#### Check for updates

#### OPEN ACCESS

EDITED BY Chengzhong Yang, Chongqing Normal University, China

REVIEWED BY Florian Mauduit, University of California, Davis, United States Edit Eszterbauer, HUN-REN Veterinary Medical Research Institute, Hungary

\*CORRESPONDENCE Itziar Estensoro itziar.estensoro@csic.es

RECEIVED 13 February 2025 ACCEPTED 05 June 2025 PUBLISHED 27 June 2025

#### CITATION

Estensoro I, López-Gurillo B, Tafalla C, Craig R, Atkinson S, Sitjà-Bobadilla A and Bartholomew J (2025) Better the devil you know: sympatric fish host copes better than allopatric with a myxozoan parasite. *Front. Cell. Infect. Microbiol.* 15:1576014. doi: 10.3389/fcimb.2025.1576014

#### COPYRIGHT

© 2025 Estensoro, López-Gurillo, Tafalla, Craig, Atkinson, Sitjà-Bobadilla and Bartholomew. This is an open-access article distributed under the terms of the Creative Commons Attribution License (CC BY). The use, distribution or reproduction in other forums is permitted, provided the original author(s) and the copyright owner(s) are credited and that the original publication in this journal is cited, in accordance with accepted academic practice. No use, distribution or reproduction is permitted which does not comply with these terms.

## Better the devil you know: sympatric fish host copes better than allopatric with a myxozoan parasite

Itziar Estensoro<sup>1\*</sup>, Beatriz López-Gurillo<sup>1</sup>, Carolina Tafalla<sup>2</sup>, Ryan Craig<sup>3</sup>, Stephen Atkinson<sup>3</sup>, Ariadna Sitjà-Bobadilla<sup>1</sup> and Jerri Bartholomew<sup>3</sup>

<sup>1</sup>Fish Pathology Group, Instituto de Acuicultura Torre de la Sal, Consejo Superior de Investigaciones Científicas (IATS, CSIC), Castellón, Spain, <sup>2</sup>Biotechnology Department, Instituto Nacional de Investigación y Tecnología Agraria y Alimentaria (INIA), Consejo Superior de Investigaciones Científicas (CSIC), Madrid, Spain, <sup>3</sup>Department of Microbiology, Oregon State University, Corvallis, OR, United States

Ceratomyxosis due to the myxozoan parasite Ceratonova shasta affects salmonids, causing severe enteritis leading to hemorrhaging and necrosis. The waterborne parasite stages penetrate host gills and reach the fish intestine through the bloodstream. Steelhead trout (Oncorhynchus mykiss) populations from C. shasta endemic watersheds (sympatric) are less susceptible than populations from non-endemic watersheds (allopatric). We investigated the immune effectors behind these different susceptibilities. Both steelhead trout strains were exposed to C. shasta genotype O. Intestinal tissue and serum samples of control and exposed sympatric and allopatric fish were taken at 8, 15, 22, 29, 57, and 183 days post exposure. Gills were taken at 1 and 8 days post exposure. Parasite abundance and histopathology were evaluated on tissue sections of fish that tested PCR+ for the parasite. Zap70<sup>+</sup> T cells, IgT<sup>+</sup> B cells, and IgD<sup>+</sup> B cells were quantified, and the presence of specific IgM was evaluated from fish sera by immunohistochemistry. Parasite counts were significantly lower and limited to a shorter duration in sympatric fish. The initial intestinal inflammatory response in both fish strains was characterized by hyperplasia of the lamina propria-submucosa and epithelial infiltration of lymphocytes. Remarkably, hyperplasia was resolved earlier in sympatric fish, whereas in allopatric fish, hyperplasia was not resolved by the end of the experiment, coinciding with high intestinal parasite counts and sporogenesis. An increase of intestinal Zap70<sup>+</sup> T cells occurred before IgT<sup>+</sup> B cells peaked, earlier in sympatric than in allopatric fish. Low numbers of intestinal  $IgD^+$  B cells were detected in both strains. In gills, an early increase of Zap70<sup>+</sup> T cells was observed in both fish strains at 1 day post exposure, and an increase of IgD<sup>+</sup> B cells occurred only in the allopatric strain. Specific circulating IgM was detected much later in both fish strains, only at 57 and 183 days post exposure. In conclusion, sympatric steelhead trout restricted intestinal proliferation of C. shasta genotype

0 more effectively than allopatric fish, probably due to an earlier T cell response triggering a stronger IgT-based mucosal adaptive response in the intestine. Specific circulating IgM appeared later in both sympatric and allopatric fish, probably also contributing systemic protection.

#### KEYWORDS

*Ceratonova shasta*, genotype 0, intestinal inflammatory response, steelhead trout strains, lymphocytes, specific IgM, IgT, IgD



Overview of infection and response kinetics of sympatric (SYM, green) and allopatric (ALLO, yellow) steelhead trout exposed to *Ceratonova shasta* G0, during a 183-day experimental parasite challenge. Red triangles pointing up represent increase of a given parameter.

#### **1** Introduction

The myxozoan parasite *Ceratonova shasta* is endemic across the Pacific Northwest of the U.S. and Canada, and comprises at least three genotypes that affect different species of salmon and trout (Atkinson and Bartholomew, 2010b, a; Stinson and Bartholomew, 2012). Much of the research on *C. shasta* host–parasite interactions has focused on the more virulent genotypes, which affect endangered and threatened salmon populations: genotype I (GI), with specificity for chinook salmon (*Oncorhynchus tshawytscha*) (Hurst and Bartholomew, 2012; Bjork et al., 2014), and genotype II

(GII), which has a broader host range with coho salmon (*Oncorhynchus kisutch*) as the primary host (Barrett and Bartholomew, 2021; Barrett et al., 2021; Taggart-Murphy et al., 2021).

Genotype 0 (G0) causes non-lethal infections in rainbow trout/ anadromous steelhead trout (*O. mykiss*) exclusively. G0 infections are rarely documented in *O. mykiss* native to waters where the parasite is endemic (sympatric). The ability of rainbow trout native to rivers where *C. shasta* is absent (allopatric) to survive G0 infections supports the low virulence of this genotype (Atkinson and Bartholomew, 2010b). In contrast, allopatric salmonid strains are highly susceptible to infections by GI (chinook salmon) and GII (coho salmon, rainbow trout) and have served as infection models because of the low dose required to cause disease and mortality (Atkinson and Bartholomew, 2010a, b; Bjork and Bartholomew, 2010; Hurst and Bartholomew, 2012; Bjork et al., 2014).

Because of its low virulence, G0 infections have received less attention than GI and GII. However, there are several aspects of this genotype that make it unique and a compelling target for research. Evolutionarily, the ancestral host-parasite relationship for C. shasta was likely O. mykiss and G0 (Breyta et al., 2020), and understanding this relationship may provide insight into how these hosts and parasites co-exist. In contrast to the acute and lethal infections observed with GI and GII infections, G0 causes a chronic infection that culminates in long-term proliferation of the parasite in the intestinal lumen (Taggart-Murphy et al., 2021). This suggests a mode of transmission that would allow continuous parasite dissemination throughout the migratory range of the host, in contrast to a transmission model for GI and GII that relies largely on the death of the fish host, and therefore a point source dissemination (Bartholomew et al., 2022). This also suggests that O. mykiss has evolved parasite-specific defenses, and identification of these traits could inform selection of C. shasta-resistant fish strains and enhancement of protective host responses.

To better understand how C. shasta G0 interacts with O. mykiss, we challenged a strain of steelhead trout sympatric with C. shasta (i.e., adapted host) and an allopatric strain (i.e., non-adapted "susceptible" host) with a known parasite dose in the laboratory. We characterized host-parasite dynamics in these adapted and non-adapted hosts focusing on the parasite distribution and proliferation kinetics and on the host immune response at the portal of entry, the gills, and the main parasite target, the intestine. Histopathological inflammatory alterations and central immune effectors involved in the mounting of the adaptive immune response were studied. Among the cellular effectors, we analyzed activated T cells, which mediate B cell activation and antibody production upon specific antigen recognition (Laing and Hansen, 2011), and B cells, responsible for the specific immune response. We focused our B cell analysis on the IgT<sup>+</sup> subset, since IgT is the teleost-exclusive immunoglobulin isotype, specialized in mucosal immunity, and on the IgD<sup>+</sup> subset, since although little is known about its tissue distribution and dynamics upon infection (Holzer et al., 2021), IgD has also been associated with mucosal responses in fish (Castro et al., 2014; Perdiguero et al., 2019; Herranz-Jusdado et al., 2023). Generally, IgM<sup>+</sup> B cells are most abundant in hematopoietic organs and blood of teleosts, although they are also present in mucosal tissues. Therefore, soluble IgM prevails in blood and was evaluated here in the steelhead sera.

#### 2 Materials and methods

#### 2.1 Fish

Strains of steelhead were obtained from hatcheries in Oregon, USA, where *C. shasta* is endemic (sympatric) or absent (allopatric):

allopatric winter steelhead from Alsea Hatchery (44°25'N, 123°33' W; ALLO; allopatric strain) and sympatric summer steelhead from Round Butte Hatchery (44°60'N, 121°27'W; SYM; sympatric strain). Fish were transported in coolers with aerated, UVsterilized well water ~13°C and held at the Oregon State University (OSU) John L. Fryer Aquatic Animal Health Laboratory (AAHL) in 380-L circular tanks on flow-through UVsterilized well water. At the start of exposure, ALLO fish had an average weight of 23.8 g and a fork length of 13.7 cm; SYM fish had an average weight of 31.5 g and a fork length of 14.3 cm.

## 2.2 Infection of annelid host and harvest of G0 for fish exposures

Populations of the annelid host Manayunkia occidentalis were collected at two sites in the Klamath River, CA, USA (42°07'03.3"N 122°03'03.8"W and 41°51'35.6"N 122°34'00.3"W) (Alexander et al., 2016). Annelids were held at the AAHL in two, 60-L mesocosm tanks partially filled with gravel/cobble substrate and supplied with flow-through, UV-sterilized Willamette River water as a nutrition source. The annelids were held for 12 months to purge any existing parasite. During this time, effluent was sampled weekly (1 L), filtered (at 5 µm), and analyzed by qPCR (Hallett and Bartholomew, 2006). Once samples of effluent consistently tested negative for C. shasta, annelids were inoculated with C. shasta myxospores from intestines and fecal matter of rainbow trout infected with G0 by exposure in the Klamath River (Alexander et al., 2016). Prior to adding myxospores to the mesocosm, a sample was sequenced (Atkinson et al., 2018) to confirm that only G0 was present. Annelids were dosed with G0 myxospores weekly for 12 wk. During this time, samples of effluent were collected weekly, following a 24-h static period, and analyzed by qPCR to quantify G0 actinospore production.

Seeding of mesocosms with infected tissue was ceased 10 d prior to fish exposure to determine infectious dose in water samples and minimize contribution from myxospore stages. For a week prior to exposure, mesocosms were maintained in static conditions with aeration. Approximately 10 L of water was removed from each mesocosm every 24 h, and the tanks were refilled with UV sterilized Willamette River water. For the exposure, ~130 L of water harvested from the mesocosms were stored in an aerated, 190-L polyethylene container. Parasite spores were up to 1 wk old at the time of exposure to fish. Three 1-L samples analyzed by qPCR measured an average density of 75 C. *shasta* actinospores/L.

#### 2.3 Fish infection

Fish exposures were randomized across 16 100-L tanks, with 4 tanks for exposure of each steelhead strain and 4 tanks containing controls of each strain. 130 L of harvested mesocosm water (containing 75 C. *shasta* actinospores/L) was divided into 8 aerated exposure tanks (16.5 L per tank; ~1238 actinospores/ tank). An equivalent amount of 13°C UV-sterilized well water

was added to 8 aerated control tanks. 104 fish from each strain were randomly assigned to exposure and control tanks with 4 replicates of 16 fish exposed per strain (~78 actinospores/fish) and 4 replicates of 10 control fish per strain. Fish were held in the aerated bath exposures for 24 h, after which 13°C UV-sterilized well water flow was resumed. Temperature of flow through water was gradually increased to 18°C over 3 d. Fish were held for 183 d, fed daily, and monitored for disease signs and mortality.

#### 2.4 Sample collection

Sampling of exposed and control fish occurred on 1, 8, 15, 22, 29, 57, and 183 days post-exposure (dpe). One fish from each control group and two fish from each exposure group were selected randomly and killed using MS222 (0.1 g/L; Sigma-Aldrich). Blood was drawn from the caudal vessels with a heparinized syringe, and transferred to 2-mL tubes. The blood was refrigerated overnight and then centrifuged at  $3000 \times g$  for 5 min. Serum was drawn off using a pipette, transferred to 0.5-mL vials, and frozen at  $-20^{\circ}$ C for determination of specific antibodies.

Fish were dissected immediately after blood collection, and 3 transverse sections of intestine were removed. The first (posterior) and third (anterior) sections were cut to ~0.5 cm and placed into a 2-mL tube containing RNAlater, refrigerated overnight, and then divided into  $2 \times 2$ -mL tubes and frozen—one tube for transcriptome analysis, the other for qPCR. The second, middle section of intestine, was cut to 2–3 cm and placed in buffered formalin for 48 h before transfer to 70% ethanol for histology. Tools used for dissections were soaked in H<sub>2</sub>O<sub>2</sub> for 3–5 min and then rinsed in deionized water between samples.

At 1 and 8 dpe, two gill arches were removed from each fish. One was placed in buffered formalin for 48 h before transfer to 70% ethanol for histology; the other was placed in a 2-mL tube and frozen for qPCR.

For mortalities that occurred during the study, skin scrapings and intestinal swabs were examined microscopically for external parasites and presence of *C. shasta* myxospores, respectively. Tissue samples of the intestine were taken for PCR assay.

#### 2.5 PCR and qPCR

Fish intestinal tissue samples were digested using a modified "boiled-crude" method of Palenzuela et al. (1999): incubation at 56°C for 1–2 h with 195  $\mu$ L buffer ATL (Qiagen) and 5  $\mu$ L proteinase K (20 mg/mL) to digest tissue, followed by heat denaturation at 85°C for 15 min, and then dilution 1:100 prior to assay by PCR (Cs1479F GCATCACCTGCTCTGAGAAGAGTGG, Cs2067R GGTCTTCATCGATGTTTTTGCCGAGG) (Atkinson and Bartholomew, 2010a; Atkinson et al., 2018). Parasite genotype was determined by sequencing PCR products using an ABI BigDye Terminator Cycle Sequencing Kit v3.1 and ABI3730 Genetic Analyzer (Applied Biosystems, Foster City, CA, USA) at the Center for Quantitative Life Sciences, CQLS, OSU (Atkinson and Bartholomew, 2010b). Fish gill samples were processed using the Qiagen DNeasy Animal Tissue Kit (Qiagen) using the manufacturer's instructions. qPCR was used to determine presence and amount of parasite in gill samples and in 1-L water samples, and was performed using a modified *C. shasta* SSU rDNA qPCR assay (Cs1034F CCAGCTTGAGATTAGCTCGGTAA, Cs1104R CCCCGGAA CCCGAAAG, TaqMan Probe Cs1058T 6FAM-CGAGCCAAGTT GGTCTCTCCGTGAAAAC-TAMRA) (Hallett et al., 2012; Alama-Bermejo et al., 2019). For gill tissue samples, we used the same DNA amount per reaction (134 ng) and ran a four-point 10-fold dilution standard curve of a purified PCR product to calculate SSU rDNA copy numbers of the parasite. All samples were run in triplicate, with a positive *C. shasta* sample as an interplate calibrator, and a no-template control. The corresponding formalin-fixed samples of fish tissue that tested positive for *C. shasta* DNA by PCR or qPCR were selected for histopathological assessment.

#### 2.6 Histopathological analysis

Formalin-fixed gill and intestine samples were routinely dehydrated and processed for paraffin histology. One gill and two intestinal cross-sections were cut at 4 µm, stained with Giemsa, and then used to evaluate parasite prevalence and intensity of infection. Intensity was scored from 1 to 6 based on the number of parasite stages observed in the entire tissue section: 1 = 1-25 parasite stages; 2 = 26-50 parasite stages; 3 = 51-100 parasite stages; 4 = 101-150parasite stages; 5 = 151-200 parasite stages; 6 = >200 parasite stages. In intestines, proliferative trophozoites that infiltrated into the lamina propria–submucosa or epithelium, free luminal proliferative trophozoites, and free luminal disporoblasts were scored separately.

Intestinal thickness was measured from the outer *muscularis* layer to the epithelium in intestinal cross-sections, including *muscularis* layers, *stratum granulosum*, *stratum compactum*, and lamina propriasubmucosa. Intestinal hyperplasia was evaluated by measuring the surface area of the lamina propria–submucosa in intestine crosssections. Lymphocyte epithelial infiltration in the intestine was scored ranging from 0 (absence) to 3 (very abundant: 25–30 intraepithelial lymphocytes [IELs] per microscope field at 500× magnification).

Observations were performed with a Leitz Dialux22 (Leica, Hesse, Germany) light microscope, and images were taken with an Olympus DP70 Camera (Olympus, Tokyo, Japan). Thickness and surface measurements were performed with ImageJ.

#### 2.7 Lymphocyte immunohistochemistry

Paraffin-embedded tissue sections were collected on Superfrost Plus slides (Menzel-Gläser, Braunschweig, Germany) for immunolabeling of lymphocytes: Zap70+ T cells, IgT<sup>+</sup> B cells, and IgD<sup>+</sup> B cells were analyzed. Details on antibodies are included in Table 1.

Washing steps between incubations consisted of successive 5-min immersions in TTBS (20 mM Tris-HCl, 0.5 M NaCl, 0.05% Tween 20, pH 7.2) and TBS (20 mM Tris-HCl, 0.5 M NaCl, pH 7.2), and incubations were made in a humid chamber at room temperature

Primary antibody	Origin	Producing species	Concentration (µg/mL)	Epitope retrieval buffer	Incubation conditions
Anti-human Zap70 *	Cell signaling	Rabbit	1:50 <sup>§</sup>	Sodium citrate, pH 6	ON, 4°C
Anti-trout IgT *	Ballesteros et al. (2013)	Mouse	3	Tris-EDTA, pH 9	ON, 4°C
Anti-trout IgD *	Ramírez-Gómez et al. (2012)	Mouse	5	Tris-EDTA, pH 9	1 h, RT
Anti-trout IgM	Ballesteros et al. (2013)	Mouse	10	Tris-EDTA, pH 9	1 h, RT

TABLE 1 Primary antibodies used for immunolabeling of trout lymphocytes (\*) and parasite stages.

ON, overnight; RT, room temperature; <sup>§</sup>manufacturer's instructions.

unless otherwise stated. Tissue sections were routinely deparaffinized and hydrated, before endogenous peroxidase was quenched in 0.3% H<sub>2</sub>O<sub>2</sub> for 30 min. Heat-induced epitope retrieval was carried out by submerging the slides in the corresponding buffer solution and then heating them in a microwave oven for 5 min at 800 W and 5 min at 450 W. Tissues were incubated with primary antibodies diluted in 1% BSA in TBS, then with the corresponding biotinylated secondary antibody (horse anti-mouse or goat anti-rabbit), and then with avidin-biotin-peroxidase complex (ABC) following the manufacturer's instruction (Vector Laboratories, CA, USA). Finally, bound peroxidase was visualized after 2 min incubation with 3,3'diaminobenzidine tetrahydrochloride chromogen (DAB, Sigma-Aldrich, MO, USA), and the reaction was stopped with deionized water. Slides were counterstained with Gill's hematoxylin, dehydrated, and mounted with di-N-butyl phthalate in xylene. Negative controls omitted the primary antibodies, secondary antibody, and ABC. Zap70<sup>+</sup>, IgT<sup>+</sup>, and IgD<sup>+</sup> immunoreactive lymphocytes were counted in 10 random microscope fields at ×500 magnification for each fish (2 control ALLO, 2 control SYM, 4 exposed ALLO, and 4 exposed SYM), and means were calculated for each fish and tissue.

Specific circulating IgM against C. shasta in the sera of the ALLO and SYM fish was detected immunohistochemically on parasite stages of intestinal sections with high infection intensity (score 6) in an indirect immunohistochemical sandwich ELISA. Briefly, endogenous peroxidase was quenched in tissue sections as described above. Sections were then incubated with pooled serum samples (N = 4) of each time point and strain for exposed fish, and 1 unexposed control serum pool (N = 4) of each strain. 16 serum pools (7 exposure times  $\times$  2 fish strains and 1 unexposed control  $\times$  2 fish strains) were tested, diluted 1:25 in 1% BSA. Parasite-bound specific Ig of sera was immunolabeled as described above, i.e., sections were incubated with mouse anti-trout IgM or IgT antibodies, horse antimouse biotinylated antibody, ABC, and DAB, and slides were then counterstained and mounted. The presence and intensity of immunoreactivity in the different parasite stages were evaluated, and representative images were taken.

#### 2.8 Transmission electron microscopy

Four intestine samples of infected ALLO steelhead trout were fixed immediately in 2.5% glutaraldehyde in 0.1 M sodium cacodylate buffer at pH 7.2 for 24 h at 4°C and washed in the same buffer. Post-fixation was carried out for 1 h in 1% osmium tetroxide in the same buffer, and samples were dehydrated and embedded in Spurr's resin. Ultrathin sections were contrasted in uranyl acetate and viewed using a Hitachi HT7800 transmission electron microscope (TEM) operating at an accelerating voltage of 120 kV.

#### 2.9 Statistics

Data on the observed intestinal histopathological alterations (intestinal thickness, hyperplasia, and IEL scoring) and lymphocyte counts in gills and intestines (Zap70<sup>+</sup> T cells, IgT<sup>+</sup> B cells, and IgD<sup>+</sup> B cells) were analyzed for statistical significance. First, Student ttests were performed to compare SYM vs. ALLO within the control unexposed experimental fish. Since no differences were found (Supplementary Table 1), these data were pooled into a single control unexposed experimental group for the subsequent analyses. Secondly, differences were analyzed between data of control unexposed, exposed SYM, and exposed ALLO experimental groups within each time point by one-way analysis of variance (ANOVA-I) followed by Student-Newman-Keuls test, and for intestines also along the six time points. Data that failed the normality or equal variance test were analyzed with Kruskal-Wallis ANOVA-I on ranks followed by Dunn's method. Data on lymphocyte counts in gills were compared by Student *t*-test or by Mann-Whitney U sum test for non-normally distributed data. In addition, data of the different intestinal alterations and lymphocyte counts were pooled regardless of the infection time for the three experimental groups and compared by ANOVA-I followed by Student-Newman-Keuls test, or by Kruskal-Wallis ANOVA-I on ranks followed by Dunn's method for non-normal distributed data.

Data on parasite intestinal infection intensity for the different parasite stages (intratissue proliferative trophozoites, luminal proliferative trophozoites, and luminal disporoblasts) were analyzed for differences between SYM and ALLO fish within each time point by Student *t*-test or by Mann–Whitney *U* sum test for non-normal distributed data, and along the six time points by ANOVA-I followed by Student–Newman–Keuls test, or by Kruskal–Wallis ANOVA-I on ranks followed by Dunn's method for non-normal distributed data. SigmaPlot v14.5 software (Systat Software Inc., CA, USA) was used for statistical analyses, and the significance level was set at P < 0.05.

#### 2.10 Ethical approval

Fish exposures were conducted at the AAHL under Animal Care and Use (ACUP) permit #21-0141.

#### **3** Results

## 3.1 Infection outcome: molecular detection and intestinal parasite counts

Infection was detected in the gills of exposed fish by qPCR as early as 1 dpe, in both fish strains. When intestines were first sampled at 8 days post-exposure, only one ALLO sample was positive; by 15 dpe, all exposed fish were PCR positive. Fish remained infected throughout the 183 days of the experiment. In the gill sections of 1 dpe and 8 dpe fish, no C. shasta stages could be found. In intestines, however, proliferative parasite stages were found from 15 dpe onward (Figures 1, 2; Supplementary Table 2). Low numbers of proliferative trophozoites migrating through the lamina propria-submucosa or epithelium were observed in SYM fish from 15 dpe to 57 dpe, whereas in ALLO fish, they were present from 15 dpe to 183 dpe (Figures 1A, 2). Two different developmental stages of C. shasta, namely, proliferative trophozoites and disporoblasts with mature spores, were observed in the intestinal lumen of exposed fish, either free or attached to the mucosal surface (Figures 1B, 2). Infection intensity with such luminal trophozoites was highest in SYM fish at 29 dpe and decreased later on. In ALLO fish, highest counts of luminal trophozoites occurred also at 29 dpe and were sustained until 183 dpe, a time point at which C. shasta trophozoites were significantly more numerous in ALLO than in SYM. Parasite disporoblasts appeared first at 29 dpe in both trout strains, though with significantly higher numbers in ALLO than in SYM. After 29 dpe, *C. shasta* disporoblasts persisted only in ALLO fish (Figures 1B, 2). Limited mortality occurred following exposure in both exposed SYM and ALLO groups but appeared to be non-specific and not related to *C. shasta* infection. All control fish were negative for *C. shasta*.

#### 3.2 Histopathological observations

In gills, no significant histopathological alterations were observed. In intestines, an overall increase in thickness was observed from 22 dpe onward in exposed steelhead trout due significant hyperplasia of the lamina propria-submucosa (Figures 3A, B). Thickness of the underlying intestinal layers (muscularis, stratum granulosum, and stratum compactum) did not change (Supplementary Figure 1). Hyperplasia was significant in SYM fish earlier than in ALLO, in which it became significant only from 29 dpe onward. By 57 dpe, intestinal hyperplasia peaked in ALLO fish, but had subsided in SYM fish (Figures 2, 3B). Overall, the increased intestinal thickness due to the hyperplasia of the lamina propria-submucosa was significant in both strains, regardless of the infection timing, when compared to control unexposed (CTRL) fish (Supplementary Figures 2A, B). The inflammatory response in the intestines of exposed fish included a significant infiltration of lymphocytes in the intestinal epithelium (Figures 2, 3C; Supplementary Figure 2C). This increase of intraepithelial lymphocytes (IELs) occurred in both fish strains and was significant in SYM fish from 15 dpe on and in ALLO fish from 22 dpe onward. Sections of CTRL intestines are shown in Supplementary Figure 3 for comparison.



#### FIGURE 1

*Ceratonova shasta* infection intensity in exposed sympatric (SYM) and allopatric (ALLO) steelhead trout intestines along the parasite challenge. (A) Proliferative trophozoites infiltrated in the lamina propria–submucosa or epithelium. (B) Free luminal parasite stages: proliferative trophozoites and disporoblasts. Infection intensity score is the number of parasite stages observed in the entire intestinal cross-section: 1 = 1-25; 2 = 26-50; 3 = 51-100; 4 = 101-150; 5 = 151-200; 6 = 200-300. Lowercase and uppercase letters indicate significant differences among sampling times within SYM or ALLO fish, respectively. Asterisks indicate significant differences between SYM and ALLO fish within a time post-exposure. P < 0.05. Scale bars =  $20 \, \mu$ m.



observed at 29 dpe, including both proliferative trophozoites (D<sub>1</sub>) and disporoblasts with mature spores (D<sub>3</sub>). Intraepithelial proliferative trophozoites (D<sub>2</sub>) were also present. At 57 dpe, intraepithelial proliferative trophozoites and lymphocytes (E<sub>1</sub>), and luminal proliferative trophozoites and disporoblasts with mature spores (E<sub>2</sub>) were noted. At 183 dpe, intraepithelial and luminal proliferative trophozoites (F<sub>1</sub>), severe high intraepithelial lymphocyte infiltration (F<sub>2</sub>), and luminal disporoblasts with mature spores (F<sub>3</sub>) were found. Black arrowheads = luminal proliferative trophozoites; white arrowheads = luminal disporoblasts with mature spores; black arrows = intratissue proliferative trophozoites; asterisks = intraepithelial lymphocyte infiltration. Giemsa stained sections. Scale bars in (A–C) and (D–F) = 50 µm; scale bars in A<sub>1</sub>–A<sub>3</sub>, B<sub>1</sub>–B<sub>2</sub>, C<sub>1</sub>–C<sub>3</sub>, D<sub>1</sub>–D<sub>3</sub>, E<sub>1</sub>–E<sub>2</sub>, and F<sub>1</sub>–F<sub>3</sub> = 20 µm.

#### 3.3 Lymphocyte quantification

Different lymphocyte types were labeled and quantified in the gill and intestinal tissues of the experimental steelhead trout (summarized in Supplementary Figures 2D–F, 4). Numbers of Zap70 immunoreactive T cells were significantly higher in the gills of exposed SYM and ALLO fish at 1 dpe (Figure 4A). These T cells were found spread in the gill filament and lamellae

(Figures 4C–E). In the intestines of both SYM and ALLO exposed fish, T cell numbers were significantly higher than in control fish at 15 dpe (Figure 4B; Supplementary Figure 2D). T cell abundance peaked in exposed SYM intestines at 22 dpe, whereas in exposed ALLO, it peaked later at 29 dpe (Figures 4F–H). By 183 dpe, intestinal numbers of T cells in exposed fish were restored to the levels of control fish in both fish strains. In control intestines, constitutive numbers of T cells remained unchanged, and T cells



were found mostly restricted to the base of the epithelium, whereas in exposed animals, they appeared invading the lamina propria– submucosa and infiltrated in the epithelium.

IgT immunoreactive B cells in the gills were scarce, and their numbers did not change significantly over time (Figure 5A) but were significantly higher in SYM than in ALLO upon *C. shasta* exposure (Supplementary Figure 4B). Interestingly, the IgT immunolabel was noted at the surface of gill lamellae of control and exposed SYM fish but was absent in ALLO gills (Figures 5C–E). In intestines, a consistent increase of IgT<sup>+</sup> B cells was observed in exposed SYM fish from 29 dpe onward and in exposed ALLO only at 57 dpe (Figure 5B). Only few dispersed IgT<sup>+</sup> B cells were observed in the intestines of control and exposed ALLO fish, while in exposed SYM intestines, IgT<sup>+</sup> B cell aggregates were observed in the lamina propria–submucosa and abundant IgT<sup>+</sup> B cells infiltrated the epithelial layer (Figures 5F–H). Overall, SYM fish presented the highest IgT<sup>+</sup> B cell numbers in intestines, and in ALLO fish, a less intense but still significant increase of IgT<sup>+</sup> B cells was observed (Supplementary Figure 2E).

In addition to B cell labeling, IgT immunolabeling was also observed on the surface of luminal intestinal parasite stages, indicating that those parasite stages were IgT-coated (Figure 6A). This label was, however, inconsistent among the different parasite stages within the same intestinal sample and among SYM and ALLO fish.

IgD immunoreactive B cells were the scarcest lymphocyte type in both tissues. However,  $IgD^+$  B cell numbers in gills were significantly higher in exposed ALLO fish than in the other experimental groups (Figures 7A, C-E; Supplementary Figure 4C). At the intestinal level, higher numbers of  $IgD^+$  B cells were registered only in exposed SYM fish at 22 and 29 dpe (Figure 7B; Supplementary Figure 2F), and these formed cell aggregates in the lamina propria–submucosa (Figures 7F–H). No IgD immunolabeling was observed on parasite stages (Figure 6B).

#### 3.4 Specific circulating antibodies

Parasite labeling with specific IgT of the tested serum pools was negative for all times and both fish strains. By contrast, parasite stages were specifically labeled with serum IgM of *C. shasta*-exposed SYM and ALLO fish at 57 and 183 dpe (Figure 8). Differences in



Zap70 immunoreactive T cells (Zap70-IR; brown stain) in intestines (A, C-E) and gills (B, F-H) of control unexposed (CTRL; C, F) and *Ceratonova shasta* exposed sympatric (SYM; D, G) and allopatric (ALLO; E, H) steelhead trout. (A, B) Different lowercase and uppercase letters stand for significant differences among sampling times within SYM or ALLO fish, respectively. Asterisks stand for significant differences between two experimental groups within a time post-exposure (P < 0.05). Microphotographs of sampling times highlighted in dashed boxes are represented in (C-H): Intestinal sections correspond to 29 days post-exposure (dpe) and gill sections correspond to 1 dpe; each insert shows an area of the non-lamellar filament surface. Hematoxylin stained sections. Scale bars = 20 µm.

parasite immunoreactivity were noted depending on the stage and its location. Only proliferative trophozoites were detected by the circulating IgM (not disporoblasts), and in general, luminal trophozoites attached to the mucosal surface of the intestine were the less immunoreactive ones, compared to free luminal trophozoites and to migrating intratissue trophozoites. Immunolabel was apparently located on the surface of *C. shasta* and in the cytoplasm of secondary daughter cells. Parasite pseudopodia were intensely labeled, especially in the luminal trophozoites. Label intensity was more intense with SYM sera than with ALLO sera only for some parasite stages.

#### 3.5 Ultrastructural observations

Only proliferative trophozoites of *C. shasta* were detected in the intestinal samples analyzed by TEM. These stages consisted of a primary cell encompassing up to four secondary daughter cells (Figure 9). Intraepithelial infiltration of lymphocytes was evident in the intestines, and trout lymphocytes were observed surrounding parasite stages, in direct contact with their primary cell (Figures 9A–D). The parasites' primary cells presented pseudopodia and cytoplasmic protrusions extending toward or in between host cells (Figures 9C, H, O), and the primary cell



IgT immunoreactive B cells (IgT-IR; brown stain) in intestines (A, C-E) and gills (B, F-H) of control unexposed (CTRL; C, F) and *Ceratonova shasta* exposed sympatric (SYM; D, G) and allopatric (ALLO; E, H) steelhead trout. (A, B) Lowercase and uppercase letters indicate significant differences among sampling times within SYM or ALLO fish, respectively. Asterisks indicate significant differences between two experimental groups within a time post-exposure (P < 0.05). Microphotographs of sampling times highlighted in dashed boxes are represented in (C–H): Intestinal sections correspond to 29 days post-exposure (dpe) and gill sections correspond to 1 dpe. Hematoxylin stained sections. Scale bars = 20  $\mu$ m.

membrane contacting lymphocytes or other host cells presented numerous ridges and blebbing (Figures 9C, F, G). The primary cell cytoplasm contained lipid inclusions and numerous vesicles, whereas the cytoplasm of developing secondary cells was densely packed with ribosomes, abundant rough endoplasmic reticulum (RER), and mitochondria (Figures 9C, F, G, I, M, S). The nuclei of the secondary cells showed a large eccentric nucleolus (Figures 9E, L, R). Free parasite stages were also observed in the intestinal lumen surrounded by undifferentiated cell debris (Figures 9J, N, P), and some also presented extended pseudopodia (Figure 9O).

#### 4 Discussion

The genotype 0 lineage of *Ceratonova shasta* has the ability to proliferate and transmit without causing the death of its fish host (Atkinson and Bartholomew, 2010a; Alama-Bermejo et al., 2019; Taggart-Murphy et al., 2021) and thus offers a model to investigate parasite tolerance through the factors involved in both myxozoan virulence and host resistance. Graphical Abstract summarizes the main findings of this study, in which we investigated the host– parasite relationship from the perspective of the host. In the current



Ig immunoreactivity on intestinal *Ceratonova shasta* stages. (A) IgT immunolabeling observed on luminal parasite stages in a sympatric steelhead host. Note the surface labeling on parasite trophozoites attached to the intestinal mucosa (arrowheads). (B) Absence of IgD immunolabeling on parasite stages in a sympatric steelhead host. Note the light positive label on the intestinal brush border. Scale bars = 10  $\mu$ m.

infection challenge, *C. shasta* G0 did not cause mortality in either ALLO or SYM hosts. In the gills, although *C. shasta* G0 was not observed by histology and no histopathological signs were observed, PCR+ fish were detected from both SYM and ALLO steelhead groups at 1 dpe, consistent with previous observations that the gills are the sites of initial parasite invasion (Bjork and Bartholomew, 2010). Earlier studies (Bjork and Bartholomew, 2010; Bjork et al., 2014) concluded that resistance to *C. shasta* GI invasion did not occur in the gills of either SYM or ALLO chinook salmon, and parasite morphology, replication, and migration from the gills into the intestine appeared similar between susceptible ALLO rainbow trout and resistant SYM chinook salmon.

# 4.1 Intestinal parasite proliferation and sporogenesis are restricted in the sympatric fish

In the intestine, both fish strains were PCR+ from 15 dpe onward throughout the entire study. Differences in infection dynamics were observed between strains, and these differences influenced the period when sporulation occurred. In SYM steelhead trout, trophozoite proliferation was lower and sporogenesis seemed occasional, compared to the ALLO steelhead. These observations may suggest that *C. shasta* G0 has adjusted its life cycle in its SYM host through their long co-evolutionary history, that the SYM host has developed a differential, perhaps more effective immune response against the parasite, or that both of these occur and complement each other.

Early sporogenesis in resistant anadromous hosts supports completion of the parasite's life cycle while hosts are still in the river ecosystem and spores can thus encounter the annelid host. Nevertheless, proliferative trophozoites persisting in the intestinal lumen of SYM fish might provide an opportunity for the parasite to remain dormant until some triggering occurs for sporulation. This strategy also entails parasite adaptation to a coelozoic lifestyle, which is generally less pathogenic for the host. The long host-parasite co-evolution of genotype 0 has been assumed from its high specificity for the putative ancestral hosts, low virulence, and phylogenetic reconstruction and genetic structure models (Breyta et al., 2020; Bartholomew et al., 2022). Under these circumstances, the introduction of ALLO fish stocks in water bodies endemic for C. shasta G0 might have provided a new host scenario in which parasite proliferation and sporogenesis might be unleashed. Transcriptomic, proteomic, and epitope analysis may help to explain the underlying mechanisms for the differential expression of parasite virulence factors and life stage-specific markers in SYM and ALLO steelhead.

There is no previous data on the differential infection dynamics of C. shasta G0 in SYM and ALLO fish hosts, although studies with genotypes I and II have provided interesting results. Upon exposure to a lower actinospore dose of the more virulent C. shasta GIIR, resistant SYM steelhead effectively contained parasite proliferation at 7 dpe, whereas in susceptible ALLO steelhead, parasite replication was still exponentially increasing at 21 dpe (Barrett and Bartholomew, 2021). In those fish, C. shasta GIIR stages invaded the intestinal mucosa and submucosa, as was also reported for C. shasta GI in SYM and ALLO chinook salmon (Bjork and Bartholomew, 2010). In contrast, in this study, proliferation of C. shasta G0 continued apparently unrestricted through the 183 days of the study only in the intestinal lumen of SYM and ALLO steelhead, and sporogenesis was limited to the early infection (29 dpe) only in SYM hosts whereas it continued in ALLO until the end of the experiment. A previous study evidenced that C. shasta G0 differed from the other genotypes in having a low proliferation rate, lower spore production, and less active parasite stages (Alama-Bermejo et al., 2019). Cell movement of motile myxozoan stages has been attributed to tissue invasion and immune evasion strategies (EL-Matbouli et al., 1995; Alama-Bermejo et al., 2012; Hartigan et al., 2016). Indeed, the virulent genotypes (I and II) which proliferate in the lamina propria-submucosa and epithelium rely on their high motility to evade host immune factors of the gut-associated immune tissue (GALT), as completion of their life cycle depends on rapid proliferation and spore release, before host death (Alama-Bermejo et al., 2019, 2020). However, high motility may not be crucial for the success of C. shasta G0 infections, which differ from the other genotypes in their high luminal prevalence and low tissue invasion.

# 4.2 Intestinal hyperplasia is a sign of inflammatory dysregulation in allopatric fish

The intestinal histopathology of ALLO steelhead trout upon *C. shasta* G0 infection involved mild submucosal inflammation with



IgD immunoreactive B cells (IgD-IR; brown stain) in intestines (A, C–E) and gills (B, F–H) of control unexposed (CTRL; C, F) and *Ceratonova shasta* exposed sympatric (SYM; D, G) and allopatric (ALLO; E, H) steelhead trout. (A, B) Lowercase letters indicate significant differences among sampling times within SYM fish. Asterisks indicate significant differences between two experimental groups within a time post-exposure (P < 0.05). Microphotographs at sampling times highlighted in dashed boxes are represented in (C–H): Intestinal sections correspond to 29 days post-exposure (dpe) and gill sections correspond to 8 dpe. Hematoxylin stained sections. Scale bars = 20 µm.

moderate infiltration of immune cells and minimal epithelial damage (Taggart-Murphy et al., 2021). In the present study, two main characteristic inflammatory signs were observed in the intestine upon exposure to *C. shasta* G0, regardless of the fish strain: submucosal hyperplasia and intraepithelial infiltration of lymphocytes. The submucosal hyperplasia overlapped with the presence of proliferative trophozoites migrating through the intestinal layers in both host strains. Thus, hyperplasia in SYM fish seemed mostly resolved from 29 dpe onward. In ALLO fish, however, the onset of the hyperplasia was delayed, and the condition persisted until 183 dpe, pointing to an inflammatory dysregulation when compared to SYM. Previous studies have also

attributed extensive intestinal inflammation and lymphocytic infiltration to the proliferation of *C. shasta* GIIR stages within the tissue accompanied by an upregulation of proinflammatory cytokine expression (Taggart-Murphy et al., 2021). However, epithelial disruption and total breakdown of the intestinal barrier, as reported during *C. shasta* GIIR infection in ALLO steelhead trout (Barrett and Bartholomew, 2021), did not occur in either ALLO or SYM hosts upon exposure to *C. shasta* G0.

The inflammatory response protects the fish host by inducing repair mechanisms through recruitment of inflammatory filtrates and proliferation of local immune effectors (Soliman and Barreda, 2023). However, inappropriate regulation may result in an excessive



inflammatory response and pathogenesis as reported from some myxozoan infections (Sitjà-Bobadilla et al., 2015; Holzer et al., 2021). Fine tuning of the regulatory response during hostparasite co-evolution may be responsible for the histopathological differences observed in the intestines of ALLO and SYM steelhead during C. shasta G0 infection. This immune regulation/ dysregulation is particularly evident in two intestinal myxozoan genera that have been the focus of research due to their devastating impact on aquaculture fish species, Ceratonova and Enteromyxum. The adaptive immune response of gilthead seabream (Sparus aurata) resistant to reinfection with Enteromyxum leei displayed a post-inflammatory gene expression profile in the intestine involving downregulation of  $il1\beta$  and  $hsp90\alpha$  and upregulation of il10(Picard-Sánchez et al., 2019). This might hint toward an early scenario of inflammation resolution occurring only during C. shasta G0 infections, in which the inflammatory response elicited by any parasite stages migrating through the tissue is effectively suppressed or regulated by the fish hosts. Thus, a stronger and earlier postinflammatory profile would be expected in SYM intestines than in ALLO. Once more, further studies on the differential transcriptomics of SYM and ALLO steelhead are needed to identify subtle differences in their regulatory response at the intestinal level, with special focus on regulatory T cell markers and anti-inflammatory cytokines, as well as on innate factors.

## 4.3 Intestinal T cell response is triggered earlier in the sympatric fish

T cell activation is an early event in the adaptive immune response, occurring upon specific antigen recognition through the T cell receptor, which encompasses recruitment and activation of the Zap70 kinase. Zap70 kinase is considered a pan T cell marker, expressed on CD8<sup>+</sup> cytotoxic T cells (CTLs) as well as on CD4<sup>+</sup> helper T cells, mediators of B cell activation and antibody production through cytokine release (Laing and Hansen, 2011). Intestinal T cell recruitment/proliferation was evidenced in the exposed fish by Zap70 immunolabeling. An increase in T cells was detected in both host strains when trophozoites were first detected in the intestinal tissue. Zap70<sup>+</sup> T cells were mostly consistent with intraepithelial T lymphocytes in their morphology and location, but we cannot discount the presence of CTLs, capable of directly killing parasite cells. For now, we have demonstrated a T cell response against C. shasta G0, but we cannot discern to what extent the later T cell response in ALLO hosts contributes to slower parasite clearing in the intestinal tissue and to the subsequent unimpeded luminal parasite proliferation beginning at 29 dpe. The analysis of transcriptional profiles of T cell subsets will aid in understanding the T cell response that takes place in the intestines of SYM vs. ALLO steelhead trout after C. shasta G0 exposure, as



Ceratonova shasta proliferative trophozoites in allopatric Oncorhynchus mykiss intestines at 3-4 months post exposure. In lower magnification images, parasite stages are colored green and fish lymphocytes are colored red. (A) C. shasta trophozoite located in the intestinal epithelium, in close contact with fish lymphocytes. (B) Higher magnification of the parasite in (A) showing three secondary (s) cells within the primary (p) cell. (C) Higher magnification of the area in the upper left dashed square in (B) showing the contact between host cell and p cell with abundant cytoplasmic vesicles (white arrowhead). Note the ribosome-packed cytoplasm and abundant rough endoplasmic reticulum (RER, black arrowhead) in the s cell. (D) Higher magnification of the area in the upper right dashed square in (B) showing the p cell cytoplasm with abundant cytoplasmic vesicles (white arrowheads) and electrodense lipid droplets (black arrowheads). Note the p cell cytoplasmic protrusions in the dashed square, which is magnified in (E). (F) Higher magnification of the area in the lower dashed square in (B) showing the contact area between p parasite cell and host lymphocyte. Note the p cell with pseudopod (black arrowhead) reaching between host cells and abundant cytoplasmic vesicles (white arrowhead). Within the p cell, the s cell harbors abundant cytoplasmic ribosomes. The dashed square is magnified in (G), showing the irregular contact surface between the p cell and a lymphocyte, with numerous ridges and blebbings (black arrowhead). (H) Higher magnification of the area in the middle-dashed square in (B) showing the ribosomepacked cytoplasm and abundant RER (black arrowhead) and mitochondria (white arrowheads) in the s cell. (I) Intestinal cross-section with a C. shasta luminal stage and an intraepithelial stage close to the basal membrane. (J) Higher magnification of the lower dashed square in (I) showing a trophozoite with four internal s cells. (K) Higher magnification of the dashed square in (J) showing detail of the junction between trophozoite and epithelial basal membrane (black arrowhead) of the host intestine. (L) Higher magnification of the upper dashed square in (I) showing detail of the luminal stage membrane in contact with luminal host cell debris. (M) Intestinal cross-section with luminal trophozoite and abundant infiltrated intraepithelial lymphocytes. (N) Higher magnification of the right dashed square in (M) showing detail of an intraepithelial lymphocyte. (O) Higher magnification of the left dashed square in (M) showing trophozoite in the intestinal lumen. Note the dashed square magnified in (P) showing parasite's pseudopod contacting luminal host cell debris (black arrowhead). (Q) Intraepithelial trophozoite consisting of three s cells within the p cell. Note the abundant RER (black arrowhead) and the mitochondria (white arrowheads) in s cells, and the cytoplasmic lipid inclusions (black arrow) and vesicles (white arrows) in the p cell. (R) Higher magnification of the dashed square in (Q) showing details of RER (black arrowhead) in the ribosome-packed cytoplasm of the s cell and mitochondria (white arrowheads) in the neighboring host cell, p, primary cell; h, host cell; s, secondary cell; N<sub>s</sub>, nucleus of the s cell. Scale bars in (A, I, M) = 10  $\mu$ m; in (B, F, J, L, N–Q) = 1  $\mu$ m; in (C–E, G, H, K, R) = 100 nm.

previously reported for CTL signature gene expression in *C. shasta* GIIR infection (Barrett and Bartholomew, 2021) and in resistant gilthead seabream to *E. leei* reinfection (Picard-Sánchez et al., 2019).

The available literature on T cell responses to fish myxozoans is limited. However, activation of the IFN\gamma-signaling pathway mediated by T<sub>H1</sub> cells seems a common response in resistance against myxozoan infections. An effective T<sub>H1</sub> response was assumed from the ALLO rainbow trout resistant to C. shasta G0 infection due to the intestinal  $ifn\gamma$  upregulation (Taggart-Murphy et al., 2021) and steelhead trout resistant to the virulent C. shasta GIIR (Barrett and Bartholomew, 2021). Thus, early specific C. shasta recognition might be an essential aspect for immune resistance. The studied enteromyxosis models also provide a range of pathogenesis and immune responses in naïve vs. resistant gilthead seabream and turbot (Scophthalmus maximus) against E. leei and Enteromyxum scophthalmi, respectively. In the highly susceptible naïve turbot, exhaustion and downregulation of T<sub>H1</sub>, T<sub>H17</sub>, and CTL gene markers (Robledo et al., 2014) follow the initial T<sub>H1</sub> response (Ronza et al., 2016), whereas in naïve gilthead seabream with a less severe disease course, the T cell response encompasses a strong intestinal recruitment of Zap70<sup>+</sup> and Tbet<sup>+</sup> T lymphocytes along with expression profiles of CTL, T<sub>H1</sub>, and T<sub>H2</sub> responses (Piazzon et al., 2018). Furthermore, gilthead seabream resistance against enteromyxosis was associated with mucosal recruitment of T cells from the head kidney (Picard-Sánchez et al., 2019). Also, gibel carp (Carassius auratus gibelio) with a severe Myxobolus hunghuensis pharynx infection, displayed T cell recruitment into the infection site along with T<sub>H1</sub> and T<sub>H2</sub> transcriptional profiles (Zhao et al., 2019). Thus, among myxozoonoses, an early activation of the T cell response in mucosal infection sites is supported systemically, and its correct regulation seems critical for myxozoan resistance and disease development. A remarkable example for effective T cell activation is resistant Atlantic salmon (Salmo salar) re-exposed to the muscleinfecting Kudoa thyrsites, in which recruitment of CD8<sup>+</sup> CTLs to the infection site aligns with upregulation of CTL signature genes and B cell markers, resulting in parasite clearance and protection (Braden et al., 2018).

## 4.4 A strong intestinal IgT<sup>+</sup> B cell response leads protection in the sympatric fish

B cell and antibody responses stimulated upon T cell activation are the next adaptive immune effectors in the battle against myxozoan diseases. The mucosal B cell response was the most significant differential trait we observed between steelhead trout strains. Among the B cell subtypes we analyzed, the IgT response was the dominant one in the intestine. An intense IgT<sup>+</sup> B cell response, spread along the intestinal mucosa and the lamina propria, was triggered in SYM fish, following a strong T cell response, whereas ALLO fish presented a mild and delayed IgT<sup>+</sup> B cell response. In contrast, IgD<sup>+</sup> B cells were almost absent in the intestine of control and exposed steelhead trout. Further investigation is warranted to decipher if those are double  $IgM^+IgD^+$  B cells or if single  $IgD^+$   $IgM^-$  B cells are involved in the immune response against *C. shasta* G0. Nonetheless,  $IgM^+IgD^+$  B cells are rare in rainbow trout mucosal tissues, with most  $IgM^+$  B cells lacking IgD expression (Herranz-Jusdado et al., 2023). In any case, this study has evidenced immunologic traits of the SYM fish phenotype (early T cell activation and  $IgT^+$  cell recruitment), which might help to further establish markers for the selection of resistant fish stocks (e.g., *zap70, igt* expression).

## 4.5 Specific serum IgM appears late and does not differ among fish strains

By contrast, the specific circulating IgM response detected at 57 and 183 dpe was similar among SYM and ALLO steelhead. Thus, the lower parasite load in the intestines of SYM fish may not be attributed to the IgM response, whose contribution to host protection at the later time points seems limited. The strongest immunogenic labeling of C. shasta G0 was observed in trophozoites migrating through the intestinal layers and in luminal trophozoites, with high cell polarization and with pseudopodia that appeared involved in mucosal attachment. Interestingly, disporoblasts and mature spores seemed to lack immunogenicity, at least by IgM. While the specific IgT in sera was below detection threshold, we detected IgT<sup>+</sup> immunolabel coating luminal parasite stages in both SYM and ALLO hosts, supporting its limited systemic involvement against this parasite and the important mucosal role of IgT. However, it has been suggested that antigens in C. shasta trophozoites may mimic trout antigens to evade immune recognition (Bartholomew et al., 1989), and we cannot exclude the possibility that the observed IgT coating on C. shasta G0 intestinal stages is rather a sign of parasite masking than of immune targeting.

Chronic or prolonged C. shasta infections provide the time frame for hosts to develop an Ig response, though to what extent those antibodies provide protection is still unclear (Bartholomew et al., 2022). Yet, the mucosal role of IgT was first reported from rainbow trout intestines infected with C. shasta G0 (Zhang et al., 2010). Supporting our results, an increased transcription of secreted igm and igt was triggered at 29 dpe to C. shasta G0 in rainbow trout intestines, and this upregulation was significantly higher for igt than for igm (Taggart-Murphy et al., 2021). SYM steelhead were capable of reverting enteritis and cleared C. shasta GIIR from the intestinal tissue by 60 dpe (Barrett et al., 2021). These fish evidenced at 21 dpe a strong induction of genes related to epithelial repair and to the adaptive immune response, including a notable upregulation of Ig transcription, particularly IgT. In general, acute C. shasta infections with the virulent genotypes GIIR in steelhead and chinook salmon (Hurst et al., 2019; Barrett and Bartholomew, 2021) or GI in chinook salmon (Bjork et al., 2014) induce igm and igt expression, although susceptible ALLO fish were not able to resolve infection or parasite proliferation and the inflammatory process continued. The adaptive response appears rather late and ineffective, as mortality due to fatal enteronecrosis occurs around 20 dpe. Increased Ig expression and

recruitment of Ig<sup>+</sup> B cells to the infection site is not uncommon among myxozoan infections, pointing to the importance of the specific local immune response. This has been reported from both intestinal infection models, E. leei in gilthead seabream and E. scophthalmi in turbot (Bermúdez et al., 2006; Estensoro et al., 2012; Robledo et al., 2014; Piazzon et al., 2016; Ronza et al., 2016). Parasitespecific circulating IgM was related to protection against reinfection in both enteromyxoses (Sitjà-Bobadilla et al., 2007; Picard-Sánchez et al., 2019). Yet, protective antibodies appear late after the primary challenge of naïve fish, and in the case of the more virulent and acute turbot enteromyxosis, specific IgM prevalence in serum is low and fish mortality is high (Sitjà-Bobadilla et al., 2006, 2007). The slower progression of the chronic and less virulent enteromyxosis in gilthead seabream, however, seems to provide the time frame for fish to build up a more effective adaptive response, which might be a scenario closer to G0 ceratomyxosis studied herein. Thus, as in gilthead seabream, it will be worth studying, through secondary challenges and passive immunization, whether the more effective primary response which culminates with higher titers of parasite-specific IgM protects steelhead against reinfection by C. shasta G0 or, furthermore, if it provides cross-protection against C. shasta GIIR. The role played by resistance traits related to T cell response and IgT<sup>+</sup> B cell response in such potential cross-protection against other parasite genotypes upon a secondary challenge would also be worth studying.

Immunomodulatory evasion strategies, common among myxozoans, include increased Ig titers during an ineffective and delayed specific immune response along with B cell exhaustion (reviewed in Sitjà-Bobadilla (2008)). In fact, the expression of high Ig clonotype diversity upon polyclonal expansion of diverse Ig B cell subsets (hyperimmunoglobulinemia) was reported from the Ig repertoires of rainbow trout during the profound B cell dysregulation caused by Tetracapsuloides bryosalmonae kidney infection (Abos et al., 2018) and during E. leei intestinal infection in naïve gilthead seabream (Picard-Sánchez et al., 2020). Also, antigen expression shift and epitope masking might be used by myxozoans to evade recognition by specific antibodies (Holzer et al., 2021). Thus, it is still unclear whether the specific antibody responses against C. shasta G0 are immunoprotective and whether a differential clonal expansion among the Ig repertoire is induced in SYM and ALLO fish upon ceratomyxosis.

# 4.6 The ultrastructure of *C. shasta* G0 primary cells suggests functional specialization for migration, environment exploration, and nutrient acquisition

*C. shasta* G0 ultrastructure in intestinal sections revealed extracellular trophozoites in the paracellular space between enterocytes and in the intestinal lumen. Cell protrusions (pseudopodia) of different lengths were observed in primary cells of both, intratissue and free luminal trophozoites, reaching between host cells. Such pseudopodia are compatible with the ones reported by

Alama-Bermejo et al. (2019) from C. shasta GIIR, which were associated with the parasite's exploratory behavior in complex environments. Although C. shasta G0 ultrastructure was not studied by those authors, from the expression of motility-related factors, it was inferred that early C. shasta G0 intestinal stages were less active but performed strong directional and adhesive migration characterized by an elongated cell shape with a leading edge. We also observed such elongated parasite shapes in the intratissue stages by TEM and light microscopy. After the initial intestinal invasion, C. shasta G0 stages abandon migration by 29 dpe (Alama-Bermejo et al., 2019), which aligns with the time point at which we observed luminal parasite stages peaking. Primary cell folds and protrusions also increase the parasites' absorptive surface, aiding nutrition from the host cells. Hence, pinocytotic vesicles for nutrient acquisition have been reported, and vacuoles, lysosomes, vesicles, and lipidic droplets in primary cells were interpreted as food reserves, which were scarce in secondary and tertiary daughter cells of Enteromyxum spp. (Redondo et al., 2003; Cuadrado et al., 2008). By TEM, we also observed numerous vesicles of different sizes and densities, and electrodense lipid inclusions in primary cells, indicative of their nourishment-related activity. By contrast, secondary cells contained a remarkable high abundance of mitochondria, RER, and ribosomes, indicative of their proliferative stage and high protein synthesis activity. Further, some of the infiltrated intraepithelial lymphocytes observed by TEM had an irregular contact surface with the parasite's primary cell, which might be a sign of direct interaction between both cell types.

## 4.7 The response against *C. shasta* G0 in gills of both fish strains is succinct

Finally, the immune response in gills, the portal of entry for the parasite, deserves some attention. As soon as fish are exposed to the parasite, recognition events in both fish strains seem to trigger the T cell response, although activation seemed stronger in SYM steelhead, aligning with the observed earlier activation of the T cell response at the intestinal level. No significant differences were observed in regard to the scarce IgT<sup>+</sup> B cells in the gills, and only SYM fish showed an increasing trend in IgT<sup>+</sup> B cell numbers, which occurred immediately after parasite exposure and decreased over time. Interestingly, IgT immunolabel on the mucosal surface of gills was weak or null in ALLO fish. These observations may indicate the increased immunocompetence of SYM fish. Previous studies on the gill invasion and later dispersion of C. shasta stages have not reported histopathological findings during these early phases, and it was suggested that embedding of the early parasite stages in endothelial cells of the gill blood vessels might impede immune recognition (Bjork and Bartholomew, 2010). Also, parasite-induced immunosuppression aiding an initial host invasion was reported from the gills of SYM and ALLO steelhead at 1 dpe to C. shasta GIIR, which encompassed T cell development and activation (Barrett and Bartholomew, 2021). Thus, these previous results might depict a similar situation to the one occurring in host gills during C. shasta G0 invasion.

# 4.8 IgD<sup>+</sup> B cells present an opposite response in gills and intestine of both fish strains

Surprisingly, although IgD<sup>+</sup> B cells were the scarcest lymphocytes observed in gills, this was the only leukocyte type that increased significantly, and only in ALLO steelhead after parasite exposure. And more intriguingly, IgD<sup>+</sup> B cells were almost absent in the intestine of ALLO, while they increased in SYM host. The specific role of this Ig class in defense remains an enigma. B cells co-expressing IgD and IgM are the most abundant B cell subset in fish systemic tissues such as blood or spleen. Nonetheless, B cell subsets exclusively expressing IgD (IgD<sup>+</sup> IgM<sup>-</sup> plasmablasts) have been identified in different mucosal surfaces, including intestine, gills, and skin (Castro et al., 2014; Perdiguero et al., 2019; Herranz-Jusdado et al., 2023). Furthermore, IgD seemed involved in maintenance of commensal microbiota homeostasis (Perdiguero et al., 2019), which may explain the IgD immunolabeling observed on the intestinal brush border. From our current results, we cannot attribute the observed IgD<sup>+</sup> B cells to a specific cell subtype, but their increase after the parasite challenge seems to follow the common pattern of IgD responses against inflammatory stimuli in teleosts (Bjørgen et al., 2023; Herranz-Jusdado et al., 2023; Perdiguero et al., 2024; Karami et al., 2025).

#### **5** Conclusions

The current comparative challenge of SYM and ALLO hosts with the non-pathogenic genotype C. shasta G0 provides clues to the differential parasite dynamics in the intestine (Graphical Abstract). In SYM steelhead trout, trophozoites proliferated less and sporogenesis was occasional. Histopathology revealed that intestinal hyperplasia was resolved in SYM fish, whereas it continued for more than 180 days in ALLO fish. The successful coordination and regulation of earlier T cell activation and a stronger subsequent mucosal IgT response in SYM host hindered luminal proliferation of the parasite and reversed the intestinal inflammation earlier than in ALLO fish. In contrast to the more virulent genotypes I and II, G0 seems not to induce excessive dysregulation of intestinal T and B cells, provokes a less severe disease pattern, and develops into a chronic sublethal mainly celozoic infection probably aided by the systemic IgM protection, even in ALLO fish. Furthermore, after the long hostparasite co-evolutionary history, the limited C. shasta G0 proliferation and sporogenesis in SYM hosts might point to an adjustment of the parasite's life cycle, and the early and effective inflammatory and immune response in SYM intestines might point to a differential, more effective host strategy: both complement each other and would allow for greater parasite tolerance in the SYM fish hosts. The current results at tissue and protein levels will be validated by functional molecular studies, which will help establish resistance markers for disease control in fish stocks, and cross-protection studies with the distinct parasite genotypes will aid in the development of vaccination strategies. From a management perspective, stocking of ALLO fish into waters where G0 is present in the absence of the more virulent GII would likely result in parasite amplification and increased dissemination as a result of the longer period of parasite release. While this may not result in direct mortality of SYM hosts, these fish did show disease pathology and presumed reduced fitness before the infection resolved.

#### Data availability statement

The original contributions presented in the study are included in the article/Supplementary Material. Further inquiries can be directed to the corresponding author.

#### **Ethics statement**

The animal study was approved by Animal Care and Use Committee of the John L. Fryer Aquatic Animal Health Laboratory at the Oregon State University. The study was conducted in accordance with the local legislation and institutional requirements.

#### Author contributions

IE: Conceptualization, Data curation, Formal analysis, Investigation, Methodology, Software, Supervision, Validation, Visualization, Writing – original draft, Writing – review & editing. BLG: Formal analysis, Investigation, Writing – review & editing. CT: Investigation, Methodology, Writing – review & editing. RC: Investigation, Methodology, Writing – review & editing. SA: Writing – original draft, Writing – review & editing, Conceptualization, Investigation, Methodology, Resources. ASB: Funding acquisition, Project administration, Resources, Supervision, Writing – original draft, Writing – review & editing, Investigation, Methodology, Conceptualization. JB: Conceptualization, Formal analysis, Funding acquisition, Investigation, Methodology, Project administration, Resources, Supervision, Validation, Visualization, Writing – original draft, Writing – review & editing.

#### Funding

The author(s) declare that financial support was received for the research and/or publication of this article. This study was funded in part by the Bureau of Reclamation (Reclamation), U.S. Department of Interior. Funding was provided by Reclamation as part of its mission to manage, develop, and protect water and related resources in an environmentally and economically sound manner in the interest of the American public. Funding was provided through Interagency Agreement #R23PG00059. The views in this report are the author's and do not necessarily represent the views of Reclamation. Funding

for J. Bartholomew to work at CSIC was provided through a Fulbright Scholar Grant. Additional support was provided by the Generatitat Valenciana AICO2023 funding (CIAICO/2022/144).

#### Acknowledgments

K. Buchmann kindly provided the anti-trout IgM and anti-IgT antibodies, whereas E. Bromage provided the anti-IgD antibody used in this study. We thank J. Monfort and L. Rodríguez for their technical assistance on the histological processing, R. Milston-Clements for assistance with sampling and fish husbandry, D. Da Piedade Silva for preparation of samples for shipments, and R. Hold for preparation of the TEM samples.

#### **Conflict of interest**

The authors declare that the research was conducted in the absence of any commercial or financial relationships that could be construed as a potential conflict of interest.

#### References

Abos, B., Estensoro, I., Perdiguero, P., Faber, M., Hu, Y. F., Rosales, P. D., et al. (2018). Dysregulation of B cell activity during proliferative kidney disease in rainbow Trout. *Front. Immunol.* 9. doi: 10.3389/fimmu.2018.01203

Alama-Bermejo, G., Bron, J. E., Raga, J. A., and Holzer, A. S. (2012). 3D morphology, ultrastructure and development of *Ceratomyxa puntazzi* stages: First insights into the mechanisms of motility and budding in the Myxozoa. *PloS One* 7, e32679. doi: 10.1371/journal.pone.0032679

Alama-Bermejo, G., Meyer, E., Atkinson, S. D., Holzer, A. S., Wiśniewska, M. M., Kolísko, M., et al. (2020). Transcriptome-wide comparisons and virulence gene polymorphisms of host-associated genotypes of the Cnidarian parasite *Ceratonova shasta* in salmonids. *Genome Biol. Evol.* 12, 1258–1276. doi: 10.1093/gbe/evaa109

Alama-Bermejo, G., Holzer, A. S., and Bartholomew, J. L. (2019). Myxozoan adhesion and virulence: *ceratonova shasta* on the move. *Microorganisms* 7, 397. doi: 10.3390/microorganisms7100397

Alexander, J. D., Bartholomew, J. L., Wright, K. A., Som, N. A., and Hetrick, N. J. (2016). Integrating models to predict distribution of the invertebrate host of myxosporean parasites. *Freshw. Sci.* 35, 1263–1275. doi: 10.1086/688342

Atkinson, S. D., and Bartholomew, J. L. (2010a). Disparate infection patterns of *Ceratomyxa shasta* (Myxozoa) in rainbow trout (*Oncorhynchus mykiss*) and Chinook salmon (*Oncorhynchus tshawytscha*) correlate with internal transcribed spacer-1 sequence variation in the parasite. *Int. J. Parasitol.* 40, 599–604. doi: 10.1016/JJIPARA.2009.10.010

Atkinson, S. D., and Bartholomew, J. L. (2010b). Spatial, temporal and host factors structure the *Ceratomyxa shasta* (Myxozoa) population in the Klamath River basin. *Infection Genet. Evol.* 10, 1019–1026. doi: 10.1016/J.MEEGID.2010. 06.013

Atkinson, S. D., Hallett, S. L., and Bartholomew, J. L. (2018). Genotyping of individual *Ceratonova shasta* (Cnidaria: Myxosporea) myxospores reveals intra-spore ITS-1 variation and invalidates the distinction of genotypes II and III. *Parasitology* 145, 1588–1593. doi: 10.1017/S0031182018000422

Ballesteros, N. A., Castro, R., Abos, B., Rodríguez Saint-Jean, S. S., Pérez-Prieto, S. I., and Tafalla, C. (2013). The pyloric caeca area is a major site for IgM+ and IgT+ B cell recruitment in response to oral vaccination in rainbow trout. *PloS One* 8, e66118. doi: 10.1371/journal.pone.0066118

Barrett, D. E., and Bartholomew, J. L. (2021). A tale of two fish: Comparative transcriptomics of resistant and susceptible steelhead following exposure to *Ceratonova shasta* highlights differences in parasite recognition. *PloS One* 16, e0234837. doi: 10.1371/JOURNAL.PONE.0234837

Barrett, D. E., Estensoro, I., Sitjà-Bobadilla, A., and Bartholomew, J. L. (2021). Intestinal transcriptomic and histologic profiling reveals tissue repair mechanisms underlying resistance to the parasite *Ceratonova shasta*. *Pathogens* 10. doi: 10.3390/ pathogens10091179

#### Generative AI statement

The author(s) declare that no Generative AI was used in the creation of this manuscript.

#### Publisher's note

All claims expressed in this article are solely those of the authors and do not necessarily represent those of their affiliated organizations, or those of the publisher, the editors and the reviewers. Any product that may be evaluated in this article, or claim that may be made by its manufacturer, is not guaranteed or endorsed by the publisher.

#### Supplementary material

The Supplementary Material for this article can be found online at: https://www.frontiersin.org/articles/10.3389/fcimb.2025. 1576014/full#supplementary-material

Bartholomew, J. L., Alexander, J. D., Hallett, S. L., Alama-Bermejo, G., and Atkinson, S. D. (2022). *Ceratonova shasta*: a cnidarian parasite of annelids and salmonids. *Parasitology* 149, 1862–1875. doi: 10.1017/S0031182022001275

Bartholomew, J. L., Rohovec, J. S., and Fryer, J. L. (1989). Development, characterization, and use of monoclonal and polyclonal antibodies against the myxosporean, *Ceratomyxa shasta. J. Protozoology* 36, 397–401. doi: 10.1111/j.1550-7408.1989.tb05534.x

Bermúdez, R., Vigliano, F., Marcaccini, A., Sitjà-Bobadilla, A., Quiroga, M. I., and Nieto, J. M. (2006). Response of Ig-positive cells to *Enteromyxum scophthalmi* (Myxozoa) experimental infection in turbot, *Scophthalmus maximus* (L.): A histopathological and immunohistochemical study. *Fish Shellfish Immunol.* 21, 501–512. doi: 10.1016/j.fsi.2006.02.006

Bjørgen, H., Oaland, Ø., Midtllyng, P., Tafalla, C., Krogdahl, Å., and Koppang, E. O. (2023). IgD-transcript positive cells suggest hypersensitivity in feed-related intestinal inflammation in the Atlantic salmon. *Fish Shellfish Immunol.* 132, 108469. doi: 10.1016/j.fsi.2022.108469

Bjork, S. J., and Bartholomew, J. L. (2010). Invasion of *Ceratomyxa shasta* (Myxozoa) and comparison of migration to the intestine between susceptible and resistant fish hosts. *Int. J. Parasitol.* 40, 1087–1095. doi: 10.1016/j.ijpara.2010.03.005

Bjork, S. J., Zhang, Y. A., Hurst, C. N., Alonso-Naveiro, M. E., Alexander, J. D., Sunyer, J. O., et al. (2014). Defenses of susceptible and resistant Chinook salmon (*Onchorhynchus tshawytscha*) against the myxozoan parasite *Ceratomyxa shasta*. Fish Shellfish Immunol. 37, 87–95. doi: 10.1016/J.FSI.2013.12.024

Braden, L. M., Rasmussen, K. J., Purcell, S. L., Ellis, L., Mahony, A., Cho, S., et al. (2018). Acquired Protective Immunity in Atlantic Salmon Salmo salar against the Myxozoan Kudoa thyrsites Involves Induction of MHII $\beta^+$  CD83<sup>+</sup> Antigen-Presenting Cells. Infection Immun. 86. doi: 10.1128/iai.00556-17

Breyta, R., Atkinson, S. D., and Bartholomew, J. L. (2020). Evolutionary dynamics of *Ceratonova* species (Cnidaria: Myxozoa) reveal different host adaptation strategies. *Infection Genet. Evol.*, 78, 104081. doi: 10.1016/J.MEEGID.2019.104081

Castro, R., Bromage, E., Abós, B., Pignatelli, J., González Granja, A., Luque, A., et al. (2014). CCR7 is mainly expressed in teleost gills, where it defines an IgD<sup>+</sup>IgM<sup>-</sup> B lymphocyte subset. *J. Immunol.* 192, 1257–1266. doi: 10.4049/jimmunol.1302471

Cuadrado, M., Marques, A., Diamant, A., Sitjà-Bobadilla, A., Palenzuela, O., Álvarez-Pellitero, P., et al. (2008). Ultrastructure of *Enteromyxum leei* (Diamant, Lom, & Dyková, 1994) (Myxozoa), an enteric parasite infecting gilthead sea bream (*Sparus aurata*) and sharpsnout sea bream (*Diplodus puntazzo*). J. Eukaryotic Microbiol. 55, 178–184. doi: 10.1111/j.1550-7408.2008.00325.x

EL–Matbouli, M., Hoffmann, R. W., and Mandok, C. (1995). Light and electron microscopic observations on the route of the triactinomyxon-sporoplasm of *Myxobolus cerebralis* from epidermis into rainbow trout cartilage. *J. Fish Biol.* 46, 919–935. doi: 10.1111/J.1095-8649.1995.TB01397.X

Estensoro, I., Calduch-Giner, J. A., Kaushik, S., Pérez-Sánchez, J., and Sitjà-Bobadilla, A. (2012). Modulation of the IgM gene expression and IgM immunoreactive cell distribution by the nutritional background in gilthead sea bream (*Sparus aurata*) challenged with *Enteromyxum leei* (Myxozoa). *Fish Shellfish Immunol.* 33, 401–410. doi: 10.1016/j.fsi.2012.05.029

Hallett, S. L., and Bartholomew, J. L. (2006). Application of a real-time PCR assay to detect and quantify the myxozoan parasite *Ceratomyxa shasta* in river water samples. *Dis. Aquat. Organisms* 71, 109–118. doi: 10.3354/DAO071109

Hallett, S. L., Ray, R. A., Hurst, C. N., Holt, R. A., Buckles, G. R., Atkinson, S. D., et al. (2012). Density of the waterborne parasite *Ceratomyxa shasta* and its biological effects on salmon. *Appl. Environ. Microbiol.* 78, 3724–3731. doi: 10.1128/AEM.07801-11/SUPPL\_FILE/AEM07801-11\_SUPPLEMENTAL.PDF

Hartigan, A., Estensoro, I., Vancová, M., Bílý, T., Patra, S., Eszterbauer, E., et al. (2016). New cell motility model observed in parasitic cnidarian *Sphaerospora molnari* (Myxozoa: Myxosporea) blood stages in fish. *Sci. Rep.* 6, 39093. doi: 10.1038/srep39093

Herranz-Jusdado, J. G., Morel, E., Simón, R., Díaz-Rosales, P., and Tafalla, C. (2023). Teleost IgD<sup>+</sup>IgM<sup>-</sup> B cells in gills and skin have a plasmablast profile, but functionally and phenotypically differ from IgM<sup>+</sup>IgD<sup>-</sup> B cells in these sites. *iScience* 26, 107434. doi: 10.1016/j.isci.2023.107434

Holzer, A. S., Piazzon, M. C., Barrett, D., Bartholomew, J. L., and Sitjà-Bobadilla, A. (2021). To react or not to react: The dilemma of fish immune systems facing myxozoan infections. *Front. Immunol.* 12. doi: 10.3389/FIMMU.2021.734238/BIBTEX

Hurst, C. N., Alexander, J. D., Dolan, B. P., Jia, L., and Bartholomew, J. L. (2019). Outcome of within-host competition demonstrates that parasite virulence doesn't equal success in a myxozoan model system. *Int. J. Parasitol. Parasites Wildl.* 9. doi: 10.1016/ j.ijppaw.2019.03.008

Hurst, C. N., and Bartholomew, J. L. (2012). *Ceratomyxa shasta* genotypes cause differential mortality in their salmonid hosts. *J. Fish Dis.* 35, 725–732. doi: 10.1111/J.1365-2761.2012.01407.X

Karami, A. M., Kania, P. W., Al-Jubury, A., Stefanova, D., Krych, L., Madsen, L., et al. (2025). Gut microbiota in rainbow trout *Oncorhynchus mykiss* with different susceptibility to *Flavobacterium psychrophilum* infection. *Aquaculture* 596, 741841. doi: 10.1016/j.aquaculture.2024.741841

Laing, K. J., and Hansen, J. D. (2011). Fish T cells: Recent advances through genomics. *Dev. Comp. Immunol.* 35, 1282–1295. doi: 10.1016/J.DCI.2011.03.004

Palenzuela, O., Trobridge, G., and Bartholomew, J. L. (1999). Development of a polymerase chain reaction diagnostic assay for *Ceratomyxa shasta*, a myxosporean parasite of salmonid fish. *Dis. Aquat. Organisms* 36, 45–51. doi: 10.3354/DAO036045

Perdiguero, P., Martín-Martín, A., Benedicenti, O., Díaz-Rosales, P., Morel, E., Muñoz-Atienza, E., et al. (2019). Teleost igD<sup>+</sup>IgM<sup>-</sup> B cells mount clonally expanded and mildly mutated intestinal igD responses in the absence of lymphoid follicles. *Cell Rep.* 29, 4223–4235. doi: 10.1016/J.CELREP.2019.11.101

Perdiguero, P., Jiménez-Barrios, P., Morel, E., Abós, B., and Tafalla, C. (2024). Singlecell atlas of rainbow trout peripheral blood leukocytes and profiling of their early response to infectious pancreatic necrosis virus. *Front. Immunol.* 15. doi: 10.3389/ fimmu.2024.1404209

Piazzon, M. C., Galindo-Villegas, J., Pereiro, P., Estensoro, I., Calduch-Giner, J. A., Gómez-Casado, E., et al. (2016). Differential modulation of IgT and IgM upon parasitic, bacterial, viral, and dietary challenges in a perciform fish. *Front. Immunol.* 7. doi: 10.3389/fmmu.2016.00637

Piazzon, M. C., Estensoro, I., Calduch-Giner, J. A., Del Pozo, R., Picard-Sánchez, A., Pérez-Sánchez, J., et al. (2018). Hints on T cell responses in a fish-parasite model: *Enteromyxum leei* induces differential expression of T cell signature molecules depending on the organ and the infection status. *Parasites Vectors* 11, 443. doi: 10.1186/s13071-018-3007-1 Picard-Sánchez, A., Estensoro, I., Del Pozo, R., Piazzon, M. C., Palenzuela, O., and Sitjà-Bobadilla, A. (2019). Acquired protective immune response in a fish-myxozoan model encompasses specific antibodies and inflammation resolution. *Fish Shellfish Immunol.* 90, 349–362. doi: 10.1016/j.fsi.2019.04.300

Picard-Sánchez, A., Estensoro, I., Perdiguero, P., Del Pozo, R., Tafalla, C., Piazzon, M. C., et al. (2020). Passive immunization delays disease outcome in gilthead Sea bream infected with *Enteromyxum leei* (Myxozoa), despite the moderate changes in IgM and IgT repertoire. *Front. Immunol.* 11. doi: 10.3389/FIMMU.2020.581361/BIBTEX

Ramírez-Gómez, F., Greene, W., Rego, K., Hansen, J. D., Costa, G., Kataria, P., et al. (2012). Discovery and characterization of secretory IgD in eainbow trout: Secretory IgD is produced through a novel splicing mechanism. *J. Immunol.* 188, 1341–1349. doi: 10.4049/jimmunol.1101938

Redondo, M. J., Quiroga, M. I., Palenzuela, O., Nieto, J. M., and Álvarez-Pellitero, P. (2003). Ultrastructural studies on the development of *Enteromyxum scophthalmi* (Myxozoa), an enteric parasite of turbot (*Scophthalmus maximus* L.). Parasitol. Res. 90, 192–202. doi: 10.1007/s00436-002-0810-5

Robledo, D., Ronza, P., Harrison, P. W., Losada, A. P., Bermúdez, R., Pardo, B. G., et al. (2014). RNA-seq analysis reveals significant transcriptome changes in turbot (*Scophthalmus maximus*) suffering severe enteromyxosis. *BMC Genomics* 15, 1149. doi: 10.1186/1471-2164-15-1149

Ronza, P., Robledo, D., Bermúdez, R., Losada, A. P., Pardo, B. G., Sitjà-Bobadilla, A., et al. (2016). RNA-seq analysis of early enteromyxosis in turbot (*Scophthalmus maximus*): new insights into parasite invasion and immune evasion strategies. *Int. J. Parasitol.* 46, 507–517. doi: 10.1016/j.ijpara.2016.03.007

Sitjà-Bobadilla, A. (2008). Living off a fish: A trade-off between parasites and the immune system. Fish Shellfish Immunol. 25, 358-372. doi: 10.1016/j.fsi.2008.03.018

Sitjà-Bobadilla, A., Redondo, M. J., Bermúdez, R., Palenzuela, O., Ferreiro, I., Riaza, A., et al. (2006). Innate and adaptive immune responses of turbot, *Scophthalmus maximus* (L.), following experimental infection with *Enteromyxum scophthalmi* (Myxosporea: Myxozoa). *Fish Shellfish Immunol*. 21, 485–500. doi: 10.1016/j.fsi.2006.02.004

Sitjà-Bobadilla, A., Palenzuela, O., Riaza, A., Macias, M. A., and Álvarez-Pellitero, P. (2007). Protective acquired immunity to *Enteromyxum scophthalmi* (Myxozoa) is related to specific antibodies in *Psetta maxima* (L.) (Teleostei). *Scandinavian J. Immunol.* 66, 26–34. doi: 10.1111/j.1365-3083.2007.01942.x

Sitjà-Bobadilla, A., Schmidt-Posthaus, H., Wahli, T., Holland, J. W., and Secombes, C. J. (2015). "Fish immune responses to myxozoa," in *Myxozoan Evolution, Ecology and Development.* Eds. B. Okamura, A. Gruhl and J. Bartholomew (Springer International Publishing, Switzerland). doi: 10.1007/978-3-319-14753-6\_14

Soliman, A. M., and Barreda, D. R. (2023). The acute inflammatory response of teleost fish. *Dev. Comp. Immunol.* 146, 104731. doi: 10.1016/j.dci.2023.104731

Stinson, M. E. T., and Bartholomew, J. L. (2012). Predicted redistribution of *Ceratomyxa shasta* genotypes with salmonid passage in the eschutes River, Oregon. *J. Aquat. Anim. Health* 24, 274–280. doi: 10.1080/08997659.2012.716012

Taggart-Murphy, L., Alama-Bermejo, G., Dolan, B., Takizawa, F., and Bartholomew, J. (2021). Differences in inflammatory responses of rainbow trout infected by two genotypes of the myxozoan parasite *Ceratonova shasta*. *Dev. Comp. Immunol.* 114, 103829. doi: 10.1016/J.DCI.2020.103829

Zhang, Y. A., Salinas, I., Li, J., Parra, D., Bjork, S., Xu, Z., et al. (2010). IgT, a primitive immunoglobulin class specialized in mucosal immunity. *Nat. Immunol.* 11, 827–836. doi: 10.1038/ni.1913

Zhao, Y., Liu, X., Sato, H., Zhang, Q., Li, A., and Zhang, J. (2019). RNA-seq analysis of local tissue of *Carassius auratus gibelio* with pharyngeal myxobolosis: Insights into the pharyngeal mucosal immune response in a fish-parasite dialogue. *Fish Shellfish Immunol.* 94, 99–112. doi: 10.1016/j.fsi.2019.08.076