

REVIEW ARTICLE

RIG-I-Like Receptors (RLRs) and Toll Like Receptors (TLRs) Mediated Regulation of Type I Interferons (IFNs) Signaling In Fish

Ambika Binesh^{a*} and Kaliyamurthi Venkatachalam^a

a- Institute of Fisheries Post Graduate Studies, Tamil Nadu Dr. J. Jayalithaa Fisheries University, OMR Campus, Chennai – 603 103.

* Corresponding author's Mail ID: ambikabinesh@gmail.com and kaliyamurthi@tnfu.ac.in

ABSTRACT

Inflammatory mediators known as cytokines (class II) which comprised of Interferons (IFNs) play vital roles in host immune defense and the discovery of IFNs is considered as a pioneer in the field of immunology. Fish type I IFNs may signal to the downstream receptors in a same way as in mammals. In zebrafish, two group IFNs such as IFN1/4 mediated signaling through CRFB1 and CRFB5 receptor complex whereas IFN2/3 signal through CRFB2 and CRFB5 complex. Three families of pattern recognition receptors (PRRs) which includes retinoic acid-inducible gene 1 (RIG-I)-like receptors (RLRs), Toll-like receptors (TLRs), and cytosolic DNA sensors, are required in the type I IFN response in mammals.

Keywords: Type I IFNs; RLRs; TLRs; JAK-STAT

Received 11.05.2019

Revised 18.09.2019

Accepted 16.10.2019

How to cite this article:

A Binesh and K Venkatachalam. RIG-I-Like Receptors (RLRs) and Toll Like Receptors (TLRs) Mediated Regulation of Type I Interferons (IFNs) Signaling in Fish. Adv. Biores., Vol 10 [6] November 2019:10-14.

INTRODUCTION

Discovery of Interferons (IFNs) play an essential role in the field of biomedicine especially in immunological studies for the past sixty years. Inflammatory mediators known as cytokines (class II) which comprised of IFNs play vital roles in host immune defense [1], especially against viruses [2]. IFNs are categorised in to three types as type (I, II and III) based on their similarity in sequence, biological function and its organisation of genome [2]. Almost all nucleated cells respond to type I IFNs [3] [4]. This review attempts to summarize the recent discoveries on the type I IFNs systems in fish, which is regulated by RLRs and TLRs.

DISCOVERY AND CLASSIFICATION OF TYPE I IFNS

In 2003, three separate groups worked in zebrafish [5], green spotted pufferfish [6] and Atlantic salmon [7], identified first fish IFN gene. Many copies of type I IFN gene is present in fish similar to other vertebrate species based on their genome linkage, so the gene copy number varied from four in zebrafish to eleven in atlantic salmon [8] [9] and appear to exist in all fish species [10] [11]. Type I IFNs in fish was originally grouped into two types based on the number of cysteine residues (required indissulfide bond formation) as group I – contains two cysteine and group II – contains four cysteine residues [12]. Based on the order of discovery and location of chromosome, type I IFN gene copies are depicted by arabic numerals for instance, in zebrafish IFNs genes (four types IFNs 1-4) are classified into group I comprised of IFNs 1 and IFNs 4 whereas group II comprised of IFNs 2 and IFNs 3 [5] [8] [13]. Complex type I IFNs with subsets a, b, c, d, e and f are distinguished in salmonids [14] [9]. IFNs subtype h is the newly added and identified in perciformes [15] [16] [17].

RECEPTORS OF TYPE I IFN MEDIATED SIGNALING PATHWAY

Type I IFN-mediated signalling pathway initiated with the interaction between type I IFNs and their receptors IFNAR1 and IFNAR2 (heterodimeric receptor complex) in mammals [1]. Researchers mainly immunologists suggested that the two receptors (IFNAR1 and IFNAR2) pertain to the class II cytokine

receptor family, which is known as cytokine receptor family B (CRFB) in fish[6] [18]. There are 17 CRFB members are present in zebrafish and pufferfish revealed in genome-wide sequencing. CRFB1 and CRFB2 showed homology to mammalian IFNAR2 whereas CRFB5 is homologue to mammalian IFNAR1[18]. In zebrafish, two group IFNs such as IFN1(IFNa) and IFN4 (IFNd) mediated signaling through CRFB1 and CRFB5 receptor complex whereas IFN2/3 (IFNc) signal through CRFB2 and CRFB5 complex[8] [19].

REGULATION OF TYPE I IFN-MEDIATED SIGNALING PATHWAY

In mammals, interaction between type I IFNs and their receptors stimulates the binding of tyrosine kinase 2 (TYK2) to IFNAR1 and Janus kinase 1 (JAK1) to IFNAR2 mediated through JAK-STAT pathway[20]. Phosphorylation of STAT1 and STAT2 (Signal Transducers and Activators of Transcription) by the aforementioned kinases are dimerized and put together with IFN regulatory factor (IRF) 9 to form an IFN stimulated gene factor 3 complex (ISGF3). Then the translocation of this trimolecular complex to the nucleus activate the transcription by binding to IFN-stimulated response elements (ISREs)[21](Figure 1). Copious immunogenetics studies revealed components of JAK-STAT pathway such as TYK2, JAK1, STAT1, STAT2 and IRF9, also exist in fish[22] [23][24].

Table 1. Conservation of JAK- STAT mechanism

Species	IFNAR1 IFNAR2	TYK1 TYK2	References
Mammals	Absence of enzymatic activity	Presence of Kinase activity	[25]
Fish	Absence of enzymatic activity	Presence of Kinase activity (Atlantic salmon TYK2)	[26]
Grass carp	Absence of enzymatic activity	Presence of Kinase activity (CRFB1 and CRFB5)	[27]

Interestingly, Fish type I IFNs may signal to the downstream receptors in a same way as in mammals. Two STAT1 genes (STAT1a and STAT1b) in zebrafish show similarity with human STAT genes. All five domains of human STAT1a is identified in zebrafish STAT1a at the same time, lack of C-terminal transcriptional activation domain is observed in both human and zebrafish STAT1b[28]. Zebrafish STAT1a is able to rescue IFN-mediated growth suppression in a STAT1-deficient human cell line, thus play an important role in type I IFN mediated signalling[29] [30].In congruent with this findings, phosphorylation and translocation of Atlantic salmon STAT1a in to nucleus was observed with recombinant IFNa1treatment[31]. In orange-spotted grouper, overexpression of STAT1a show antiviral activity against iridovirus and nodavirus by upregulating the ISGs expression[30].However, gibel carp STAT1(resembles like zebrafish and human STAT1b) induce ISG and inhibit viral infection[32]. Invitro studies in mandarin fish[16]and Co-IP assay in salmon[33]revealed that the ISGF3 complex (STAT1, STAT2 and IRF9) conserved in fish. At the same time, studies show that fish IRF9 is essential for the type I IFN-mediated signalling[34][35][36].

Numerous studies suggested that three families of pattern recognition receptors (PRRs) which includes retinoic acid-inducible gene I (RIG-I)-like receptors (RLRs), Toll-like receptors (TLRs), and cytosolic DNA sensors, are required in the type I IFN response in mammals[37]and in fish[38].

RLR-MEDIATED TYPE I IFN RESPONSE

Family of cytosolic receptors that is RLRs which recognize viral RNAs with three members, including RIG-I, melanoma differentiation-associated gene 5 (MDA5) and laboratory of genetics and physiology 2 (LGP2)[39]and the downstream molecules mitochondrial antiviral signalling protein (MAVS), and TANK binding kinase 1 (TBK1) are found conserved in fish.Upon the recognition of dsRNA from viruses, RLR components RIG-I/MDA5 recruits MAVS which then become associated with TRAF3 and TBK1, leading to the phosphorylation and activation of IRF3/IRF7 to elicit type I IFN response(Figure 1). RIG-I seems to be lost in fish of Acanthopterygii whereas MDA5 and LGP2 appear to exist in all fish species[40] [41].

Notwithstanding, it remains to be determined whether there are multicopy genes STAT1 and STAT2 in a wide range of fish taxa. In addition, investigating the functional similarity and divergence of STAT1 and STAT2 fish multicopy genes in Type IFN signalling will be interesting.However, there are still two important and intriguing questions to be answered with regard to fish RLR-mediated type IFN response. First, since RIG-I appears to be lost in certain groups of fish species[40][41], clarifying the mechanism for

compensating for RIG-I deficiency in these species would be interesting. Second, much more research is needed to fully elucidate the dual roles of Type I IFN-mediated antiviral response fish LGP2 and to understand the factors that influence the functional switch of LGP2 in fish[42].

TLR-MEDIATED TYPE I IFN RESPONSE

Type I integral proteins also known as TLRs comprising an ectodomain containing ligand-rich repeats (LRRs), a transmembrane region and cytosolic Toll-IL-1 receptor (TIR) domains that mediate downstream signaling pathways[43]. TLR3, TLR19 and TLR22 are mainly involved in the activation of type I IFN response in fish[44] [45]which comprises a larger TLR collection, which shows similarity to mammalian TLRs and non-mammalian TLRs [46]. Among the three TLRs in fish, TLR3 is localized in endoplasmic reticulum and recognizes short dsRNA, TLR22 is located particularly in plasma membrane and recognizes long dsRNA and recruit TRIF to elicit type I IFN response [47] [48](Figure 1). The studies from *Takifugurubripes* commonly called fugu/puffer showed the ligand recognition and type I IFN-inducing activity of fish TLR3 and TLR22 whereas TLR19 response identified in grass carp.

In other fish species, the functional properties of these TLRs remain to be further characterized. Furthermore, although fish TLR9 may bind CpG-containing DNA as in mammals [49][50], it remains to be shown whether it can activate Type I IFN production. Future studies are also required to determine whether ssRNA can be conservatively recognized by fish TLR7 and TLR8 and to trigger type IFN response[42].

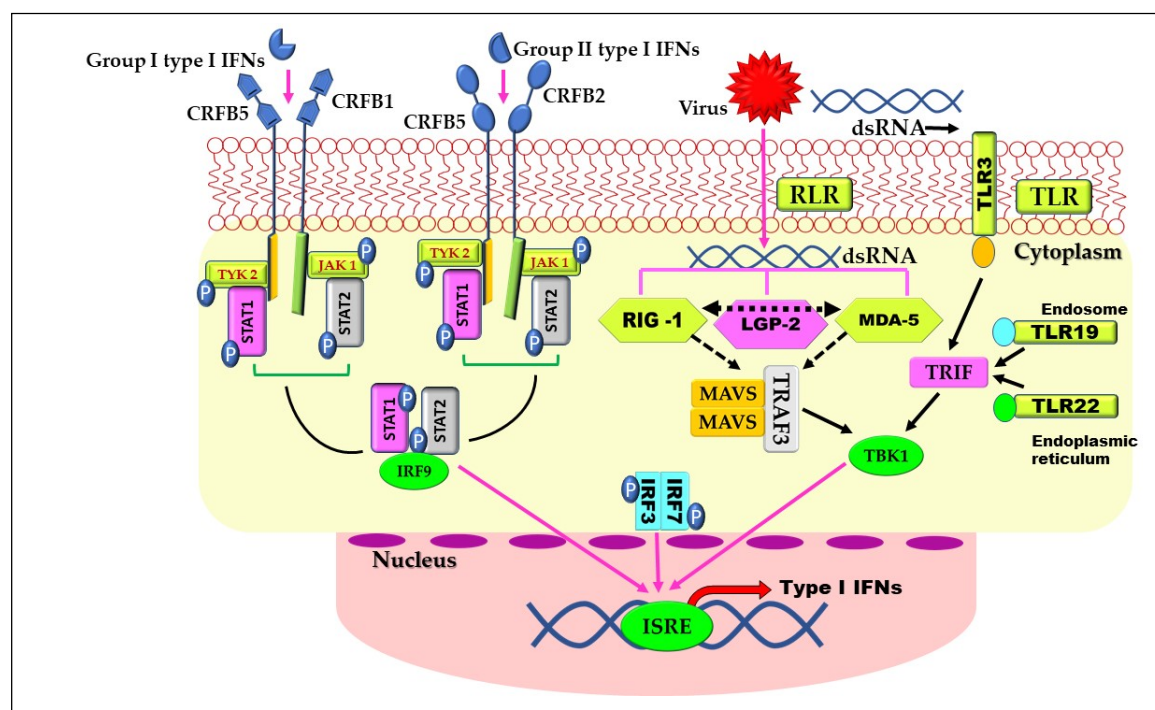


Figure 1: Signalling pathway model for type I IFNs in fish. Two groups of IFNs interact with its common receptor CRFB5 and the two different receptors CRFB1 and CRFB2. Upon ligand – receptor interaction TYK2 and JAK1 are recruited and activated, leads to the phosphorylation of STAT1 and STAT2 which become dimerized and form a trimolecular complex with ISGF3, ultimately translocates to the nucleus and binds to ISREs, thus activating the transcription. Upon the recognition of dsRNA from viruses, RLR components RIG-I/MDA5 recruits MAVS which then become associated with TRAF3 and TBK1, whereas TLR3, TLR19 and TLR22 which recruit TRIF and TBK1 leading to the phosphorylation and activation of IRF3/IRF7 to elicit type I IFN response.

CONCLUSIONS AND FUTURE PERSPECTIVE

First, since RIG-I appears to be lost in certain groups of fish species, it would be interesting to clarify the mechanism in these species to compensate for the RIG-I deficiency. Second, much more research is needed to fully elucidate the dual roles of Type I IFN-mediated antiviral response of fish LGP2 and the factors influencing the functional switch of LGP2 in fish. Future studies are also needed to determine whether fish TLR7 and TLR8 can conservatively recognize ssRNA and trigger the response of type I IFN.

CONFLICTS OF INTEREST

All authors disclose that there are no conflicts of interest that could inappropriately influence the outcome of the study.

REFERENCES

1. Binesh A, Devaraj SN, Halagowder D. (2018). Atherogenic diet induced lipid accumulation induced NF κ B level in heart, liver and brain of Wistar rat and diosgenin as an anti-inflammatory agent. *Life Sci.* ;196:28-37.
2. Pestka S, Krause CD, Walter MR. (2004). Interferons, interferon like cytokines, and their receptors. *Immunological Reviews.* 202: 8–32.
3. Lazear HM, Nice TJ, (2015). Diamond MS. Interferon-lambda: immune functions at barrier surfaces and beyond. *Immunity*;43: 15–28.
4. Teijaro JR. (2016). Type I interferons in viral control and immune regulation. *Current Opinion in Virology.* 16: 31–40.
5. Altmann SM, Mellon MT, Distel DL, Kim CH. (2003). Molecular and functional analysis of an interferon gene from the zebrafish, *Danio rerio*. *Journal of Virology.* 77: 1992–2002.
6. Lutfalla G, RoestCrollius H, Stange-Thomann N, Jaillon O, Mogensen K, Monneron D. (2003). Comparative genomic analysis reveals independent expansion of a lineage-specific gene family in vertebrates: the class II cytokine receptors and their ligands in mammals and fish. *BMC Genomics.* 4: 29.
7. Robertsen B, Bergan V, Rokenes T, Larsen R, (2003). Albuquerque A. Atlantic salmon interferon genes: cloning, sequence analysis, expression, and biological activity. *Journal of Interferon and Cytokine Research.* 23: 601–612.
8. Aggad D, Mazel M, Boudinot P, Mogensen KE, Hamming OJ, Hartmann R et al. (2009). The two groups of zebrafish virus induced interferons signal via distinct receptors with specific and shared chains. *Journal of Immunology.* 183: 3924–3931.
9. Svingerud T, Solstad T, Sun B, Nyrud ML, Kileng O, Greiner Tollersrud L et al. (2012). Atlantic salmon type I IFN subtypes show differences in antiviral activity and cell-dependent expression: evidence for high IFN β /IFN γ -producing cells in fish lymphoid tissues. *Journal of Immunology.* 189: 5912–5923.
10. Secombes CJ, Zou J. (2017). Evolution of interferons and interferon receptors. *Frontiers in Immunology.* 8: 209.
11. Robertsen B. (2018). The role of type I interferons in innate and adaptive immunity against viruses in Atlantic salmon. *Developmental and Comparative Immunology.* 80: 41–52.
12. Chang M, Nie P, Collet B, Secombes CJ, Zou J.(2009). Identification of an additional two-cysteine containing type I interferon in rainbow trout *Oncorhynchus mykiss* provides evidence of a major gene duplication event within this gene family in teleosts. *Immunogenetics.* 61: 315–325.
13. Lopez-Munoz A, Roca FJ, Meseguer J, Mulero V. (2009). New insights into the evolution of IFNs: zebrafish group II IFNs induce a rapid and transient expression of IFN-dependent genes and display powerful antiviral activities. *Journal of Immunology.* 182: 3440–3449.
14. Sun B, Robertsen B, Wang Z, Liu B.(2009). Identification of an Atlantic salmon IFN multigene cluster encoding three IFN subtypes with very different expression properties. *Developmental and Comparative Immunology.*33: 547–558.
15. Ding Y, Ao J, Huang X, Chen X. (2016). Identification of two subgroups of type I IFNs in perciform fish large yellow croaker *Larimichthys crocea* provides novel insights into function and regulation of fish type I IFNs. *Frontiers in Immunology.* 7:343.
16. Laghari ZA, Chen SN, Li L, Huang B, Gan Z, Zhou Y et al. (2018). Functional, signalling and transcriptional differences of three distinct type I IFNs in a perciform fish, the mandarin fish *Siniperca chuatsi*. *Developmental and Comparative Immunology.* 84: 94–108.
17. Milne DJ, Campoverde C, Andree KB, Chen X, Zou J, Secombes CJ. (2018). The discovery and comparative expression analysis of three distinct type I interferons in the perciform fish, meagre (*Argyrosomus sregius*). *Developmental and Comparative Immunology.* 84: 123–132.
18. Stein C, Caccamo M, Laird G, Leptin M.(2007). Conservation and divergence of gene families encoding components of innate immune response systems in zebrafish. *Genome Biology.* 8: R251.
19. Levraud JP, Boudinot P, Colin I, Benmansour A, Peyrieras N, Herbomel P et al.(2007). Identification of the zebrafish IFN receptor: implications for the origin of the vertebrate IFN system. *Journal of Immunology.* 178: 4385–4394.
20. Stark GR, Darnell JE Jr.(2012). The JAK-STAT pathway at twenty. *Immunity.* 36: 503–514.
21. Platanias LC. (2005). Mechanisms of type-I- and type-II-interferon mediated signalling. *Nature Reviews Immunology.* 5: 375–386.
22. Xu C, Evensen O, Munang'andu HM. (2016). A de novo transcriptome analysis shows that modulation of the JAK-STAT signaling pathway by salmonid alphavirus subtype 3 favours virus replication in macrophage/dendritic-like TO-cells. *BMC Genomics.* 17: 390.
23. Jin Y, Zhou T, Li N, Liu S, Xu X, Pan Y et al. (2018). JAK and STAT members in channel catfish: identification, phylogenetic analysis and expression profiling after Edwardsiella ictaluri infection. *Developmental and Comparative Immunology.* 81: 334–341.
24. Binesh A, Devaraj SN, Halagowder D. (2019). Molecular interaction of NF κ B and NICD in monocyte-macrophage differentiation is a target for intervention in atherosclerosis. *J Cell Physiol.* 234(5):7040-7050.

25. Piehler J, Thomas C, Garcia KC, Schreiber G. (2012). Structural and dynamic determinants of type I interferon receptor assembly and their functional interpretation. *Immunological Reviews*. 250: 317–334.
26. Sobhkhez M, Hansen T, Iliev DB, Skjesol A, Jorgensen JB. (2013). The Atlantic salmon protein tyrosine kinase Tyk2: molecular cloning, modulation of expression and function. *Developmental and Comparative Immunology*. 41: 553–563.
27. Hou Q, Gong R, Liu X, Mao H, Xu X, Liu D et al. (2017). Poly I: C facilitates the phosphorylation of Ctenopharyngodonidellus type I IFN receptor subunits and JAK kinase. *Fish and Shellfish Immunology*. 60: 13–20.
28. Song H, Yan YL, Titus T, He X, Postlethwait JH. (2011). The role of STAT1b in zebrafish hematopoiesis. *Mechanisms of Development*. 128: 442–456.
29. Oates AC, Wollberg P, Pratt SJ, Paw BH, Johnson SL, Ho RK et al. (1999). Zebrafish stat3 is expressed in restricted tissues during embryogenesis and stat1 rescues cytokine signalling in a STAT1-deficient human cell line. *Developmental Dynamics*. 215: 352–370.
30. Zhang J, Huang X, Ni S, Liu J, Hu Y, Yang Y et al. (2017). Grouper STAT1a is involved in antiviral immune response against iridovirus and nodavirus infection. *Fish and Shellfish Immunology*. 70: 351–360.
31. Skjesol A, Hansen T, Shi CY, Thim HL, (2010). Jorgensen JB. Structural and functional studies of STAT1 from Atlantic salmon (*Salmo salar*). *BMC Immunology*. 11: 17.
32. Yu FF, Zhang YB, Liu TK, Liu Y, Sun F, Jiang J et al. (2010). Fish virus-induced interferon exerts antiviral function through STAT1 pathway. *Molecular Immunology*. 47: 2330–2341.
33. Sobhkhez M, Skjesol A, Thomassen E, Tollersrud LG, Iliev DB, Sun B et al. (2014). Structural and functional characterization of salmon STAT1, STAT2 and IRF9 homologs sheds light on interferon signaling in teleosts. *FEBS Open Bio*. 4: 858–871.
34. Shi J, Zhang YB, Liu TK, Sun F, Gui JF. (2012). Subcellular localization and functional characterization of a fish IRF9 from crucian carp *Carassius auratus*. *Fish and Shellfish Immunology*. 33: 258–266.
35. Huang CJ, Chou CM, Lien HW, Chu CY, Ho JY, Wu Y et al. (2017). IRF9-Stat2 fusion protein as an innate immune inducer to activate Mx and interferon-stimulated gene expression in zebrafish larvae. *Marine Biotechnology*. 19: 310–319.
36. Wu Z, Wang L, Xu X, Lin G, Mao H, Ran X et al. (2017). Interaction of IRF9 and STAT2 synergistically up-regulates IFN and PKR transcription in *Ctenopharyngodonidella*. *Molecular Immunology*. 85: 273–282.
37. Wu J, Chen ZJ. (2014). Innate immune sensing and signaling of cytosolic nucleic acids. *Annual Review of Immunology*. 32: 461–488.
38. Poynter S, Lisser G, Monjo A, DeWitte-Orr S. (2015). Sensors of infection: viral nucleic acid PRRs in fish. *Biology*. 4: 460–493.
39. Loo YM, Gale M Jr. (2011). Immune signaling by RIG-I-like receptors. *Immunity*. 34: 680–692.
40. Zou J, Chang M, Nie P, Secombes CJ. (2009). Origin and evolution of the RIG-I like RNA helicase gene family. *BMC Evolutionary Biology*. 9: 85.
41. Chen SN, Zou PF, Nie P. (2017). Retinoic acid-inducible gene I (RIG-I)-like receptors (RLRs) in fish: current knowledge and future perspectives. *Immunology*. 151: 16–25.
42. Zhen Gan, Shan Nan Chen, Bei Huang, Jun Zou, Pin Nie. (2019). Fish type I and type II interferons: composition, receptor usage, production and function. *Reviews in Aquaculture*. 1–32.
43. Kawai T, Akira S. (2010). The role of pattern-recognition receptors in innate immunity: update on Toll-like receptors. *Nature Immunology*. 11: 373–384.
44. Matsuo A, Oshiumi H, Tsujita T, Mitani H, Kasai H, Yoshimizu M et al. (2008). Teleost TLR22 recognizes RNA duplex to induce IFN and protect cells from birnaviruses. *Journal of Immunology*. 181: 3474–3485.
45. Ji J, Rao Y, Wan Q, Liao Z, Su J. (2018). Teleost-specific TLR19 localizes to endosome, recognizes dsRNA, recruits TRIF, triggers both IFN and NF-kappaB pathways, and protects cells from grass carp reovirus infection. *Journal of Immunology*. 200: 573–585.
46. Pietretti D, Wiegertjes GF. (2014). Ligand specificities of Toll-like receptors in fish: indications from infection studies. *Developmental and Comparative Immunology*. 43: 205–222.
47. Sullivan C, Postlethwait JH, Lage CR, Millard PJ, Kim CH. (2007). Evidence for evolving Toll-IL-1 receptor-containing adaptor molecule function in vertebrates. *Journal of Immunology*. 178: 4517–4527.
48. Funami K, Matsumoto M, Oshiumi H, Akazawa T, Yamamoto A, Seya T. (2004). The cytoplasmic 'linker region' in Toll-like receptor 3 controls receptor localization and signaling. *International Immunology*. 16: 1143–1154.
49. Iliev DB, Skjaeveland I, Jorgensen JB. (2013). CpG oligonucleotides bind TLR9 and RRM-containing proteins in Atlantic salmon (*Salmo salar*). *BMC Immunology*. 14: 12.
50. Yeh DW, Liu YL, Lo YC, Yuh CH, Yu GY, Lo JF et al. (2013). Toll-like receptor 9 and 21 have different ligand recognition profiles and cooperatively mediate activity of CpG-oligodeoxynucleotides in zebrafish. *Proceedings of the National Academy of Sciences of the United States of America*. 110: 20711–20716.

Copyright: © 2019 Society of Education. This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.