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Temperature modulates PFAS accumulation and energy allocation in sheepshead minnows

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Abstract

Climate warming and chemical pollution shape aquatic ecosystems, yet the mechanisms underlying their combined effects remain poorly understood. We investigated how climate-relevant surface water warming alters per- and polyfluoroalkyl substances (PFAS) toxicokinetics and their effects on sheepshead minnows (*Cyprinodon variegatus*) physiological performance. Adult fish were chronically exposed to an environmentally relevant PFAS mixture (perfluorooctane sulfonate (PFOS) + perfluorooctanoate (PFOA)) under three warming scenarios. Tissue PFAS concentrations, whole-organism metabolic rates, swimming performance, reproductive parameters and somatic indices were assessed. Temperature modified PFAS tissue concentrations in a compound- and tissue-specific manner, notably promoting PFOA redistribution to eggs. Metabolic responses were temperature-dependent: at 26 °C, higher tissue PFAS concentrations were associated with elevated standard and maximum metabolic rates (SMR and MMR), maintaining aerobic scope (AS). At 28.5 °C, SMR remained stable while MMR and AS declined with rising PFAS, indicating reduced oxygen available for energetically demanding activities. Despite unchanged swimming and reproductive outputs, an increased hepatosomatic index with increasing tissue PFAS concentrations and altered PFAS distribution suggest detoxification costs. These findings indicate that warming is likely to exacerbate contaminant stress, with consequences for coastal fish population resilience and offspring development. PFAS risk assessment should account for multiple stressors under ongoing climate change.

Keywords: PFAS, temperature, fish, accumulation, metabolism, reproduction, swimming performance

Synopsis

Limited research addresses how temperature affects PFAS toxicokinetics and toxicity. This study shows that warming reshapes tissue PFAS concentrations and distribution, and influences fish energy-allocation trade-offs.

1. Introduction

Multiple stressors shape ecosystems worldwide and pose a growing threat for aquatic organisms (1). In particular, chemical pollution and climate warming can interact to increase contaminant exposure and heighten organismal sensitivity (2,3), with consequences for growth, reproduction, and behavior that may promote nonlinear ecological responses and reduce ecosystem resilience (4,5,6). The mechanisms driving these interactions remain poorly understood, especially for contaminants of emerging concern, despite their importance for anticipating contaminant risks under accelerating climate change.

Per- and polyfluoroalkyl substances (PFAS) represent a large family of synthetic contaminants for which temperature-dependent effects remain largely unknown. PFAS have been widely used since the 1950s for their unique chemical and physical properties (7). Strong carbon–fluorine bonds and fluorinated chains make them highly stable and water- and oil-repellent. Some PFAS, like perfluoroalkyl acids (PFAAs), also have a hydrophilic group, giving them surfactant-like behavior (8). These characteristics make PFAS resistant to degradation in natural conditions and allow their widespread transport and accumulation, especially in aquatic ecosystems (9,10). Perfluorooctanoate (PFOA) and perfluorooctane sulfonate (PFOS) are among the most historically widely used long-chain PFAAs and have attracted attention in the scientific, public, and regulatory communities (11). Due to their persistence, bioaccumulation, and toxicity, they are listed under Annexes A (Elimination) and B (Restriction), respectively, in the Stockholm Convention on Persistent Organic Pollutants (12,13) UNEP 2009; UNEP 2019) and are regulated in many countries (14). Although their production and use have been largely phased out since the early 2000s (11), their extreme stability and remaining sources continue to contaminate surface and groundwater. Concentrations typically range from a few ng⬚L^{-1} to low µg⬚L^{-1} (10), but much higher levels have been reported near industrial sites and legacy contamination areas; for example, up to 42,000⬚ng⬚L^{-1} of PFOA and 2,700⬚ng⬚L^{-1} of PFOS have been found in groundwater near 3M company disposal sites in Minnesota (15).

PFAA bioaccumulation and health risks in people and wildlife are well documented (9). The

amphiphilic structure of PFAAs confers a high affinity for serum albumin, fatty acid–binding proteins

(FABPs), and other endogenous proteins that influence their partitioning within organisms (16). In

fish, PFAS primarily accumulate in protein-rich tissues such as blood, liver, and kidneys (17), and can

also bind to vitellogenin, contributing to deposition in eggs and posing potential multigenerational

risks. Generally, PFAS bioaccumulation increases with carbon chain length and hydrophobicity and

depends on the nature of the functional group (18). In fishes, PFOS and PFOA induce reprotoxicity

(19), cause developmental abnormalities (20), and disrupt energy metabolism (21).

Although the physiological impacts of PFAS are increasingly recognized, little is known about how

temperature may alter their uptake, distribution, and toxicity in fish (22, 23). Temperature governs

the environmental behavior of contaminants in aquatic environments, influencing solubility, binding

affinities, and partitioning (5). For PFAS specifically, non-covalent interactions with proteins,

stabilized by van der Waals forces and hydrogen bonding, are exothermic and weaken at higher

temperatures, potentially enhancing dissociation and elimination (24, 25). In parallel, temperature is

a fundamental driver of ectotherm performance, as it sets the pace of biochemical reactions and

defines the energetic capacity available for essential functions such as growth, locomotion, and

reproduction (26, 27, 28, 29). As energy demands increase with temperature (30), increasing

metabolic rates may, in turn, enhance contaminant uptake (31). Altogether, clarifying how climate

warming and PFAS exposure interact to shape bioenergetic traits is essential for predicting energy

balance and organismal fitness (32). Whole-organism metabolic rates provide a valuable integrative

measure, reflecting oxygen consumption and the combined capacity for oxygen delivery and energy

conversion. However, research on PFAS and fish metabolic rates remains scarce (33). Critically, no

studies have addressed PFAS mixtures, and only a single study has examined temperature-

dependent effects, reporting metabolic disruptions in juvenile fish exposed solely to PFOS (34). This

gap underscores the need to evaluate PFAS toxicity through whole-organism metabolic traits that

capture energy allocation trade-offs and ecological function, particularly under varying thermal

conditions.

The overarching aim of this study was to assess how climate-driven temperature change may alter

PFAS toxicokinetics and compromise physiological performance in sheepshead minnows (*Cyprinodon*

variegatus). *Sheepshead minnows* are widely used in toxicology studies (35, 36). They are a

eurythermal species broadly distributed along the U.S. Atlantic coast and Gulf of Mexico, including

areas influenced by wastewater treatment plant (WWTPs) discharges, a PFAS source due to low

removal efficiencies (37). Consistent with this exposure context, our field sampling downstream of

WWTPs (Connecticut shore, Long Island Sound) detected PFOS/PFOA at all investigated sites and

documented sheepshead minnows, confirming exposure relevance (unpublished data). In our study,

we aimed (i) to characterize how temperature affects the tissue accumulation and distribution of a

PFAS mixture made of PFOS and PFOA, and (ii) to evaluate whether temperature modulates the

toxicity of the mixture on fish metabolic, swimming and reproductive performance. We

hypothesized that PFOS would accumulate to higher concentrations than PFOA because of its

greater protein-binding affinity and slower elimination (18), and would therefore primarily drive

PFAS-associated toxicity. We further expected temperature to influence PFAS accumulation by

altering the thermodynamics of PFAS-protein interactions and physiological processes such as

uptake and elimination (24). Lastly, we predicted that PFAS exposure would increase metabolic

oxygen demand beyond the fish's oxygen supply capacity, thereby reducing AS and limiting

performance, with the magnitude of these effects being temperature dependent. To test these

predictions, we conducted two complementary, laboratory-based chronic aqueous exposure

experiments with adult sheepshead minnows acclimated to three surface water temperatures

representative of current and future warming scenarios in the northeastern USA. By integrating

thermal and contaminant stressors within a whole-organism framework, this study advances our

understanding of how combined environmental stressors influence fish physiological function and

ecological fitness in the context of global change.

2. Materials and Methods

2.1 Fish husbandry and temperature acclimation

All fish husbandry and experimental procedures were in accordance with the University of Connecticut's Institutional Animal Care and Use Committee protocols A22-049 & A23-049. Juvenile sheepshead minnows (*Cyprinodon variegatus,* > 60 days post hatch (dph)) from Aquatic BioSystems, Inc. (Fort Collins, CO, USA) were reared to adulthood (> 150 dph) in two 380 L - 1360 L recirculating aquaculture systems (Iwaki Aquatic, Systems and Services, Holliston, MA, USA).

During husbandry, water quality (temperature, salinity, dissolved oxygen, pH) was checked three times a week, and ammonia, nitrite, and nitrate were checked weekly to ensure they remained within acceptable limits. Prior to experiment onset, sexually mature fish were acclimated for at least three weeks to their assigned temperature (Exp. 1: 24 °C, 26 °C, or 28.5 °C; Exp. 2: 24 °C or 28.5 °C) under the same monitoring regime. Fish were fed daily with standard commercial flaked food (TetraMin Tropical Flakes, Tetra, Melle, Germany), and maintained under a 14 h light:10 h dark photoperiod cycle during the holding and experimental periods.

2.2 Experimental design

We conducted two complementary experiments to test how temperature affects the tissue accumulation and distribution of PFOS and PFOA and the fitness of small-bodied fish. In the first experiment, we assessed PFAS accumulation (whole-body, muscle tissue and eggs) among temperature treatments as well as the coupled influences of PFAS and elevated temperature exposure on fish metabolism, swim performance, and reproduction. The second experiment, motivated by the results of the first, assessed the influence of temperature on PFOS and PFOA distribution among muscle, liver, ovary, and blood. Three temperatures were tested, representative of contemporary summer surface water temperatures (24°C) and those projected 50 years and 100 years in the future (26 °C and 28.5 °C) for the Long Island Sound estuary (38). We targeted a

combined PFOS+PFOA exposure concentration of 100 µg·L^{-1}, corresponding to the combined U.S.

EPA Freshwater Aquatic Life Water Quality Criteria for PFOA and PFOS (39). Details of the exposure

module, water-quality parameters during PFAS exposure, PFAS stock-solution preparation, and PFAS

concentration monitoring are provided in the Supplementary Information (Figure S1, Tables S1-S2).

Fish were randomly assigned to PFAS-exposed or non-exposed groups and one of three temperature

treatments, for six experimental conditions in experiment 1 and four conditions in experiment 2

(Figure S1). For the first experiment, 684 fish (mean lengths and weights ± standard errors (SE): 53.0

± 0.39 cm, 3.26 ± 0.07 g) were distributed across 36 aquaria (6 treatments * 6 replicates with 19 fish

per aquaria, and at a ratio of 12 females for 7 males). Experimental exposures were initiated on a

rolling basis as aquaria became available, in batches of six aquaria at a time, with each batch

comprising all six treatment conditions. A pre-exposure period of 6 days was conducted to allow

them to acclimate to the system, establish baseline egg production, and to allocate fish to exposure

groups, ensuring that any changes in reproductive output during the exposure period could be

attributed to PFAS exposure rather than natural variation (40). Each aquarium included spawning

boxes for daily egg quantification throughout the pre-exposure and exposure periods. Egg samples

were collected for PFAS analysis from the three PFAS-exposed groups at day 0 (prior to exposure),

day 14, and day 28, sampling three aquaria per treatment at each time point (n= 27 samples; 9

samples per PFAS-exposed treatment group). At the end of the 28-day exposure, two males from

each aquarium were selected for swimming and metabolic performance trials (see below). Males

were prioritized to limit variance under sample-size and logistical constraints, as metabolism varies

by sex and reproductive status. All other fish were euthanized using tricaine methane sulphonate

(MS-222; 0.5 g·L^{-1}; Syndel, USA). Four fish (two males and two females) per aquarium were collected

for biometric measurements (total length, height, width, and body weight) and dissected for skinless

muscle for PFAS analysis. Liver and gonads were weighed to calculate somatic indices. Water, eggs,

and muscle tissue were preserved at -20 °C for PFAS analyses.

181 The second experiment was performed using a new batch of fish. In total, 105 (49.2 ± 0.22 cm, 2.40

182 ± 0.04 g) fish were distributed across 12 aquaria (4 treatments * 3 replicates with 7-10 fish per

183 aquaria and at a ratio of 6-9 females for 1 male). All fish were euthanized at the end of the 28-day

184 exposure. Three females per aquarium were collected for biometric measurements (total length,

185 height, width, and body weight) and dissected for skinless muscle, liver, blood, and ovary for PFAS

186 analysis. Liver and gonads were weighed to calculate somatic indices. Water, muscle, liver, blood and

187 ovary tissue were preserved at -20°C for PFAS analyses.

188 2.3 Somatic and reproductive endpoints

189 Hepatosomatic index (HSI) and gonadosomatic index (GSI) were calculated in both experiments as:

190 $$\text{Somatic index } (\%) = \frac{\text{organ weight (g)}}{\text{body weight (g)}} \times 100 \textbf{ equation (1)}$$

191 Daily (eggs·female^{-1}·day^{-1}) and cumulative (eggs·female^{-1}) egg production in experiment 1 were

192 calculated for each aquarium on each day over the 28-day exposure period by summing egg counts

193 and normalizing by the number of females. Egg production data were expressed per female to

194 account for any discrepancies in the initial 12:7 female-to-male ratio, including any differences due

195 to mortality during the exposure period. These daily values were then averaged by treatment group

196 to visualize temporal trends. To summarize reproductive output, mean daily egg production was

197 calculated per aquarium over the full period and then averaged across groups. Finally, linear

198 regressions of daily and cumulative egg production over time were used to estimate rates of change

199 (slopes), which were also averaged by treatment group.

200 2.4 Swimming and metabolic performances protocol

201 Swimming performance and aerobic metabolic rates of individual males were determined using

202 1500 mL intermittent-flow swim tunnels (Loligo Systems, Tjele, Denmark). Water flow was calibrated

203 with digital particle tracking velocimetry and controlled by a DAQ-M data acquisition device. Oxygen

204 consumption (MO$_2$, mg O$_2$ kg^{-1} h^{-1}) was recorded by intermittent-flow respirometry (AutoResp®,

version 3.2.2) connected to a Witrox-4 oxygen meter (Loligo Systems). Probes were calibrated

weekly with a two-point, temperature-paired calibration (0% and 100% air saturation), and no signal

drift was detected during the experiment. All swimming trials were performed at the fish's

respective treatment temperature (24, 26, or 28.5 °C).

Fish were fasted for 24 h prior to the swimming test. Fish were individually transferred into the swim

chamber and acclimated by gradually increasing the water velocity over a 10-minute period. Each

fish then remained overnight in the swim chamber at a low water velocity (U = 0.7 Body Length per

second (BL s^{-1})) for habituation. Swimming and metabolic performances were assessed using a

critical swimming speed (U_{crit}) protocol, in which each fish was exposed to a stepwise increase in

water velocity, all performed at the same time of each day (e.g., 41,42,43). Swimming speeds were

increased stepwise by 0.3 BL s^{-1} after each complete respirometry cycle which consisted of a 3-

minute flush phase to maintain oxygen saturation (i.e., above 80% of air saturation), followed by a 1-

minute closed wait phase to allow flow stabilization, and a 10-minute closed oxygen measurement

phase at steady swimming speed, with the flush pump turned off. Swimming trials were terminated

when fatigue was reached, indicated by the fish's inability to swim against the current and the

adoption of a C-shaped posture on the grid located at the back of the swim chamber.

At the end of the stepwise protocol, males were removed from the swim chamber and euthanized

using MS-222 (0.5 g L^{-1}). For each male, total length, height, width, and body weight were

recorded, and the whole body was then stored at -20 °C for subsequent PFAS analysis. Swim

respirometers were cleaned between trials with bleach and rinsed three times with freshwater.

Background respiration (e.g., microbial respiration) was measured in the empty swim tunnel for at

least 20 minutes before and after each trial.

Calculation of fish metabolic rates and swimming performance.

Oxygen consumption (unscaled MO$_2$, mg O$_2$ kg^{-1} h^{-1}) of the fish was calculated using the following

equation (28):

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$$\text{Unscaled MO}_2 = \left[\left(\frac{\Delta[O_2]}{\Delta t}\right) - \left(\left(\frac{\Delta[O_2]}{\Delta t}\right)_{bact}\right)\right] \times \left(\frac{V_{resp}}{m}\right) \textbf{ equation (2)}$$

where $\Delta[O_2]/\Delta t$ (mg O_2 L^{-1} h^{-1}) is the rate of oxygen concentration decrease in the swim

respirometer over time during each MO_2 measurement period. Only slopes with a regression

coefficient greater than 0.85 were considered valid. $(\Delta[O2]/\Delta t)_{bact}$ (mg O_2 L^{-1} h^{-1}) represents the

background respiration slope, calculated as the mean of two measurements taken before and after

the swimming test in the empty chamber. V_{resp} is the respirometer volume (1.5 L) minus the volume

of the fish, and m (kg) is the fish body mass.

As fish respiration depends on animal body mass, metabolic rates were corrected to a standard body

mass of 0.1 kg using an allometric exponent of 0.68, consistent with values reported for *C.*

variegatus (e.g., 44, 45):

240
$$MO_2 = \text{Unscaled MO}_2 \times \left(\frac{m}{m_{corr}}\right)^{1-A} \textbf{ equation (3)}$$

where MO_2 (mg O_2 kg^{-1} h^{-1}) is the oxygen consumption standardized to a reference body mass m_{corr}

of 0.1 kg. Unscaled MO_2 (mg O_2 kg^{-1} h^{-1}, equation 2) is the uncorrected oxygen consumption

calculated for each fish with a body mass m (kg). A is the allometric exponent of the relationship

between metabolic rate and fish mass.

The Standard Metabolic Rate (SMR, in mg O_2 kg^{-1} h^{-1}) of each fish was estimated by plotting MO_2

against swimming speed (BL s^{-1}) and extrapolating the resulting linear regression to a null swimming

velocity (e.g., 41,42,46). The highest MO_2 recorded during the U_{crit} test was considered an estimate

of the Maximum Metabolic Rate (MMR, in mg O_2 kg^{-1} h^{-1}, e.g., 42, 47).

Aerobic scope (AS, mg O_2 kg^{-1} h^{-1}) was then calculated for each individual as the difference

between MMR and SMR (e.g., 28,41).

U_{crit} (BL s^{-1}) was calculated according to the following formula (41, 43, 48):

252
$$U_{crit} = U_t + \frac{t_1}{t} \times U_1 \textbf{ equation (4)}$$

where U_t (BL·s^{-1}) is the highest velocity that the fish maintained for a complete swimming step, t_1

(min) is the time spent at the fatigue velocity step, t (min) is the time of a complete swimming step

(i.e. 14 min) and U_1 is the last increase of the velocity before the fish fatigued (i.e., 0.3 BL·s^{-1}).

2.5 Chemical analyses

All samples (water and tissues) were analyzed for PFOS and PFOA concentrations at the University of

Connecticut's Center for Environmental Sciences and Engineering (CESE). Water samples were

processed according to U.S. EPA Method 537.1, while tissue samples were extracted and analyzed

following the protocol described in Campbell et al. (49). Analyses were conducted using an ACQUITY

UPLC system coupled to a tandem mass spectrometer (UPLC–MS/MS; Waters, Milford, MA, USA).

Standard quality assurance/quality control (QA/QC) procedures were followed, including analysis of

method blanks, duplicate samples, pre-extraction matrix spikes, and laboratory control samples. The

method reporting limits were 1.8 ng·L^{-1} for PFOA and 2.3 ng·L^{-1} for PFOS in water samples, and 0.49

ng·g^{-1} for PFOA and 0.35 ng·g^{-1} for PFOS in tissue samples. Concentrations are reported as ng·g^{-1} wet

weight for tissue samples and ng·L^{-1} for water samples.

2.6 Statistical analysis

All statistical analyses were performed in R (version 4.3.1). Selection between linear mixed models

(LMMs) and linear models (LMs) was guided by Akaike Information Criterion, Bayesian Information

Criterion, and likelihood ratio tests (using the ANOVA function). In cases where (1) the inclusion of

sex did not improve model fit, or (2) sex-related terms and interactions were not statistically

significant, the simpler model excluding sex was retained for clarity and ease of interpretation.

Diagnostic checks included visual inspection of residual and Q–Q plots to assess linearity,

homoscedasticity, and normality, supported by Shapiro–Wilk tests. When assumptions were not

met, variables were log-transformed and models refit. Mixed models were fitted using the lme4

package (lmer), and fixed effects were evaluated via Type III Wald chi-square tests using the car

package. Post hoc tests of interactions were performed with emmeans-tests with Tukey correction.

Statistical significance was defined as $p < 0.05$. The specific models used are described below.

2.6.1 Tissue concentrations, distribution, and temperature effects

To test whether tissue PFOS, PFOA, and ΣPFAS concentrations vary with temperature, LMMs were

used with temperature and PFAS treatments as fixed effects and aquarium identity as a random

effect to account for aquarium-level variation. In experiment 1, temperature was modeled either as

a continuous variable to assess linear trends (including both PFAS-exposed and non-exposed

individuals), or as a categorical factor to capture potential non-linear or threshold effects. In

experiment 2, temperature was modeled only as a categorical factor. PFAS concentrations in eggs

(Exp. 1) were not subjected to statistical analysis due to limited replication (n = 3 aquaria per PFAS

treatment group) and were considered exploratory.

To test whether temperature altered PFOS's proportional contribution to ΣPFAS within each tissue

among PFAS-exposed fish, LMMs were used in experiment 1 with temperature included as a fixed

effect and modeled either as a continuous variable or as a categorical factor, and aquarium was

included as a random effect. In experiment 2, comparisons between groups were conducted using

either Student's t-tests or Wilcoxon rank-sum tests, depending on the distribution of the data.

Temperature effects on PFAS distribution in PFAS-exposed fish (Exp. 2) were evaluated by analyzing

the percent share of PFAS concentrations by tissue and organ:blood concentration ratios. Percent

shares were modeled for each compound with LMs including temperature and tissue as fixed

effects. Organ:blood ratios were compared between temperatures using either Student's t-tests or

Wilcoxon rank-sum tests, depending on the distribution of the data.

2.6.2 Temperature effects on PFAS toxicity

To test whether fish length and weight were different between treatment groups, LMMs were used

in both experiments with temperature and PFAS treatment as fixed effects, and aquarium identity as

a random effect, while mortality was analyzed using LMs, with temperature and PFAS treatments as

fixed effects.

For the somatic indices and metabolic and swimming performances, statistical analyses were

structured in two main components. First, the main and interactive effects of temperature and PFAS

exposure on SMR, MMR, AS, U_{crit}, HSI, and GSI were quantified using LMs with temperature (24, 26,

28.5 °C) and PFAS treatment (non-exposed vs. PFAS-exposed) as categorical factors (3*2 design).

Second, to test whether individual PFAS concentrations predicts somatic indices and metabolic and

swimming performances, and whether this depends on temperature, LMs or LMMs were used using

log-transformed PFOS, PFOA, and ΣPFAS as continuous factors, and temperature as a categorical

factor. Aquarium identity was included as a random factor when appropriate. Simple per-

temperature linear regressions were then fitted to describe the slopes within each temperature

when the PFAS*temperature interaction was significant, indicating temperature-dependent PFAS

effects.

To test whether PFAS concentrations predicts daily egg production per female, and whether this

depends on temperature, LMMs were used. As a preliminary step, LMMs were fitted separately

within each temperature group during the pre-exposure period to assess potential baseline

differences in reproductive output prior to PFAS exposure. Since no significant differences were

detected at baseline, a LMM model was used during the exposure period to test the effects of

temperature, PFAS treatment (exposed vs. non-exposed), and their interaction. In addition, LMs

were used to compare the slopes of egg production over time across treatment groups, using both

daily and cumulative egg counts, to assess dynamic reproductive responses throughout the

exposure.

3. Results and Discussion

In the context of rising multi-stressor pressures on aquatic ecosystems, this study assessed how climate-driven temperature increases may alter PFAS toxicokinetics and compromise physiological performance in sheepshead minnows, a coastal species of ecological relevance. Our central question was how temperature modulates PFAS effects on fish fitness. Accordingly, in the sections that follow, we first quantify PFAS patterns and then evaluate how they vary with temperature. Experiments were conducted under environmentally relevant contaminant and temperature regimes, highlighting broader risks to aquatic health under global change.

3.1. Temperature modulation of PFAS toxicokinetics

The first objective of our study was to characterize how climate-driven temperature increases affects the tissue accumulation and distribution of PFOS and PFOA. In the first experiment, PFAS concentrations and distribution were assessed for eggs (Figure S2, Table S3), whole-body samples of swimming fish and muscle tissue (Table S4), and based on the observed patterns, a second experiment was designed to assess PFAS concentrations and distribution among muscle, blood, ovaries, and liver (Table S4).

PFOS dominance across tissues and maternal transfer.

PFAS (primarily PFOS) was detected in 87% of non-exposed fish tissues from both experiments with ΣPFAS concentrations between 8- and 70-fold lower than in PFAS-exposed fish, indicating a relatively minor source of PFAS (i.e., background contribution from source water or food; Table S4). After accounting for the low background, PFAS treatment significantly increased PFOS and PFOA concentrations in exposed vs. non-exposed fish across all tissues and temperatures (Treatment effect: all $p < 0.05$; Figure 1, Table S5). Compared with U.S. EPA monitoring data, muscle ΣPFAS in our study (Means 263–777 ng·g^{-1} ww; Table S4) were higher than national medians (11.8 ng·g^{-1} ww; PFOS-dominated; 50) yet below hotspot maxima (≤ 5,150 ng·g^{-1} ww near historical fluorochemical

facilities; 51,52). Thus, our values reflect substantial accumulation while remaining within the range

observed in environmentally impacted systems. Mirroring the PFOS-dominated pattern reported in

national monitoring, our samples were PFOS-dominated: PFOS accounted for 94–99% of ΣPFAS

across tissues and temperatures despite a 1:1 mixture (Figure 1), aligning with observations across

taxa including fish, birds and mammals (53). This dominance may be partly attributed to the stronger

protein-binding affinity of PFOS, as demonstrated in carp (*Cyprinus carpio*) by Zhong et al. (54), likely

due to stronger ionic interactions of the sulfonate vs. the carboxylate group, leading to slower

elimination and greater bioaccumulation. In line with the stronger protein-binding of PFOS described

above, the percent share of PFOS concentrations by tissue was highest in blood, next in liver, and

lower in ovary and muscle (Exp. 2; tissue effect, $p < 0.001$; Figure 2, Table S6), matching prior

evidence that PFOS accumulates preferentially in compartments rich in serum albumin and fatty

acid-binding proteins (17,55). By contrast, the share of PFOA concentrations in tissue was flatter,

with the largest share in blood and comparable shares across liver, muscle, and ovary (tissue effect:

$p < 0.001$; Exp. 2, Figure 2, Table S6). Although ovaries carried lower PFAS concentrations than blood

or liver (Figures 1-2), reproductive tissues still accumulated strongly over the exposure period. In

eggs (Exp. 1), ΣPFAS rose from 3.30–8.10 ng·g^{-1} ww at day 0 to 54.5–113 ng·g^{-1} ww at day 14 (13–34

fold higher), and further to 388–543 ng·g^{-1} ww at day 28 (48–165 fold higher from day 0; Figure S2,

Table S3). In ovaries (Exp. 2), ΣPFAS increased from 25.8–56.5 ng·g^{-1} ww in non-exposed females to

1,470–1,530 ng·g^{-1} ww in exposed females at day 28 (26–59 fold higher, treatment effect: $p = 0.032$;

Figure 1, Tables S4-5). Together with prior reports (56), our findings reinforce maternal transfer

through eggs/ovaries as a key PFAS exposure route.

Temperature shapes PFAS accumulation and maternal transfer.

Under warming, ΣPFAS varied in parallel with PFOS, reflecting PFOS dominance in tissues.

In Exp. 1, ΣPFAS and PFOS concentrations in PFAS-exposed fish declined with increasing water

temperature, moderately in whole-body (swimming males; ΣPFAS: estimate ± SE = –302 ± 150,

PFAS*temperature (continuous), $p = 0.052$; PFOS: –307 ± 150, $p = 0.049$; Figure 1, Table S5) and

significantly in muscle (both sexes; ΣPFAS: −97.7 ± 47, p = 0.045; PFOS: −98.8 ± 47, p = 0.041; Figure

1, Table S5). Muscle also showed a 28.5 °C threshold (PFAS*Temperature (factor), ΣPFAS: p = 0.021;

PFOS: p = 0.019; Table S5): ΣPFAS and PFOS at 28.5 °C were lower than at 24.0 °C (post hoc tests,

ΣPFAS: p = 0.035; PFOS p = 0.031) and 26.0 °C (ΣPFAS: p = 0.012; PFOS: p = 0.009), with no difference

between 24 and 26 °C (both p > 0.05). By contrast, for whole-body, the categorical model detected

no PFAS*temperature effects for ΣPFAS or PFOS (all p > 0.05; Table S5), although trends mirrored the

continuous analysis. The decrease in muscle and whole-body PFOS concentrations with warming is

consistent with temperature weakening of PFAS–protein binding (57) and suggests enhanced

elimination and/or redistribution of free PFOS to excretory organs. Experimental evidence on

temperature-modulated PFAS toxicokinetics in fish is extremely limited (25), though our results are

supported by observations in rainbow trout (*Oncorhynchus mykiss*), where elevated temperatures

increased PFOS clearance and hepatic redistribution, reducing muscle concentrations and decreasing

half-lives in multiple tissues (i.e., liver, brain, kidney; 25). In Exp. 2, the temperature-driven decline of

ΣPFAS and PFOS were not observed in muscle, blood, ovary and liver of PFAS-exposed females,

where ΣPFAS and PFOS remained stable with warming (all temperature effect and

PFAS*temperature interaction: p > 0.05; Figure 1, Table S5). The discrepancies in muscle PFOS

temperature responses between experiments likely reflect a plateau in muscle PFOS concentrations

(maximum binding capacity or steady state) reached at 24 °C in both experiments and maintained at

28.5 °C in Exp. 2. This interpretation is supported by comparable muscle PFOS concentrations across

Exp. 1 (24 °C) and Exp. 2 (24 °C, 28.5 °C; Table S4) and by unchanged organ-to-blood PFOS ratios

across temperatures in Exp. 2 (Figure 3, Table S7), which may indicate a redistribution plateau in

which uptake and elimination scale proportionally. A higher PFAS burden in Exp. 2 (not measured)

could have kept muscle near saturation at 28.5 °C, masking temperature effects. Further

investigations are needed to understand the temperature-related differences in muscle PFOS

concentrations, potentially including PFOS burden measurements.

In parallel, warming increased tissue PFOA concentrations in PFAS-exposed fish from both

experiments. In Exp. 1, muscle PFOA increased with temperature (both sexes; estimate ± SE = 1.71 ±

0.70; PFAS*temperature (continuous): p = 0.019; Figure 1, Table S5). Muscle also showed a 28.5 °C

threshold (PFAS*temperature (factor): p = 0.029; Table S5): PFOA at 28.5 °C exceeded 24 °C and 26

°C (post hoc tests, both p < 0.018), with no difference between 24 and 26 °C (p > 0.05). By contrast,

whole-body PFOA showed no statistically significant temperature effect (swimming males; all p >

0.05; Figure 1, Table S5). In Exp. 2 (females) warming from 24 °C to 28.5 °C significantly increased

PFOA in blood (2.4-fold; PFAS*temperature: p = 0.003) and ovary (4.3-fold; PFAS*temperature: p =

0.042), and showed non-significant increases in liver (~4.0-fold; PFAS*temperature: p = 0.139) and

muscle (~2.0-fold; PFAS*temperature: p = 0.108; Figure 1, Tables S4–S5). This change in PFOA

muscle concentrations with temperature may partly be explained by PFOS–PFOA competition for

shared and limited binding sites. While competition between PFOS and PFOA (or between two long-

chain PFAS) has not been experimentally demonstrated, such a hypothesis is supported by prior

evidence of antagonism between long- and short-chain PFAS compounds with overlapping binding

affinities (e.g., in zebrafish (*Danio rerio*); 58) and by their different polar head groups that likely

influence their binding affinities and transport kinetics (59). In Exp. 1, warming likely weakened

PFOS-protein binding, reducing muscle PFOS concentrations. This could have released common

binding sites, enabling greater PFOA association with muscle, in line with the observed increase in

muscle PFOA concentrations with warming. Whole-body PFOA concentrations remained stable,

which is consistent with redistribution toward muscle (or other tissues) rather than net

accumulation. In Exp. 2, by contrast, evidence points to PFAS saturation of binding sites in muscle at

both temperatures (i.e., plateau conditions, see above), with PFOS occupying shared sites. This could

explain that PFOA did not increase significantly with temperature in muscle of PFAS-exposed

females.

Overall, warming altered PFAS accumulation across tissues in PFAS-exposed fish in a compound-

specific manner: In Exp. 1, PFOS decreased with warming while PFOA increased. In Exp. 2, PFOA

increased whereas PFOS remained stable with warming. Nonetheless, across both experiments the

mixture composition shifted similarly with warming: the PFOS fraction of ΣPFAS declined with

temperature (i.e., PFOA's share increased), while PFOS remained dominant (all p < 0.05; Figure 1,

Table S5).

As warming increased PFOA's share, more PFOA was directed to reproductive tissues: in PFAS-

exposed females, the ovary:blood PFOA ratio was higher at 28.5 °C than at 24 °C ($p = 0.034$; Figure 3;

Table S7), whereas liver:blood and muscle:blood ratios were unchanged. To our knowledge, these

data provide novel experimental evidence that maternal PFAS transfer is temperature-sensitive,

adding to a growing body of work linking maternal PFAS transfer to adverse developmental

outcomes in offspring (60). Recent work in zebrafish, for example, demonstrated transgenerational

effects of PFAS exposure, where PFOA, PFOS, and their mixture altered larval behavior and induced

widespread changes in gene expression in the F1 generation (60). These findings underscore the

importance of understanding maternal transfer mechanisms under warming conditions and their

potential consequences for reproductive success, early-life health, and long-term ecological impacts.

Evidence on temperature-dependent PFAS toxicokinetics in fish is scarce, and to our knowledge, our

work provides the first experimental evidence that warming reconfigures tissue-specific

accumulation under co-exposure to a PFAS mixture (PFOS + PFOA) in fish, including reproductive

tissues, reshaping mixture composition toward a greater PFOA share.

Figure 1: PFAS concentrations (PFOS, PFOA, ΣPFAS; ng·g⁻¹) and PFOS contribution (%) in fish tissues across temperatures.

Panels are arranged left to right by compound (PFOS, PFOA, ΣPFAS) plus the PFOS fraction of ΣPFAS; and top to bottom by

tissue, with experiment 1 at the top (whole-body, then muscle) and experiment 2 below (muscle, blood, ovary, liver).

Crossbars represent group means ± SE. Circles represent individual fish (open circles = non-exposed, filled circles = PFAS-

exposed). Temperature groups are represented by color: 24°C in blue and 28°C in black. In experiment 1, lines show

linear mixed models (LMMs) fits with PFAS and temperature as continuous predictors. Lines are plotted only when the

interaction is significant with shaded 95% confidence intervals (solid = PFAS-exposed; dashed = non-exposed). In

experiment 2, PFAS concentrations were analyzed by LMMs with temperature as a factor. Where interactions are

significant, letters indicate pairwise differences (emmeans-tests with Tukey correction), and *p-values* for the main effect of

PFAS treatment are annotated when significant. For PFOS contribution panels, asterisks indicate significant differences

from Student's t-test or Wilcoxon test. Sample size: Exp. 1 - non-exposed fish: *n* = 9–12, PFAS-exposed fish: *n* = 17–24

(muscle) or 7–10 (whole-body); Exp. 2 - *n* = 8–9 fish per group. *Note:* the y-axis is scaled higher for blood and truncated for

muscle to enhance visibility.

Figure 2: Percent share of PFAS concentrations by tissue (PFOS, PFOA, ΣPFAS; %) in PFAS-exposed fish across

temperatures in experiment 2. Percent shares were calculated as the concentration in a given tissue divided by the sum of

concentrations in blood, liver, muscle, and ovary from the same fish. Crossbars represent group means ± SE. Circles

represent individual fish. Temperature groups are represented by color: 24°C in blue and 28°C in black. Letters indicate

pairwise differences among tissues (emmeans-tests). Sample size: n = 8–9 fish per group.

.

Figure 3: PFAS organ:blood concentration ratios (PFOS, PFOA, ΣPFAS) in fish from PFAS-exposed groups across temperatures in experiment 2. Ratios were calculated for muscle, liver and ovary relative to blood concentrations. Crossbars represent group means ± SE. Circles represent individual fish. Temperature groups are represented by color: 24□°C in blue and 28□°C in black. Asterisk indicate a significant difference between groups (Student's t-test). Sample size: n = 8–9 fish per group.

3.2. Temperature modulates PFAS exposure effects on fish fitness

The second objective of our study was to evaluate whether climate-driven temperature modulates

the toxicity of the mixture on fish metabolic, swimming and reproductive performance. To this end,

we used two complementary approaches: (i) full-factorial analyses of PFAS*temperature effects

(results for metabolic rates, somatic indices, body length/mass , and U_{crit} are provided in the SI;

Figures S3-S4-S5, Tables S8-S11-S13-S14), and (ii) analyses of the relationships between internal

PFAS concentrations and physiological traits across temperatures (Figures 4-5-6, Tables S9-S10-S12-

S15-S16). Here, we focus on the internal concentration–trait relationships, as these most directly

inform temperature-dependent toxicity.

End-century warming does not constrain aerobic capacity in unexposed minnows.

Aerobic scope represents the portion of oxygen supply that an individual can allocate to support

energy-demanding activities beyond basic maintenance and is widely considered ecologically

relevant (26, 29, 41, 61). In our study, we found no significant difference in AS (405.2 to

482.2 mg·kg^{-1}·h^{-1}), as well as in SMR (101.5 to 108.7 mg·kg^{-1}·h^{-1}) and in MMR (512.3 to

572.6 mg·kg^{-1}·h^{-1}) between 24 °C, 26 °C, and 28.5 °C in unexposed fish, indicating stable metabolic

performance across this temperature range (Figure S3, Table S8). The temperatures tested in our

study fall within an intermediate thermal range for *S. minnows*, between the cooler (21 °C) and

warmer (32 °C) conditions explored in previous studies. For instance, Kirby et al. (48) found a

significant increase of MMR and AS at 21 °C compared to 32 °C, suggesting that aerobic capacity

improves toward the upper end of this tested range, although no intermediate temperatures were

tested to assess where the peak occurs. By contrast, Jung et al. (62) found no significant difference in

RMR, MMR, or AS between 25 °C and 30 °C, indicating a performance plateau over this range. Our

lack of significant differences between 24 °C and 28.5 °C aligns with this plateau pattern, implying

that these temperatures fall within or near the species' optimal thermal window for maintaining

aerobic scope. Note that the values reported in the present study are lower than those in Kirby et al.

(48) and Jung et al. (62) because our data were allometrically scaled to account for individual body

mass, a step not applied in the comparative studies. Our results therefore highlight that projected

end-century warming within 24–28.5 °C is unlikely to constrain aerobic performance in unexposed

fish; this conclusion does not account for the multiple co-exposures typical of natural environments

(1).

Temperature modifies PFAS impacts on metabolic performance.

Whole-body PFOA concentrations showed no association with AS, SMR, or MMR and no interactions

with temperature (all $p > 0.05$; Figure 4, Table S9). By contrast, metabolic traits were associated with

whole-body ΣPFAS and PFOS, and these associations were temperature-dependent (all

PFAS*temperature: $p > 0.017$ across AS, SMR, MMR; Figure 4, Table S9). At 24 °C, PFOS and ΣPFAS

were not related to AS, SMR, or MMR (all $p > 0.475$; Table S10). At 26 °C, SMR and MMR increased

with PFOS and with ΣPFAS (all $p < 0.037$, all $R^2 > 0.34$; Table S10), such as AS showed no association

with PFAS concentrations (PFOS: $p=0.386$; ΣPFAS: $p=0.400$). These patterns indicate a PFAS-

related rise in oxygen demand at 26 °C that was matched by greater oxygen-supply capacity (higher

SMR/MMR), thereby maintaining aerobic scope. This compensatory response, along with the slightly

lower whole-body PFAS concentration relative to 24 °C (Figure 1), suggests added energetic costs of

active elimination (and other constraints), even though AS was maintained. Indeed, in fish, PFAS are

primarily eliminated via renal and biliary excretion, similar to other vertebrates (55). In mammals,

ATP-binding cassette (ABC) transporters actively mediate PFAS excretion (63), and these

transporters are highly conserved and expressed in fish detoxification organs, including liver, kidney,

and gills (64,65). Recent work in Atlantic cod (*Gadus morhua*) shows that exposure to single PFOS,

PFOA, and PFNA, or their mixture (total ~138 ug.L^{-1}), significantly modulates ABC transporter gene

expression in liver tissue (66). Together, these findings suggest that active, ATP-dependent transport

likely contributes to PFAS elimination in fish and may add to the energetic cost of detoxification,

although this cost remains to be quantified.

At 28.5 °C, PFAS concentrations were not related to SMR (PFOS: p = 0.934; ΣPFAS: p = 0.993; Table

S10), while both MMR and AS declined with increasing PFOS (MMR: p = 0.005, R^2 = 0.46; AS: p =

0.008, R^2 = 0.43) and ΣPFAS (MMR: p = 0.006, R^2 = 0.46; AS: p = 0.008, R^2 = 0.43). These patterns

indicate that the compensatory capacity observed at 26 °C failed at 28.5 °C, resulting in a mismatch

between oxygen demand and supply, as reflected by the decrease in MMR and aerobic scope,

despite a reduced whole-body ΣPFAS concentration (Figure 1). Physiological constraints that were

likely present but compensated for at 26 °C may have become critical, preventing fish from

maintaining sufficient oxygen uptake and delivery. PFOS exposure has been shown to alter

mitochondrial enzyme activity (e.g., citrate synthase, cytochrome c oxidase; 67), downregulate key

quality control genes (*pink1*, *fis1*), and reduce basal and ATP-linked oxygen consumption rates (68) in

fish species exposed to 0.1-1 mgL⁻¹. Such mitochondrial impairments could limit how much oxygen

can be used to support high metabolic demands at the whole-organism level, helping to explain the

decline in MMR and aerobic scope observed at 28.5 °C in our study. Furthermore, PFOS and PFOA

have been shown to cause structural damage to gill tissue, reducing oxygen diffusion efficiency

(67,69), and to induce cardiac oxidative stress and dysfunction that may constrain oxygen transport

capacity (70). Finally, lower ΣPFAS concentrations at 28.5 °C (Figure 1) may reflect reduced uptake

and/or enhanced passive loss rather than more efficient active elimination, given the limited oxygen

supply, and demonstrates that thermal stress reduces physiological tolerance, amplifying the impact

of a lower contaminant burden.

To our knowledge, few studies have quantified whole-organism metabolic rates in PFAS-exposed fish

(33, 34), and prior to ours none have tested the effects of a PFAS mixture on metabolic rates. Xia et

al. (33) reported no significant effects on MMR (SMR not assessed) in juvenile goldfish (*Carassius*

auratus) exposed for 48 h to 0.5 mg.L⁻¹ of PFOS, with effects only at the highest doses tested (32.0

mg.L⁻¹). Building on this, Xia et al. (34) demonstrated that temperature can change the dose-

response threshold for PFOS effects on resting metabolic rate in juvenile *Spinibarbus sinensis*: at

higher temperatures, lower doses triggered an increase in RMR (lowest observed effect

concentration: 5 mg.L⁻¹ at 18 °C vs. 0.8 mg.L⁻¹ at 28 °C after 40 days). However, no significant

effect of PFOS on MMR or aerobic scope was observed across temperature. In accordance with Xia

et al. (34), our results also demonstrate that a lower PFAS concentrations can become more

metabolically costly at higher temperatures. However, we found that thermal stress did not increase

SMR at higher temperature but instead constrained the ability to maintain peak oxygen supply

suggesting that specific metabolic endpoints and sensitivity thresholds may vary across species.

Figure 4: Relationships between whole-body PFAS concentrations (log-transformed PFOS, PFOA, ΣPFAS; ng·g⁻¹) and

metabolic traits (SMR, MMR, and AS; mg O₂ kg⁻¹ h⁻¹) or critical swimming speed (U_crit; BL s⁻¹) across temperatures in

experiment 1. Panels are arranged left to right by compound (PFOS, PFOA, ΣPFAS) and top to bottom by trait (SMR, MMR,

AS, U_{crit}). Circles represent individual fish. Temperature groups are represented by color: 24°C in blue, 26°C in orange,

and 28.5°C in black. Lines show linear mixed models (LMMs) fits with shaded 95% confidence intervals and with PFAS as a

continuous predictor and temperature as a factor. Where interactions are significant, *p-values* are reported. Sample size: n

= 16-18 fish per group.

Hepatic somatic index reveals sex- and context-dependent PFAS effects.

HSI showed context-dependent responses to PFAS. In experiment 1, the relationships between

muscle PFAS concentrations and HSI differed by sex (ΣPFAS*sex: p = 0.033; PFOS*sex: p = 0.029;

Figure 5, Table S12): females exhibited higher HSI with increasing muscle ΣPFAS concentrations,

driven by PFOS (ΣPFAS: slope = 0.081 ± 0.04; PFOS: slope = 0.082 ± 0.04, p < 0.05), whereas males

showed no association. PFOA was unrelated to HSI (ΣPFOA*sex: p = 0.185; Table S12). There was no

evidence that temperature modulated these relationships (PFAS*temperature, all p > 0.05). The

PFOS-driven HSI increase in females in Exp. 1 mirrors field observations: Piva et al. (71), reported

higher HSI in *Squalius cephalus* from highly contaminated freshwater sites compared to less polluted

locations, attributing it to hepatic hyperplasia and hypertrophy, likely reflecting an adaptive

response to elevated PFAS accumulation in the liver. The liver may increase in mass to enhance

detoxification capacity. Similar structural liver changes (vacuolization, steatosis, and cell

proliferation) have been well documented in mammals and zebrafish chronically exposed to PFAS

(72,73). In experiment 2 (females only), HSI was not related to muscle ΣPFAS, PFOS, or PFOA (all p >

0.065; Figure 5, Table S12). We did not observe PFAS*temperature interactions, though temperature

had an overall effect (lower HSI at 28.5 °C than at 24 °C; all p < 0.002 across PFOS, PFOA, ΣPFAS). The

lack of an HSI–PFOS association in experiment 2, despite a positive relationship in experiment 1, may

reflect reduced statistical power given the smaller sample size (n = 8–9 fish per group), and/or higher

internal PFAS exposure in experiment 2 that was not captured by muscle PFOS. Because muscle

PFOS at 24 °C was similar between experiments, it likely served as a weak surrogate for the hepatic

dose (the proximate driver of HSI) if true internal exposure was higher in experiment 2. In addition,

the lower baseline HSI at 28.5 °C may have compressed the response range, further flattening the

relationship. Taken together, our results suggests that PFAS, particularly PFOS, can induce liver

enlargement in females. This may reflect compensatory detoxification responses, although hepatic

alterations remain possible.

Figure 5: Relationships between muscle PFAS concentrations (log-transformed PFOS, PFOA, ΣPFAS; ng·g⁻¹) and somatic

indices (HSI and GSI; %) across temperatures in both experiments. Panels are arranged left to right by compound (PFOS,

PFOA, ΣPFAS); and top to bottom by somatic index (HSI, GSI), with experiment 1 on the upper row and experiment 2 on the

lower row for each index. Circles (females) and triangles (males) represent individual fish. Temperature groups are

represented by color: 24°C in blue, 26°C in orange, and 28.5°C in black. Lines show linear models (LMs) fits with shaded

95% confidence intervals and with PFAS as a continuous predictor and temperature as a factor (solid lines = females,

dashed lines = males). Sample size: Exp. 1 - n = 23-24 fish per sex and per temperature group; Exp. 2 - n = 17-18 females per

temperature group.

PFAS and temperature did not elevate mortality.

Mortality was low overall (Exp. 1: 41/684, 6%; Exp. 2: 15/105, 14%) and did not differ by PFAS

treatment across temperatures in either experiment (all $p > 0.251$), indicating no PFAS- or

temperature-related increase in mortality. Most deaths involved females (> 70%), consistent with

aggressive male-female interactions reported during the experiment, this pattern was accentuated

in Exp. 2 by greater size asymmetry between sexes, increasing female susceptibility to male

aggression.

PFAS-temperature interactions on ecological performance.

A lower aerobic scope should constrain energetically demanding activities. Accordingly, we

examined ecologically relevant performance metrics, starting with swimming performance,

quantified in males as critical swimming speed (U_{crit}, Figure 4), a standard integrative index of

aerobic capacity and overall fitness in fish. U_{crit} were not associated with whole-body PFAS

concentrations, or with temperature, and there were no significant interactions (all $p > 0.05$ across

PFOS, PFOA and ΣPFAS; Figure 4, Table S15). Overall, U_{crit} values in our fish (4.7–5.5 BL·s⁻¹ across 24–

28.5°C) were within published range for the same species (e.g., 42), though slightly lower than

those reported by Kirby et al. (48), likely reflecting methodological and scaling differences. The

absence of U_{crit} responses, despite temperature-dependent effects on AS/MMR is contrary to our

expectations because the stepwise U_{crit} test recruits aerobic slow muscle fibers and typically covaries

with aerobic metabolic performance in teleosts (e.g., mahi-mahi (*Coryphaena hippurus*), (74);

multiple damselfish species, (75); golden grey mullet (*Chelon auratus*), (43)). However, this

decoupling between U_{crit}, MMR, and AS aligns with prior work with the same species as our study. In

sheepshead minnows, Kirby et al. (48) found that acclimation to 32 °C increased MMR and AS

relative to 21 °C, yet U_{crit} did not increase. They hypothesized that sheepshead minnow's skeletal

muscle exhibits high thermal plasticity (including flexible nerve stimulation, contractile kinetics, force

generation, and enzyme activity), that may partly explain why this species shows consistent

swimming performance across a wide range of temperatures. Under PFAS exposure, Xia et al. (33)

similarly reported unchanged U_{crit} in goldfish despite elevated MMR at 32 mg.L^{-1} of PFOS, likely due

to reduced swimming efficiency caused by gill damage or disrupted glucose metabolism. Thus, U_{crit}

may be relatively robust to PFAS concentrations and temperatures tested in our study, any effects

on aerobic capacity could have been offset by compensatory locomotor strategies.

As reproduction is one such energy-demanding function and key component of fitness, both GSI

(Figure 5) and egg production (Figure 6) were also assessed to evaluate potential sublethal impacts

on reproductive investment. Mean daily egg production per female was not influenced by PFAS

exposure and temperature, and there were no significant interactions (all $p > 0.095$; Figure 6C, Table

S16). Similarly, egg production remained stable over time, with no acceleration or decline

attributable to PFAS exposure or temperature (slopes of daily and cumulative eggs production: all p

> 0.05; Figure 6D-E, Table S16). GSI was not associated with muscle PFAS concentration, and no

PFAS*temperature interactions were detected (both experiments; all $p ≥ 0.065$ across PFOS, PFOA,

and ΣPFAS; Figure 5, Table S12). In females from Exp. 2, however, GSI varied with temperature

overall (all $p ≤ 0.002$; Figure 5, Table S12), being lower at 28.5 °C than at 24 °C. In Exp. 1, GSI did not

differ between sexes (data pooled; note that females had an overall greater GSI in the factorial

model, Sex: $p = 0.005$; Table S11) and showed no overall temperature effect (Table S12). These

findings align with those of Gust et al. (76), who found no significant effect of PFOS (0.1 to 100

μg·L^{-1}) on parental or first-generation egg production in zebrafish. Conversely, other studies have

reported adverse reproductive outcomes. For instance, Suski et al. (77) exposed sexually mature

fathead minnows (*Pimephales promelas*) to PFOS for 42 days, reporting a reduction in male GSI at

44 µg·L⁻¹ and decreased fecundity in females at 140 µg·L⁻¹. Similarly, Kang et al. (78) reported

reduced egg production in Japanese medaka (*Oryzias latipes*) following co-exposure to 10 mg·L⁻¹

PFOA and 1 mg·L⁻¹ PFOS for 21 days. These discrepancies likely reflect differences in PFAS

concentrations, exposure duration, compound identity (single vs. mixture), and species-specific

sensitivity. Although we detected no measurable effects on reproductive output or GSI in our fish,

we cannot exclude the possibility of delayed or transgenerational effects. For instance, Lee et al. (21)

exposed *O. latipes* to a mixture of PFOS, PFOA, PFBS, and PFNA at 0.5 µg·L⁻¹ over three generations

(238 days), and reported significant reproductive alterations (e.g., inhibition of hatchability,

induction of VTG expression).

Figure 6: Egg production across temperatures and PFAS exposure conditions in experiment 1. (A) Daily egg production

(eggs·female⁻¹·day⁻¹) and (B) cumulative egg production (eggs·female⁻¹) per female over the 28-day exposure period. (C)

Mean daily egg production per female (eggs·female⁻¹·day⁻¹). Rate of (D) daily and (E) cumulative egg production per

female, estimated as the slope of daily and cumulative egg counts over time (Δ eggs·female⁻¹·day⁻¹). Points and crossbars

represent group means ± SE. Circles represent individual fish (open circles = non-exposed, filled circles = PFAS-exposed).

Temperature groups are represented by color: 24 °C in blue, 26 °C in orange, and 28.5 °C in black. Sample size: n = 5-6

aquaria   per   group.

By combining thermal stress and chronic PFAS exposure, this study demonstrates that predicted

climate warming can profoundly reshape contaminant dynamics and alter physiological performance

in a key coastal fish species. To our knowledge, this is the first experimental demonstration that

warming modifies both the toxicokinetics (internal distribution and maternal transfer) and the

physiological toxicity of a PFAS mixture composed of PFOS and PFOA. Temperature altered PFAS

accumulation in a compound- and tissue-specific manner, shifting the mixture toward PFOA.

Notably, warming promoted PFOA redistribution toward reproductive tissues and increased

maternal transfer to eggs. Importantly, beyond a thermal threshold, fish lose their ability to fully

compensate for the metabolic cost of contamination, resulting in reduced aerobic capacity. Although

swimming and reproductive outputs were not impaired under the tested conditions, the observed

physiological alterations, such as liver enlargement and changes in tissue distribution, point to

hidden energetic costs associated with detoxification. These results indicate that even exposure

conditions under environmentally relevant concentrations combined with warming temperatures

may increase the vulnerability of estuarine fish to contaminant stress, with implications for the

resilience and ecological fitness of coastal fish populations and potentially for the development and

survival of their offspring.

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Margot Grimmelpont: Methodology – Conceptualization, Investigation, Formal analysis, Writing –

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Anthony Provotas: Chemical analysis; **Jessica Brandt:** Methodology – Conceptualization, Writing –

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Data Availability Statement

The dataset underlying this study has been deposited in Zenodo (DOI:

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