02$) of mice revealed that diet influenced these outcomes in a sex-specific manner. For instance, feed intake was 18, 21, and 18 % greater in FO-, BO-, and EO-fed males, respectively, compared with CO-fed males (3.48, 3.57, and 3.48 g *v.* 2.94 g, respectively, $P < 0.05$; Fig. 1(a)). Similarly, weight gain was 34, 52, and 42 % greater in FO-, BO-, and EO-fed males, respectively, compared with CO-fed males (28.7, 32.5, and 30.3 g *v.* 21.4 g, respectively, $P < 0.05$; Fig. 1(b)). Yet, these comparative differences were not observed for feed efficiency (Fig. 1(c)). Of note, we also observed considerable heterogeneity in weight (online Supplementary Fig. 1) and weight gain (Supplementary Fig. 2) in both male and female CD-1[®] mice in response to varying dietary fat quality. In sum, diet influenced feeding behaviour and weight gain differently in males *v.* females.

Parameters of glucose tolerance and insulin sensitivity

The final blood glucose measurement of ITT was 10 % lower in EO-fed mice than in CO-fed mice ($P < 0.05$), an indicator that EO-fed mice exhibited improved peripheral insulin sensitivity (Table 2). In addition, ITT AUC tended to be lower in EO-fed mice (7 %) compared with CO-fed mice ($P = 0.07$). By all metrics used to assess glucose homeostasis in this study, except percent change from baseline in blood glucose during ITT, glucose tolerance and insulin sensitivity were greater in females compared with males, regardless of diet ($P \leq 0.001$; Table 3). Unexpectedly, when examining the effect of time (i.e. age) on parameters of glucose homeostasis, GTT AUC was found to be 21 % lower at 12 months of experimental feeding than at all other time points ($P < 0.05$; Table 4), indicating that glucose tolerance improved with age. To investigate this finding further, we examined the interaction between the effects of diet, sex, and time on GTT AUC ($P = 0.001$) and final blood glucose

measurement of GTT ($P < 0.001$); at 3 months of experimental feeding, FO-, BO-, and EO-fed males had an 81, 90, and 56 % greater GTT AUC, respectively, than CO-fed males (28 565, 30 085, and 24712 *v.* 15816, respectively, $P < 0.05$; Fig. 2(a)). Yet, this phenotype disappeared 3 months later, and by 12 months of experimental feeding, FO- and EO-fed males had a 33 % and 26 % lower GTT AUC, respectively, compared with CO-fed males (14 704 and 16 230 *v.* 22 043, respectively, $P < 0.05$; Fig. 2(a)). Notably, no diet-induced changes over time were detected in females. Similar results were observed for the final blood glucose measurement of GTT (Fig. 2(b)), but no interaction between the effects of diet, sex, and age was found for ITT AUC (Fig. 2(c)). Compared with CO-fed males, EO-fed males had a lower final blood glucose measurement of ITT at 6 months (28 % difference, 64 *v.* 46 mg/dl, respectively, $P < 0.05$) and at 9 months (32 % difference, 76 *v.* 52 mg/dl, respectively, $P < 0.05$) of experimental feeding. However, there were no differences at 12 months of experimental feeding (Fig. 2(d)). Taken together, long-term supplementation of dietary fat with sources of unique FA compositions, particularly fish and echium oil, may beneficially modulate glucose homeostasis in a sex- and age-dependent manner.

Correlations of parameters of glucose homeostasis with tissue fatty acids

Spearman correlations were performed to assess the relationship between parameters of glucose homeostasis at 12 months of experimental feeding and dietary FA content measured in insulin-sensitive tissues at time of harvest (14 months of age), i.e. liver (Fig. 3(a) and 3(b)), muscle (Fig. 3(c) and 3(d)) and adipose tissue (Fig. 3(e) and 3(f)). Due to the pronounced sex differences observed in measurements of metabolic health in mice, all correlations for this study were performed separately for males and females. In liver, muscle, and adipose tissue of males, fish-derived FAs were negatively associated with GTT AUC and final blood glucose measurement of GTT ($\rho = -0.36$ – -0.42 , $P < 0.05$), indicating that fish-derived FAs beneficially modulate glucose tolerance. Similar results were found for muscle tissue, but not liver or adipose tissue, of female mice ($P < 0.05$; Fig. 2(d)). Yet, unexpectedly, the content of fish-derived FAs in adipose tissue of females was positively correlated with ITT AUC ($\rho = 0.37$) and final blood glucose measurement of ITT ($\rho = 0.41$, $P < 0.05$). Overall, diet-derived FAs in tissue may play a role in glucose tolerance and insulin sensitivity; however, this effect may be more pronounced in males than females.

Correlations of parameters of glucose homeostasis with colonic bacterial composition

In males, the abundance of the colonic genera *Clostridium* and *Oscillospira* was negatively correlated with fasting glucose ($\rho = -0.52$ and -0.45 , respectively, $P < 0.05$; Fig. 4(a)). In addition, observed genera (an index of α diversity) were negatively associated with fasted insulin ($\rho = -0.60$) and HOMA-IR ($\rho = -0.56$), while *Akkermansia*, *Bacteroides*, and *Parabacteroides* were all positively associated with AUC GTT ($\rho = 0.44$ – 0.52) in males ($P < 0.05$). In females, abundance of *Ruminococ-*

Table 2. Daily feed intake, feed efficiency, body weight, and parameters of glucose homeostasis of CD-1® mice fed one of four experimental isoenergetic high-fat (40 % fat of total energy) diets

	Diet*												P value						
	CO			FO			BO			EO			D	S	T	D × S	D × T	S × T	D × S × T
	Mean	SEM	n	Mean	SEM	n	Mean	SEM	n	Mean	SEM	n							
Daily feed intake (g)	3.14	0.07	79	3.29	0.07	78	3.23	0.08	79	3.32	0.05	80	–	0.07	<0.001	0.004	<0.001	0.02	0.005
Feed efficiency†	0.007	0.000	78	0.009	0.001	77	0.007	0.001	73	0.008	0.001	78	–	<0.001	<0.001	–	–	<0.001	0.07
Body weight (g)	42.8	1.2	78	49.2	1.8	77	48.0	1.9	79	47.4	1.7	81	0.09	<0.001	<0.001	0.009	0.02	0.004	0.09
Body weight gain (g)	17.4	1.0	78	23.3	1.5	77	21.9	1.5	79	21.3	1.4	81	–	<0.001	<0.001	0.02	0.03	0.004	–
Fed glucose (mg/dl)	98	2	79	104	4	78	102	3	79	104	4	79	–	<0.001	0.02	–	0.009	0.002	0.03
Fed insulin (ng/ml)	3.71	0.52	70	5.72	0.69	69	5.66	0.68	73	6.19	0.72	71	–	<0.001	<0.001	–	–	<0.001	–
Fasted glucose (mg/dl)	88	3	76	84	3	76	91	4	77	84	3	79	–	<0.001	–	–	–	0.09	–
Fasted insulin (ng/ml)	1.29	0.23	57	1.23	0.22	59	1.59	0.32	60	1.85	0.28	58	–	0.001	–	–	–	<0.001	–
HOMA-IR‡	9.57	1.89	57	7.49	1.24	55	11.71	1.89	59	11.76	1.70	57	–	<0.001	–	–	–	<0.001	–
Area under the curve (GTT)	17 696	999	73	17 583	877	74	20 664	1349	75	16 921	847	76	0.09	<0.001	<0.001	–	0.001	0.001	0.001
Blood glucose (120 min of GTT)	104	7	72	104	6	74	122	10	75	97	6	76	–	<0.001	<0.001	–	0.01	0.03	<0.001
Area under the curve (ITT)	3716	193	61	3807	161	58	3843	250	60	3446	148	58	0.07	<0.001	–	0.04	–	0.03	–
% change from baseline (ITT)§	32.5	2.9	61	25.5	2.5	58	29.1	2.8	60	32.0	2.23	58	–	–	–	–	0.06	0.03	–
Blood glucose (60 min of ITT)	48 ^a	3	59	51 ^a	2	56	50 ^a	3	56	43 ^b	2	58	0.03	<0.001	–	0.02	–	0.02	0.009

CO, diet composed of 100 % USA fat blend; FO, diet composed of 70 % USA fat blend and 30 % fish (menhaden) oil; BO, diet composed of 70 % USA fat blend and 30 % dairy (butter oil) fat; EO, diet composed of 70 % USA fat blend and 30 % echium oil. D, diet; S, sex; T, time (month of data collection); GTT, glucose tolerance test; ITT, insulin tolerance test.

* Values are expressed as mean ± standard error of the mean and are collapsed by sex and time. Data were collected at 3, 6, 9, and 12 months of experimental feeding. Means without a common letter differ ($P < 0.05$).

† Feed efficiency = total weight gain (g)/total feed intake (kcal).

‡ HOMA-IR, homeostatic model of assessment of insulin resistance = (glucose_{0 min} × insulin_{0 min})/405.

§ % change from baseline = ((Final blood glucose – initial blood glucose)/initial blood glucose) × 100.

Effects of dietary fat, sex and age on health

Table 3. Daily feed intake, feed efficiency, body weight, and parameters of glucose homeostasis of male and female CD-1® mice

	Sex*						P value						
	Male			Female			D	S	T	D × S	D × T	S × T	D × S × T
	Mean	SEM	n	Mean	SEM	n							
Daily feed intake (g)	3.37	0.04	160	3.12	0.05	156	–	0.07	<0.001	0.004	<0.001	0.02	0.005
Feed efficiency†	0.010 ^a	0.000	153	0.005 ^b	0.000	153	–	<0.001	<0.001	–	–	<0.001	0.07
Body weight (g)	57.6 ^a	0.7	159	35.9 ^b	0.9	156	0.09	<0.001	<0.001	0.009	0.02	0.004	0.09
Body weight gain (g)	28.3 ^a	0.7	159	13.5 ^b	0.9	156	–	<0.001	<0.001	0.02	0.03	0.004	–
Fed glucose (mg/dl)	116 ^a	3	158	88 ^b	1	157	–	<0.001	0.02	–	0.009	0.002	0.03
Fed insulin (ng/ml)	8.65 ^a	0.42	155	1.29 ^b	0.20	128	–	<0.001	<0.001	–	–	<0.001	–
Fasted glucose (mg/dl)	101 ^a	3	157	72 ^b	1	151	–	<0.001	–	–	–	0.09	–
Fasted insulin (ng/ml)	2.20 ^a	0.20	140	0.42 ^b	0.07	94	–	0.001	–	–	–	<0.001	–
HOMA-IR‡	15.47 ^a	1.21	136	2.34 ^b	0.46	92	–	<0.001	–	–	–	<0.001	–
Area under the curve (GTT)	23231 ^a	772	152	12998 ^b	359	146	0.09	<0.001	<0.001	–	0.001	0.001	0.001
Blood glucose (120 min of GTT)	138 ^a	6	151	75 ^b	2	146	–	<0.001	<0.001	–	0.01	0.03	<0.001
Area under the curve (ITT)	4901 ^a	150	94	2918 ^b	71	143	0.07	<0.001	–	0.04	–	0.03	–
% change from baseline (ITT)§	20.4	1.8	94	36.0	1.7	143	–	–	–	–	0.06	0.03	–
Blood glucose (60 min of ITT)	65 ^a	2	94	35 ^b	1	135	0.03	<0.001	–	0.02	–	0.02	0.009

D, diet; S, sex; T, time (month of data collection); GTT, glucose tolerance test; ITT, insulin tolerance test.

* Values are expressed as mean ± standard error of the mean and are collapsed by diet and time. Means without a common letter differ ($P < 0.05$). Data were collected at 3, 6, 9, and 12 months of experimental feeding.

† Feed efficiency = total weight gain (g)/total feed intake (kcal).

‡ HOMA-IR, homeostatic model of assessment of insulin resistance = (glucose_{0 min} × insulin_{0 min})/405.

§ % change from baseline = ((Final blood glucose – initial blood glucose)/initial blood glucose) × 100.

Table 4. Daily feed intake, feed efficiency, body weight, and parameters of glucose homeostasis of CD-1® mice at 3, 6, 9 and 12 months of experimental feeding

	Month*												P value							
	3			6			9			12			D	S	T	D × S	D × T	S × T	D × S × T	
	Mean	SEM	n	Mean	SEM	n	Mean	SEM	n	Mean	SEM	n								
Daily feed intake (g)	3.19 ^{bc}	0.07	81	3.33 ^a	0.07	79	3.32 ^{ab}	0.06	79	3.15 ^c	0.06	77	–	0.07	<0.001	0.004	<0.001	0.02	0.005	
Feed efficiency†	0.011 ^a	0.001	80	0.008 ^b	0.000	77	0.006 ^c	0.000	76	0.005 ^d	0.000	73	–	<0.001	<0.001	–	–	<0.001	0.07	
Body weight (g)	39.1 ^d	1.3	80	46.8 ^c	1.6	80	49.6 ^b	1.7	79	52.3 ^a	1.8	76	0.09	<0.001	<0.001	0.009	0.02	0.004	0.09	
Body weight gain (g)	13.3 ^d	1.0	80	20.9 ^c	1.3	80	23.6 ^b	1.4	79	26.3 ^a	1.5	76	–	<0.001	<0.001	0.02	0.03	0.004	–	
Fed glucose (mg/dl)	99 ^{ab}	2	80	110 ^a	5	79	96 ^b	2	79	103 ^{ab}	3	77	–	<0.001	0.02	–	0.009	0.002	0.03	
Fed insulin (ng/ml)	3.54 ^b	0.44	81	4.66 ^a	0.46	68	8.20 ^a	0.93	76	4.82 ^b	0.46	58	–	<0.001	<0.001	–	–	<0.001	–	
Fasted glucose (mg/dl)	83	3	79	87	4	79	92	4	79	85	2	71	–	<0.001	–	–	–	0.09	–	
Fasted insulin (ng/ml)‡	0.76	0.09	65	1.41	0.23	72	2.03	0.34	64	2.05	0.36	33	–	0.001	–	–	–	<0.001	–	
HOMA-IR‡,§	5.21	0.78	63	9.69	1.56	71	13.19	1.84	63	15.23	2.81	31	–	<0.001	–	–	–	<0.001	–	
Area under the curve (GTT)	18309 ^a	1040	77	20419 ^a	1242	75	18722 ^a	965	76	15210 ^b	788	70	0.09	<0.001	<0.001	–	0.001	0.001	0.001	
Blood glucose (120 min of GTT)	104 ^a	7	77	122 ^a	10	74	108 ^a	7	76	94 ^b	5	70	–	<0.001	<0.001	–	0.01	0.03	<0.001	
Area under the curve (ITT)	3753	243	58	3437	170	57	3796	156	62	3816	197	60	0.07	<0.001	–	0.04	–	0.03	–	
% change from baseline (ITT)	31.6	3.0	58	33.3	2.3	57	28.4	2.7	62	26.1	2.5	60	–	–	–	–	0.06	0.03	–	
Blood glucose (60 min of ITT)	48	3	55	44	3	57	50	2	60	49	3	57	0.03	<0.001	–	0.02	–	0.02	0.009	

D, diet; S, sex; T, time (month of data collection); GTT, glucose tolerance test; ITT, insulin tolerance test.

* Values are expressed as mean ± standard error of the mean and are collapsed by diet and sex. Means without a common letter differ ($P < 0.05$).

† Feed efficiency = total weight gain (g)/total feed intake (kcal).

‡ Results of statistical analysis similar when data from month 12 is excluded.

§ HOMA-IR, homeostatic model of assessment of insulin resistance = (glucose_{0 min} × insulin_{0 min})/405.

|| % change from baseline = ((Final blood glucose – initial blood glucose)/initial blood glucose) × 100.

Effects of dietary fat, sex and age on health

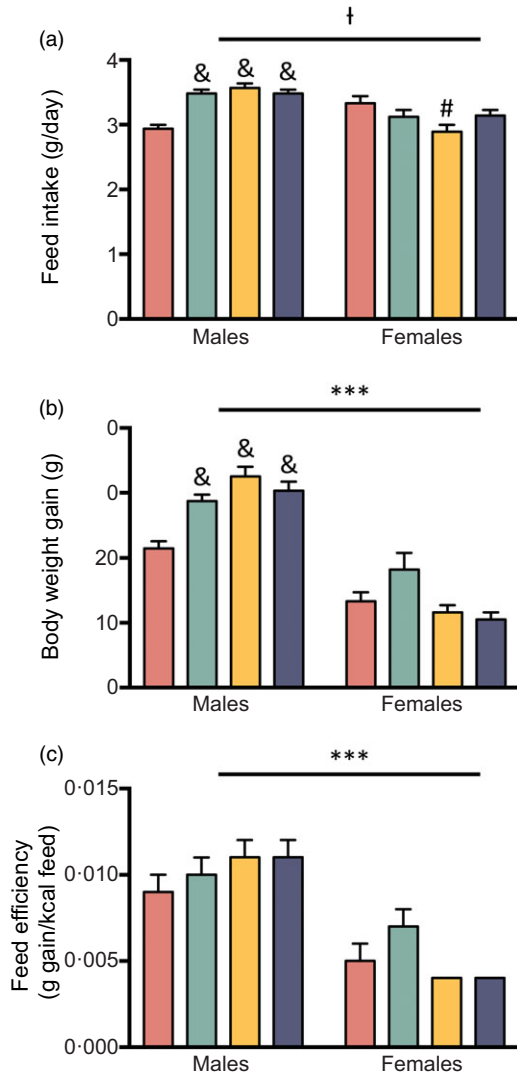


Fig. 1. Feed intake (a), body weight gain (b), and feed efficiency (c) of male and female of CD-1[®] mice fed one of four experimental isoenergetic high-fat (40 % fat of total energy) diets with fat content consisting of either 100 % USA fat blend (CO) or 70 % USA fat blend and 30 % fish oil (FO), dairy fat (BO), or echium oil (EO), respectively. Values are expressed as mean \pm standard error of the mean from data collected at 3, 6, 9, and 12 months of experimental feeding. † = $P = 0.07$, males v. females collapsed by diet and time. *** $P < 0.001$, males v. females collapsed by diet and time. & = $P < 0.05$, FO-, BO-, or EO-fed males v. CO-fed males collapsed by time. # = $P < 0.05$, BO-fed females v. CO-fed females collapsed by time. Data were analysed via linear mixed model with an unstructured covariance, specifying diet, sex, and time as fixed effects and body weight as covariate when appropriate. ■, CO; ■, FO; ■, BO; ■, EO.

cus and *Turicibacter* was negatively correlated with fed glucose and fed insulin, respectively ($\rho = -0.55$ and -0.56 , respectively, $P < 0.05$; (Fig. 4 (b)). Abundance of *Lachnospirillum*, *Parabacteroides*, and *Roseburia* and observed genera were positively associated with final blood glucose measurement of ITT in females ($\rho = 0.52$ – 0.67 , $P < 0.05$). Furthermore, *Eubacterium* positively correlated with fed insulin in both male and female mice ($\rho = 0.45$ and 0.67 , respectively, $P < 0.05$). Collectively, abundance and diversity of specific colonic bacteria were associated with metabolic health in a sex-specific fashion.

Assessment of plasma estradiol concentration

To evaluate whether females were reproductively senescent at the end of the study, we compared circulating E2 concentrations in a subset of female mice at a time of sexual maturity and again when aged (i.e., 5 and 14 months of age, respectively). As expected, plasma E2 concentrations were lower as female mice aged ($P < 0.01$; online Supplementary Fig. 3).

Discussion

Ageing is a significant risk factor in the development of metabolic diseases such as T2D, yet, there is a lack of effective lifestyle prevention strategies to protect against age-related metabolic impairments. Research shows that diet, and specifically dietary fat quality, can significantly influence glucose homeostasis, however, whether or not life-long consumption of beneficial FAs can attenuate glucose intolerance and insulin resistance during ageing is not clear. Our study therefore aimed to evaluate whether long-term supplementation of a diet with fish oil, dairy fat, or echium oil attenuates age-related glucose intolerance and insulin resistance induced by consumption of a FA composition mixture representative of the average U.S. American diet in out-bred mice.

A principal finding of our work is the prominent sexual divergence in the metabolic response of mice to diets varying in fat quality. Diet *per se* had a minimal impact on parameters of glucose homeostasis, but when males and females were assessed separately, we observed that dietary fat quality influenced glucose tolerance notably in males but not females. In particular, consumption of echium oil or fish oil had a protective effect against deterioration of glucose tolerance in males. Echium oil is a seed oil that may have unique health properties due to its high content of PUFAs, particularly α -linolenic acid and stearidonic acid (both *n*-3 FAs), as well as γ -linolenic acid (*n*-6 FA)⁽³¹⁾. Studies in rodents^(42–44) and humans^(45,46) indicate that the primary physiologic benefits of echium oil are its lowering effect on cholesterol and/or triacylglycerol blood concentrations, likely by modulating transcription of hepatic genes related to lipogenesis^(42,43) and inflammation⁽⁴³⁾. Of note, Kavanagh *et al.*⁽³⁰⁾ showed that dietary echium oil promoted glucose disposal in aged, hyperglycaemic and hyperinsulinaemic monkeys, which aligns with our work showing that EO-fed mice, regardless of sex, lowered blood glucose following an insulin bolus. We and others have shown that consumption of echium oil-derived stearidonic acid results in an increased tissue content of downstream long-chain *n*-3 FAs (e.g. EPA)^(31,47), which are known to impart health benefits by modulating atherogenic risk factors^(19,48–50). Additionally, research also suggests that γ -linolenic acid is another FA constituent in echium oil that favourably influences lipid metabolism⁽⁴³⁾. Fish, fish oil, and *n*-3 FA supplements have been extensively studied as nutritional interventions to protect against metabolic diseases^(19,48–50). In rodents, research suggests that supplementation of marine oils rich in long-chain *n*-3 FAs to a high-fat diet (45–60 % of total energy) can protect against cardiometabolic risk factors^(51–53). Loehfelm *et al.*⁽⁵³⁾ showed that a high-fat (60 % of total energy) diet enriched with mussel oil prevented diet-induced weight gain

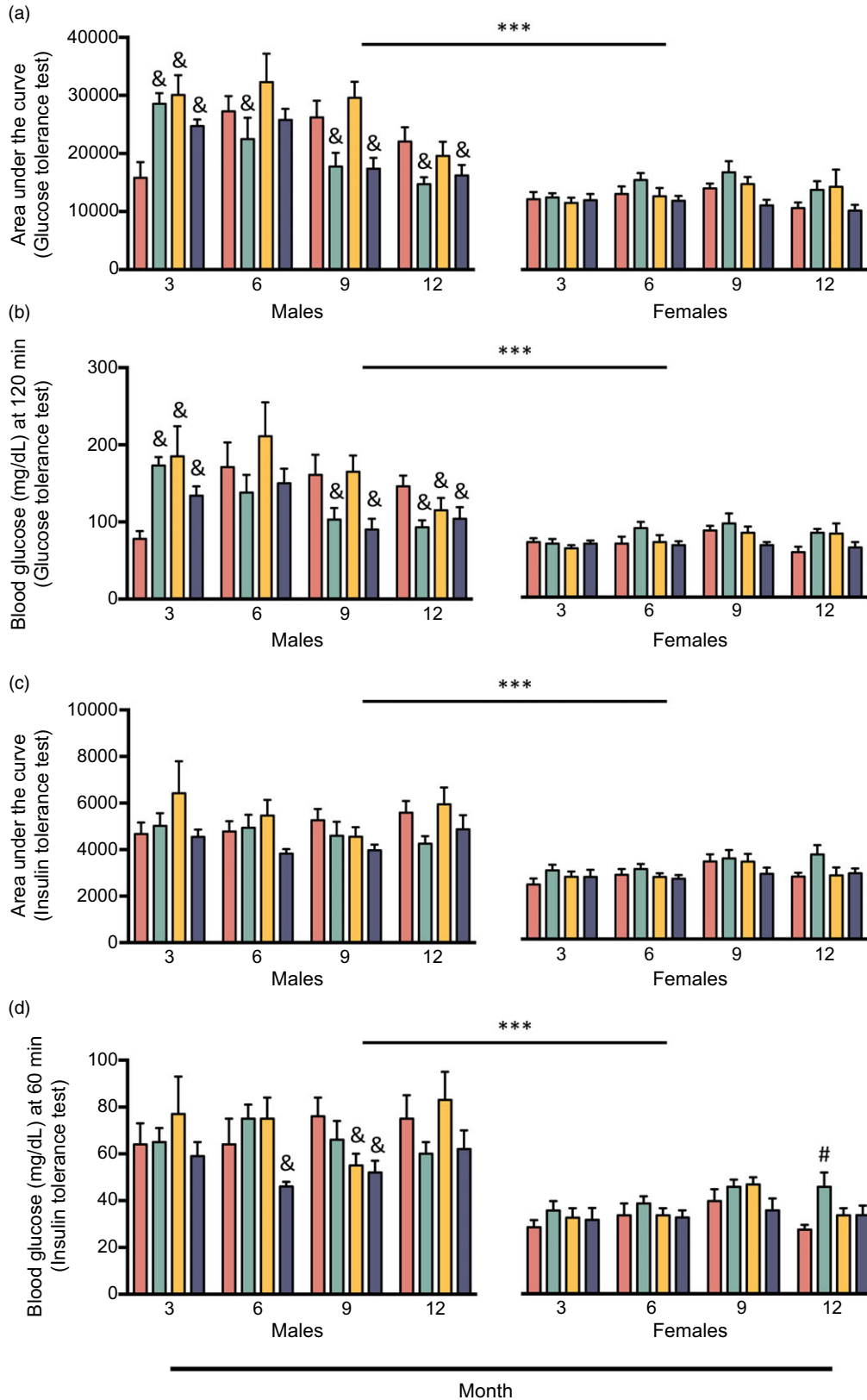


Fig. 2. Area under the curve of glucose tolerance test (a), final blood glucose measurement of glucose tolerance test (b), area under the curve of insulin tolerance test (c), and final blood glucose measurement of insulin tolerance test (d) of male and female CD-1[®] mice at 3, 6, 9, and 12 months of feeding one of four experimental isoenergetic high-fat (40 % fat of total energy) diets with fat content consisting of either 100 % USA fat blend (CO) or 70 % USA fat blend and 30 % fish oil (FO), dairy fat (BO), or echium oil (EO), respectively. Values are expressed as mean \pm standard error of the mean. *** $P < 0.001$, males v. females collapsed by diet and time. & = $P < 0.05$, FO-, BO-, or EO-fed males v. CO-fed males within each respective month. # = $P < 0.05$, FO-fed females v. CO-fed females within the respective month. Data were analysed via linear mixed model with an unstructured covariance, specifying diet, sex, and time as fixed effects and body weight as covariate when appropriate. ■, CO; ■, FO; ■, BO; ■, EO.

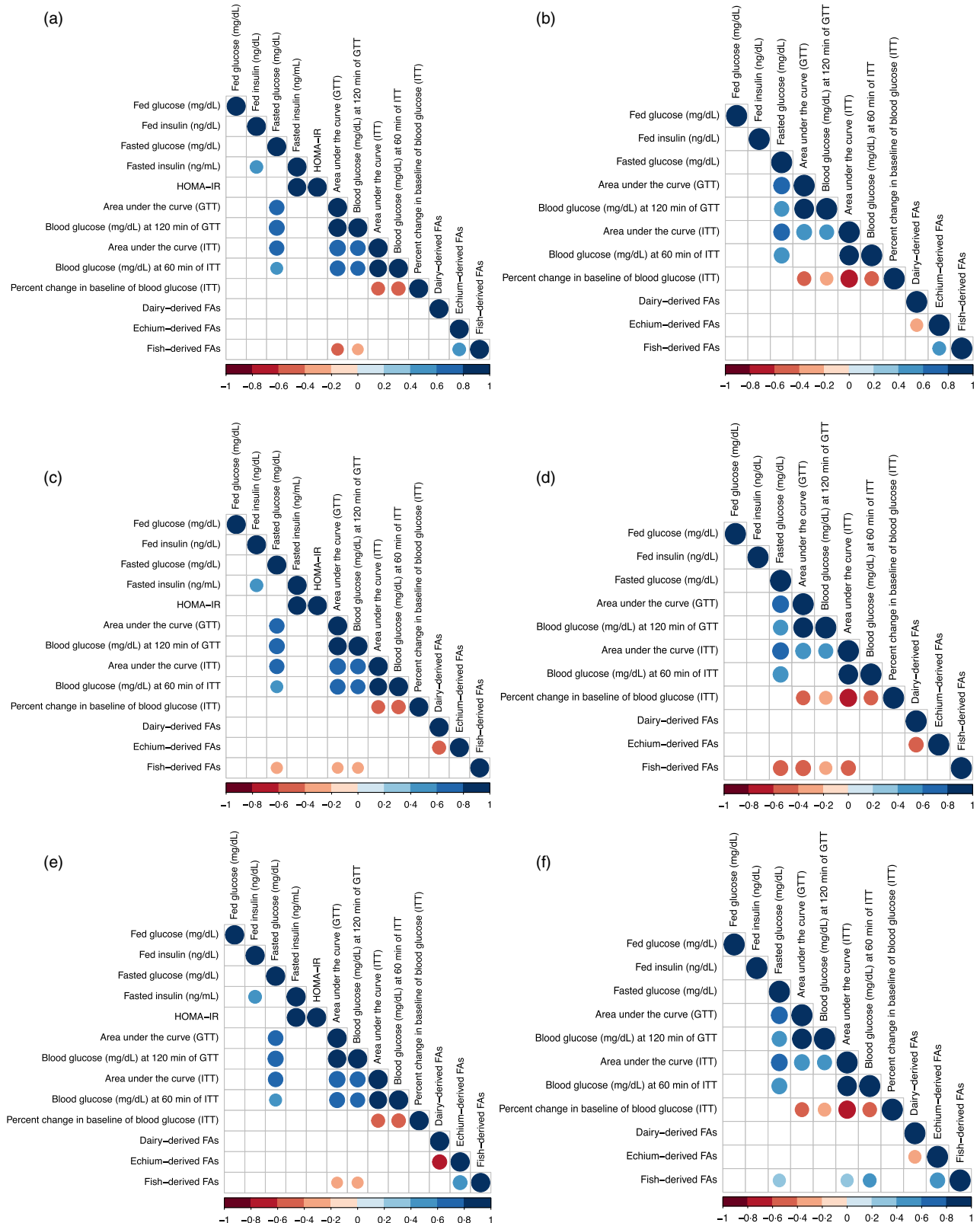


Fig. 3. Spearman correlation matrices between metabolic parameters determined at 12 months of experimental feeding and fatty acids measured in liver tissue of male (a), liver tissue of female (b), muscle tissue of male (c), muscle tissue of female (d), adipose tissue of male (e), and adipose tissue of female (f) CD-1[®] mice at harvest (14 months of age). A positive correlation (closer to 1) is signified by a darker shade of blue; a negative correlation (closer to -1) is signified by a darker shade of red ($P < 0.05$). HOMA-IR, homeostatic model of assessment of insulin resistance = $(\text{glucose}_{0 \text{ min}} \times \text{insulin}_{0 \text{ min}}) / 405$. GTT, glucose tolerance test. ITT, insulin tolerance test. Percent change in baseline = $(\text{Final blood glucose} - \text{initial blood glucose}) / \text{initial blood glucose} \times 100$. FAs, fatty acids. Dairy-derived FAs include the sum of 15:0, 16:1 n-7, 17:0, 18:1 n-7, and 18:1 n-9, n-11. Echium-derived FAs include the sum of 18:3 n-6, c6, c9, c12 (n-6), 18:3 n-3, c9, c12, c15 (n-3), and 18:4 n-3, c6, c9, c12, c15 (n-3). Fish-derived FAs include the sum of 20:5 n-3, c5, c8, c11, c14, c17 (n-3), 22:5 n-3, c7, c10, c13, c16, c19 (n-3), and 22:6 n-3, c4, c7, c10, c13, c16, c19 (n-3). Variables of fasted insulin and HOMA-IR were not included in analysis with females due to the low number of observations.

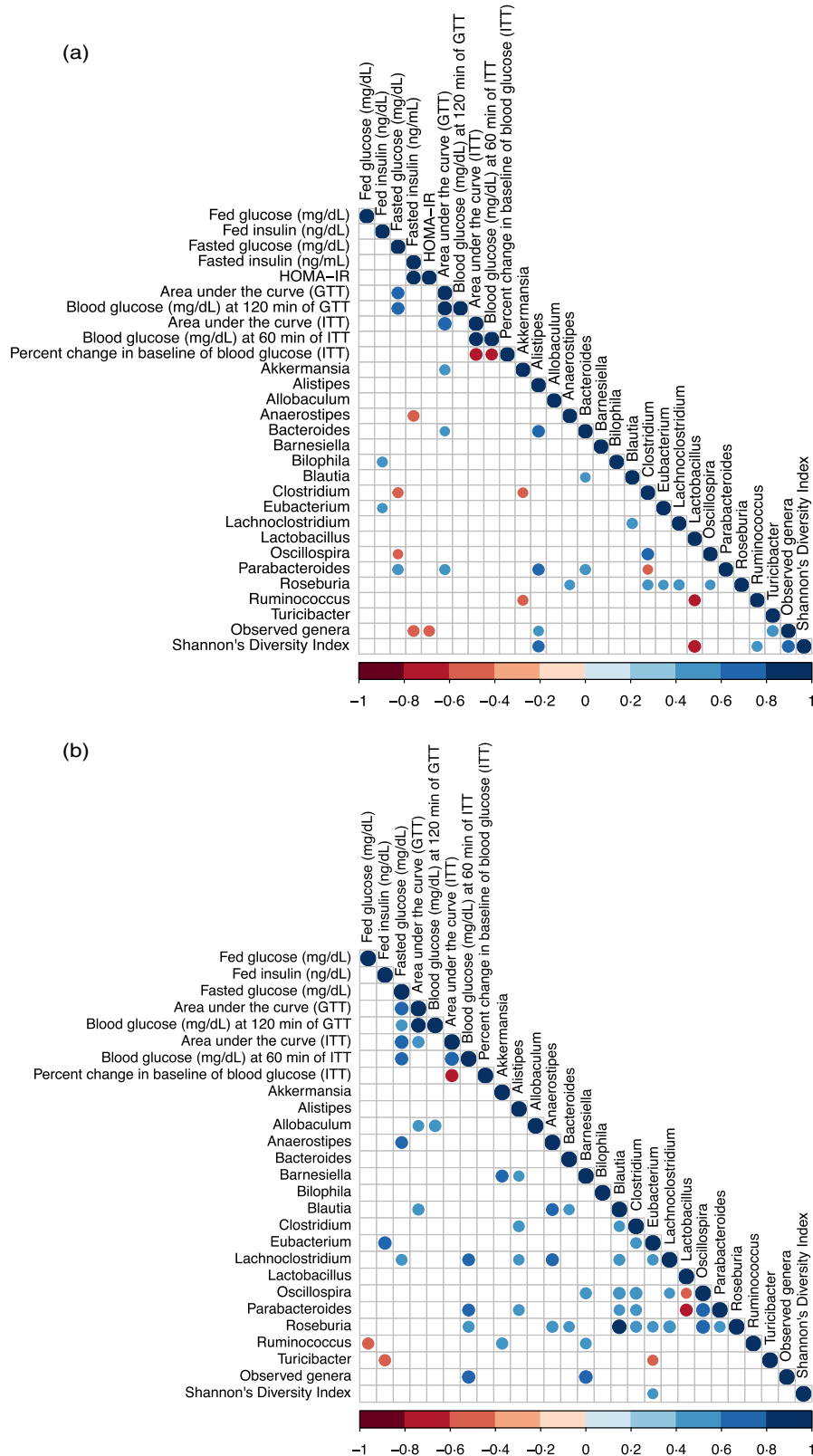


Fig. 4. Spearman correlation matrices between metabolic parameters determined at 12 months of experimental feeding, diversity indices of colonic bacteria measured before harvest (13.5 months of age), and abundance by counts of colonic bacterial genera (mean relative abundance > 1%) measured before harvest (13.5 months of age) of male (a) and female (b) CD-1[®] mice. A positive correlation (closer to 1) is signified by a darker shade of blue; a negative correlation (closer to -1) is signified by a darker shade of red ($P < 0.05$). HOMA-IR, homoeostatic model of assessment of insulin resistance = $(\text{glucose}_{0 \text{ min}} \times \text{insulin}_{0 \text{ min}}) / 405$. GTT, glucose tolerance test. ITT, insulin tolerance test. Percent change in baseline = $((\text{Final blood glucose} - \text{initial blood glucose}) / \text{initial blood glucose}) \times 100$. Variables of fasted insulin and HOMA-IR were not included in analysis with females due to the low number of observations.

in young mice (~10–12 weeks of age) after 1 week of feeding, and either attenuated weight-gain or decreased weight in lean or obese, respectively, aged mice (~12 months of age) after 4 weeks of feeding. However, observational studies in humans assessing the effect of fish oil consumption on T2D risk are highly controversial⁽¹⁸⁾. For example, research has generally observed that fish oil intake is associated with an increased risk of T2D in USA populations⁽¹⁸⁾. Here, we demonstrate via a controlled experiment that fish oil supplementation to a base diet with a FA composition reflecting the average U.S. American FA composition can improve metabolic predictors of T2D risk in males.

Males have a higher prevalence of T2D than females⁽¹⁰⁾; however, the differential influence of dietary fat quality on T2D risk by sex is not well characterised. Studies assessing dietary FA content and composition and metabolic health outcomes using animal models have often used only males or females^(43,44,51,54), while human trials may not explicitly test the effect of sex^(45,46,55,56) (e.g. sex included as a covariate in statistical analysis). In our study, sex was the most consistent effect on parameters of glucose tolerance and insulin sensitivity, even when considering known risk factors of diet and age. While previous work underscores that sex hormones play an important role in T2D risk^(57–60), the biological mechanisms underlying the observed differences in metabolic health of male and female mice in our work are not clear. Although mice are not considered elderly until 18 months of age, reproductive senescence begins between the age of 10 and 15 months⁽⁶¹⁾, and as a result estrogen levels in females begin to decline⁽⁶²⁾. Similarly, our work shows that E2 concentrations had declined in aged female mice (i.e. 14 months of age) compared with female mice during sexual maturity. Nevertheless, for the duration of the study and regardless of age, females surprisingly remained more protected against T2D risk than males. Therefore, the significance of circulating sex hormones in the context of our study, particularly as mice aged, should be carefully considered. To fully understand the implications of our work, more long-term studies are warranted, particularly focused on the intersection of dietary fat quality, sex and sex hormones, and T2D risk in a frank elderly population.

We also observed several sex-specific correlations between metabolic parameters and FA content in tissues. Overall, this was expected, as we have observed in this cohort of mice that the distribution of unique FAs derived from fish, dairy, and echium in tissues differed between males and females (these results have been discussed extensively previously⁽³¹⁾). Notably, we found that the content of fish-derived FAs (i.e. EPA, docosapentaenoic acid, and DHA) in liver, muscle, and adipose tissue correlated with improved glucose tolerance in males, with similar results found in females but in skeletal muscle only. This points to a relationship between diet-derived FAs from fish oil and from echium oil, as we have shown previously in this cohort of mice that echium oil consumption increases tissue content of long-chain *n*-3 FAs⁽³¹⁾. These results align with the current consensus that fish-derived *n*-3 FAs are potent regulators of tissue-specific gene expression and function (as reviewed in⁽⁶³⁾). For example, EPA and DHA can beneficially regulate membrane fluidity and signalling pathways when incorporated into the

membrane phospholipids, as well as promote homeostasis by modulating gene expression for lipid metabolism and inflammation. In this study, no correlations between the content of diet-derived FAs in liver tissue and measurements of glucose tolerance or insulin sensitivity were observed, while fish-derived FA content in adipose tissue was associated with indices of insulin resistance, in female mice. However, the physiologic relevance of these findings is ambiguous, as females were at very low risk of metabolic abnormalities.

No correlations between dairy-derived FAs in tissues and metabolic outcomes were found. While this is likely explained by the modest metabolic improvements exhibited in mice supplemented with dairy fat, it is striking that BO-fed males were significantly less glucose intolerant than CO-fed males after 12 months of experimental feeding. It is challenging to contextualise that these results via other rodent studies, as dietary supplementation with dairy fat in other work has shown mixed results on metabolic parameters^(54,64–66) that are likely more reflective of the respective study designs (e.g. animal model, diet formulation). For example, when male C57BL/6 mice were fed a low-fat chow or a high-fat diet (10 and 37.5% of total energy, respectively) with fat composed of milk fat, lard, or safflower oil, modest changes in markers of inflammation in adipose tissue were found in mice fed milk fat, whereas markers of inflammation in adipose tissue were significantly enhanced in mice fed safflower oil⁽⁵⁴⁾. An important consideration, however, is that the dietary fat of these murine diets was derived from a single fat source (i.e. 100% milk fat). In general, our results more closely support several epidemiological studies showing that dairy fat intake is negatively associated with T2D^(67,68) and related metabolic impairments^(23,25), particularly in elderly populations⁽⁶⁹⁾. One possible explanation for this is that the preparation of our murine diets reflects the macronutrient profile (in percent energy) and the FA composition of an U.S. American diet, which has been suggested to lead to the most accurate results in terms of mimicking human physiology⁽⁷⁰⁾. Yet, it is important to note that murine and epidemiological studies cannot completely account for all exposures (or lifestyle factors) occurring simultaneously in the complex lives of humans; thus, more research is needed to understand the physiologic effects of dairy-derived FAs on health.

Gut microbiota are increasingly recognised as a critical factor for health maintenance and disease pathogenesis^(71–75), and studies have established that diet is integral to this relationship^(51,54,76). Our study suggests that colonic bacterial genera may differentially impact metabolic outcomes in mice after feeding a controlled diet long-term. For example, an index of α diversity (i.e. observed genera) was associated with measurements of insulin sensitivity (in males only), consistent with previous studies^(77–79). Unexpectedly, however, we found a positive association between the abundance of *Akkermansia* and glucose intolerance (i.e. AUC GTT) in male mice and the abundance of *Eubacterium* and insulin resistance in both male and female mice. *Akkermansia muciniphila*, a recently discovered genus and species⁽⁸⁰⁾, is a mucin-degrading bacterium largely observed as beneficial to systemic metabolism in rodent^(81,82) and human studies^(73,83,84). Likewise, certain



Eubacterium spp. are gaining recognition as potential targets for microbiome-centric therapeutic strategies to improve health in humans⁽⁸⁵⁾. In particular, daily oral dosing of *Eubacterium ballii* for 4 weeks has been reported to improve markers of insulin sensitivity in obese, diabetic *db/db* mice⁽⁸⁶⁾. While these relationships between gut bacteria and metabolic risk factors observed in our work are surprising, it is noteworthy that the field of gut microbiota and human disease is still in its infancy. For example, *Eubacterium* is a very diverse genus both in its phylogeny and functional characteristics, and consensus of the definition of the *Eubacterium* genus is ongoing⁽⁸⁵⁾. Moreover, a survey of the literature establishes that the association of bacterial taxa and host phenotype varies widely across studies^(87,88) and that more controlled trials are needed to fully define the function of specific bacteria in isolation and as part of the collective microbial population. To that end, future research utilising metagenomics and metatranscriptomics to evaluate functional pathways of gut bacteria may resolve the direct or indirect role of gut bacterial composition and their derived metabolites on the net effectiveness of specific diets to mitigate disease risk.

Mice are an adequate but imperfect model organism to study the effectiveness of nutritional interventions on disease risk in humans, as the murine gastrointestinal tract has certain differences when compared with that of humans⁽⁸⁹⁾. Yet, an advantage of our experimental approach was the utilisation of an outbred mouse stock to mimic the genetic heterogeneity of the human population. In addition, by using a relatively short-lived mouse model, we were capable of examining diet effects over several different life stages. However, due to progressive morbidity in our study population, we did not study the effects of dietary fat quality in a frank elderly population; hence, more work is needed to contextualise our research in a human population over 65 years of age. Importantly, our work is limited in its characterisation of the onset of reproductive senescence in female mice during the study, which was assessed at only two timepoints. In addition, estrous cycle was not monitored or coordinated by research personnel, as the significant amount of time and labour required in such a large cohort and long duration of our study (~14 months) is not practical. Nevertheless, our study demonstrates the need for future studies evaluating differences in males and females to focus on the role of sex hormones in glucose tolerance and insulin sensitivity in response to dietary interventions, both during sexual maturity and throughout ageing and reproductive quiescence. Lastly, since dietary sources of unique FAs were incorporated with a diet typical of an U.S. American diet, our results may not be generalisable to other populations with distinct, complex, and varying dietary patterns.

Our work demonstrates that long-term supplementation of dietary fat with sources of unique FAs, notably fish oil and echium oil, can beneficially modulate glucose homeostasis during ageing in a sex-dependent manner. Our study sheds light on the sex-specific role of tissue FA content and colonic bacterial composition on metabolic health. Overall, our findings underscore the importance of sex in formulating future dietary recommendations to mitigate development of metabolic diseases such

as T2D risk through adulthood. Moving forward, clinical trials examining the metabolic impact of long-term dietary fat quality in males and females across different age groups are needed to validate our findings.

Acknowledgements

The authors would like to thank Keara McElroy-Yaggy for assistance during sample collection and animal procedures.

This work was supported by an Armin Grams Memorial Research Award (J.K. and T.L.J., <http://med.uvm.edu/>) from the Center on Aging at the UVM Robert Larner, M.D. College of Medicine and a USDA-NIFA Hatch Fund (J.K., accession number: 1006628; <https://nifa.usda.gov/>). The aforementioned funders had no role in the design, analysis, or writing of this article.

The authors' responsibilities were as follows – T. L. J. and J. K.: formulated the research questions and designed research project; T. L. J., J. K., and A. L. U.: conducted research; A. L. U.: analysed data; T. L. J., J. K., and A. L. U.: interpreted the findings; T. L. J., J. K., and A. L. U.: wrote paper. All authors read and approved the final manuscript.

There are no conflicts of interest.

Supplementary material

For supplementary material accompanying this paper visit <https://doi.org/10.1017/S0007114521004001>

References

1. Roberts AW, Ogunwole SU, Blakeslee L, *et al.* (2018) The Population 65 Years and Older in the United States: 2016 American Community Survey Reports. <https://www.census.gov/content/dam/Census/library/publications/2018/acs/ACS-38.pdf> (accessed October 2020).
2. He W, Goodkind D & Kowal P (2016) International Population Reports, P95/16–1, an Aging World: 2015. U.S. Gov. Publ. Off. <https://www.census.gov/content/dam/Census/library/publications/2016/demo/p95-16-1.pdf> (accessed October 2020).
3. USA Census Bureau (2018) The population 65 years and older in the United States. <https://www.census.gov/library/publications/2018/acs/acs-38.html> (accessed October 2020).
4. De Tata V (2014) Age-related impairment of pancreatic beta-cell function: pathophysiological and cellular mechanisms. *Front Endocrinol* **5**, 138.
5. Jackson RA (1990) Mechanisms of age-related glucose intolerance. *Diabetes Care* **13**, 9–19.
6. Enge M, Arda HE, Mignardi M, *et al.* (2017) Single-cell analysis of human pancreas reveals transcriptional signatures of aging and somatic mutation patterns. *Cell* **171**, 321–330.e14.
7. Palmer AK, Gustafson B, Kirkland JL, *et al.* (2019) Cellular senescence: at the nexus between ageing and diabetes. *Diabetologia* **62**, 1835–1841.
8. CDC Diabetes Risk Factors (2021) <https://www.cdc.gov/diabetes/basics/risk-factors.html> (accessed October 2020).
9. International Diabetes Federation Type 2 Diabetes (2020) <https://www.idf.org/aboutdiabetes/type-2-diabetes> (accessed October 2020).
10. Saeedi P, Petersohn I, Salpea P, *et al.* (2019) Global and regional diabetes prevalence estimates for 2019 and projections

- for 2030 and 2045: results from the International Diabetes Federation Diabetes Atlas. *Diabetes Res Clin Pract* **157**, 107843.
11. Eddy DM, Schlessinger L & Kahn R (2005) Clinical outcomes and cost-effectiveness of strategies for managing people at high risk for diabetes. *Ann Intern Med* **143**, 251–264.
 12. Katula JA, Vitolins MZ, Morgan TM, *et al.* (2013) The healthy living partnerships to prevent diabetes study: 2-year outcomes of a randomized controlled trial. *Am J Prev Med* **44**, S324.
 13. Costa B, Barrio F, Cabré JJ, *et al.* (2012) Delaying progression to type 2 diabetes among high-risk Spanish individuals is feasible in real-life primary healthcare settings using intensive lifestyle intervention. *Diabetologia* **55**, 1319–1328.
 14. Ramachandran A, Snehalatha C, Mary S, *et al.* (2006) The Indian Diabetes Prevention Programme shows that lifestyle modification and metformin prevent type 2 diabetes in Asian Indian subjects with impaired glucose tolerance (IDPP-1). *Diabetologia* **49**, 289–297.
 15. Knowler WC, Barrett-Connor E, Fowler SE, *et al.* (2002) Reduction in the incidence of type 2 diabetes with lifestyle intervention or metformin. *N Engl J Med* **346**, 393–403.
 16. Andersen CJ & Fernandez ML (2013) Dietary strategies to reduce metabolic syndrome. *Rev Endocr Metab Disord* **14**, 241–254.
 17. Jump DB (2002) The biochemistry of *n*-3 polyunsaturated fatty acids. *J Biol Chem* **277**, 8755–8758.
 18. Wallin A, Di Giuseppe D, Orsini N, *et al.* (2012) Fish consumption, dietary long-chain *n*-3 fatty acids, and risk of type 2 diabetes: Systematic review and meta-analysis of prospective studies. *Diabetes Care* **35**, 918–929.
 19. Natto ZS, Yaghmour W, Alshaeri HK, *et al.* (2019) *n*-3 fatty acids effects on inflammatory biomarkers and lipid profiles among diabetic and cardiovascular disease patients: a systematic review and meta-analysis. *Sci Rep* **9**, 1–10.
 20. Månsson HL (2008) Fatty acids in bovine milk fat. *Food Nutr Res* **52**, 1821.
 21. Unger AL, Bourne DE, Walsh H, *et al.* (2020) Fatty acid content of retail cow's milk in the Northeastern USA—what's in it for the consumer? *J Agric Food Chem* **68**, 4268–4276.
 22. Mozaffarian D, De Oliveira Otto MC, Lemaitre RN, *et al.* (2013) Trans-palmitoleic acid, other dairy fat biomarkers, and incident diabetes: the multi-ethnic study of atherosclerosis (MESA). *Am J Clin Nutr* **97**, 854–861.
 23. Kratz M, Marcovina S, Nelson JE, *et al.* (2014) Dairy fat intake is associated with glucose tolerance, hepatic and systemic insulin sensitivity, and liver fat but not β -cell function in humans. *Am J Clin Nutr* **99**, 1385–1396.
 24. Ardisson Korat A V, Li Y, Sacks F, *et al.* (2019) Dairy fat intake and risk of type 2 diabetes in 3 cohorts of USA men and women. *Am J Clin Nutr* **110**, 1192–1200.
 25. Santaren ID, Watkins SM, Liese AD, *et al.* (2014) Serum penta-decanoic acid (15:0), a short-term marker of dairy food intake, is inversely associated with incident type 2 diabetes and its underlying disorders. *Am J Clin Nutr* **100**, 1532–1540.
 26. Mahendran Y, Ågren J, Uusitupa M, *et al.* (2014) Association of erythrocyte membrane fatty acids with changes in glycemia and risk of type 2 diabetes. *Am J Clin Nutr* **99**, 79–85.
 27. Lee TC, Ivester P, Hester AG, *et al.* (2014) The impact of polyunsaturated fatty acid-based dietary supplements on disease biomarkers in a metabolic syndrome/diabetes population. *Lipids Health Dis* **13**, 1–11.
 28. Heine RJ, Mulder C, Popp-Snijders C, *et al.* (1989) Linoleic-acid-enriched diet: long-term effects on serum lipoprotein and apolipoprotein concentrations in insulin sensitivity in noninsulin-dependent diabetic patients. *Am J Clin Nutr* **49**, 448–456.
 29. Wanders AJ, Blom WAM, Zock PL, *et al.* (2019) Plant-derived polyunsaturated fatty acids and markers of glucose metabolism and insulin resistance: a meta-analysis of randomized controlled feeding trials. *BMJ Open Diabetes Res Care* **7**, 585.
 30. Kavanagh K, Flynn DM, Jenkins KA, *et al.* (2013) Stearidonic and γ -linolenic acids in echium oil improves glucose disposal in insulin resistant monkeys. *Prostaglandins Leukot Essent Fat Acids* **89**, 39–45.
 31. Unger A, Jetton T & Kraft J (2021) Tissue and circulating fatty acids as biomarkers to evaluate long-term fat intake are tissue and sex dependent in CD-1 mice. *J Nutr* **151**, 1779–1790.
 32. Johnson MM, Swan DD, Surette ME, *et al.* (1997) Dietary supplementation with γ -linolenic acid alters fatty acid content and eicosanoid production in healthy humans. *J Nutr* **127**, 1435–1444.
 33. Charles River CD-1® IGS Mouse (2021) <https://www.criver.com/products-services/find-model/cd-1r-igs-mouse?region=3611> (accessed September 2020).
 34. Maita K, Hirano M, Harada T, *et al.* (1988) Mortality, major cause of morbidity, and spontaneous tumors in CD-1 mice. *Toxicol Pathol* **16**, 340–349.
 35. Kilkeny C, Browne W, Cuthill I, *et al.* (2010) Improving bio-science research reporting: the ARRIVE guidelines for reporting animal research. *J Pharmacol Pharmacother* **1**, 94.
 36. US Department of Health and Human Services 2015–2020 Dietary Guidelines (2015) [health.gov.https://health.gov/dietaryguidelines/2015/guidelines/](https://health.gov/dietaryguidelines/2015/guidelines/) (accessed October 2019).
 37. Ervin RB, Wright JD, Wang C-Y, *et al.* (2004) Dietary intake of fats and fatty acids for the USA population: 1999–2000. *Adv Data* **348**, 1–6.
 38. Unger AL, Eckstrom K, Jetton TL, *et al.* (2019) Colonic bacterial composition is sex-specific in aged CD-1 mice fed diets varying in fat quality. *PLOS ONE* **14**, e0226635.
 39. Unger AL, Eckstrom K, Jetton TL, *et al.* (2020) Facility-dependent metabolic phenotype and gut bacterial composition in CD-1 mice from a single vendor: a brief report. *PLOS ONE* **15**, e0238893.
 40. IACUC Sample Size Calculations. <https://www.bu.edu/researchsupport/compliance/animal-care/working-with-animals/research/sample-size-calculations-iacuc/> (accessed April 2020).
 41. Gupta D, Jetton TL, LaRock K, *et al.* (2017) Temporal characterization of β cell-adaptive and -maladaptive mechanisms during chronic high-fat feeding in C57BL/6NTac mice. *J Biol Chem* **292**, 12449–12459.
 42. Zhang P, Boudyguina E, Wilson MD, *et al.* (2008) Echium oil reduces plasma lipids and hepatic lipogenic gene expression in apoB100-only LDL receptor knockout mice. *J Nutr Biochem* **19**, 655–663.
 43. Shewale S V, Boudyguina E, Zhu X, *et al.* (2015) Botanical oils enriched in *n*-6 and *n*-3 FADS2 products are equally effective in preventing atherosclerosis and fatty liver. *J Lipid Res* **56**, 1191–1205.
 44. Forrest LM, Boudyguina E, Wilson MD, *et al.* (2012) Echium oil reduces atherosclerosis in apoB100-only LDLrKO mice. *Atherosclerosis* **220**, 118–121.
 45. Surette ME, Edens M, Chilton FH, *et al.* (2004) Dietary Echium oil increases plasma and neutrophil long-chain (*n*-3) fatty acids and lowers serum triacylglycerols in hypertriglyceridemic humans. *J Nutr* **134**, 1406–1411.
 46. Kuhn K, Fuhrmann C, Köhler M, *et al.* (2014) Dietary echium oil increases long-chain *n*-3 PUFAs, including docosapentaenoic acid, in blood fractions and alters biochemical markers for cardiovascular disease independently of age, sex, and metabolic syndrome. *J Nutr* **144**, 447–460.



47. Kuhnt K, Weiß S, Kiehntopf M, *et al.* (2016) Consumption of echium oil increases EPA and DPA in blood fractions more efficiently compared to linseed oil in humans. *Lipid Health Dis* **15**, 1–11.
48. Harris WS (1996) *n*-3 Fatty acids and lipoproteins: comparison of results from human and animal studies. *Lipids* **31**, 243–252.
49. Kabir M, Skurnik G, Naour N, *et al.* (2007) Treatment for 2 mo with *n*-3 polyunsaturated fatty acids reduces adiposity and some atherogenic factors but does not improve insulin sensitivity in women with type 2 diabetes: a randomized controlled study. *Am J Clin Nutr* **86**, 1670–1679.
50. De Luis DA, Conde R, Aller R, *et al.* (2009) Effect of *n*-3 fatty acids on cardiovascular risk factors in patients with type 2 diabetes mellitus and hypertriglyceridemia: an open study. *Eur Rev Med Pharmacol Sci* **13**, 51–55.
51. Caesar R, Tremaroli V, Kovatcheva-Datchary P, *et al.* (2015) Crosstalk between gut microbiota and dietary lipids aggravates WAT inflammation through TLR signaling. *Cell Metab* **22**, 658–668.
52. Cui C, Li Y, Gao H, *et al.* (2017) Modulation of the gut microbiota by the mixture of fish oil and krill oil in high-fat diet-induced obesity mice. *PLOS ONE* **12**, e0186216.
53. Loehfelm A, Rizwan MZ & Tups A (2021) A New Zealand green-lipped mussel oil-enriched high-fat diet exhibits beneficial effects on body weight and metabolism in mice. *Br J Nutr* **125**, 972–982.
54. Huang EY, Leone VA, Devkota S, *et al.* (2013) Composition of dietary fat source shapes gut microbiota architecture and alters host inflammatory mediators in mouse adipose tissue. *J Parenter Enter Nutr* **37**, 746–754.
55. Zheng JS, Lin M, Imamura F, *et al.* (2016) Serum metabolomics profiles in response to *n*-3 fatty acids in Chinese patients with type 2 diabetes: a double-blind randomised controlled trial. *Sci Rep* **6**, 1–8.
56. Sawada T, Tsubata H, Hashimoto N, *et al.* (2016) Effects of 6-month eicosapentaenoic acid treatment on postprandial hyperglycemia, hyperlipidemia, insulin secretion ability, and concomitant endothelial dysfunction among newly-diagnosed impaired glucose metabolism patients with coronary artery disease. *Cardiovasc Diabetol* **15**, 1–14.
57. Bonds DE, Lasser N, Qi L, *et al.* (2006) The effect of conjugated equine oestrogen on diabetes incidence: the Women's Health Initiative randomised trial. *Diabetologia* **49**, 459–468.
58. Kanaya AM, Herrington D, Vittinghoff E, *et al.* (2003) Glycemic effects of postmenopausal hormone therapy: the heart and estrogen/progestin replacement study: a randomized, double-blind, placebo-controlled trial. *Ann Intern Med* **138**, 1–9.
59. Muka T, Nano J, Jaspers L, *et al.* (2017) Associations of steroid sex hormones and sex hormone-binding globulin with the risk of type 2 diabetes in women: a population-based cohort study and meta-analysis. *Diabetes* **66**, 577–586.
60. Ding EL, Song Y, Manson JE, *et al.* (2007) Plasma sex steroid hormones and risk of developing type 2 diabetes in women: a prospective study. *Diabetologia* **50**, 2076–2084.
61. Dutta S & Sengupta P (2016) Men and mice: relating their ages. *Life Sci* **152**, 244–248.
62. Brinton RD (2012) Minireview: translational animal models of human menopause: challenges and emerging opportunities. *Endocrinology* **153**, 3571–3578.
63. Deckelbaum RJ, Worgall TS & Seo T (2006) *n*-3 fatty acids and gene expression. *Am J Clin Nutr* **83**, 1520S–1525S.
64. Prieto I, Hidalgo M, Segarra AB, *et al.* (2018) Influence of a diet enriched with virgin olive oil or butter on mouse gut microbiota and its correlation to physiological and biochemical parameters related to metabolic syndrome. *PLOS ONE* **13**, e0190368.
65. Rolland V, Roseau S, Fromentin G, *et al.* (2002) Body weight, body composition, and energy metabolism in lean and obese Zucker rats fed soybean oil or butter. *Am J Clin Nutr* **75**, 21–30.
66. Hidalgo M, Prieto I, Abriouel H, *et al.* (2014) Effect of virgin and refined olive oil consumption on gut microbiota. Comparison to butter. *Food Res Int* **64**, 553–559.
67. Yakoob MY, Shi P, Willett WC, *et al.* (2016) Circulating biomarkers of dairy fat and risk of incident diabetes mellitus among men and women in the USA in two large prospective cohorts. *Circulation* **133**, 1645–1654.
68. Imamura F, Fretts A, Marklund M, *et al.* (2018) Fatty acid biomarkers of dairy fat consumption and incidence of type 2 diabetes: a pooled analysis of prospective cohort studies. *Plos Med* **15**, e1002670.
69. Babio N, Becerra-Tomás N, Martínez-González MÁ, *et al.* (2015) Consumption of yogurt, low-fat milk, and other low-fat dairy products is associated with lower risk of metabolic syndrome incidence in an elderly mediterranean population. *J Nutr* **145**, 2308–2316.
70. Whelan J & Whelan J (2020) Conversion of dietary polyunsaturated fats between humans and rodents: a review of allometric scaling models. *Prostaglandins Leukot Essent Fat Acids* **158**, 102094.
71. Turnbaugh PJ, Ley RE, Mahowald MA, *et al.* (2006) An obesity-associated gut microbiome with increased capacity for energy harvest. *Nature* **444**, 1027–1031.
72. Ley RE, Turnbaugh PJ, Klein S, *et al.* (2006) Microbial ecology: human gut microbes associated with obesity. *Nature* **444**, 1022–1023.
73. Allin KH, Tremaroli V, Caesar R, *et al.* (2018) Aberrant intestinal microbiota in individuals with prediabetes. *Diabetologia* **61**, 810–820.
74. Gilbert JA, Quinn RA, Debelius J, *et al.* (2016) Microbiome-wide association studies link dynamic microbial consortia to disease. *Nature* **535**, 94–103.
75. Galicia-Garcia U, Benito-Vicente A, Jebari S, *et al.* (2020) Pathophysiology of type 2 diabetes mellitus. *Int J Mol Sci* **21**, 1–34.
76. Menni C, Zierer J, Pallister T, *et al.* (2017) *n*-3 fatty acids correlate with gut microbiome diversity and production of N-carbamylglutamate in middle aged and elderly women. *Sci Rep* **7**, 11079.
77. Candela M, Biagi E, Soverini M, *et al.* (2016) Modulation of gut microbiota dysbioses in type 2 diabetic patients by macrobiotic Ma-Pi 2 diet. *Br J Nutr* **116**, 80–93.
78. Tong X, Xu J, Lian F, *et al.* (2018) Structural alteration of gut microbiota during the amelioration of human type 2 diabetes with hyperlipidemia by metformin and a traditional chinese herbal formula: a multicenter, randomized, open label clinical trial. *MBio* **9**, e02392–e02417.
79. Menni C, Zhu J, Le Roy CI, *et al.* (2020) Serum metabolites reflecting gut microbiome α diversity predict type 2 diabetes. *Gut Microbes* **11**, 1632–1642.
80. Derrien M, Vaughan EE, Plugge CM, *et al.* (2004) Akkermansia muciniphila gen. nov., sp. nov., a human intestinal mucin-degrading bacterium. *Int J Syst Evol Microbiol* **54**, 1469–1476.
81. Everard A, Belzer C, Geurts L, *et al.* (2013) Cross-talk between Akkermansia muciniphila and intestinal epithelium controls diet-induced obesity. *Proc Natl Acad Sci USA* **110**, 9066–9071.
82. Plovier H, Everard A, Druart C, *et al.* (2017) A purified membrane protein from Akkermansia muciniphila or the pasteurized bacterium improves metabolism in obese and diabetic mice. *Nat Med* **23**, 107–113.



83. Zhang X, Shen D, Fang Z, *et al.* (2013) Human gut microbiota changes reveal the progression of glucose intolerance. *PLoS One* **8**, e71108.
84. de la Cuesta-Zuluaga J, Mueller NT, Corrales-Agudelo V, *et al.* (2017) Metformin is associated with higher relative abundance of mucin-degrading Akkermansia muciniphila and several short-chain fatty acid-producing microbiota in the gut. *Diabetes Care* **40**, 54–62.
85. Mukherjee A, Lordan C, Ross RP, *et al.* (2020) Gut microbes from the phylogenetically diverse genus Eubacterium and their various contributions to gut health. *Gut Microbes* **12**, 1802866.
86. Udayappan S, Manneras-Holm L, Chaplin-Scott A, *et al.* (2016) Oral treatment with Eubacterium hallii improves insulin sensitivity in db/db mice. *Npj Biofilms Microbiomes* **2**, 1–10.
87. He Y, Wu W, Zheng HM, *et al.* (2018) Regional variation limits applications of healthy gut microbiome reference ranges and disease models. *Nat Med* **24**, 1532–1535.
88. Gurung M, Li Z, You H, *et al.* (2020) Role of gut microbiota in type 2 diabetes pathophysiology. *EBioMedicine* **102**, 590.
89. Nguyen TLA, Vieira-Silva S, Liston A, *et al.* (2015) How informative is the mouse for human gut microbiota research? *Dis Model Mech* **8**, 1–16.