Dietary Protein Source Influence on Body Size and Composition in Growing Zebrafish

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Abstract

The importance of nutritional components on growth and body composition outcomes has been demonstrated in multiple model organisms. Although zebrafish (*Danio rerio*) have an established role in research laboratories for its utility in understanding developmental biology and genetics, the influence of diet composition on basic growth outcomes is less well demonstrated. In the current study, four protein sources were tested in isolation using isonitrogenous diets or combined using a defined lab diet. Fish ($n \approx 60$ /group) were group housed ($n \leq 10$ fish/1.8 L tank) and fed *ad libitum* three times daily for 12 weeks. Fish were assessed for effects on length, body weight, and body composition (lean and fat mass). Individuals fed wheat gluten protein were significantly shorter in length, with significantly lower body weight and lean mass in both male and female fish, although percent body fat was high compared with other diets. Casein-fed fish similarly had significantly reduced body length, body weight, and lean and fat mass in both male and female fish, with a low percent body fat compared with other diets (leanest). Fish protein hydrolysate-fed fish had significantly lower lean mass and a high percent body fat, whereas soy protein isolate diet performed similarly to a mixed-protein control diet for all measured outcomes. These results suggest that the protein source, with accompanying amino acid ratios or additional protein source differences, has a significant impact on growth and body composition outcomes in zebrafish when fed in a semipurified, defined diet background.

Introduction

LTHOUGH ZEBRAFISH (Danio rerio) have been used exten-Asively in a number of scientific disciplines (e.g., developmental biology, genetics, toxicology), their utility as a model of nutrition research has been limited, partly because of the lack of standardized, defined diets and husbandry conditions.¹ This is particularly surprising given the extensive knowledge of developmental biology, the sequenced genome, available genetic models, and forward genetic screening capabilities, as well as the economy of scale allowing significantly larger sample sizes to be investigated for less cost than other common vertebrate models. Although there are multiple commercial diets that have been used in zebrafish studies, defined diets like those used in rodent research have only recently been reported.^{2,3} These studies have demonstrated that formulated diets of defined composition were sufficient for growth and survival promotion compared with commercial diets.

The influence of diet composition and quantity on growth, metabolism, disease, and longevity is well recognized among many species. A general theory for mammals has developed, which proposes "diets that promote growth and early maturation are inversely associated with health and longevity," supported by diet composition studies and alterations in growth hormone signaling.^{4–7} Conversely, diets that delay growth or maturation, such as calorie restriction, reduce disease incidence, and increase longevity.^{8,9} However, observations from different fish species do not clearly support an inverse relationship between growth and longevity.^{10,11} Although it is commonly ascribed that caloric consumption is the main determinant of the growth, health, and longevity outcomes, it also appears that dietary composition (particularly the protein amount, source, and individual amino acids—e.g., methionine) may directly/indirectly contribute to health and longevity outcomes.^{12,13}

Beyond basic diet macronutrient composition, there is an increasing awareness of the influence of essential dietary nutrients (e.g., amino acids) on growth and disease onset, metabolism, feeding behavior, and even longevity in a variety of animal models.^{12,14–19} While the essential amino acid

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requirements may be predicted based on other nutrition studies using teleost models,^{20–23} few nutrient requirement studies have been formally presented in the research literature for zebrafish.^{3,24,25} Nevertheless, although zebrafish nutrition and obesity research is still developing compared with rodent models,²⁶ the information reported from decades of rodent nutrition research may be used to rapidly test dietary compositions for a variety of growth and eventual health-related outcomes.

Dietary macronutrient content including carbohydrate, fat, and protein is commonly reported and discussed as a relative percentage of total calories. Research in a number of animal models has shown that the source of dietary protein, independent of the relative caloric contribution, can have a significant impact on growth, reproduction, and disease/ longevity.¹³ Protein sources are also reported to influence feed conversion efficiency, or a measure of how much weight is gained per calorie eaten, through alterations in energy uptake, utilization, metabolism, and/or food intake patterns.²⁷⁻²⁹ Certain protein sources, such as milk or egg, are described as growth promoting, whereas plant-based protein sources are sometimes described as growth limiting because of limitations of essential amino acids.^{30,31} This inference can sometimes be a technical distinction, in that even growth-limiting diets can permit weight gain, just at a lower rate than growthpromoting diets when provided in ad libitum rations. However, in the area of aging and disease, both diets and genetic alterations that are growth promoting often result in earlyonset metabolic and age-related disease, along with increased mortality.^{9,32} A series of experiments were performed using four common protein sources (singularly or in combination) used in formulated diets to assess differences in growth including length, weight, and body composition to better understand the nutritional requirements of zebrafish within the laboratory research environment and determine the influence of various protein sources used in laboratory diets.

Materials and Methods

Zebrafish (AB strain) were obtained from the Aquatic Animal Research Core at University of Alabama at Birmingham (UAB). Embryos were collected randomly from a mass spawn of adult zebrafish (10 males and 10 females for each of the two cohorts, fed previously the mixed-protein diet described below and in Table 1). Embryos were transferred to Petri dishes (~50/dish) and incubated for 5 days at 28.5°C. At 5 days postfertilization (dpf), hatched larvae were transferred to 1.8 L tanks with \sim 2 cm of static water and fed rotifers (*Bran*chionus plicatilis) ad libitum enriched with Nannochloropsis (RotiGrow Omega, Reed Mariculture) three times daily until day 28. At 10 dpf water flow was initiated for all tanks as a slow drip. At day 28, fish were randomly distributed into 1.8 L tanks representing one of five diet treatments (n = 10 individuals per tank, five tanks per diet treatment). A photograph of all fish in each tank was recorded with a Nikon DS FIL or Nikon D-70, and the length of each fish was determined by NIS Elements 3.1 image analysis from a digital image. Average lengths were evaluated by analysis of variance (ANOVA) to ensure no significant differences among the treatments at the beginning of the experiment. All zebrafish were maintained at 28°C and $1500 \,\mu\text{S/cm}$ conductivity in a recirculating system (Aquaneering, Inc.). Flow rates were adjusted to provide at least two water changes per hour within each tank. Municipal tap water was filtered through 5 μ m sediment filter, followed by charcoal, reverse osmosis, and a cation/anion exchange resin (Kent Marine) before the addition of synthetic sea salts (Instant Ocean) to obtain final conductivity for the system water source. Sodium bicarbonate was used to maintain pH of the system

	Amount included (g/100 g total)						
Ingredient	MIX	FPH	CAS	SOY	WG		
Fish protein hydrolysate (82%) ^{a,b}	18.20	59.00					
Casein (vita-free) (96%) ^{a,c}	22.75		51.00				
Soy protein isolate (92%) ^{a,d}	4.55			52.00			
Wheat gluten (80%) ^{a,e}	9.10				60.00		
Dextrin	18.25	13.85	21.85	20.85	12.85		
Soy lecithin	4.00	4.00	4.00	4.00	4.00		
Canthaxanthin	2.31	2.31	2.31	2.31	2.31		
Ascorbylpalmitate	0.04	0.04	0.04	0.04	0.04		
Vitamin premix BML-2	4.00	4.00	4.00	4.00	4.00		
Mineral mix BTm	3.00	3.00	3.00	3.00	3.00		
Betaine	0.15	0.15	0.15	0.15	0.15		
Potassium phosphate monobasic	1.15	1.15	1.15	1.15	1.15		
Alginate	5.38	5.38	5.38	5.38	5.38		
Cholesterol	0.12	0.12	0.12	0.12	0.12		
Menhaden oil	4.67	4.67	4.67	4.67	4.67		
Corn oil	2.33	2.33	2.33	2.33	2.33		

TABLE 1. COMPOSITION OF DIETS

^aProtein content by percentage.

^bThe Scoular Company, Sopropeche–C.P.S.P. 90.

^cMP Biomedicals, Cat no. 904798.

^dMP Biomedicals, Cat no. 905456.

^eSigma Aldrich, Cat no. G5004.

MIX, mixed; FPH, fish protein hydrolysate; CAS, casein; SOY, soy protein isolate; WG, wheat gluten.

water (adjusted daily to maintain \sim 7.4 pH). At least 20% of system water was exchanged weekly. System water quality including total alkalinity (~50 ppm CaCO₂) and total hardness $(\sim 200 \text{ ppm CaCO}_2)$ have been previously measured but were not systematically obtained over the entire course of the experiment. Tanks were maintained on the same conditioned water system throughout the experiment, but rotated across rack positions at weekly intervals to reduce environmental confounding from noise, light, vibration, or other unidentified sources. Tanks were siphoned weekly to remove any uneaten feed or debris. Total ammonia nitrogen, nitrite, and nitrate were measured colorimetrically (Mars Fish Care, Inc.), and all but nitrate remained below detectable limits.

Diets

Diet ingredient information is provided in Table 1. Each diet was produced with a single, common base mix (all ingredients minus the protein) using chemically defined ingredients. To this base mix was added one of four protein sources or a combination of all four (control diet). The control diet contained a mixed-protein source including fish protein hydrolysate (FPH; Scoular Company), casein (CAS, vitamin-free; Affymetrix-USB), soy protein isolate (SOY; MP Biomedicals), and wheat gluten (WG; MP Biomedicals). Each experimental diet contained only a single-protein source with adjustments to the carbohydrate component (dextrin) to calorically compensate for nitrogen differences between individual protein sources. Diets were mixed in an orbital mixer (Kitchen Aid) and then extruded with a Kitchen Aid extruder (KPEXTA) and were air-dried to $\sim 10\%$ moisture content before storage. Full diet analyses for proximate composition, amino acid composition, and mineral content were performed by MVTL Laboratories, Inc., and the results are provided in Table 2. For feeding, diets were ground to a powder (250-500 µm sieved) and measured aliquots (approx. $\geq 5\%$ of body weight) were dispensed at ~900, 1300, and 1700 hr during the light cycle in surplus quantities. Fish

TABLE 2. PROTEIN SOURCE DIETS: COMPOSITIONAL ANALYSIS							
	MIX	FPH	CAS	SOY	WG		
Proximate analysis							
Moisture (%) ^{a,b}	11.52, 11.33	10.97, 10.72	10.30, 10.18	10.08, 9.94	9.58, 9.45		
Fat (%) ^{a,c}	12.34, 12.05	15.86, 15.65	10.47, 10.34	10.33, 10.19	11.10, 10.90		
Fiber (%) ^a	1.60, 1.58	1.77, 1.67	1.41, 1.60	1.65, 1.67	1.73, 1.85		
Protein (%) ^{a,d}	44.90, 44.40	45.30, 45.19	44.50, 43.97	43.90, 43.58	49.50, 49.09		
Ash (%) ^a	6.84, 6.85	8.37, 8.34	7.89, 7.79	6.86, 6.90	5.95, 5.98		
Amino acids ^e							
Cysteine (%)	0.400	0.400	0.190	0.530	1.010		
Methionine (%)	1.080	1.160	1.320	0.560	0.730		
Lysine (%)	2.930	3.200	3.610	2.700	0.660		
Alanine (%)	1.894	2.915	1.427	1.932	1.249		
Arginine (%)	2.090	2.748	1.642	3.187	1.547		
Aspartic Acid (%)	3.284	3.849	3.322	4.996	1.509		
Glutamic Acid (%)	9.631	5.626	10.120	8.388	18.060		
Glycine (%)	2.072	3.984	0.876	1.848	1.622		
Isoleucine (%)	2.023	1.709	2.394	2.132	1.763		
Leucine (%)	3.631	2.866	4.354	3.500	3.293		
Serine (%)	2.140	1.804	2.461	2.103	2.177		
Threonine (%)	1.706	1.744	1.917	1.612	1.217		
Valine (%)	2.424	2.019	3.016	2.196	1.868		
Histidine (%)	1.120	0.940	1.337	1.110	0.975		
Phenylalanine (%)	2.130	1.573	2.387	2.325	2.600		
Tyrosine (%)	1.877	1.574	2.372	1.473	1.491		
Taurine (%)	0.098	0.367	0.010	0.010	0.010		
Tryptophan (%)	0.506	0.374	0.620	0.582	0.467		
<i>Minerals</i> ^{e,f}							
Calcium (%)	0.81	0.86	0.80	0.85	0.80		
Copper (ppm)	11.26	8.94	8.00	20.01	13.98		
Iron (ppm)	123.60	125.30	79.21	139.50	99.34		
Magnesium (ppm)	440.30	642.80	259.90	530.20	564.10		
Manganese (ppm)	106.60	105.00	106.40	109.80	115.50		
Phosphorus (%)	1.31	1.38	1.41	1.39	1.10		
Potassium (%)	0.82	1.24	0.58	0.76	0.66		
Sodium (%)	0.91	1.29	0.67	1.15	0.82		
Zinc (ppm)	41.78	43.18	40.73	36.51	41.21		

^{a,b}Duplicate measures.

^cFat by ethyl ether extraction.

^dProtein = $N \times 6.25$.

ePercent by weight.

^fParts per million.

were observed to consume the diets at each level of the water column.

Growth measures

Similar to length measures at baseline for randomization, all fish were subsequently photographed in clear-bottom containers at several intervals (10, 13, and 16 weeks postfertilization). Digital images were used to measure individual fish body lengths (from the tip of the mouth to the base of the caudal fin) within a tank and averaged within the diet treatment against a reference length standard in all images. All lengths were recorded to the nearest 0.01 mm. Final body weight was measured at study completion on tricaine methanesulfonate (15 mg/L)-anesthetized fish before body composition assessment. Anesthetized fish were quickly blotted to remove surface moisture and weighed to the neared 0.0001 g.

Body composition assessment

Body composition measurement (lean and fat mass) was performed at study completion by quantitative magnetic resonance (QMR) using the EchoMRI 3-in-1 system and the tissue probe holder. Tricaine (MS-222; 300 mg/L) was used to rapidly immobilize the fish in a small volume of water (also containing MS-222; 300 mg/L) for the duration of the QMR measure. Anesthetic induction was performed on single fish $(\sim 2-5 \text{ min})$ to monitor swimming motion, loss of equilibrium, cessation of opercular movement, and heart contractions.³³ QMR scans were performed using the appropriate tissue sample holder (with MS-222 containing water) and high precision setting, requiring $\sim 3 \min \text{ per scan}$. Body composition measures include lean mass (or fat free mass) and fat mass reported to the nearest 0.001 g. After the scans, fish were subsequently revived by submersion in fresh system water and gentle perfusion of the gills using a disposable pipette.

Euthanasia

At study completion, fish were euthanized by rapid submersion in ice-cold water with MS-222 $(300 \text{ mg/L})^{33}$ and the carcass stored at -80° C until disposal.

Data analyses

Data are reported as means and standard error (SE) of the mean. Length (at each of the three measurement time points), final weight, fat, lean/fat free mass, and percent fat were separately compared by ANOVA using SAS v.9.1, adjusting for cohort and/or body weight as covariates (ANCOVA) when significance between groups or relationships were observed. Data and analyses were stratified by sex, to accommodate significant differences between sexes. Observed significant differences ($p \le 0.05$) were further tested using Duncan's multiple range *post hoc* test for among-groups differences (p < 0.05).

Results

Fish consumed each of the five diets with no overt feeding differences observed during the course of the study. All diets initially floated and were consumed by fish at the surface of the water, but subsequently sank (given enough time) and

TABLE 3. LENGTH (MM) BY COHORT AND GROUP

Cohort	Group	Ν	10 weeks*	13 weeks	16 weeks
1	MIX FPI CAS SOY WG	26 25 25	$\begin{array}{c} 16.94 \ (0.59)^{a} \\ 15.46 \ (0.80)^{a,b} \\ 15.66 \ (0.81)^{a,b} \\ 16.46 \ (0.46)^{a} \\ 14.27 \ (0.65)^{b} \end{array}$	$\begin{array}{c} 19.57 \ (0.55)^{a} \\ 17.72 \ (0.82)^{a,b} \\ 17.58 \ (0.83)^{a,b} \\ 18.20 \ (0.45)^{a} \\ 15.99 \ (0.61)^{b} \end{array}$	$\begin{array}{c} 22.83 \ (0.57)^{a} \\ 21.31 \ (0.77)^{a,b} \\ 20.99 \ (0.84)^{a,b} \\ 20.65 \ (0.35)^{b} \\ 18.80 \ (0.51)^{c} \end{array}$
2	MIX FPI CAS SOY WG	35 35 34	$\begin{array}{c} 16.39 \ (0.60)^a \\ 15.64 \ (0.62)^{a,b} \\ 15.12 \ (0.77)^{a,b} \\ 17.04 \ (0.58)^a \\ 13.99 \ (0.69)^b \end{array}$	$\begin{array}{c} 19.74 \ (0.65)^{a,b} \\ 18.70 \ (0.76)^{a,b} \\ 17.87 \ (0.83)^{b,c} \\ 20.51 \ (0.63)^{a} \\ 16.23 \ (0.79)^{c} \end{array}$	$\begin{array}{c} 23.74 \ (0.62)^a \\ 23.36 \ (0.61)^a \\ 22.11 \ (0.77)^{a,b} \\ 23.80 \ (0.38)^a \\ 20.60 \ (0.70)^b \end{array}$

Values shown are mean (standard error of the mean). Different letters indicate significant differences between groups at p < 0.05 (Duncan multiple range *post hoc* test).

*Age denoted in weeks postfertilization.

were also consumed from the floor of the tanks. Despite the fact that all fish were fed ad libitum (in excess) three times per day, lengths were significantly different among protein sources at 10, 13, and 16 weeks (Table 3 and Fig. 1). Mean length was significantly smaller in the WG group compared with all others at each time point and study end in both cohorts (Table 3 and Fig. 1), with CAS fish having an intermediate length compared with the remaining groups (MIX, FPH, SOY; Table 3 and Fig. 1). Similarly, final body weight was lowest in the WG group (males and females), with small but significant differences present across the other groups when considering both sexes (MIX \approx SOY>FPH>CAS>WG; Table 4 and Fig. 2). A significant, positive relationship was observed for body weight with both lean and fat mass measures for all fish (Fig. 3). Additionally, female fish as a whole were heavier (p < 0.01), with a higher fat mass (p < 0.01) than males (data not shown). There was a significant difference between groups for fat mass (p < 0.0001) and lean mass (p < 0.001), with CAS and WG fish having the lowest mean fat mass (Table 4 and Fig. 4), and WG fish having the lowest lean mass amounts

25 a,b MIX FPH SOY 20 WG a.b -ength (mm) 15 10 5 0 WK 10 WK 13 WK 16

FIG. 1. Mean body length (mm) of zebrafish at 10, 13, and 16 weeks postfertilization fed either the mixed (MIX), fish protein hydrolysate (FPH), casein (CAS), soy protein isolate (SOY), or wheat gluten (WG) as the protein source in the diet. Different letters indicate between-group differences at p < 0.05.

DIETARY PROTEIN SOURCE AND GROWTH IN ZEBRAFISH

Cohort	Group	Sex	N	Weight (mg)	Fat (mg)	Lean (mg)	% Fat
1	MIX	М	13	204.4 (19.8) ^a	26.7 (2.3) ^a	164.2 (19.2) ^a	14.86 (0.81)
		F	11	$178.6(24.2)^{a}$	$26.1 (3.4)^{a,b}$	155.5 (17.4) ^{a,b}	14.50 (0.83)
	FPH	Μ	13	163.7 (14.4) ^{a,b}	25.2 (1.6) ^a	127.6 (12.1) ^{a,b,c}	17.25 (0.88)
		F	10	145.9 (22.0) ^{a,b}	23.8 (3.0) ^{a,b}	109.0 (15.6) ^{b,c}	19.02 (1.25)
	CAS	Μ	13	138.5 (20.0) ^{b,c}	14.9 (1.5) ^b	115.9 (16.2) ^{b,c}	13.53 (1.80)
		F	6	162.3 (25.5) ^a	$17.8(2.4)^{\rm b}$	132.0 (17.6) ^{a,b}	11.92 (0.38)
	SOY	Μ	14	168.8 (12.8) ^{a,b}	$23.6 (1.8)^{a}$	138.4 (10.9) ^{a,b}	14.65 (0.63)
		F	10	212.0 (21.7) ^a	$32.7 (3.8)^{a}$	176.2 (16.7) ^a	15.55 (0.79)
	WG	Μ	18	107.3 (7.6) ^c	17.0 (1.2) ^b	93.6 (6.5) ^c	15.61 (0.54)
		F	6	79.4 (16.4) ^b	17.0 (1.6) ^b	70.0 (13.5) ^c	21.06 (2.10)
2	MIX	Μ	20	206.1 (16.9) ^{a,b}	27.6 (2.1) ^{a,b}	176.4 (15.6) ^a	13.90 (0.44)
		F	14	320.9 (16.2) ^a	45.1 (5.9) ^{a,b}	266.2 (19.6) ^a	14.16 (1.04)
	FPH	Μ	21	190.6 (15.4) ^b	27.5 (2.2) ^{a,b}	146.9 (13.5) ^{a,b}	16.29 (0.61)
		F	9	302.9 (54.4) ^a	56.8 (18.0) ^a	250.0 (76.8) ^a	18.38 (0.14)
	CAS	Μ	22	186.7 (16.2) ^b	$21.3(1.7)^{b}$	166.8 (15.2) ^{a,b}	11.65 (0.43)
		F	5	149.0 (23.0) ^b	$16.8(2.78)^{c}$	115.3 (15.1) ^b	12.59 (0.59)
	SOY	Μ	26	243.0 (13.3) ^a	$30.0 (1.5)^{a}$	186.0 (9.0) ^a	13.91 (0.32)
		F	7	281.1 (28.2) ^a	47.7 (8.6) ^a	230.0 (45.4) ^{a,b}	17.27 (0.27)
	WG	Μ	21	140.3 (18.4) ^c	$21.8(2.9)^{b}$	132.8 (17.1) ^b	15.17 (1.15)
		F	6	153.0 (22.5) ^b	19.4 (1.8) ^{b,c}	124.6 (26.9) ^b	14.70 (1.57)

TABLE 4. FINAL BODY WEIGHT AND COMPOSITION BY COHORT AND GROUP

Values shown are mean (standard error of the mean). Different letters indicate significant differences between groups within a given sex and cohort at p < 0.05 (Duncan multiple range *post hoc* test).

in both sexes (Table 4 and Fig. 5), with these differences persisting after controlling for body weight (data not shown). CAS and FPH fish had a lower lean mass than the MIX controls (Table 4 and Fig. 5) after co-varying for body weight (data not shown). Similarly, the relative fat amount (% total fat) was significantly different between groups (p < 0.001) with CAS-fed fish having the leanest (lowest% fat) body composition and the FPH and WG having the highest percent fat (Table 4, data not shown).

Discussion

This study demonstrates a significant impact of dietary protein source, despite *ad libitum* feeding protocols and isonitrogenous diets, on growth outcomes, including length, body weight, and body composition, using the zebrafish model. Additionally, multiple single-protein source diets performed as well as a mixed-protein source diet for length and weight outcomes, with more complex body composition (lean and fat mass) responses observed among groups. WG protein was deficient for growth, producing fish in both sexes that were shorter in length, with significantly lower body weight and lean mass, although percent body fat was high compared with other diets. CAS-fed fish similarly had significantly lower body length, body weight, and lean and fat mass in both male and female fish, with the lowest percent body fat (leanest) compared with the other protein sources. FPH-fed fish, a presumably high-quality, replete protein

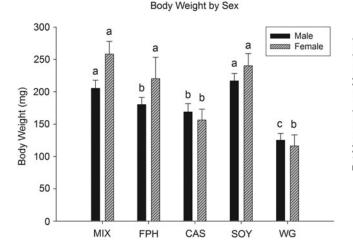


FIG. 2. Mean group body weight of zebrafish at 16 weeks postfertilization for males and females. Different letters indicate between-group differences within a given sex at p < 0.05.

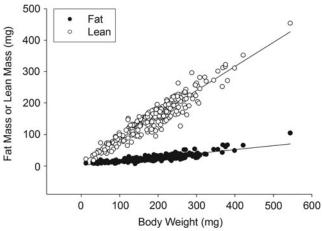


FIG. 3. Relationship between body weight and fat mass or lean mass of zebrafish at 16 weeks postfertilization across all diet groups. Lean mass $r^2 = 0.8993$, p < 0.001; fat mass $r^2 = 0.7447$, p < 0.001.

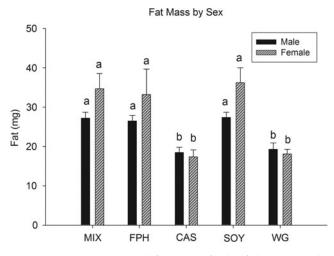


FIG. 4. Mean group total fat mass of zebrafish at 16 weeks postfertilization for males and females. Different letters indicate between-group differences within a given sex at p < 0.05.

source, had significantly lower lean mass and a high percent body fat, whereas SOY-fed fish resembled mixed-protein controls for all measured outcomes. Despite females being heavier and possessing a larger fat mass than males when all groups were combined, the significant differences between groups for growth outcomes were related to diet treatment and not sex-specific.

All diets were fed as a sole source of nutrients (no additional live feeds or commercial supplements) in a uniform physical state from 4 weeks postfertilization for the duration of the study. Additionally, the base formulation (excluding the protein sources) was kept consistent among groups, pointing to the significant impact dietary protein and potentially amino acid ratios may have on growth and body composition outcomes in zebrafish. The diets were formulated to maintain a similar nitrogen content (as presented in Table 2).

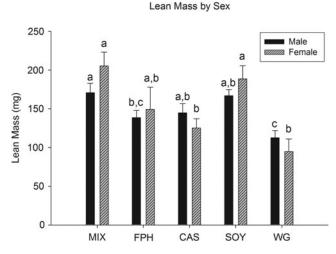


FIG. 5. Mean group total lean mass of zebrafish at 16 weeks postfertilization for males and females. Different letters indicate between-group differences within a given sex at p < 0.05.

However, because of the varying proportion of nitrogen present in the different dietary protein sources, an additional dietary caloric source had to be modified to approximate isocaloric levels between test diets. In this case, the carbohydrate content was manipulated to balance between diets because of its caloric density and the desire to measure body composition outcomes (lean and fat content), which may be expected to change with the increasing or decreasing dietary lipid content. Thus, we cannot formally exclude the possibility that manipulation of the dietary carbohydrate content also influences the measures growth outcomes,³⁴ although previous findings would suggest that dietary carbohydrate does not affect growth outcomes at those levels.³⁵ Diets with similar carbohydrate content and protein content but different protein sources (see Tables 1 and 2: SOY vs. CAS) still exhibited significant differences in multiple growth outcomes, most likely attributable to the protein source contribution to the individual diets.

While the individual, essential amino acid requirements of zebrafish have not been systematically tested, assuming similar requirements to other fish,^{20,23} it may not be surprising that the wheat-fed fish exhibited the lowest growth, provided the known limitations of lysine in wheat.^{23,36–38} Previous research with soy protein also might suggest that limitation of essential, sulfur-containing amino acids (methionine and cysteine) might be expected to limit growth outcomes,28,39,40 but the fish in this study responded well with similar growth outcomes to the MIX control diet. One explanation for this difference may reside in the high protein content of the diet $(\sim 45\%)$, which could offset any limitation of specific amino acids proportionally with a given protein source because of the high absolute mass amount present. Alternatively, dietary betaine, serving as a methyl donor, may promote production of methionine. Progressive reductions in the protein content of the diet (<45%) should be tested to determine minimal requirements for essential amino acids.

An additional limitation of this and the majority of nutrition studies is the diet complexity and proportional nature of nutrients. One example is the amino acid differences among diets. Since diets were formulated to be isonitrogenous, any elevation in a single amino acid must be compensated by a reduction in one or more amino acids to maintain the overall nitrogen balance (see Table 2). This complexity limits the ability to confidently ascribe the given growth outcomes to a single nutrient (amino acid) as the proportion of compensating changes in other nutrients may equally have attributed to the observed difference. Additionally, it is not yet known how feeding in excess of dietary needs three times a day may influence these and other outcomes. Accurate determination of food intake in an aquaculture system such as this is challenging, particularly on an individual fish basis. Given the observed variance among individuals within a single tank, it would be valuable to know whether food intake differences account for any of the variability. Similarly, with essential nutritional elements, it would be beneficial to know if limitation invokes a hypophagic response because of a perceived dietary inferiority or a hyperphagic response to increase overall nutrient intake to meet any given individual limiting requirement.

The diets performed well for growth measures, including length and overall body weight. Overall length, weight, and percent lipid were similar to the findings of Kaushik *et al.*,³

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who report a length of 23 mm at 9 weeks of age with a total body lipid of $\sim 10\%$.³ Another recent work reports fish that appear to be significantly larger than those in the present study, with a final length of \sim 33 mm and a body weight of 350 mg.²⁴ Whether strain differences partially account for these growth differences has not been evaluated. Given the larger starting and final lengths,²⁴ it would be of interest to determine if dietary composition differences produced similar growth responses in another zebrafish strain background or if earlier initiation of formulated feeds altered the growth parameters. Additionally, tank densities were different between the two studies, with the current study having three times the density of the previous work (e.g., current: ~5.5 fish/L vs. ~1.6 fish/L).24 Although the zebrafish model has garnered much recent attention for its po-tential use in lipid and obesity research,⁴¹⁻⁴⁷ few studies have reported overall body composition outcomes. In the present work, the dietary protein source significantly impacted lean and fat mass outcomes. Across treatments the fish retained a relatively lean phenotype, with the CAS-fed fish being the leanest. Although the dietary protein amount has been shown to significantly influence food intake, body weight, and body composition outcomes (low proteinincreased intake with greater body fat vs. high proteindecreased intake with greater body lean), the comparison of different protein sources has had mixed results regarding their effectiveness for desirable body composition alterations (fat reduction and lean accretion), particularly when provided in a purified, nonsupplemented form. Given the observed difference in body weight and composition, it would be of interest to know if physiological responses paralleled these growth outcomes, with alterations in endocrine profiles, metabolic responses, and reproductive output, which were not fully measured in this study. As the diets were isonitrogenous, the body composition response with changes in lean and fat mass between protein sources raises questions about the influence of amino acid ratios or other nonnutritive compounds that may accompany the protein sources for their molecular impact on both the host organism (zebrafish) as well as the microbiome inhabiting the gut.⁴⁸ Future studies may assess nutrient-specific alteration in the microbioata with diet and body composition outcomes.

The data presented demonstrate a significant impact of the protein source on growth outcomes in zebrafish using a semipurified diet formulation. Finding which nutrient differences are responsible for the differential growth outcomes as well as determining minimal requirements for the zebrafish model will be important for diet standardization. Finally, these results suggest that body composition, both lean and fat mass, is significantly influenced by the protein source and should be carefully considered in future studies that may utilize the zebrafish as a model of diet-induced obesity or lipid metabolism.

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Author Disclosure Statement

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