

Adaptive Immune System in Fish

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How to cite

Kordon, A.O., Pinchuk, L., Karsi, A. (2022). Adaptive Immune System in Fish. *Turkish Journal of Fisheries and Aquatic Sciences*, 22(4), TRJFAS20235. <http://doi.org/10.4194/TRJFAS20235>

Article History

Received 07 July 2021

Accepted 06 November 2021

First Online 25 November 2021

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Keywords

Adaptive immunity

T cells

B cells

Abstract

The immune system of all jawed vertebrates is composed of two major subsystems, the innate (non-specific) and adaptive (specific) immune system. The innate immune system is the first to respond to infectious agents; however, it does not provide long-lasting protection. The adaptive immune system is activated later and responds to pathogens with specificity and memory. The main components of the adaptive immune system, including T cell receptors (TCRs), major histocompatibility complex (MHC), immunoglobulins (Igs), and recombination-activating gene (RAG) arose in the first jawed fish (cartilaginous and teleost fish). This review explores and discusses components of the adaptive immune system in teleost fish and recent developments in comparative immunology. Similar to mammals, the adaptive immune system in teleost fish is divided into two components: cellular-mediated responses and humoral-mediated responses. T cells, the principal elements of cellular-mediated adaptive immune responses, differentiate into effector helper T (Th) cells or effector cytotoxic T cells (CTLs). The central elements involved in the differentiation of Th subsets in mammals, cytokines and master transcription factors, have also been identified in teleost fish. In addition, each subset of Th cells, defined with a particular cytokine to control the immune responses, has been described in teleost fish. Similar to mammals, CTLs contribute to cellular cytotoxicity in teleost fish. B cells are central players in humoral-mediated adaptive immunity by producing opsonizing, neutralizing and complement-binding antibodies and inducing antibody-dependent cellular cytotoxicity (ADCC). Three classes of antibodies named IgM, IgD, and IgT/Z have been characterized in teleost fish. The presence of an adaptive immune system and consequent immune memory in teleost fish allows vaccination, the most appropriate method for disease control in aquaculture. Immunological studies in fish provide a comprehensive assessment of the fish immune system, which is crucial for understanding the evolution of the mammalian immune system.

Introduction

All jawed vertebrates, including cartilaginous and teleost (bony) fish, share essential fundamentals of the immune system, consisting of two main parts: innate (non-specific) and adaptive (specific) immune responses. Innate immunity, the first line of the defense

system, recognizes pathogens in a non-specific manner and creates a fast reaction to eliminate the microbes, but does not provide long-term protection (Turvey & Broide, 2010). If the innate immune system is insufficient to clear the infectious agents, adaptive immunity is activated by non-specific immunity and interferes with pathogens by reacting with specificity

and long-lasting protection (memory) (Rauta et al., 2012). The adaptive immune system is composed of highly specialized cells, T- and B-lymphocytes, and proteins that destroy and inhibit the growth of invaders (Spiering, 2015). In contrast to innate immunity, adaptive immune responses rely on the highly diverse antigen-specific receptors expressed on T- and B-cells by a few hundred germ-line-encoded gene elements that are assembled by somatic hypermutation and recombination mechanisms of the variable (V), diversity (D), and joining (J) gene segments (Chaplin, 2010; Litman et al., 2010; Tonegawa, 1983). Therefore, the adaptive immune system possesses exquisite specificity for many different pathogens. Another key property of adaptive immunity is memory, as the adaptive immune system generates memory cells that provide long-lived specific immunity, thus playing a crucial role in protection against recurrent infections with a quick and efficient response to the same pathogens (Tangye & Tarlinton, 2009).

Cartilaginous and teleost (bony) fish are the earliest vertebrate group containing basic principles of immunity similar to mammals, despite certain differences between them. Although teleost fish include approximately 50% of all vertebrate species, most of the studies in fish immunology have been defined over the last decades (Voff, 2005). Previously, we highlighted recent knowledge on the innate immune responses and antigen-presenting cells (APCs) in teleost fish (Kordon et al., 2018). In this review, we describe the major components of the adaptive immune system in teleost fish.

Adaptive Immunity in Fish

The major specific components of the adaptive immune system, such as T cell receptors (TCRs), the major histocompatibility complex (MHC), immunoglobulins (Igs), and the recombination-activating gene (RAG), appeared roughly 450-500 million years ago in the first jawed fish (cartilaginous and teleost fish) (Table 1) (Brazeau & Friedman, 2015; Flajnik & Kasahara, 2010). Like in mammals, adaptive immunity in teleost fish is divided into two main components: cellular and humoral responses (Figure 1) (Biller-Takahashi J. D. & C., 2014). T cells, key elements of cellular adaptive immunity, mature in the thymus and differentiate into effector cells, including cytotoxic T cells (CTLs) or helper T (Th) cells (Nakanishi et al., 2015). Although B cells, key elements of humoral adaptive immunity, are derived from bone marrow in mammals, they are produced in the anterior kidney of teleost fish and differentiate into plasma cells that produce antibodies (Carsetti, 2000; Zwollo et al., 2005). Antibodies can be present in two different forms: a soluble form, known as immunoglobulins (Igs), and a membrane-bound form, known as B cell receptor (BCR) (Smith et al., 2019).

The BCR and TCR, expressed on B and T cells, are antigen-specific receptors formed by V(D)J recombination and somatic hypermutation (Nemazee, 2000). Rearrangement of V(D)J genes in different combinations generates a wide-range diversity of antigen-receptor specificities (Carmona et al., 2016). Proteins encoded by recombination-activating genes (RAG1 and RAG2) mediate V(D)J gene rearrangement by recognition of recombination signal sequences (RSSs) and cleavage of target DNA (Carmona & Schatz, 2017; Fugmann, 2001; Schatz et al., 1989). V(D)J gene segments are present in teleost fish, such as zebrafish, rainbow trout, and fugu (Mashoof & Criscitiello, 2016). In addition, the expression of RAG1 and RAG2 genes have been identified in teleost fish, and the V(D)J gene rearrangement in teleost fish is dependent on RAG proteins (Dickerson & Findly, 2017; Greenhalgh & Steiner, 1995; Lee et al., 2014; Willett et al., 1997). Moreover, somatic hypermutation in mammals is initiated by an enzyme, activation-induced cytidine deaminase (AID), that is also expressed in teleost fish (Barreto et al., 2005). In addition to the formation of TCRs and BCRs, these mechanisms induce an irreversible change in the DNA of each cell, and the progenies of these cells inherit the genes that encode the same receptor specificity, including memory T and B cells that provide long-lived specific immunity (Firdaus-Nawi & Zamri-Saad, 2016).

In addition to antigen specificity, immunological memory is defined as the fundamental feature of the adaptive immune system. The majority of specific immune cells die after clearance of the invading pathogens, but a small percentage of cells give rise to long-lived memory cells that mediate a rapid and protective immune response against previously encountered pathogens (Pennock et al., 2013; Youngblood et al., 2017). Effective immunization relies on the combination of antigen specificity and memory in adaptive immunity. Like mammals, memory T and B cells have been identified in teleost fish (Scapigliati et al., 2018). The proliferation of memory T cells has been described in immunized carp by IL-10 modulation (Piazzon et al., 2015).

Cell-Mediated Responses in Adaptive Immunity

T cells play a fundamental role in cell-mediated responses of adaptive immunity by either involving the regulation of other leukocytes functions or directly killing infected host cells. T cells are developed in the thymus; therefore, they are also called thymocytes. The transcription factors for T cell development in teleost fish, such as *ikaros* (expressed by early T-cells) and *lck* (expressed later in development) seem to be similar in mammals (Willett et al., 2001). In addition to molecular mechanisms, cellular mechanisms of T cell development are conserved in jawed vertebrates (Boehm et al., 2012; Boehm & Swann, 2014; Hirano et al., 2011; Langenau &

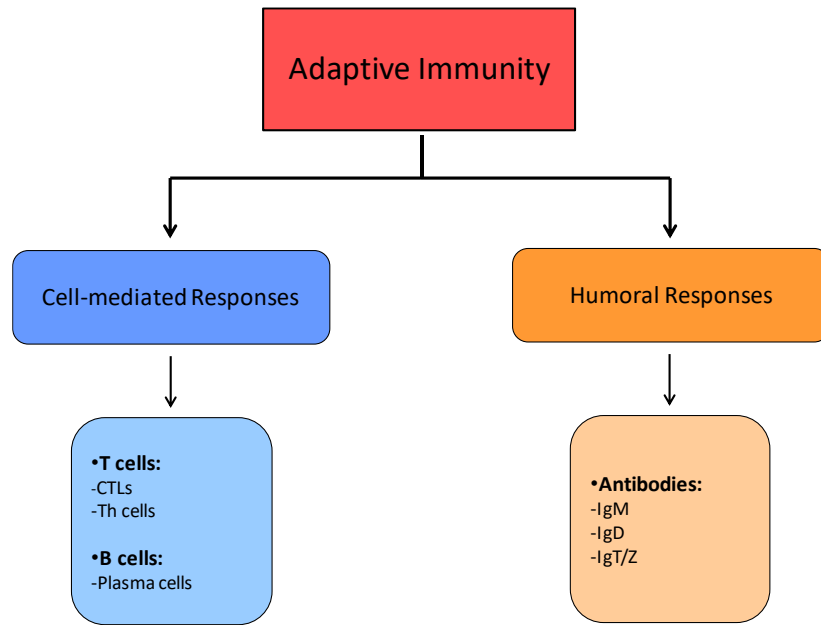


Figure 1. The components of adaptive immunity in teleost (bony) fish. The adaptive immune system of teleost fish consists of two major components: 1) Cell-mediated responses, and 2) Humoral responses.

Table 1. Comparative assessment of general adaptive immune components between teleost fish and mammals

	Teleost (Bony) Fish	Mammals
T and B cells	+	+
BCR	+	+
TCR	+	+
Mechanism of Diversity	V(D)J recombination	V(D)J recombination
RAG genes	+	+
AID	+	+
SHM	+	+
Germinal Center	-	+
CSR	-	+
Ig Isotypes	IgM, IgD, and IgT/Z	IgM, IgD, IgG, IgE, and IgA

Zon, 2005). Recently, the development of T cells in teleost fish was comprehensively reviewed elsewhere (Bajoghli et al., 2019). Mature T-cells are present in lymphoid tissues of teleost fish, such as thymus, kidney, spleen, and mucosa-associated lymphoid tissues, including the intestine, skin, and gills (Nakanishi et al., 2015).

All T cells possess a TCR, formed by RAG-mediated V(D)J gene rearrangement and recognize specific antigens. In mammals, TCR is a type I transmembrane glycoprotein with a short cytoplasmic tail and is associated with the CD3 complex, a transmembrane protein complex with an intracellular signaling domain (immunoreceptor tyrosine-based activation motif (ITAM)), which forms the TCR:CD3 complex (Birnbaum et al., 2014; Love & Hayes, 2010). The structure of the TCR:CD3 complex is conserved between mammals and teleost fish (Castro et al., 2011; Øvergård et al., 2009; Shang et al., 2008). Moreover, T cells in mammals are

present mainly in two sublineages based on the nature of heterodimeric receptor chains with TCR: $\alpha\beta$ -TCR or $\gamma\delta$ -TCR (Smith et al., 2019). Most of the T cells express $\alpha\beta$ -TCR found in blood and lymphoid organs, whereas $\gamma\delta$ -T cells represent a small fraction of total T cell population in mammals (~5 %) (Carding & Egan, 2002; Janeway et al., 1988). Similar to mammals, TCR genes encode the TCR α , β , γ , and δ chains in teleost fish, such as channel catfish, zebrafish, and Atlantic salmon (Bilal et al., 2018; Castro et al., 2011; Fischer et al., 2002; Haire et al., 2000; Hordvik et al., 2004; Nam et al., 2003; Wilson et al., 1998). Also, the $\gamma\delta$ -T cells constitute 7-20% of total lymphocytes in blood and lymphoid tissues of zebrafish (Wan et al., 2017).

Although $\gamma\delta$ -T cells recognize antigens directly, $\alpha\beta$ -T cells recognize peptides of antigens bound to major histocompatibility complex (MHC) molecules (Konigshofer & Chien, 2006). The $\alpha\beta$ -T cells in mammals are further divided into two major populations based on

their function, cytotoxic CD8+ T cells (CTLs) and helper CD4+ T (Th) cells. CD8+ T cells are activated by peptides derived from intracellular antigens and presented by MHC class I molecules, and their function is to kill infected host cells. Helper CD4+ T cells are stimulated by peptides derived from extracellular antigens and presented by MHC class II molecules, and they regulate the responses of other leukocytes (Andersen et al., 2006; Banchereau & Steinman, 1998; Kordon et al., 2018). The functional roles of both T cells and T cell-related molecules, CD4, CD8, MHC class I and II, have been described in teleost fish (Fischer et al., 2013; Koppang et al., 2010; Nakanishi et al., 2015; Hideaki Toda, Yasutaka Saito, et al., 2011).

Major Histocompatibility Complex and Activation of Naïve T Cells

Major histocompatibility complex (MHC) class I and II molecules play a pivotal role in adaptive immunity. MHC class I molecules are composed of two domains ($\alpha 1$ and $\alpha 2$), forming antigen binding platform, and one membrane-spanning domain ($\alpha 3$). The microglobulin domain ($\beta 2$) is bound non-covalently to the membrane-spanning domain (Chen et al., 2010; Chen et al., 2017). On the other hand, the binding region of MHC class II molecules are formed by $\alpha 1$ and $\beta 1$ domains, and two membrane-spanning domains ($\alpha 2$ and $\beta 2$) are present in MHC class II molecule (Chen et al., 2010; Chen et al., 2017; Yamaguchi & Dijkstra, 2019). Although both MHC class I and II genes were described in most teleost fish and are conserved in jawed vertebrates, some teleost fish, such as Atlantic cod lack MHC class II genes (Smith et al., 2019; Star et al., 2011). However, more genes-related to MHC class I molecules were identified in Atlantic cod to compensate for the absence of MHC class II counterparts compared to other teleost fish (Smith et al., 2019; Star & Jentoft, 2012; Star et al., 2011). Also, the heterodimer complexes and the conserved structure for peptide-ligand binding of MHC molecules in teleost fish are similar to those found in mammals (Smith et al., 2019; Yamaguchi & Dijkstra, 2019).

In mammals, MHC class I molecule is expressed by all nucleated cells and presents the peptides from intracellular antigens to CD8+ T cells (Cytotoxic T cells) (Hewitt, 2003). However, MHC class II molecule presents the peptides processed from endocytosed/phagocytosed antigens (extracellular antigens) on only professional APCs, dendritic cells, macrophages, and B cells to CD4+ T cells (helper T cells) (Cho & Roche, 2013). Similar to mammals, MHC class II molecules are expressed by all professional APCs of teleost fish (Yamaguchi & Dijkstra, 2019)(ref). However, MHC class I molecules are ubiquitously expressed and detected in multiple tissues, including spleen and anterior kidney (Grimholt et al., 2002; Smith et al., 2019). Also, the roles of teleost MHC class I and II molecules are similar to those in mammals. For

example, upon antigen stimulation, the expression of both MHC class I and II genes were upregulated in teleost fish similarly to mammals (Scharsack et al., 2007). Also, the elevated expression of MHC class I and II genes increased the survival and resistance of Atlantic salmon in *Aeromonas salmonicida* infection (Kjøglum et al., 2008; Smith et al., 2019). Furthermore, increases in expression of MHC class II and CD4 genes correlated with enhanced adaptive immune responses to *Edwardsiella ictaluri* (*E. ictaluri*) live attenuated vaccine strains in channel catfish lymphoid organs (Kordon, Abdelhamed, et al., 2019; Kordon et al., 2021).

In addition, the interaction between TCR and the MHC: peptide complex on APCs lack enough signal to activate fully naïve T cells in mammals. Therefore the priming of T cells requires the combination of three different signals delivered in sequence: 1) antigen recognition (TCR:MHC), 2) costimulation, and 3) cytokines (Figure 2) (Sckisel et al., 2015). The co-stimulatory signal is generated by the interaction of a co-stimulatory factor, CD28, expressed on T cells and B7.1 (CD80) and B7.2 (CD86) ligands found on APC. As a third signal, cytokines secreted by APCs determine the differentiation of activated T cell into a particular effector T cell subset (Muñoz-Wolf & Lavelle, 2018; Nace et al., 2012). Similar to mammals, co-stimulatory molecules (CD28, B7.1, and B7.2) have been identified in several teleost fish, such as rainbow trout and zebrafish (Bernard et al., 2006; Hansen et al., 2009; Sugamata et al., 2009). Cytokines for T cell differentiation have been determined in teleost fish, similar to mammals (Ashfaq et al., 2019).

Effector CD4+ T helper Cells

CD4+ T cells accomplish numerous crucial functions in mammals by stimulating cell-mediated immunity of macrophages, granulocytes (neutrophils, eosinophils, and basophils), CD8+ T cells, and B cells to produce antibodies through the production of several cytokines (Reiner, 2007). Similar functions of CD4+ T cells have been described in teleost fish: in particular, the effective bacterial killing triggered by CD4+ T cells in ginbuna carp (Nayak & Nakanishi, 2013). Moreover, the upregulation of cytokine, IL-12, related to CD4+ T cells was documented in fugu during extracellular parasitic infection (Yoshiura et al., 2003). Although the mammalian CD4 molecule contains four Ig-like domains (D1-D4), teleost fish possess two CD4 genes: CD4-1 containing four Ig-like domains and CD4-2 containing two or three Ig-like domains (Ashfaq et al., 2019; Maddon et al., 1987). Recently, we showed the increased expression of both CD4-1 and CD4-2 genes in the lymphoid tissues of channel catfish infected with *E. ictaluri* strains (Kordon, Abdelhamed, et al., 2019; Kordon et al., 2021).

Upon activation, the naïve CD4+ T cells differentiate into specific subsets named Th1, Th2, Th9, Th17, Tfh cells, and Tregs, with each subset defined by a

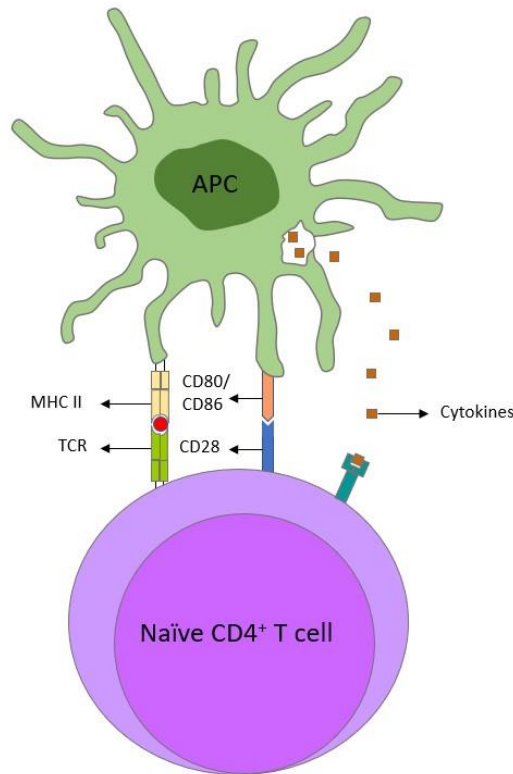


Figure 2. Three signals delivered by APCs to prime naïve T cells. **Signal 1** is generated from the interaction between MHC: peptide complex and TCR and required for activation of naïve T cells. **Signal 2** is a co-stimulatory signal (CD80/CD96:CD28) delivered by the same APC for survival and proliferation of the T cell. **Signal 3** is a cytokine signal. Cytokines are secreted by APC and other innate lymphocytes for differentiation of the T cells into a particular subtype.

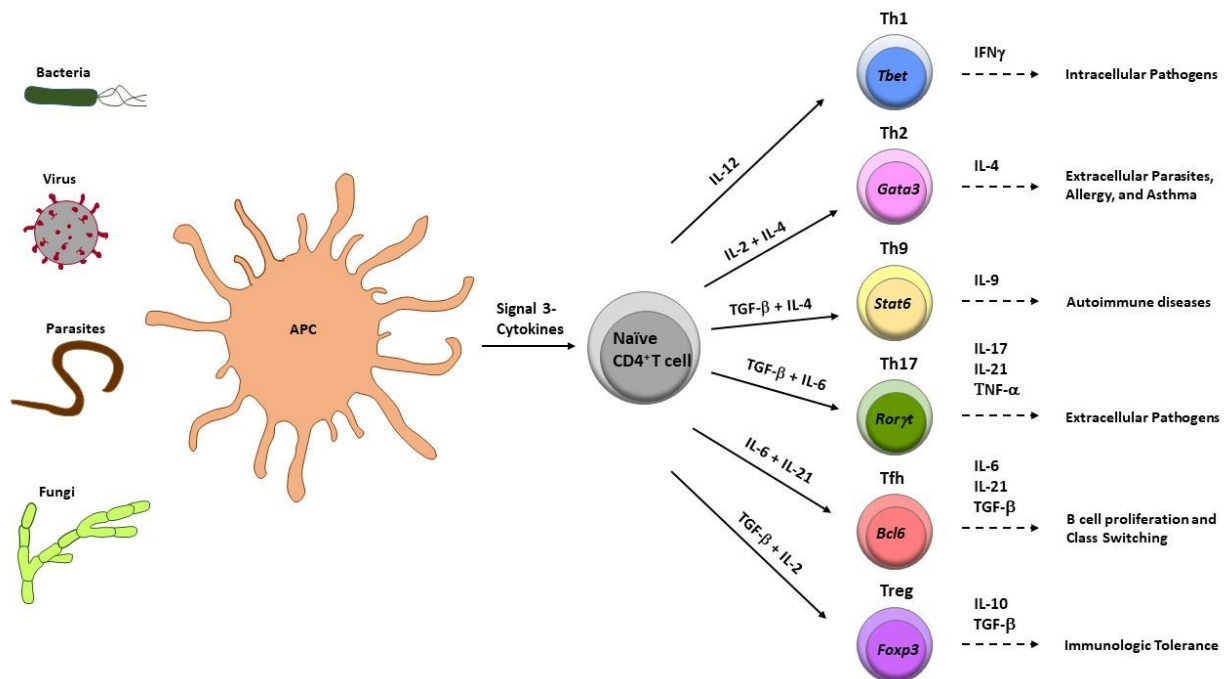


Figure 3. Pathogen-induced differentiation of naïve CD4+ T cell into different subsets. Followed by interaction with an APC infected with different pathogens (bacteria, viruses, parasites, or fungi), Th cells may differentiate into several effector cell subsets depending on the cytokine signal (solid arrows). Each Th type secretes a unique set of cytokines (dashed arrows) and promotes the pathogen-specific immune responses.

particular cytokine profile control over the immune responses (Figure 3) (Annunziato & Romagnani, 2009; Dardalhon et al., 2008; Saito et al., 2010; Wan & Flavell, 2009). In mammals, APCs activated through the pattern recognition receptors produce large amounts of interleukin 12 (IL-12) that induce natural killer cells (NK) to secrete interferon γ (IFN- γ), thus initiating Th1 differentiation (Luckheeram et al., 2012; Trinchieri et al., 2003). Also, several transcription factors, including the T-box transcription factor (T-bet), signal the transducer and activator of transcription-1 (STAT-1), and STAT-4 are involved in full differentiation of the Th1 cells (Ashfaq et al., 2019). In teleost fish, multiple isoforms of IL-12 and IFN- γ are key cytokines for Th1 differentiation (Nascimento et al., 2007; Yoshiura et al., 2003). In addition, the T-bet gene has been characterized in teleost fish, such as zebrafish, rainbow trout, and grass carp, and T-bet is considered to be a master transcription factor for both fish and mammals (S. Mitra et al., 2010; Wang et al., 2013; Wang, Holland, et al., 2010). The Th1 cells mount an immune response to intracellular pathogens by the release of IFN- γ that activates macrophages and enhances their phagocytic activity (Luckheeram et al., 2012). The Th1 cells in teleost fish generate one or two forms of IFN- γ , IFN- γ and IFN- γ rel (Smith et al., 2019). In *E. ictaluri* infection, we reported elevated expression levels of the IFN- γ gene, correlated with CD4-1 and CD4-2 genes' elevated expression in the lymphoid organs of channel catfish (Kordon, Abdelhamed, et al., 2019; Kordon et al., 2021).

The IL-2 and IL-4 are critical cytokines responsible for Th2 differentiation through the regulation of transcription factors containing STAT-5, STAT-6, and GATA-3 (Ashfaq et al., 2019; Zhu et al., 2001). Similarly, two genes of both IL-2 and IL-4 have been identified in teleost fish (Díaz-Rosales et al., 2009; Li et al., 2007). In addition to key cytokines, the master transcription factor GATA-3 has been described in distinct fish species, including zebrafish, salmonids, and grass carp (Kumari et al., 2009; Neave et al., 1995; Wang et al., 2013). The Th2 cells are effective against extracellular parasites, inducing the degranulation of cells and mast cells, and the production of antibodies by producing key cytokines including IL-4, IL-5, IL-9, IL-10, IL-13 (Walker & McKenzie, 2018). Similar to mammals, Th2-related immune responses were detected in fish gills and skin with a higher expression of IL-4/13A and GATA-3 against parasites (Takizawa, Koppang, et al., 2011).

Th17 cells, another Th cell lineage, are involved in the elimination of extracellular pathogens, including bacteria and fungi. The main cytokines IL-6, IL-21, IL-23, and transforming growth factor-beta (TGF- β) drive the differentiation of Th17 cells, along with regulators including retinoic acid receptor-related orphan receptors gamma (ROR γ) and alpha (ROR α) and the signal transducer and activator of transcription 3 (STAT-3) (Zhu et al., 2010). The major elements required for

mammalian Th17 cell segregation pathway are also found in teleost fish, containing signature cytokines (IL-6, IL-21, IL-23, and TGF- β) and the master transcription factor ROR γ (Ashfaq et al., 2019). Th17 cells in mammals produce IL-17 (IL-17A, IL-17F), IL-21, and IL-22, which mediate inflammatory responses, inducing the recruitment of inflammatory cells, such as neutrophils to site of infection, and the production of pro-inflammatory cytokines, such as IL-1, IL-6, and TNF α (Luckheeram et al., 2012). Multiple isoforms of IL-17A/F genes (IL-17A/F1, IL-17A/F2, and IL-17A/F3) have been characterized in teleost fish, such as zebrafish, rainbow trout, and fugu (Gunimaladevi et al., 2006; Kono et al., 2008; Monte et al., 2013). Additionally, IL-17A/F2 stimulated antibacterial defenses by inducing the expression of antibacterial peptides and pro-inflammatory cytokines, IL-6 and IL-8, in rainbow trout (Secombes et al., 2011). Moreover, Th17-like immune response has been identified in zebrafish mucosal tissues after live attenuated *Vibrio anguillarum* infection (Zhang et al., 2014).

Regulatory T cells (Treg) regulate the maintenance of immunologic tolerance to self and foreign antigens. The key cytokines, TGF- β and IL-10, play a vital role in the differentiation of Treg cells through the main transcription factor, forkhead box P3 (FoxP3) (Yuan & Malek, 2012). Similar to mammals, distinct isoforms of TGF- β and IL-10-type genes are also present in fish (Ashfaq et al., 2019; Zhan & Jimmy, 2000). In addition to key cytokines, the master regulator FoxP3 has been identified in teleost fish (Suman Mitra et al., 2010; Wang, Monte, et al., 2010). Mammalian Tregs produce the same effector cytokines, TGF- β and IL-10, that suppress pro-inflammatory responses after the clearance of pathogens to prevent tissue damage, and maintain peripheral tolerance (Ouyang et al., 2011). The IL-10 produced by Tregs has been identified in most teleost fish species, and its expression was highly elevated at 2-5 weeks post-infection (Piazzon et al., 2017). Also, Tregs in teleost fish preserve peripheral tolerance in the skin of rainbow trout (Leal et al., 2016).

In mammals, follicular helper T cells (Tfh) regulate the proliferation of B cells and Ig class switching, especially in the germinal centers of lymphoid tissues. The major cytokines, IL-6 and IL-21, and transcription factor Bcl6 are involved in differentiation process of Tfh cells (Nurieva & Chung, 2010). The effector Tfh cells secrete IL-6, IL-21, and TGF- β to perform their function (Srivastava et al., 2018). In contrast to mammals, teleost fish lack germinal centers and Ig isotype switching. Recently, Th9 cells were defined as a subgroup of CD4+ T cells contributing to immune responses against intestinal worms and autoimmune diseases, including multiple sclerosis (MS), inflammatory bowel disease (IBD), rheumatoid arthritis (RA), systemic lupus erythematosus (SLE), psoriasis, and cancer, and they exhibit potent anti-tumor properties (Chen et al., 2020; Srivastava et al., 2018). The cytokines TGF- β and IL-4 and

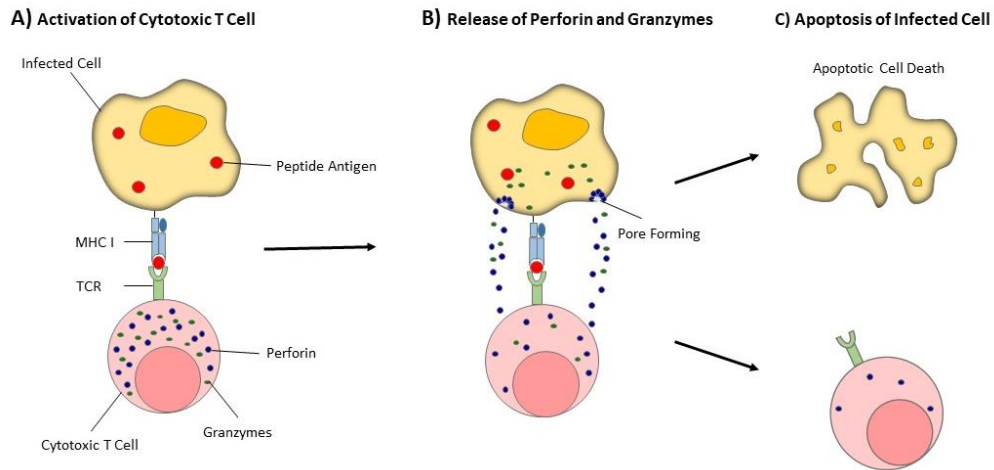


Figure 4. Clearance of infected host cells by effector CD8⁺ T (Cytotoxic T) cells in teleost fish. Cytotoxic T cells bind by their TCRs that recognize peptide: MHC class I complex on the infected host cells (A). Cytotoxic T cells secrete perforin and granzymes, that induce the formation of pores and release of cytotoxins in the cytoplasm of infected host cell (B). The infected host cells undergo apoptotic cell death, and cytotoxic T cells are detached from infected cells (C).

master transcriptional factor STAT-6 are responsible for Th9 cell differentiation (Chen et al., 2020). The significant cytokine IL-9 is secreted by Th9 cells (Srivastava et al., 2018). On the other hand, two subpopulations of CD4⁺ T cells, T_{fh} and Th9 cells, have not yet been detected in teleost fish species.

Effector CD8⁺ Cytotoxic T Cells

CD8⁺ T cells play a crucial role in immune defense against intracellular pathogens, especially virus infection. IL-12 released from the APC is the key cytokine driving the differentiation of mammalian CD8⁺ T cells through the master transcription regulators, including T-bet and eomesodermin (Eomes) (Ahuja et al., 2007; Intlekofer et al., 2005). In teleosts, homologues of IL-12 subunits have been reported (Yoshiura et al., 2003). Similarly, *T-bet* has been characterized in many teleost fish species, and gene expression analysis showed that T-bet triggers the activation of CD8⁺ T cells. Also, Eomes has been identified in teleost fish, such as zebrafish, Atlantic salmon, and rainbow trout (Yamaguchi et al., 2019). Furthermore, the highest level of Eomes expression was observed in rainbow trout CD8⁺ T cells (Takizawa et al., 2014). The CD8 molecule in mammals is present in two forms: a homodimer of two α chains or a heterodimer of α and β chains (Cole & Gao, 2004). Both α and β chains of the CD8 molecule have also been described in most species of teleost fish (Forlenza et al., 2008; Takizawa, Dijkstra, et al., 2011). Furthermore, we demonstrated that an intracellular pathogen, *E. ictaluri* strains, induced the upregulation of both CD8 α and CD8 β gene expressions in the lymphoid tissues of channel catfish (Kordon, Abdelhamed, et al., 2019; Kordon et al., 2021).

Mammalian CD8⁺ T cells induce the apoptosis of infected host cells by two pathways: the secretory and non-secretory mechanisms. The secretory pathway is characterized by the release of granular toxins, including perforin and serine proteases known as granzyme, while the non-secretory pathway requires the engagement of target-cell death receptors, such as Fas, located on the CD8⁺ T cell surface (Jaime-Sanchez et al., 2020). The secretory pathway in teleost fish is similar to that of mammals (Figure 4). A perforin-like molecule has been identified in multiple teleost species (Hwang et al., 2004; Praveen et al., 2004). CD8 α ⁺ cells in ginbuna crucian carp kill virus-infected cells by using perforin, and perforin inhibitor, concanamycin A, inhibited the killing function of CD8 α ⁺ lymphocytes in channel catfish and ginbuna crucian carp (H. Toda et al., 2011; Zhou et al., 2001). These studies showed that perforin induces a similar pathway of killing in teleost fish (Figure 4). In addition to perforin, granzyme with a similar structure to that of mammals has been identified in teleost fish (Figure 4) (Huang et al., 2010; Praveen et al., 2004). The granzyme B-like molecule was involved in the cytotoxicity of CD8⁺ T cells in fish (Hideaki Toda, Takeshi Yabu, et al., 2011). Moreover, highly elevated levels of granzyme mRNA was detected in CD8⁺ T cells of ginbuna crucian carp in *Edwardsiella tarda* (*E. tarda*) infection (Matsuura et al., 2014). Granzyme A-like serine protease in ginbuna crucian carp contributed cell-mediated immunity through cytotoxic activity (Matsuura et al., 2016). Furthermore, the non-secretory pathway has been studied in teleost fish. For instance, the FasL-like molecule, apoptosis ligand, has been characterized in several teleost species, such as channel catfish, tilapia, and gilthead sea bream (Cuesta et al., 2003; Long et al., 2004; Ma et al., 2014). The FasL-like protein in tilapia

Table 2. General structure of Ig isotypes in teleost fish

Ig Isotypes	Teleost (Bony) Fish	Mammals
IgM	Tetramer No J chain Disulfide bonds-associated Serum	Pentamer J chain-associated Serum
	Mucosal in the absence of IgT/Z	
IgD	Transmembrane Secreted: No V domain C domain ranging from 2 to 16	Monomer
	Unique to teleost fish Mucosal Tetramer in mucosal tissue Monomer in serum	Not present

resulted in cytotoxicity to the Hela cells through apoptosis (Ma et al., 2014).

Teleost CD8⁺ T cells are involved in cell-mediated cytotoxicity that control intracellular pathogens. The population of CD8⁺ cells killed virus-infected target cells in fish (Somamoto et al., 2013; Hideaki Toda, Takeshi Yabu, et al., 2011; Yamaguchi et al., 2019). Increased cytotoxic activity of CD8⁺ T cells with elevated numbers of CD8⁺ cells contribute to the elimination of *E. tarda*-infected cells and bacteria in the spleen and kidney of fish (Yamasaki et al., 2013). Furthermore, adoptive transfer of CD8⁺ lymphocytes from carp hematopoietic necrosis virus-infected fish to naïve recipients provided effective protection in ginbuna crucian carp at the latter virus challenge (Somamoto et al., 2013).

Humoral-Mediated Responses in Adaptive Immunity

B cells mediate the humoral immunity branch of the adaptive immune system by secreting antibodies. Antibodies are divided into two groups: secreted soluble form, immunoglobulins (Igs) or antibodies, or B cell receptors (BCR), membrane-bound form. Mammalian Igs are composed of two heavy chains (IgH) and two light chains (IgL) that are engaged by disulfide bonds resulting in a 'Y' shape molecule (Chiu et al., 2019). Both IgH and IgL chains are comprised of one variable region, called the Fab region (fragment, antigen-binding) that give specificity to the antibody for antigen binding, and one or more constant domains, called Fc region (fragment, crystallizable) that determine the effector functions of the antibody (Smith et al., 2019). Antibodies induce the neutralization, internalization, and elimination of pathogens, as well as antibody-dependent cellular cytotoxicity (ADCC) through Fc receptor-bearing effector cells (Forthal, 2014; Teillaud, 2001). Antibodies also activate a complement cascade that mediates internalization of complement-coated pathogens by phagocytic cells and lysis of pathogens by membrane attack complex formation (Forthal, 2014). The structure and function of Igs in teleost fish are similar to those in mammals (Mashoof & Criscitiello, 2016). Three classes

of Igs have been identified in teleost fish: IgM, IgD, and IgZ/T (Table 2) (Mashoof & Criscitiello, 2016).

The BCR in mammals consists of the membrane-bound antibody associated with the Ig- α /Ig- β heterodimer, known as CD79a/b, that is crucial for signal transduction because of the presence of ITAM domain (Treanor, 2012). Like TCR, antigen specificity of BCR and Igs are formed by V(D)J gene rearrangement mediated by RAG genes (Nguyen et al., 2016). Similar to higher vertebrates, the IgH gene in teleost fish is arranged by the configuration of multiple V(D)J gene segments, followed by constant (C) segments (V-D-J-C). However, the IgL chain is arranged in a repeating set of V-J-C segments (Bao et al., 2010; N. Danilova et al., 2005; J. Hikima et al., 2011).

In mammals, bone marrow is the main hematopoiesis site that produces B cells. However, teleost fish lack bone marrow, and B cells are developed in the anterior kidney, the main site of hematopoiesis in teleosts. Mammalian mature B cells are found in secondary lymphoid tissues, such as spleen and lymph nodes. Although lymph nodes are not present in teleost fish, mature B cells are found in the spleen and posterior kidney of teleost fish (Zwollo et al., 2005). Also, B cells are found in distinct tissues, such as the lamina propria of intestine and in the epithelium of skin and gill (Salinas, 2015; Salinas et al., 2011). Upon recognition of an antigen, B cells differentiate into short-lived plasma cells (SLPC) formed in extra-follicular sites of secondary lymphoid tissues, with a life span of 3-5 days, or long-lived plasma cells (LLPC) formed in germinal centers of secondary lymphoid organs, with a life span of several months to a lifetime (Nutt et al., 2015). A similar differentiation pattern of B cells to plasma cells has been described in teleost fish (Ye et al., 2011; Zwollo et al., 2005). The plasma cells of teleost fish are divided into SLPCs formed in the spleen and LLPCs located only in the anterior kidney of fish (Kaattari et al., 2005; Ye et al., 2011).

The germinal center (GC) is a microenvironment where somatic hypermutation and class-switch recombination reaction for antibody diversification and affinity maturation occur (De Silva & Klein, 2015).

Activation-induced cytidine deaminase (AID) mediates somatic hypermutation to increase the affinity of antibodies and mediates class-switch recombination to generate antibodies with specialized effector functions (Leeman-Neill et al., 2018). Hence, long-lived plasma cells produce high-affinity class-switched antibodies, whereas short-lived cells express low-affinity IgM antibodies (Khodadadi et al., 2019). While teleost fish lack GCs in immune tissues, melano-macrophage centers are formed by numerous macrophages in lymphoid tissues, such as the spleen, and have a similar function as GCs (Agius & Roberts, 2003). Furthermore, the AID enzyme of teleost fish that mediates somatic hypermutation and catalyzes class-switch recombination in mammalian B cells *in vitro* was first identified in channel catfish (Barreto et al., 2005; Magor, 2015; Saunders & Magor, 2004). Despite the presence of AID in teleost fish, they still lack class-switch recombination due to the IgH gene structure, thus resulting in low efficient affinity maturation response compared to mammals (Wakae et al., 2006).

Similar to T cells, B cells differentiate into memory B cells in GCs of lymphoid tissues after activation by antigenic interaction (as discussed in detail in ref. (Akkaya et al., 2020)). Also, teleost fish have developed immunological memory that provides a rapid and effective response to a pathogen previously encountered. For example, the formation of memory B cells in rainbow trout resulted in a faster and larger magnitude of secondary response to trinitrophenylated-keyhole limpet hemocyanin (Arkoosh & Kaattari, 1991). In addition, antibody affinity was higher in salmonids during the secondary response to the same antigen because of memory B cells (Ma et al., 2013). Moreover, vaccination with attenuated Viral Hemorrhagic Septicemia rhabdovirus (VHSV) induced the generation of memory B cells that elevated titers of serum IgM in rainbow trout in response to the same antigen (Magadan et al., 2018). Therefore, vaccination is a prophylactic method to protect teleost fish against economically devastating pathogens for several years post-immunization due to immunological memory (Findly R. C. et al., 2013; Firdaus-Nawi & Zamri-Saad, 2016).

In addition to the crucial role of B cells in adaptive immunity, two subsets of mammalian B cells, B-1 cells and marginal zone (MZ) B cells, contribute to innate immunity, including cytokine production, phagocytic capability, antigen presentation, and intracellular killing (Sunyer, 2013; Zouali & Richard, 2011). Similar to B-1 and MZ B cells of mammals, B cells in teleost fish are capable of phagocytosis and microbial killing of ingested bacteria. The phagocytic ability of B cells has been described in rainbow trout, and the formation of phagolysosome was observed in these fish (Li J et al., 2006). Moreover, zebrafish B cells showed strong phagocytic ability for particulate and soluble antigens and could present antigens to T cells like mammalian B cells (Zhu et al., 2013). Furthermore, the phagocytic

capacity of B cells isolated from the anterior kidney and peripheral blood was higher than neutrophils in Atlantic cod (Øverland et al., 2010). Previously, it was reported that large amounts of B cells in catfish blood were phagocytic (Esteban et al., 2015). Recently, our research group demonstrated both active uptake of *E. ictaluri* strains by anterior kidney B cells in catfish at 30° and 4° C and phagosome and/or phagolysosome formation in the cytoplasm of B cells (Kordon, Kalindamar, et al., 2019; Kordon et al., 2020). Also, internalized *E. ictaluri* strains were destroyed by catfish B cells at 30° C, but not at 4° C (Kordon et al., 2020).

Immunoglobulin M

IgM is known as the most ancient class of antibody found in all jawed vertebrates with unique features (Flajnik, 2002). In mammals, secreted IgM is present as a pentameric form associated with the joining (J) chain (Keyt et al., 2020). Immunoglobulin M, the first identified in plasma and the most prevalent in teleost fish, is described in two different forms: secreted and transmembrane form (Mashoof & Criscitiello, 2016). The secreted form of IgM is multimerized into a tetrameric form polymerized by interchain disulfide bonds due to the absence of J chain (Table 2) (Castro & Flajnik, 2014). However, some teleost fish species, such as rainbow trout, have monomer IgM in their serum (Elcombe et al., 1985; Wilson & Warr, 1992). In rainbow trout, the binding affinity of both monomeric and tetrameric IgM is similar, but the tetrameric form of IgM is more efficient to activate the complement system (Elcombe et al., 1985). In addition, the transmembrane form of IgM in B cells is one domain shorter than the secreted form, due to alternative splicing (Sahoo et al., 2008). Moreover, IgM, the only isotype in teleost fish, has two sub-types identified in Atlantic salmon and brown trout (Hordvik et al., 2002). Similar to mammals, teleost IgM contributes to both innate and adaptive immune responses, including complement activation, induced opsonization, lysis of pathogens and mediating cellular cytotoxicity through ADCC (Boshra et al., 2004; Smith et al., 2019). Also, IgM regulates agglutination for phagocytosis that results in the clearance of pathogens (Ye et al., 2013). In addition to plasma, IgM was detected in mucosal tissues, such as the skin and intestine of fish (Peleteiro & Richards, 1988; Zhang Y et al., 2010). Immunization induces an elevated titer of IgM in fish serum, but results in a weak improvement in affinity maturation compared to mammals (Kaattari et al., 2002; Ye et al., 2013). Recently, our research team detected significantly increased IgM levels in channel catfish serum following vaccination with *E. ictaluri* strains (Kordon et al, unpublished observation).

Immunoglobulin D

Immunoglobulin D is the second class of Igs identified in fish that shares sequence similarity with IgD

in mammals (Wilson M et al., 1997). However, teleost fish contain unique features of IgD due to the many forms (from 2 to 16) of IgD constant domains (Table 2) (Edholm et al., 2011; Parra et al., 2016; F. Ramirez-Gomez et al., 2012). In addition, the only transmembrane form of IgD has been found in teleost fish, with the exception of channel catfish, Japanese puffer, and rainbow trout, which have two forms, transmembrane and secretory (J.-i. Hikima et al., 2011; Francisco Ramirez-Gomez et al., 2012). Secreted IgD (which lacks the V domain) binds directly to Fc receptor of basophils functioning as a pattern recognition receptor. Thus, it can induce inflammatory responses, including the production of pro-inflammatory cytokines, antimicrobial, opsonizing, and B cell-activating factors (Chen et al., 2009; Edholm et al., 2010). Immunoglobulin D in teleost fish is mainly found in serum and also expressed in the anterior and posterior kidneys, spleen, and gill (Bengtén & Wilson, 2015). Furthermore, the level of IgD was higher in the gill of rainbow trout compared to IgM level (Smith et al., 2019).

Immunoglobulin T/Z

IgT/Z, the only teleost fish-specific Ig, was first identified in rainbow trout (IgT) and in zebrafish (IgZ) (Table 2) (Nadia Danilova et al., 2005; Hansen et al., 2005). IgT/Z functions similar to mammalian IgA (Mashoof et al., 2014). The IgT/Z level in the gut of rainbow trout was much higher (63 times) than that in serum, whereas the concentration of IgM in the serum was much higher compared to IgT/Z (Zhang Y et al., 2010). The IgT/Z in the gut of teleost fish contributes to immune responses against intestinal parasites and bacteria (Zhang Y et al., 2010). For example, the number of IgT+ B cells elevated in the gut of rainbow trout after parasitic intestinal infection, although the number of IgM+ B cells did not change in the same tissue (Zhang Y et al., 2010). In addition to IgT+ B cells, the IgT concentration was increased in the gut of surviving rainbow trout. However, the titer of parasite-specific IgM was higher in the sera of surviving animals (Zhang Y et al., 2010). Moreover, IgT+ B cells in teleost fish were found in skin-associated lymphoid tissue and secrete IgT into skin mucus (Xu et al., 2013). IgT is present in fish serum as a monomer but has a tetramer form in the gut mucus (Zhang Y et al., 2010). Some teleost fish species, such as channel catfish, lack IgT; therefore, IgM is the main Ig class in both serum and mucosal immunity (Findly R. C et al., 2013; Maki & Dickerson, 2003; Zhao et al., 2008).

Conclusion

Recent findings in fish immunology show that significant progress has been made in the mechanistic understanding of fish immune responses. Two branches of the fish immune system, innate and adaptive immunity, provide important information for

understanding the evolution of the immune system. Studies of fish with new reagents and powerful sequencing and knockout procedures will provide a valuable framework for assessment of the highly complex mammalian adaptive immune responses. In addition, the costs of fish infections to aquaculture can be overwhelming, making the failure of immunity a major risk for commercial fish farming. The presence of an adaptive immune system and, consequently, immune memory in teleost fish allows vaccination, the most appropriate method for disease control in aquaculture. In addition to vaccination, different approaches have been developed to control infectious diseases in aquaculture, such as high-quality diets including probiotics, prebiotics, and medicinal plants and treatment with antibiotics (Chinabut & Puttinaowarat, 2005). Furthermore, farm-level biosecurity precautions, including egg disinfection, water treatments, clean food, traffic control, intense quarantine measures, removal and disposal of mortalities, are preventive approaches to keep the facility safe before any disease outbreaks (Assefa & Abunna, 2018).

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