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**Dissecting the immune pathways stimulated following injection vaccination of rainbow trout (*Oncorhynchus mykiss*) against enteric redmouth disease (ERM)**

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24

**Abstract**

25

26 Enteric redmouth disease (ERM or yersiniosis) is one of the most important diseases of  
27 salmonids and leads to significant economic losses. It is caused by the Gram-negative  
28 bacterium *Yersinia ruckeri* but can be controlled by bacterin vaccination. The first  
29 commercial ERM vaccine was licenced in 1976 and is one of the most significant and  
30 successful health practices within the aquaculture industry. Although ERM vaccination  
31 provides complete protection, knowledge of the host immune response to the vaccine and the  
32 molecular mechanisms that underpin the protection elicited is limited. In this report, we  
33 analysed the expression in spleen and gills of a large set of genes encoding for cytokines,  
34 acute phase proteins (APPs) and antimicrobial peptides (AMPs) in response to ERM  
35 vaccination in rainbow trout, *Oncorhynchus mykiss*. Many immune genes in teleost fish are  
36 known to have multiple paralogues that can show differential responses to ERM vaccination,  
37 highlighting the necessity to determine whether all of the genes present react in a similar  
38 manner. ERM vaccination immediately activated a balanced inflammatory response with  
39 correlated expression of both pro- and anti-inflammatory cytokines (eg IL-1 $\beta$ 1-2, TNF- $\alpha$ 1-3,  
40 IL-6, IL-8 and IL-10A etc.) in the spleen. The increase of pro-inflammatory cytokines may  
41 explain the systemic upregulation of APPs (eg serum amyloid A protein and serum amyloid  
42 protein P) and AMPs (eg cathelicidins and hepcidin) seen in both spleen and gills. We also  
43 observed an upregulation of all the  $\alpha$ -chains but only one  $\beta$ -chain (p40B2) of the IL-12 family  
44 cytokines, that suggests specific IL-12 and IL-23 isoforms with distinct functions might be  
45 produced in the spleen of vaccinated fish. Notably the expression of Th1 cytokines (IFN- $\gamma$ 1-  
46 2) and a Th17 cytokine (IL-17A/F1a) was also up-regulated and correlated with enhanced  
47 expression of the IL-12 family  $\alpha$ -chains, and the majority of pro- and anti-inflammatory  
48 cytokines, APPs and AMPs. These expression profiles may suggest that ERM vaccination  
49 activates host innate immunity and expression of specific IL-12 and IL-23 isoforms leading  
50 to a Th1 and Th17 biased immune response. A late induction of Th2 cytokines (IL-4/13B1-2)  
51 was also observed, that may have a homeostatic role and/or involvement in antibody  
52 production. This study has increased our understanding of the host immune response to ERM  
53 vaccination and the adaptive pathways involved. The early responses of a set of genes  
54 established in this study may provide essential information and function as biomarkers in  
55 future vaccine development in aquaculture.

56

57 **Key words:** Enteric redmouth disease, vaccination, cytokine, acute phase protein,  
58 antimicrobial peptide, gene expression, T helper 1, T helper 17, spleen, rainbow trout,

## 59 1. Introduction

60 Enteric redmouth disease (ERM, yersiniosis) is one of the most important diseases of  
61 salmonids and leads to significant economic losses [1-2]. The disease is caused by *Yersinia*  
62 *ruckeri*, a Gram-negative rod-shaped enterobacterium, first isolated from rainbow trout  
63 *Oncorhynchus mykiss* in the Hagerman Valley of Idaho, USA in the 1950s [3-5]. ERM  
64 infected rainbow trout show a general septicaemia with an inflammatory response in all  
65 tissues. The gills are the first route of entry for infection, with the spleen a major secondary  
66 lymphoid organ associated with the response [1,6].

67

68 Vaccination as a means of controlling ERM is one of the most significant and successful  
69 health practices within the aquaculture industry, and has helped to reduce the use of  
70 environmental unfriendly antibiotics to control this bacterial disease [7]. The first commercial  
71 fish vaccine for ERM was licensed in 1976, and was a bacterin prepared from formalin-killed  
72 whole cells of *Y. ruckeri*. A simple immersion is effective but intraperitoneal (ip) injection  
73 provides superior and long lasting protection against ERM [8]. However, ERM outbreaks  
74 have occurred recently due to the emergence of atypical biotypes of *Y. ruckeri* [2]. This issue  
75 can be resolved by including new isolates in the improved vaccine preparation, as shown with  
76 AquaVac(®) RELERA™ that contains both biotypes 1 and 2 of *Y. ruckeri* and provides  
77 better protection [8-9]. Moreover, distinct strains of *Y. ruckeri* are present in the environment  
78 and can evolve with the introduction of susceptible hosts and vaccination of salmonids in  
79 aquaculture [10-11]. As the production of global aquaculture continues to increase it is likely  
80 that vaccines against other bacterial disease will encounter similar issues and will need the  
81 development of improved formulations.

82

83 The development of fish vaccines has been largely empirical, based on whether a formulation  
84 is effective at increasing survival post-disease challenge [12]. This is unsatisfactory from  
85 both ethical and scientific perspectives. There is a clear need to establish methods to improve  
86 fish vaccine development, such as pre-screening of candidate vaccines in the early  
87 development phase and the quality control of vaccines. The immune system is a network of  
88 specialized cell types and tissues that communicates via cytokines and direct contact, to  
89 orchestrate specific types of responses to effect protection [13]. It is known in mammals that

90 different vaccines induce distinct transcriptional signatures, representing the highly  
91 specialized defence mechanisms that can be elicited to cope with the different pathogens and  
92 insults a host may encounter. In this context, the early innate responses to vaccines are likely  
93 critical instructors for the development of adaptive immunity at later time points [14]. For the  
94 development of new vaccines, it is crucial to determine molecular signatures of vaccine-  
95 induced immune responses to gain a better understanding of the pathways involved and  
96 mechanisms that underpin protection, as well as to predict vaccine performance.

97

98 As a highly efficacious vaccine, the ERM vaccine provides a useful model for the  
99 investigation of fish immune responses to vaccines and bacterial diseases. Thus past studies  
100 have examined immune gene responses to ERM after vaccination [15-16] or after vaccination  
101 and challenge [9,17-18]. However, only a relatively small number of genes were investigated  
102 in these studies and often in the absence of information of the paralogues present [15-18].  
103 With the recent release of salmonid genomes [19-20] and the identification of a large number  
104 of cytokine genes including the paralogues for IL-1 $\beta$  [21], TNF- $\alpha$  [22], IL-17A/F [23-24], IL-  
105 4/13 [25] and subunits of the IL-12 family [26-29] in salmonids, it is timely to revisit the  
106 early immune responses to ERM vaccination.

107

108 Hence, in this study ERM vaccination was used as a model to investigate the early cytokine  
109 responses in rainbow trout in two major and relevant immune organs, the spleen and gills.  
110 The expression of acute phase proteins (APPs) and anti-microbial peptides (AMPs) was also  
111 examined. We found that intraperitoneal injection vaccination induces an early balanced  
112 expression of pro- and anti-inflammatory cytokines and adaptive cytokines in the spleen, with  
113 a heightened expression of APPs and AMPs in both spleen and gills.

114

115

## 116 2. Materials and Methods

### 117 2.1 Fish maintenance and rearing condition

118 Apparently healthy rainbow trout with no history of infection were purchased from the Mill  
119 of Elrich Trout Fishery (Aberdeenshire, Scotland, UK) and maintained in 1-m-diameter  
120 fibreglass tanks with recirculating freshwater at 14±1°C at the Scottish fish immunology  
121 research centre, the University of Aberdeen, UK. At least 10 fish from each batch were  
122 screened for potential bacterial infection by taking head kidney swabs and growing on tryptic  
123 soy broth (Sigma, UK) agar plates. No bacterial growth was seen. Fish were fed (2%  
124 biomass) twice a day with a commercial diet (EWOS) and given two weeks for acclimation  
125 prior to vaccination.

126

### 127 2.2 Fish vaccination

128 The commercial vaccine AquaVac<sup>TM</sup> ERM (MSD Animal Health), a formalin-inactivated  
129 bacterin containing not less than 5 x 10<sup>9</sup> cells per ml of *Y. ruckeri* (Hagerman strain type 1),  
130 was used in this study. The vaccine trial was described previously [30]. Briefly, a group of 24  
131 fish (mean ± SEM = 48.8 ± 1.5) were vaccinated by intraperitoneal injection (ip) of 0.1 ml of  
132 vaccine following manufacturer's instructions. The same number of fish were injected with  
133 phosphate buffer saline (PBS) as the control. The vaccinated and control groups were kept in  
134 separate 1-m-diameter fibreglass tanks in a single recirculating freshwater system at 14±1°C.  
135 Fish handling and experimental protocols comply with the Guidelines of the European Union  
136 Council (2010/63/EU) for the use of laboratory animals, and were carried out under UK  
137 Home Office project licence PPL 60/4013, approved by the ethics committee at the  
138 University of Aberdeen.

### 139 2.3 Sampling, total RNA extraction and cDNA synthesis

140 Six fish from both the vaccinated and control groups were killed at 1, 3, 7 and 14 days post  
141 vaccination. Spleen and gills were taken from each fish and homogenised separately in TRI  
142 reagent (Sigma, UK). Total RNA extraction and cDNA synthesis was as described previously  
143 [17, 31]. The synthesised cDNA samples were diluted in TE buffer (10 mM Tris-HCl, 1mM  
144 EDTA, pH8.0) and stored at -20°C ready for real-time PCR analysis.

145

## 146 **2.4 Gene expression analysis by real-time PCR**

147 The expression of a set of genes for acute phase proteins (APPs), antimicrobial peptides  
148 (AMPs) and cytokines using real time PCR were performed as described previously [25,32-  
149 33]. Briefly, the PCR amplification was performed using a LightCycler® 480 Instrument II  
150 (Roche Applied Science) and 384 multiwell plates in a 10 µl reaction using SYBR® Green I  
151 (Invitrogen™, Carlsbad, USA) and IMMOLASE™ DNA Polymerase (Bioline, UK), and  
152 expression levels calculated using the 'LightCycler® 480 software version 1.5. Elongation  
153 factor-1α (EF-1α), a house keeping gene, was used as an internal control. The sequences of  
154 primers used, and the DDBJ/ENA/GenBank accession number of the sequence the primers  
155 were designed against are listed for each gene in **Table 1**. At least one of each primer pair  
156 was designed to cross an intron and were tested to ensure that PCR products could only be  
157 amplified from cDNA samples and not from genomic DNA. The cp value (the crossing point  
158 at which the fluorescence crosses the threshold, means+SEM) of EF-1α was 11.31+0.25  
159 (spleen samples) and 10.44+0.11 (gill samples). The expression of each gene was first  
160 normalized to that of EF-1α, and expressed as a fold change relative to the expression level of  
161 control fish at the same time points.

162

## 163 **2.5 Data and statistical analysis**

164 For the statistical analysis, all of the gene expression data were first calculated as arbitrary  
165 units after normalization to the expression level of the house keeping gene EF-1α. Then the  
166 data were log<sub>2</sub> transformed to improve the normality of real-time quantitative PCR  
167 measurements, as described previously [31]. One way-ANOVA and Bonferroni post hoc tests  
168 were used to analyse all of the expression data using IBM SPSS statistics 22 software (SPSS  
169 Inc., Chicago, IL, USA). Differences between vaccinated and control groups for each time  
170 point were considered statistically significant at  $p < 0.05$ . In addition, we also undertook a  
171 correlation analysis of vaccine modulated genes using the Spearman rank order correlation  
172 test to look for associations between expression patterns. The gene expression levels in spleen  
173 samples at day 1 and day 3 (N=12) were used for this analysis, since the majority of gene  
174 expression changes seen were during this period.

175

### 176 3. Results

177 The expression of 63 genes, including the house keeping gene EF-1 $\alpha$ , 11 APPs and AMPs, 46  
178 cytokine genes and 5 master transcription factors, were analysed by real-time RT-PCR in this  
179 study. To give an indication of relative transcription level in the spleen and gills,  $\Delta$ Cp that is  
180 the Cp of the target gene minus that of EF-1 $\alpha$ , were provided for the control fish at day 1 in  
181 **Table 1**.

182

#### 183 3.1 ERM-vaccination activates an immediate expression of APP and AMP genes in both 184 spleen and gills

185

186 APPs, eg serum amyloid A protein (SAA) and serum amyloid protein P (SAP), and AMPs,  
187 eg cathelicidins (CATH) and hepcidin, are evolutionarily conserved effector molecules of the  
188 innate immune system that have important roles in the resolution of infection and activation  
189 of the adaptive immune response [34-35] The expression of some APPs and AMPs has been  
190 shown to be induced by *Y. ruckeri* infection previously [36]. Thus their expression was  
191 examined first in response to ERM vaccination. Trout SAA was highly expressed in spleen  
192 and gills ( $\Delta$ Cp=8, **Table 1**). Its expression was significantly induced in spleen by ERM  
193 vaccination at day 1, 3 and 7, and peaked at day 1 (89-fold) and in the gills at day 1 (18-fold,  
194 **Fig. 1A**). One of the SAP paralogues, SAP1 that was lowly expressed constitutively, was also  
195 induced in spleen at day 1 and 3 (**Fig. 1B and C**).

196

197 The constitutive expression of the AMPs, CATH1, CATH2 [35] and Hepcidin [34] was  
198 relatively high in spleen and gills ( $\Delta$ Cp=9 to 16, **Table 1**). CATH1 expression was induced at  
199 day 1 and day 3 in spleen but was refractory in gills after ERM vaccination (**Fig. 1D**).  
200 CATH2 expression was highly induced and peaked at day 1 in both spleen (385 fold) and  
201 gills (32 fold) and the heightened expression lasted until day 7 in the spleen and day 3 in the  
202 gills after vaccination (**Fig. 1E**). Hepcidin expression was also induced at day 1 in both  
203 spleen (43 fold) and gills (59 fold, **Fig. 1F**) by ERM vaccination. The expression of other  
204 AMPs, including LEAP2A [37] and  $\beta$ -defensins [38] was refractory in both spleen and gills  
205 (**Table 2 and S1**).

206

#### 207 3.2 ERM-vaccination induces an early correlated upregulation of pro- and anti- 208 inflammatory cytokine expression in the spleen



209 A successful vaccine is expected to activate the innate immune system to express pro- and  
210 anti-inflammatory cytokines. Several such cytokines have been investigated previously [15-  
211 16]. However, many other cytokines discovered recently, especially the paralogues, have not  
212 been examined after ERM vaccination. Therefore, a large number of cytokine genes  
213 including all the known paralogues have been investigated in this study. Three active IL-1 $\beta$   
214 paralogues are present in salmonids [21]. IL-1 $\beta$ 1 and IL-1 $\beta$ 3 have relatively high constitutive  
215 expression in spleen and gills, compared to IL-1 $\beta$ 2 (**Table 1**). The expression of both IL-1 $\beta$ 1  
216 and IL-1 $\beta$ 2 was induced at day 1 and 3, and peaked at day 1 in the spleen (**Fig. 2A-B**), but  
217 was not modulated in the gills (**Table S1**), of vaccinated fish. IL-1 $\beta$ 3 expression was  
218 refractory (**Table 2 and S1**). In terms of fold change, IL-1 $\beta$ 2 expression was more inducible  
219 (peaked at 49 fold) than IL-1 $\beta$ 1 (peaked at 17 fold). At least three TNF- $\alpha$  paralogues are  
220 present in rainbow trout [22]. All the three genes (TNF- $\alpha$ 1-3) were up-regulated in spleen at  
221 day 1, with TNF- $\alpha$ 2 remaining elevated at day 3 in the spleen, but no changes were seen in  
222 the gills after vaccination (**Fig. 2C-E**). The other pro-inflammatory cytokines investigated  
223 included IL-8 [39], and three IL-6 family members (IL-6, IL-11 and M17) [40-41]. The  
224 expression of all these genes was induced and peaked at day 1 in the spleen (**Fig. 2F-I**) after  
225 vaccination but in gills only IL-6 expression was induced to a small extent (3 fold) at day  
226 1(**Table S1**).

227 Several genes with regulatory roles, including two paralogues of IL-10 [42] TGF- $\beta$ 1 [43] and  
228 FoxP3 [44], and the novel fish IL-1 family member nIL-1Fm [45] were also investigated. IL-  
229 10A expression was up-regulated from days 1 and 3 and peaked at day 1 (301 fold) (**Fig. 2J**)  
230 in the spleen but was not modulated in the gills of vaccinated fish, however IL-10B  
231 expression was refractory in both tissues (**Table 2 and S2**). The expression of nIL-1Fm was  
232 also induced at day 1 in the spleen by vaccination (**Fig. 2K**). The expression of both TGF- $\beta$ 1  
233 paralogues was refractory in both spleen and gills after vaccination (**Table 2 and S1**).  
234 However, the expression of the two trout master transcription factors Fox3A and Fox3B,  
235 important for mammalian TGF- $\beta$ 1 expression in mammals, was decreased at day 1 in the gills  
236 and spleen respectively (**Fig. 2L, Table 2 and S1**).

237 Correlation analysis of vaccination-modulated gene expression in the spleen at day 1 and 3  
238 revealed that the expression of the major pro-inflammatory cytokines (IL-1 $\beta$ 1, IL-1 $\beta$ 2, TNF-  
239  $\alpha$  paralogues, IL-6, IL-8, IL-11 and M17) and anti-inflammatory cytokines (IL-10A and nIL-  
240 1Fm) was highly correlated (**Table 3**). Their expression was also correlated with that of APPs  
241 (SAA) and AMPs (CATH1, CATH2 and hepcidin) (Supplementary **Table. S2**).

242

### 243 **3.3 ERM-vaccination activates an early Th1 and Th17-type response but a later Th2-** 244 **type response in the spleen**

245 Activation of specific Th responses is important for vaccine-mediated immunity [46]. The  
246 expression of the key Th-response specifying cytokines has only focused on IFN- $\gamma$ 1 after  
247 ERM vaccination to date [15-16]. With the recent success of identification of the complete  
248 repertoire of these cytokines, including three IL-4/13 paralogues and six IL-17A/F paralogues  
249 [24-25] in rainbow trout, we examined, for the first time, the repertoire of Th response related  
250 cytokines after ERM vaccination. The Th1 specifying cytokines, IFN- $\gamma$ 1 and IFN- $\gamma$ 2 were  
251 induced at day 1 (8 fold) in the spleen of vaccinated fish (**Fig. 3A-B**). Two IFN- $\gamma$  inducing  
252 cytokines, IL-18 [47] and IL-21 [31] were also significantly induced to some extent at day 1  
253 in the spleen, but not in the gills, of the vaccinated fish (**Table 2 and S1**). Of the six potential  
254 Th17 cytokines in rainbow trout, the expression of IL-17A/F1A, 2A and 3 was upregulated at  
255 day 1 in spleen, with IL-17A/F2A remaining elevated at day 3, but no changes were apparent  
256 in the gills of vaccinated fish (**Fig. 3E-G**). However, IL-17A/F1B, 2B and IL-17N (or IL-  
257 17A/F4) were refractory (**Table 2 and S1**). In terms of inducibility, IL-17A/F1A (105 fold)  
258 and 17A/F3 (19 fold) were more responsive than IL-17A/F2A (7 fold). Three potential Th2  
259 cytokines, IL-4/13A, IL-4/13B1 and IL-4/13B2 are present in salmonids [25]. A small but  
260 significant induction of the expression of IL-4/13B1 (4 fold) and IL-4/13B2 (3 fold) was  
261 found at day 3 only in the spleen of vaccinated fish (**Fig. 3J-K**). IL-4/13A expression was  
262 refractory (**Fig. 3I**), as was the expression of the master transcription factors (T-bet, GATA3  
263 and ROR $\gamma$ ) [48-49] and the T cell cytokine IL-2 [50] (**Fig. 3C, D, H, and L**) in the spleen. Of  
264 these genes only ROR $\gamma$  at day 14 was modulated in the gills but again to a very small degree  
265 (**Table S2**).

266

267 The expression of Th1 cytokines (IFN- $\gamma$ 1 and 2) and the Th17 cytokines (IL-17A/F1A and  
268 IL-17A/F3) was highly correlated ( $R=0.72-0.90$ ,  $p<0.001$ ) but there was a lack of correlation  
269 with the other Th17 cytokine IL-17A/F2A, and the Th2 cytokine IL-4/13B1, the  
270 predominantly expressed IL-4/13B isoform (**Table 4**). The expression of IFN- $\gamma$ 1, IFN- $\gamma$ 2 and  
271 IL-17A/F1A was also correlated to IL-21, the majority of pro-inflammatory cytokines (IL-  
272 1 $\beta$ 1-2, TNF- $\alpha$ 1-3, IL-6, IL-8, IL-11 and M17), anti-inflammatory cytokines (IL-10A and nIL-  
273 1Fm), APPs and AMPs (**Table S2**).

274

### 275 **3.4 ERM-vaccination induces specific isoforms of the IL-12 family members in the** 276 **spleen**

277 IL-12 family cytokines play key roles in immunity bridging the innate and adaptive immune  
278 systems. Each cytokine consists of an  $\alpha$ -chain (p19, p28 and p35), and a  $\beta$ -chain (p40 and  
279 EBI3) in mammals. The orthologues of mammalian p35 and p40 are increased in teleosts  
280 fish, and salmonids in particular, due to the whole genome duplication events in these  
281 lineages [25]. Rainbow trout has three isoforms of p35 and p40, and two p28 isoforms, that  
282 potentially make additional IL-12 family cytokines [51]. The constitutive expression of  $\alpha$ -  
283 chains (p35A1, p35A2, p35B1, p19, p28A and p28B) was lower ( $\Delta C_p=16-22$ , **Table 1**) than  
284 that of the  $\beta$ -chains (P40A, P40B2, p40C and EBI3,  $\Delta C_p=9-17$ , **Table 1**). The expression of  
285 all the  $\alpha$ -chains was significantly increased in spleen and peaked at day 1 (13.6-fold for  
286 p35A1, 37.5-fold for p35A2, 58.7-fold for p35B1, 1502.0-fold for p19, 52.6-fold for p28A  
287 and 11.1-fold for p28B), with upregulated levels remaining at day 3 except for p35A1 and  
288 p28B, in the spleen (**Fig. 4A-E**). No modulation of these genes occurred in the gills and  
289 expression of the  $\beta$ -chains was refractory in both spleen and gills, with the exception of  
290 p40B2 that was up-regulated 3.4-fold at day 1 in the spleen (**Fig. 5F, Table 2 and S1**). These  
291 expression profiles suggest that specific isoforms of IL-12 and IL-23 could be increased in  
292 the spleen after ERM vaccination.

293

294 The upregulated expression of the  $\alpha$ -chains in the spleen was significantly correlated and also  
295 correlated to that of Th1 (IFN- $\gamma$ 1/2) and Th17 (IL-17A/F1A) markers (**Table 4**). The  
296 increased expression of most of the IL-12 family  $\alpha$ -chains was also correlated to the pro-and  
297 anti-inflammatory cytokines (IL-1 $\beta$ 1-2, TNF- $\alpha$ 1-3, IL-6, IL-8, IL-11, M17, IL-21, nIL-1Fm  
298 and IL-10A), AMPs (CATH1, CATH2 and hepcidin) and APPs (SAA and SAP1, **Table S2**).

299

### 300 **3.5 Other cytokine genes**

301 Other cytokine genes examined included IL-15 [52], two IL-17C paralogues [53], IL-20 [54],  
302 IL-22 [55] and IL-34 [56] and were refractory in both spleen and gills in response to ERM

303 vaccination (**Table 2 and S1**). IL-17D, which was lowly expressed in the spleen, was  
304 inhibited at day 3 in the spleen of vaccinated fish (**Table 2**).

305

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## 306 4. Discussion

307 Through the analysis of the expression of a large number of immune genes after ERM  
308 injection vaccination in rainbow trout, we observed a systematic activation of anti-microbial  
309 defences in both spleen and gills, but a specific activation of inflammatory cytokines and  
310 specific IL-12 family members that leads to a Th1/Th17 biased immune activation in the  
311 spleen.

312

### 313 4.1 Activation of pro- and anti-inflammatory cytokines in the spleen after ERM 314 vaccination

315 In this study, we comparatively examined immune gene expression in spleen and gills after  
316 ERM vaccination. The choice of spleen over head kidney was because transcriptome changes  
317 mainly happen transiently in spleen after bacterin vaccination in fish as shown by microarray  
318 analysis [57]. The gill tissue was chosen because it is a major lymphoid tissue in salmonids  
319 [58] and an important immune organ relevant to bacterial infections such as ERM [1, 2, 6].  
320 Vaccination with a bacterin mimics a bacterial infection thereby activating the innate immune  
321 system and initiating an adaptive immune response without the development of severe  
322 disease. Thus, a number of genes known to be activated by *Y. ruckeri* infection (**Table 5**)  
323 were found to be activated by ERM vaccination in this study. However, key differences of  
324 the host responses between ERM vaccination and *Y. ruckeri* infection exist. First, only a  
325 subset of genes are commonly activated by both ERM vaccination and infection. For  
326 example, the major pro-inflammatory cytokines IL-1 $\beta$ 1-2, TNF- $\alpha$ 1-3, IL-6, IL-8, IL-17A/F  
327 (1A, 2A and 3) and IFN- $\gamma$ 1, and anti-inflammatory cytokine IL-10A were activated by both  
328 vaccination and infection. In contrast, IL-2, IL-17C1-2 and TGF- $\beta$ 1A were up-regulated only  
329 by infection and were refractory to vaccination (**Table 5**). Secondly, the kinetics of gene  
330 activation was different. For example, the activation of gene expression by ERM vaccination  
331 was rapid and peaked at day 1 for most of the genes, whilst during infection the activation of  
332 inflammatory genes such as IL-1 $\beta$ , TNF $\alpha$ , and SAA occurs later [36]. Thirdly, the activation  
333 of pro- and anti-inflammatory cytokine expression happened mainly in the spleen by  
334 vaccination but also in the gills by infection (**Table 5**). Although direct comparisons of  
335 transcript levels between different experiments could be complicated by multiple factors,  
336 such as the age and life history of the fish, and the pathogen strains used, these comparisons

337 do suggest that different pathways have been activated by vaccination and pathogenic  
338 challenge.

339

340 The differences between ERM vaccination and pathogenic *Y. ruckeri* infection observed  
341 could be attributed to (1) dose effects, (2) factors released during infection [2,3] but absent in the  
342 vaccine preparation, (3) different stress responses due to the damage caused by pathogenic  
343 infection and vaccination, and (4) the evasion mechanisms of the pathogen. The ERM  
344 vaccination used a high dose, ie equivalent to  $5 \times 10^8$  cfu/fish, whilst infection studies typically  
345 use lower doses, eg  $5 \times 10^5$  cfu/fish [17, 45]. It is well known that the host immune response  
346 (eg cytokine transcript levels) is dose (pathogen load) dependent [36]. Thus, after ERM  
347 vaccination we observed an immediate activation of a set of APPs, AMPs, and pro- and anti-  
348 inflammatory cytokines in the spleen. However, after low(er) dose infection, the bacteria  
349 multiply and spread in different target tissues, with associated tissue damage likely  
350 responsible for the relatively late kinetics of gene expression and the broad activation of  
351 immune genes in multiple tissues (eg in both the spleen and gills). Some of the differences,  
352 however, might be due to immune evasion of the pathogen. For example, IL-4/13B1-2 were  
353 shown to be induced *in vitro* in HK cells by bacterin exposure and *in vivo* in the spleen after  
354 vaccination. In contrast, their expression was inhibited in gills by *Y. ruckeri* infection [25].

355

356 Despite the differences, the expression of the major pro-inflammatory cytokines (IL-1 $\beta$ 1, IL-  
357 1 $\beta$ 2, and TNF- $\alpha$  paralogues, IL-6, IL-8, IL-11 and M17) and anti-inflammatory cytokines  
358 (IL-10A and nIL-1Fm) was highly upregulated in the spleen (but not in the gills) immediately  
359 after ERM vaccination (**Figs. 2-3**). These changes are highly correlated (**Table 3**), suggesting  
360 an effective activation of the innate immune system was established with a balanced  
361 inflammatory response.

362

#### 363 **4.2 Activation of IL-12 family cytokines may lead to a Th1/Th17 biased adaptive** 364 **immune response**

365 Among a wide range of cytokines, the IL-12 family (IL-12, IL-23, IL-27 and IL-35) has  
366 unique structural, functional, and immunological characteristics that have made this family as

367 important immunological playmakers. Each IL-12 family member is composed of an  $\alpha$ -chain  
368 (p19, p28 and p35) and a  $\beta$ -chain (p40 and Ebi3). Whilst IL-12 (p35/p40), IL-23 (p19/p40)  
369 and IL-27 (p28/EBI3) are secreted by activated antigen presenting cells (APC) during antigen  
370 presentation to naïve T cells, IL-35 (p35/EBI3) is a product of regulatory T and B cells in  
371 mammals [26, 60-61]. IL-12 and IL-23 are proinflammatory or prostimulatory cytokines,  
372 whereas IL-27 and IL-35 are inhibitory cytokines. With our recent success in cloning and  
373 characterizing all of the subunits of the IL-12 family in salmonids [26-29], this is the first  
374 study to examine their involvement in a fish vaccination model. Although the constitutive  
375 expression of  $\alpha$ -chains (p35A1, p35A2, p35B1, p19, p28A and p28B) was low, their  
376 expression was significantly increased in the spleen of vaccinated fish. The expression of the  
377  $\beta$ -chain p40B2 was also up-regulated, but not p40B1, p40C or EBI3. This expression profiles  
378 suggest that specific isoforms of IL-12 and IL-23 containing p40B2 could be produced in the  
379 spleen after ERM vaccination.

380

381 CD4<sup>+</sup> T cells, also known as T-helper (Th) cells, play an important role in orchestrating  
382 adaptive immune responses to pathogens and vaccines. When naïve CD4<sup>+</sup> T cells recognize a  
383 foreign antigen-derived peptide presented in the context of major histocompatibility complex  
384 (MHC) class II on APCs, they undergo massive proliferation and differentiation into distinct  
385 Th cell subsets such as Th1, Th2, Th17 and induced T-regulatory (iTreg) cells in mammals  
386 [46]. Whilst IL-12 and IL-23 are critical for Th1 and Th17 cell development, respectively,  
387 IL-35 has immunosuppressive effects that are mediated through regulatory T and B cells. IL-  
388 27 displays both pro- and anti-inflammatory activities. It promotes the differentiation of Th1  
389 and IL-10-producing Tr1-like regulatory cells, but inhibits Th2 and Th17 [62]. Whilst the  
390 extent to which the mammalian Th cell paradigm is conserved is still unclear in fish [46], it  
391 provides a framework to investigate the immune responses to vaccination in fish [63-64]. The  
392 expression of Th1 cytokines (IFN- $\gamma$ 1 and 2), Th17 cytokines (IL-17A/F1A, 2A and 3), and  
393 Th2 cytokines (IL-4/13B1 and 2) was increased in the spleen after ERM vaccination. It is  
394 notable that the increased expression of IFN- $\gamma$ 1 and 2, and IL-17A/F1A was earlier and  
395 higher than Th2 cytokines (**Fig. 4**). Furthermore, their upregulated expression was  
396 significantly correlated to that of the  $\alpha$ -chains of the IL-12 family, and pro-/anti-inflammatory  
397 cytokines in the spleen of the vaccinated fish (**Table 4**). It is known that rainbow trout IL-12  
398 isoforms can induce IFN- $\gamma$  expression [26]. The correlated expression of IL-12 and IL-23  
399 with Th1 and Th17 cytokines and pro-anti-inflammatory cytokines may suggest that ERM  
400 vaccination activates a balanced inflammatory response with the expression of IL-12 and IL-



401 23 leading to a Th1 and Th17 biased immune response in the spleen. The ability of IL-23 to  
402 induce Th17 cytokine expression remains to be proven since no bioactivity analysis of this  
403 cytokine has been performed in any fish species.

404

405 In mammals the differentiation of Th cell subsets and expression of lineage specifying  
406 cytokines depend on the induction of lineage-specific master transcription factors, including  
407 T-bet for Th1, GATA3 for Th2, and ROR $\gamma$ t for Th17 [45, 65]. Although the expression of  
408 Th1, Th2 and Th17 cytokines was found increased in the spleen of ERM vaccinated fish, the  
409 expression of T-bet, GATA3 and ROR $\gamma$  was refractory. However, the lack of transcriptional  
410 factor expression change at a tissue level doesn't exclude their role in the regulation of  
411 cytokine gene expression in a specific cell type. Indeed, the master transcription factors can  
412 be expressed in other cell types in addition to Th cells, and their expression can coexist in the  
413 same cell and is dynamic and quantitative [65]. Thus at the mixed tissue level, changes of  
414 gene expression in a specific cell type(s) may be diluted by the presence of other abundant  
415 cell types or by the changes in other cell types. The possibility to isolate relevant fish  
416 leucocyte populations, such as CD4<sup>+</sup> or CD8<sup>+</sup> cells, and study their responses has become  
417 possible in fish recently using antibodies [66-68] or transgenic fish [69]. For example in trout  
418 infected with *Y. ruckeri* 4 days earlier, upregulation of Th1 (IL-2, IFN- $\gamma$ ) and Th17 cytokines  
419 (IL-17A/F1a, IL-21, IL-22) is apparent in splenic CD4-1<sup>+</sup>/CD4-2<sup>+</sup> cells and CD4-1<sup>-</sup>/CD4-2<sup>+</sup>  
420 cells [70]. Such studies will undoubtedly be directed towards elucidating the responses in  
421 vaccinated fish in the near future.

#### 422 **4.3 Systematic activation of anti-microbial defences by ERM vaccination**

423 APPs are an integral part of the acute phase response. They are secreted by the liver in  
424 response to a variety of injuries and can also be expressed in extrahepatic tissues. APPs  
425 favour the systemic regulation of defence, coagulation, proteolysis, and tissue repair [62].  
426 AMPs are integral components of innate immunity, and one of the first lines of host defence  
427 against bacterial infection. The up-regulation of APPs and AMPs in both spleen and gills  
428 suggests a systemic activation of innate immune system by ERM vaccination, that provide  
429 non-specific protection after vaccination.

430

431 It is known that many proinflammatory cytokines can induce the expression of APPs and  
432 AMPs in mammals and in fish. In rainbow trout, IL-6 induces hepcidin and CATH2 but not



433 CATH1 in the macrophage cell line RTS-11[32]. In contrast TNF- $\alpha$  induces hepcidin and  
434 CATH1 but not CATH2 in HK macrophages [22], whilst IL-1 $\beta$  can induce the expression of  
435 both CATH1 and CATH2 (un-published results). Adaptive cytokines such as IL-4/13 can  
436 also induce APPs (eg SAP1), and AMPs (eg hepcidin, and CATH1 but not CATH2) in HK  
437 cells [25]. The expression of a major APP (eg SAA) and AMPs (eg hepcidin and CATH2)  
438 was positively correlated with the expression of major pro-inflammatory cytokines (**Table**  
439 **S2**), suggesting that the activation and release of pro-inflammatory cytokines may lead to the  
440 induction of APPs and AMPs following vaccination. In agreement with this notion, IL-6, that  
441 specific induces CATH2 and hepcidin, was the only up-regulated proinflammatory cytokine  
442 in gills after ERM vaccination, and may account for the high levels of expression of these  
443 two AMPs (**Fig. 1**).

444

#### 445 **4.4 Differential expression of paralogous genes**

446 Many immune genes in teleost fish are known to have multiple paralogues, especially in  
447 salmonids that have undergone an additional whole genome duplication event. For example,  
448 there are three genes for IL-1 $\beta$  and TNF- $\alpha$  in salmonids [21-22] and three each of the p35 and  
449 p40 genes, that potentially could make 9 heterodimeric IL-12 isoforms with different  
450 functions [26-27]. Thus, it is necessary to determine whether all of the genes present react in  
451 a similar manner. In this study it is clear that major differences occur between different  
452 paralogues in response to vaccination. The biggest differences were seen when one of the  
453 paralogues was responsive and the other not, as with IL-1 $\beta$ 1 and 2 vs IL-1 $\beta$ 3, IL-4/13A vs  
454 IL-4/13B1 and 2, IL-10A vs IL-10B, IL-12 p40B2 vs p40B1 and p40C, IL-17A/F1A vs IL-  
455 17A/F1B, IL-17A/F2A vs IL-17A/F2B, and SAP1 vs SAP2. Differential responses of  
456 paralogues have been seen in responses to PAMPs [22,33], infection [24,26] and cytokine  
457 stimulation [25-26]. These differences likely reflect differences in the promoters, with some  
458 of the paralogues becoming more or less responsive to particular signalling pathways,  
459 perhaps in particular cell types, or genes that are being pseudogenised. The differential  
460 responses of IL-12 p40 paralogues is of particular interest. Two isoforms of rainbow trout  
461 rIL-12 have been made as recombinant proteins that differ in the p40 chain (ie p40B or p40C).  
462 These proteins can induce IFN- $\gamma$  expression in HK cells but only the isoform containing  
463 p40C was able to also induce IL-10 [26]. The induction of all  $\alpha$ -chains but only one  $\beta$ -chain  
464 (ie p40B2) suggests that specific IL-12 and IL-23 isoforms are produced after ERM

465 vaccination that may have different functions (as seen with IL-12 isoforms) critical for a  
466 Th1/Th17 biased response.

467

#### 468 **4.5 Implications for vaccine development in fish**

469 Using an efficacious bacterial model vaccine, this study has revealed that host innate  
470 immunity is activated by ERM vaccination as evidenced by the correlated upregulation of  
471 pro- and anti-inflammatory cytokines in the spleen and the systemic increase of APPs and  
472 AMPs. Specific IL-12 members are induced that may drive the Th1/Th17 biased immune  
473 responses observed. As an efficacious vaccine must activate innate immunity and initiate  
474 specific adaptive pathways, the early responses of the set of genes studied here may provide  
475 essential information and function as biomarkers in future vaccine development for fish,  
476 potentially allowing a screening method for vaccine candidates and formulations before more  
477 expensive mortality testing.

#### 478 **4.6 Conclusions**

479 In summary, ERM vaccination immediately activates a balanced inflammatory response with  
480 correlated expression of both pro- and anti-inflammatory cytokines in the spleen. The  
481 increase of pro-inflammatory cytokines may lead to the systemic upregulation of APPs and  
482 AMPs in both spleen and gills. We also observed an upregulation of all the IL-12 cytokine  
483 family  $\alpha$ -chains, but only one  $\beta$ -chain (p40B2) which suggests specific IL-12 and IL-23  
484 isoforms with distinct functions might be produced in the spleen of vaccinated fish. Notably  
485 the expression of Th1 cytokines (IFN- $\gamma$ 1-2) and a Th17 cytokine (IL-17A/F1A) were up-  
486 regulated and correlated to that of the IL-12 family  $\alpha$ -chains, the majority of pro- and anti-  
487 inflammatory cytokines, APPs and AMPs. These expression profiles may suggest that ERM  
488 vaccination activates host innate immunity and expression of specific IL-12 and IL-23  
489 isoforms leading to a Th1 and Th17 biased immune response. This study has increased our  
490 understanding of the host immune response to ERM vaccination and the adaptive pathways  
491 involved. The early responses seen may provide useful biomarkers for future vaccine  
492 development in aquaculture.

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## 500 6. References

- 501 [1] Tobback E, Decostere A, Hermans K, Haesebrouck F, Chiers K. *Yersinia ruckeri*  
502 infections in salmonid fish. J Fish Dis. 2007; 30: 257-268.
- 503 [2] Kumar G, Menanteau-Ledouble S, Saleh M, El-Matbouli M. *Yersinia ruckeri*, the  
504 causative agent of enteric redmouth disease in fish. Vet Res. 2015; 46:103.
- 505 [3] Furones MD, Rodgers CJ, Munn CB. *Yersinia ruckeri*, the causal agent of enteric  
506 redmouth disease (ERM) in fish. Annu Rev Fish Dis. 1993; 3: 105-125.
- 507 [4] Wheeler RW, Davies RL, Dalsgaard I, Garcia J, Welch TJ, Wagley S, Bateman KS,  
508 Verner-Jeffreys DW. *Yersinia ruckeri* biotype 2 isolates from mainland Europe and the  
509 UK likely represent different clonal groups. Dis Aquat Organ. 2009; 84: 25-33.
- 510 [5] Bastardo A, Ravelo C, Romalde JL. A polyphasic approach to study the intraspecific  
511 diversity of *Yersinia ruckeri* strains isolated from recent outbreaks in salmonid culture.  
512 Vet Microbiol. 2012; 160: 176-182.
- 513 [6] Tobback E, Decostere A, Hermans K, Ryckaert J, Duchateau L, Haesebrouck F, et al.  
514 Route of entry and tissue distribution of *Yersinia ruckeri* in experimentally infected  
515 rainbow trout *Oncorhynchus mykiss*. Dis Aquat Organ. 2009; 84: 219-228.
- 516 [7] Bravo S, Midtlyng PJ. The use of fish vaccines in the Chilean salmon industry 1999-  
517 2003. Aquaculture. 2007; 270: 36-42.
- 518 [8] Chettri JK, Deshmukh S, Holten-Andersen L, Jafaar RM, Dalsgaard I, Buchmann I.  
519 Comparative evaluation of administration methods for a vaccine protecting rainbow trout  
520 against *Yersinia ruckeri* O1 biotype 2 infections. Vet Immunol Immunopathol. 2013; 145:  
521 379-85.
- 522 [9] Deshmukh S, Kania PW, Chettri JK, Skov J, Bojesen AM, Dalsgaard I, Buchmann K.  
523 Insight from molecular, pathological, and immunohistochemical studies on cellular and  
524 humoral mechanisms responsible for vaccine-induced protection of rainbow trout against  
525 *Yersinia ruckeri*. Clin Vaccine Immunol. 2013; 20: 1623-41.
- 526 [10] Barnes AC, Delamare-Deboutteville J, Gudkovs N, Brosnahan C, Morrison R,  
527 Carson J. Whole genome analysis of *Yersinia ruckeri* isolated over 27 years in Australia  
528 and New Zealand reveals geographical endemism over multiple lineages and recent  
529 evolution under host selection. Microb Genom. 2016; 2: e000095.
- 530 [11] Ormsby MJ, Caws T, Burchmore R, Wallis T, Verner-Jeffreys DW, Davies RL.  
531 *Yersinia ruckeri* Isolates Recovered from Diseased Atlantic Salmon (*Salmo salar*) in

- 532 Scotland Are More Diverse than Those from Rainbow Trout (*Oncorhynchus mykiss*) and  
533 Represent Distinct Subpopulations. *Appl Environ Microbiol.* 2016; 82: 5785-5794.
- 534 [12] Midtlyng PJ. Chapter 6 Methods for measuring efficacy, safety and potency of fish  
535 vaccines. Adams A (Ed), *Fish vaccines*. Springer Basel, 2016, p119-142.
- 536 [13] Furman D, Davis MM. New approaches to understanding the immune response to  
537 vaccination and infection. *Vaccine.* 2015; 33: 5271-5281.
- 538 [14] Maertzdorf J, Kaufmann SH, Weiner J 3rd. Molecular signatures for vaccine  
539 development. *Vaccine.* 2015; 33: 5256
- 540
- 541
- 542 -61.
- 543 [15] Raida MK, Buchmann K. Temperature-dependent expression of immune-relevant  
544 genes in rainbow trout following *Yersinia ruckeri* vaccination. *Dis Aquat Organ.* 2007;  
545 77: 41-52.
- 546 [16] Raida MK, Buchmann K. Bath vaccination of rainbow trout (*Oncorhynchus mykiss*  
547 Walbaum) against *Yersinia ruckeri*: effects of temperature on protection and gene  
548 expression. *Vaccine.* 2008; 26: 1050-1062.
- 549 [17] Harun NO, Wang T, Secombes CJ. Gene expression profiling in naïve and  
550 vaccinated rainbow trout after *Yersinia ruckeri* infection: Insights into the mechanisms of  
551 protection seen in vaccinated fish. *Vaccine.* 2011; 29: 4388-4399.
- 552 [18] Bridle AR, Koop BF, Nowak BF. Identification of surrogates of protection against  
553 yersiniosis in immersion vaccinated Atlantic salmon. *PLoS One.* 2012; 7: e40841.
- 554 [19] Berthelot C, Brunet F, Chalopin D, Juanchich A, Bernard M, Noël B, Bento P, Da  
555 Silva C, Labadie K, Alberti A, Aury JM, Louis A, Dehais P, Bardou P, Montfort J,  
556 Klopp C, Cabau C, Gaspin C, Thorgaard GH, Boussaha M, Quillet E, Guyomard R,  
557 Galiana D, Bobe J, Volff JN, Genêt C, Wincker P, Jaillon O, Roest Crollius H, Guiguen  
558 Y. The rainbow trout genome provides novel insights into evolution after whole-genome  
559 duplication in vertebrates. *Nat Commun.* 2014; 5:3657.
- 560 [20] Lien S, Koop BF, Sandve SR, Miller JR, Kent MP, Nome T, Hvidsten TR, Leong JS,  
561 Minkley DR, Zimin A, Grammes F, Grove H, Gjuvsland A, Walenz B, Hermansen RA,  
562 von Schalburg K, Rondeau EB, Di Genova A, Samy JK, Olav Vik J, Vigeland MD, Caler  
563 L, Grimholt U, Jentoft S, Våge DI, de Jong P, Moen T, Baranski M, Palti Y, Smith DR,  
564 Yorke JA, Nederbragt AJ, Tooming-Klunderud A, Jakobsen KS, Jiang X, Fan D, Hu Y,  
565 Liberles DA, Vidal R, Iturra P, Jones SJ, Jonassen I, Maass A, Omholt SW, Davidson  
566 WS. The Atlantic salmon genome provides insights into rediploidization. *Nature.* 2016;  
567 533: 200-205.
- 568 [21] Husain M, Bird S, van Zwieten R, Secombes CJ, Wang T. Cloning of the IL-1 $\beta$ 3  
569 gene and IL-1 $\beta$ 4 pseudogene in salmonids uncovers a second type of IL-1 $\beta$  gene in  
570 teleost fish. *Dev Comp Immunol.* 2012; 38: 431-446.

- 571 [22] Hong S, Li R, Xu Q, Secombes CJ, Wang T. Two types of TNF- $\alpha$  exist in teleost  
572 fish: phylogeny, expression, and bioactivity analysis of type-II TNF- $\alpha$ 3 in rainbow trout  
573 *Oncorhynchus mykiss*. J Immunol. 2013; 191: 5959-5972.
- 574 [23] Monte MM, Wang T, Holland JW, Zou J, Secombes CJ. Cloning and  
575 characterization of rainbow trout interleukin-17A/F2 (IL-17A/F2) and IL-17 receptor A:  
576 expression during infection and bioactivity of recombinant IL-17A/F2. Infect Immun.  
577 2013; 81: 340-353.
- 578 [24] Wang T, Jiang Y, Wang A, Husain M, Xu Q, Secombes CJ. Identification of the  
579 salmonid IL-17A/F1a/b, IL-17A/F2b, IL-17A/F3 and IL-17N genes and analysis of their  
580 expression following in vitro stimulation and infection. Immunogenetics. 2015; 67: 395-  
581 412.
- 582 [25] Wang T, Johansson P, Abós B, Holt A, Tafalla C, Jiang Y, Wang A, Xu Q, Qi Z,  
583 Huang W, Costa MM, Diaz-Rosales P, Holland JW, Secombes CJ. First in-depth analysis  
584 of the novel Th2-type cytokines in salmonid fish reveals distinct patterns of expression  
585 and modulation but overlapping bioactivities. Oncotarget. 2016; 7: 10917-10946.
- 586 [26] Wang T, Husain M, Hong S, Holland JW. Differential expression, modulation and  
587 bioactivity of distinct fish IL-12 isoforms: Implication towards the evolution of Th1-like  
588 immune responses. Eur J Immunol. 2014; 44: 1541-1551.
- 589 [27] Wang T, Husain M. The expanding repertoire of the IL-12 cytokine family in teleost  
590 fish: Identification of three paralogues each of the p35 and p40 genes in salmonids, and  
591 comparative analysis of their expression and modulation in Atlantic salmon *Salmo salar*.  
592 Dev Comp Immunol. 2014; 46: 194-207.
- 593 [28] Jiang Y, Husain M, Qi Z, Bird S, Wang T. Identification and expression analysis of  
594 two interleukin-23 $\alpha$  (p19) isoforms, in rainbow trout *Oncorhynchus mykiss* and Atlantic  
595 salmon *Salmo salar*. Mol Immunol. 2015; 66: 216-228.
- 596 [29] Husain M, Martin SAM, Wang T. Identification and characterisation of the IL-27  
597 p28 subunits in fish: cloning and comparative expression analysis of two p28 paralogues  
598 in Atlantic salmon *Salmo salar*. Fish Shellfish Immunol. 2014; 41: 102-112.
- 599 [30] Benedicenti O, Wang T, Wangkahart E, Milne DJ, Holland JW, Collins C, Secombes  
600 CJ. Characterisation of arginase paralogues in salmonids and their modulation by immune  
601 stimulation/ infection. Fish Shellfish Immunol. 2017; 61:138-151.
- 602 [31] Wang T, Diaz-Rosales P, Costa MM, Campbell S, Snow M, Collet B, Martin SAM,  
603 Secombes CJ. Functional characterization of a nonmammalian IL-21: rainbow trout  
604 *Oncorhynchus mykiss* IL-21 upregulates the expression of the Th cell signature cytokines  
605 IFN- $\gamma$ , IL-10, and IL-22. J Immunol. 2011; 186: 708-721.
- 606 [32] Costa MM, Maehr T, Diaz-Rosales P, Secombes CJ, Wang T. Bioactivity studies of  
607 rainbow trout (*Oncorhynchus mykiss*) interleukin-6: effects on macrophage growth and  
608 antimicrobial peptide gene expression. Mol Immunol. 2011; 48: 1903-1916.
- 609 [33] Wangkahart E, Scott C, Secombes CJ, Wang T. Re-examination of the rainbow trout  
610 (*Oncorhynchus mykiss*) immune response to flagellin: *Yersinia ruckeri* flagellin is a  
611 potent activator of acute phase proteins, anti-microbial peptides and pro-inflammatory  
612 cytokines *in vitro*. Dev Comp Immunol. 2016; 57: 75-87.



- 613 [34] Douglas SE, Gallant JW, Liebscher RS, Dacanay A, Tsoi SC. Identification and  
614 expression analysis of hepcidin-like antimicrobial peptides in bony fish. *Dev Comp*  
615 *Immunol.* 2003; 27: 589-601.
- 616 [35] Chang CI, Zhang YA, Zou J, Nie P, Secombes CJ. Two cathelicidin genes are  
617 present in both rainbow trout (*Oncorhynchus mykiss*) and Atlantic salmon (*Salmo salar*).  
618 *Antimicrob Agents Chemother.* 2006; 50: 185-195.
- 619 [36] Wiens GD, Vallejo RL. Temporal and pathogen-load dependent changes in rainbow  
620 trout (*Oncorhynchus mykiss*) immune response traits following challenge with biotype 2  
621 *Yersinia ruckeri*. *Fish Shellfish Immunol.* 2010; 29: 639-647.
- 622 [37] Zhang YA, Zou J, Chang CI, Secombes CJ. Discovery and characterization of two  
623 types of liver-expressed antimicrobial peptide 2 (LEAP-2) genes in rainbow trout. *Vet*  
624 *Immunol Immunopathol.* 2004; 10: 259-269.
- 625 [38] Casadei E, Wang T, Zou J, González Vecino JL, Wadsworth S, Secombes CJ.  
626 Characterization of three novel beta-defensin antimicrobial peptides in rainbow trout  
627 (*Oncorhynchus mykiss*). *Mol Immunol.* 2009; 46: 3358-3366.
- 628 [39] Laing KJ, Zou JJ, Wang T, Bols N, Hirono I, Aoki T, Secombes CJ. Identification  
629 and analysis of an interleukin 8-like molecule in rainbow trout *Oncorhynchus mykiss*.  
630 *Dev Comp Immunol.* 2002; 26: 433-444.
- 631 [40] Wang T, Holland JW, Bols N, Secombes CJ. Cloning and expression of the first  
632 nonmammalian interleukin-11 gene in rainbow trout *Oncorhynchus mykiss*. *FEBS J.*  
633 2005; 272: 1136-1147.
- 634 [41] Wang T, Secombes CJ. Identification and expression analysis of two fish-specific  
635 IL-6 cytokine family members, the ciliary neurotrophic factor (CNTF)-like and M17  
636 genes, in rainbow trout *Oncorhynchus mykiss*. *Mol Immunol.* 2009; 46: 2290-2298.
- 637 [42] Harun NO, Costa MM, Secombes CJ, Wang T. Sequencing of a second interleukin-  
638 10 gene in rainbow trout *Oncorhynchus mykiss* and comparative investigation of the  
639 expression and modulation of the paralogues in vitro and in vivo. *Fish Shellfish Immunol.*  
640 2011; 31:107-117.
- 641 [43] Maehr T, Costa MM, Vecino JLG, Wadsworth S, Martin SAM, Wang T, Secombes  
642 CJ. Transforming growth factor- $\beta$ 1b: a second TGF- $\beta$ 1 paralogue in the rainbow trout  
643 (*Oncorhynchus mykiss*) that has a lower constitutive expression but is more responsive to  
644 immune stimulation. *Fish Shellfish Immunol.* 2013; 34: 420-432.
- 645 [44] Wang T, Monte MM, Huang W, Boudinot P, Martin SAM, Secombes CJ.  
646 Identification of two FoxP3 genes in rainbow trout (*Oncorhynchus mykiss*) with  
647 differential induction patterns. *Mol Immunol.* 2010; 47: 2563-2574.
- 648 [45] Wang T, Bird S, Koussounadis A, Holland JW, Carrington A, Zou J, Secombes CJ.  
649 Identification of a novel IL-1 cytokine family member in teleost fish. *J Immunol.* 2009;  
650 183: 962-974.
- 651 [46] Wang T, Secombes CJ. The cytokine networks of adaptive immunity in fish. *Fish*  
652 *Shellfish Immunol.* 2013; 35: 1703-1718.

- 653 [47] Zou J, Bird S, Truckle J, Bols N, Horne M, Secombes C. Identification and  
654 expression analysis of an IL-18 homologue and its alternatively spliced form in rainbow  
655 trout (*Oncorhynchus mykiss*). Eur J Biochem. 2004; 271: 1913-1923.
- 656 [48] Wang T, Holland JW, Martin SAM, Secombes CJ. Sequence and expression analysis  
657 of two T helper master transcription factors, T-bet and GATA3, in rainbow trout  
658 *Oncorhynchus mykiss* and analysis of their expression during bacterial and parasitic  
659 infection. Fish Shellfish Immunol. 2010; 29: 705-715.
- 660 [49] Monte MM, Wang T, Costa MM, Harun NO, Secombes CJ. Cloning and expression  
661 analysis of two ROR- $\gamma$  homologues (ROR- $\gamma$ 1 and ROR- $\gamma$ 2) in rainbow trout  
662 *Oncorhynchus mykiss*. Fish Shellfish Immunol. 2012; 33: 365-374.
- 663 [50] Díaz-Rosales P, Bird S, Wang T, Fujiki K, Davidson WS, Zou J, Secombes CJ.  
664 Rainbow trout interleukin-2: Cloning, expression and bioactivity analysis. Fish Shellfish  
665 Immunol. 2009; 27:414-422.
- 666 [51] Secombes CJ, Wang T, Bird S. Chapter 5 Vertebrate cytokines and their evolution.  
667 D. Malagoli (Ed.), The Evolution of the Immune System: a Balance between  
668 Conservation and Diversity, Elsevier Press (2016), p87-150
- 669 [52] Wang T, Holland JW, Carrington A, Zou J, Secombes CJ. Molecular and functional  
670 characterization of IL-15 in rainbow trout *Oncorhynchus mykiss*: a potent inducer of IFN-  
671  $\gamma$  expression in spleen leukocytes. J Immunol. 2007; 179: 1475-1488.
- 672 [53] Wang T, Martin SA, Secombes CJ. Two interleukin-17C-like genes exist in rainbow  
673 trout *Oncorhynchus mykiss* that are differentially expressed and modulated. Dev Comp  
674 Immunol. 2010; 34: 491-500
- 675 [54] Wang T, Díaz-Rosales P, Martin SA, Secombes CJ. Cloning of a novel interleukin  
676 (IL)-20-like gene in rainbow trout *Oncorhynchus mykiss* gives an insight into the  
677 evolution of the IL-10 family. Dev Comp Immunol. 2010; 34: 158-67.
- 678 [55] Monte MM, Zou J, Wang T, Carrington A, Secombes CJ. Cloning, expression  
679 analysis and bioactivity studies of rainbow trout (*Oncorhynchus mykiss*) interleukin-22.  
680 Cytokine 2011; 55: 62-73.
- 681 [56] Wang T, Kono T, Monte MM, Kuse H, Costa MM, Korenaga H, Maehr T, Husain  
682 M, Sakai M, Secombes CJ. Identification of IL-34 in teleost fish: differential expression  
683 of rainbow trout IL-34, MCSF1 and MCSF2, ligands of the MCSF receptor. Mol  
684 Immunol. 2013; 53: 398-409.
- 685 [57] Jiang J, Miyata M, Chan C, Ngoh SY, Liew WC, Saju JM, Ng KS, Wong FS, Lee  
686 YS, Chang SF, Orbán L. Differential transcriptomic response in the spleen and head  
687 kidney following vaccination and infection of Asian seabass with *Streptococcus iniae*.  
688 PLoS One. 2014; 9: e99128.
- 689 [58] Haugarvoll E, Bjerås I, Nowak BF, Hordvik I, Koppang EO. Identification and  
690 characterization of a novel intraepithelial lymphoid tissue in the gills of Atlantic salmon. J  
691 Anat. 2008; 213: 202-209.

- 692 [59] Raida MK, Buchmann K. Development of adaptive immunity in rainbow trout,  
693 *Oncorhynchus mykiss* (Walbaum) surviving an infection with *Yersinia ruckeri*. Fish  
694 Shellfish Immunol. 2008; 25: 533-541.
- 695 [60] Sun L, He C, Nair L, Yeung J, Egwuagu CE. Interleukin 12 (IL-12) family  
696 cytokines: Role in immune pathogenesis and treatment of CNS autoimmune disease.  
697 Cytokine. 2015; 75: 249-255.
- 698 [61] Vignali DA, Kuchroo VK. IL-12 family cytokines: immunological playmakers. Nat.  
699 Immunol 2012;13:722-728.
- 700 [62] Schrödl W, Büchler R, Wendler S, Reinhold P, Muckova P, Reindl J, Rhode H.  
701 Acute phase proteins as promising biomarkers: Perspectives and limitations for human  
702 and veterinary medicine. Proteomics Clin Appl. 2016; 10: 1077-1092.
- 703 [63] Zhang H, Shen B, Wu H, Gao L, Liu Q, Wang Q, Xiao J, Zhang Y. Th17-like  
704 immune response in fish mucosal tissues after administration of live attenuated *Vibrio*  
705 *anguillarum* via different vaccination routes. Fish Shellfish Immunol. 2014; 37: 229-38.
- 706 [64] Zhang H, Fei C, Wu H, Yang M, Liu Q, Wang Q, Zhang Y. Transcriptome profiling  
707 reveals Th17-like immune responses induced in zebrafish bath-vaccinated with a live  
708 attenuated *Vibrio anguillarum*. PLoS One. 2013; 8: e73871.
- 709 [65] Fang D, Zhu J. Dynamic balance between master transcription factors determines the  
710 fates and functions of CD4 T cell and innate lymphoid cell subsets. J Exp Med. 2017;  
711 214: 1861-1876.
- 712 [66] Toda H, Saito Y, Koike T, Takizawa F, Araki K, Yabu T, Somamoto T, Suetake H,  
713 Suzuki Y, Ototake M, Moritomo T, Nakanishi T. (2011). Conservation of characteristics  
714 and functions of CD4 positive lymphocytes in a teleost fish. Dev. Comp. Immunol. 35:  
715 650-660.
- 716 [67] Castro R, Takizawa F, Chacara W, Lunazzi A, Dang TH, Koellner B, Quillet E, Six A,  
717 Fischer U, Boudinot P. (2013). Contrasted TCR $\beta$  diversity of CD8+ and CD8- T cells in  
718 rainbow trout. PLoS One 8: e60175.
- 719 [68] Kono T, Korenaga H. 2013. Cytokine gene expression in CD4 positive cells of the  
720 Japanese pufferfish, *Takifugu rubripes*. PLoS ONE 8: e66364.
- 721 [69] Dee CT, Nagaraju TR, Athanasiadis EI, Gray C, Fernandez del Ama L, Johnston SA,  
722 Secombes CJ, Cvejic A, Hurlstone AFL. (2016). CD4-Transgenic zebrafish reveals tissue-  
723 resident Th2- and regulatory T cell-like populations and diverse mononuclear phagocytes.  
724 J. Immunol. 197: 3520-3530.  
725
- 726 [70] Takizawa F, Magadan S, Parra D, Xu Z, Korytar T, Boudinot P, Sunyer JO (2016).  
727 Novel teleost CD4-bearing cell populations provide insights into the evolutionary origins  
728 and primordial roles of CD4+ lymphocytes and CD4+ macrophages. J. Immunol.  
729 196:4522-4535.  
730
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733 **Figure legend**

734 **Fig. 1 Modulation of the expression of APP and AMP genes in the spleen and gills by**  
735 **ERM vaccination.** Two groups of rainbow trout were vaccinated by ip injection with  
736 AquaVac ERM (red bars) or PBS as control (blue bars). The fish were killed at days 1, 3, 7  
737 and 14, and spleen and gills collected for gene expression analysis by real-time RT-PCR (as  
738 described in the Materials and Methods). Modulated expression was expressed as a fold  
739 change calculated as the mean expression levels in vaccinated fish normalized to that of time-  
740 matched controls in the same tissue. The means + SEM of six fish are shown. The relative  
741 significance of a Bonferroni post hoc test after a significant one way-ANOVA between the  
742 vaccinated and control groups at the same time point is shown above/within the bars as:  
743 \* $p < 0.05$ , and \*\*\* $p < 0.001$ .

744 **Fig. 2 Modulation of the expression of pro- and anti-inflammatory cytokines and**  
745 **Foxp3b in spleen by ERM vaccination.** Two groups of rainbow trout were vaccinated by ip  
746 injection with AquaVac ERM (red bars) or PBS as control (blue bars). The fish were killed at  
747 day 1, 3, 7 and 14, and spleen and gills collected for gene expression analysis by real-time  
748 PCR (as described in the Materials and Methods). Modulated expression was expressed as a  
749 fold change calculated as the mean expression levels in vaccinated fish normalized to that of  
750 time-matched controls in the same tissue. The means + SEM of six fish are shown. The  
751 relative significance of a Bonferroni post hoc test after a significant one way-ANOVA  
752 between the vaccinated and control groups at the same time point is shown above/within the  
753 bars as: \* $p < 0.05$ , and \*\*\* $p < 0.001$ .

754 **Fig. 3 Modulation of the expression of genes associated with T helper cells in spleen by**  
755 **ERM vaccination.** Two groups of rainbow trout were vaccinated by ip injection with  
756 AquaVac ERM (red bars) or PBS as control (blue bars). The fish were killed at day 1, 3, 7  
757 and 14, and spleen and gills collected for gene expression analysis by real-time PCR (as  
758 described in the Materials and Methods). Modulated expression was expressed as a fold  
759 change calculated as the mean expression levels in vaccinated fish normalized to that of time-  
760 matched controls in the same tissue. The means + SEM of six fish are shown. The relative  
761 significance of a Bonferroni post hoc test after a significant one way-ANOVA between the  
762 vaccinated and control groups at the same time point is shown above/within the bars as:  
763 \* $p < 0.05$ , \*\* $p < 0.01$  and \*\*\* $p < 0.001$ .

764 **Fig. 4 Modulation of the expression of subunits of IL-12 family in spleen by ERM**  
765 **vaccination.** Two groups of rainbow trout were vaccinated by ip injection with AquaVac  
766 ERM (red bars) or PBS as control (blue bars). The fish were killed at day 1, 3, 7 and 14, and  
767 spleen and gills collected for gene expression analysis by real-time PCR (as described in the  
768 Materials and Methods). Modulated expression was expressed as a fold change calculated as  
769 the mean expression levels in vaccinated fish normalized to that of time-matched controls in  
770 the same tissue. The means + SEM of six fish are shown. The relative significance of a  
771 Bonferroni post hoc test after a significant one way-ANOVA between the vaccinated and  
772 control groups at the same time point is shown above/within the bars as: \* $p < 0.05$ , and  
773 \*\*\* $p < 0.001$ .

774

**Table 1 Primers used for expression analysis by real-time PCR**

Gene	$\Delta$ cp*(Spleen)	$\Delta$ cp*(Gills)	Forward (5' to 3')	Reverse (5' to 3')	Acc. No.
<b>House-keeping gene</b>					
EF-1 $\alpha$	0**	0**	CAAGGATATCCGTCGTGGCA	ACAGCGAAACGACCAAGAGG	AF498320
<b>Acute phase proteins and antimicrobial peptides</b>					
SAA	8.16	8.34	GGTGAAGCTGCTCAAGGTGCTAAAG	GCCATTACTGATGACTGTTGCTGC	AM422447
SAP1	20.43	21.17	GCTGTATGGTGACCTTCAAGATCTCTC	GCGTTTGACAACAACAAATCATTGTGTC	X99385
SAP2	15.08	13.91	GGTTGTTATGCTGAACATCAAGATCTCTC	CCACCTTTGATTGCATACACAGATT	EZ763346
CATH1	9.39	9.14	ACCAGCTCCAAGTCAAGACTTTGAA	TGTCCGAATCTTCTGCTGCAA	AY594646
CATH2	11.77	11.00	ACATGGAGCGAGAAGTTCAGAAGA	GAGCCAAACCCAGGACGAGA	AY542963
Hepcidin	11.73	16.19	GCTGTTCTTTCTCCGAGGTGC	GTGACAGCAGTTGCAGCACCA	CA369786
LEAP2A	12.19	12.91	GGTTCCTGGTGTCTGCTGCT	AGTGGCCACCCTGCAAAT	AY362186
$\beta$ -defensin-1	17.99	19.57	CTGGTTTTCTATTGCTTAATGTTGTGG	GAAATGAGAAACACAGCACAAGAATCC	AM282655
$\beta$ -defensin-2	17.77	20.22	ATGGGAGACTGGGTTTGGT	ACGCAAAGCACAGCATTTAATCT	FM212656
$\beta$ -defensin-3	17.96	24.42	GGCTCTTTAGTCATTGCTTGTGGAATAC	CAGCATACATTGGCCATGTACA	FM212657
$\beta$ -defensin-4	20.58	18.13	TGGTGCTCTCGCTTCTTGG	TGGGCGACACAGCATACAATC	FM212658
<b>Cytokines</b>					
IL-1 $\beta$ 1	10.26	11.51	CCTGGAGCATCATGGCGTG	GCTGGAGAGTGTGTGGAAGACATATAG	AJ278242
IL-1 $\beta$ 2	20.89	14.09	GAGCGCAGTGAAGTGTGG	AGACAGGTTCAAATGCACCTTTATGGT	AJ245925
IL-1 $\beta$ 3	11.18	11.77	CTG AAG GCC GTC ACA ATC CA	CTGGTCCCTTACAGCGCTCCAA	AM181685
niL-1Fm	10.04	9.41	CCCATTCTCGTGACACCAG	CTGGACGACCTGGAGAGTGACT	AJ555869
IL-2	13.45	14.43	TGATGTAGAGGATAGTTGCATTGTTGC	GAAGTGTCCGTTGTGCTGTTCTC	AM422779
IL-4/13A	13.96	12.12	ACCACCACAAAGTCAAGGAGTTCT	CACCTGGTCTTGGCTCTTCAAC	FN820501
IL-4/13B1	17.21	13.28	GAGATTCATCTACTGCAGAGGATCATGA	GCAGTTGGAAGGGTGAAGCTTATTGTA	HG794522
IL-4/13B2	18.55	15.87	GAGACTCATCTATTGCGTATGATCATCG	TGCAGTTGGTTGGATGAACTTATTGTA	HG794523
IL-6	14.45	18.56	GGGAGAAAATGATCAAGATGCTCGT	GCAGACATGCCTCCTTGTGG	DQ866150
IL-8	9.58	8.72	AGAGACACTGAGATCATTGCCAC	CCCTCTTCATTGTTGTTGGC	AJ310565
IL-10A	16.55	13.68	GGATTCTACACCCTTGAAGAGCCC	GTCGTGTTGTTCTGTGTTCTGTTGT	AB118099
IL-10B	16.90	15.50	GGGATCTAGACCACATCAAGAGTCC	GATGGGAGATTTAAAGTTGTGTGTTCC	FR691804
IL-11	10.86	14.38	CTCTCGCTGCTATTGGCCA	TCTCGAATGCATGTTCCCAATAGAT	AJ535687
M17	12.14	10.58	GTGGACCTCTTAAAAACATACAAGCTCAG	GGATGGTGGCTGAAGTCTGTCTG	FM866399
IL-12 p35A1	16.62	17.51	GGAACACCACATTCAGTGAGAGTGC	CGTCTGCAACTGTGAGGAAGGAT	HE798148
IL-12 p35A2	20.73	19.77	GGAACACCACATTCAGTGAGAGTGA	CAACTGTGAGGAAGACACCCA	HG917950
IL-12 p35B1	21.49	18.58	TGCCAAACGCCAAGCTTTATTTTG	GCTGTTGAGTGCCTTTGGCTTTTGG	HG917951
IL-12 p40B1	17.14	9.77	CCCTTCTACATCCGAGAAATAGTGAAC	GTTGGTTTCACTTATAAACACCTTTTCCCTT	HE798149
IL-12 p40B2	14.14	9.46	CCGTCTACATACGAGAAATAGTGGAGA	TCAGAGTACAGCTTTCCCTGG	HG917952
IL-12 p40C	12.14	12.13	TTAAAGACAACGGAAAGGAGGAGC	CCTCCCGTAACCACATTTTCC	AJ548830
IL-23 p19	22.01	16.44	ACCTAAGAGCAGATTCAATGCCTTG	TCTTCCAGCTCTTCACTTCTCTG	KP410548
IL-27 p28A	19.88	18.47	GCAGCTGCTCAGGAGATATAAGGAGG	TCTCTCAGGTATGCTGGGTTTGG	HG794528
IL-27 p28B	22.39	22.06	GCAGCTGCTCAGGAGATATAAGGAGGA	GCTGCTCTGTGTTCCACCTTATCCAC	HG794529
EBI3	17.27	17.42	ACATCGCCACCTACAGTATGAAAGG	GGGTCCGGCTTCACAATGT	AJ620467
IL-15	7.95	7.69	TGGAATTGCTTCATAATATTGAGCTGCC	TGGTAGTTATCTGTGACCGACATGTCCTC	AJ628345
IL-17A/F1A	24.36	13.42	CAAACGTACACTTTTTGATGGTGCTG	GGGACTCATCATAGGTGGTGTGGT	KJ921977
IL-17A/F1B	18.92	20.46	CTCCAGTCTTTGACGGTGCTG	GGTTGTAATAGGCTGTGGAATGGAA	KJ921978
IL-17A/F2A	20.94	11.97	CACCTGGACCTGAAAAGCAC	GGCCACAGACAGGAAGGAGG	AJ580842
IL-17A/F2B	22.62	18.05	CCCTGGACCTGAAAACCCAT	GGCCACGGACAGGAAGGTA	KJ921979
IL-17A/F3	18.89	11.41	CTGGTGCTGGGTCTGATCATGT	GGTCTCATCGTATGTGTCGCTGATG	KJ921980
IL-17A/F4/N	25.20	18.46	AGAACTAACATGCAACAGCTCCA	CGGTTCAAGTCAATTTTTCCACGTA	KJ921981
IL-17C1	22.60	15.21	CTGGCGGTACAGCATCGATA	GAGTTATATCCATAATCTTCTGATCCGGC	FM955455
IL-17C2	16.84	16.70	CTGGCGGTACAGCATCGATA	CAGAGTTATATGCATGATGTTGGC	FM955456
IL-17D	19.30	11.10	GAAGAAATCCTCGAGCAGATGTTTG	GGGTGCTGGGAGATCCTGTATG	AJ580843
IL-18	9.02	10.01	GAGCAATGCAAAGCAGATGATTG	CATGTTTTGAGCAGCAATGTAGTC	AJ556990
IL-20	>25	19.37	CAAGAACCTGAGGCAATGCTACTG	TCTCTATAGCTTTTACTGCTGCCG	FN386780
IL-21	16.45	15.00	AAAGTTATCAAAAACCTCAACACCGAA	CCAGTCTACTGATGGCCTTTTGAAG	FM883702
IL-22	18.05	12.20	GAAGGAACACGGCTGTGCTATTAAC	GATCTAGGCGTGCACAGAAAGTC	AM748538
IL-34	9.97	9.31	AGGCAGAAACGTAACATGAAACACA	TCCACCCTCGCCCTCAGCTT	FN820429
IFN $\gamma$ 1	12.41	12.16	CAAACCTGAAAGTCCACTATAAGATCTCCA	TCCTGAATTTTCCCCTTGACATATTT	AJ616215
IFN $\gamma$ 2	14.43	14.00	CAAACCTGAAAGTCCACTATAAGATCTCCA	GGTCCAGCCTCTCCCTCAC	FM864345
TGF- $\beta$ 1A	8.13	9.90	CTCACATTTTACTGATGTCACCTTCTGT	GGACAACCTGCTCCACCTTGTG	OMY7836
TGF- $\beta$ 1B	7.47	8.72	CATGTCCATCCCCAGAAT	GGACAACCTGTTCCACCTTGTGT	FN822750
	13.09	13.29	TGTGTGGGTCCTCTTAATAGCAGGTC	CCTCAATTTTCACTGCTGATGTTGA	AJ277604
TNF- $\alpha$ 2	16.03	14.35	CTGTGTGGGCTTCTTAAATAGCAGCTT	CATTCCGTCCTGCATCGTTGC	AJ401377
TNF- $\alpha$ 3	14.94	16.51	GCTGCACCTTCTTTACCAAGAAACAAG	CCACTGAGGACTTGAATCACCATAGGT	HE798544
<b>Master transcription factors</b>					
T-bet	11.09	14.89	GGTAACATGCCAGGGAACAGGA	TGGTCTATTTTTAGCTGGGTGATGCTG	FM863825
GATA3	11.12	5.29	CCAAAAACAAGTCTATTGTCAGAAGG	TGGTGAGAGGTCGGTTGATATTGTG	FM863826
ROR $\gamma$ ***	9.04	8.61	ACAGACCTTCAAAGCTCTTGGTTGTG	GGGAAGCTTGGACACCATCTTTG	FM883712
Foxp3A	9.78	10.42	CCCAGAACCAGGTTGGAGTGT	TGACGGACAGCTTCTTCCA	FM883710
Foxp3B	9.09	9.99	TCTTGCCCACTACTCATCCC	TGACGGACAGCTTCTTCCA	FM883711

**Notes**

\* $\Delta$ cp is the average cp value of the target gene minus that of the house-keeping gene EF-1 $\alpha$  in the control fish at day 1 post vaccination. A higher  $\Delta$ cp value indicates a lower expression level.

\*\*The average cp of EF-1 $\alpha$  is 11.31+0.25 (spleen) and 10.44+0.11 (gills).

\*\*\*Primers amplify both ROR $\gamma$ a and ROR $\gamma$ b genes.

**Table 2** Fold change of transcript expression of studied genes in spleen after ERM vaccination not shown in Figs 1-

4. Asterisks indicate significant differences between vaccinated fish and controls as \*\* $p < 0.01$  and \*\*\* $p < 0.001$  (One-way-ANOVA with Bonferroni correction).

Gene	D1	D3	D7	D14
LEAP2A	0.58	0.53	0.81	0.22
$\beta$ -defensin-1	0.14	0.27	0.27	0.21
$\beta$ -defensin-2	2.57	0.35	0.17	1.07
$\beta$ -defensin-3	1.25	0.43	0.46	0.41
$\beta$ -defensin-4	1.68	0.75	0.73	0.48
IL-1 $\beta$ 3	1.11	1.22	0.95	0.92
IL-10B	1.33	0.67	0.59	0.62
IL-12 p40B1	1.21	0.84	0.66	1.41
IL-12 p40C	1.08	0.77	1.34	1.56
IL-27 p28B	11.06***	2.16	0.44	0.99
EBI3	0.75	0.48	1.45	1.38
IL-15	1.65	0.99	0.72	0.86
IL-17A/F1b	5.90	4.44	3.09	0.58
IL-17A/F2b	1.41	3.00	0.38	1.78
IL-17A/F4/N	15.95	0.97	3.15	4.52
IL-17C1	2.93	2.00	1.21	0.68
IL-17C2	1.48	2.00	0.82	1.52
IL-17D	0.50	0.29**	0.48	0.32
IL-18	1.72***	1.10	0.70	0.82
IL-20	1.66	0.24	0.04	0.37
IL-21	2.51**	0.61	1.18	1.49
IL-22	1.97	2.24	0.84	1.02
IL-34	1.59	0.52	0.69	1.04
TGF- $\beta$ 1A	1.42	1.07	0.83	0.81
TGF- $\beta$ 1B	2.43	0.74	0.58	0.48
Foxp3A	0.43***	0.74	1.26	0.76

Table 3. The Spearman's rho correlation coefficient (R) and the 2-tailed significance (p) between gene expression levels of the pro and anti-inflammatory cytokines in the spleen of vaccinated fish at day 1 and day 3 post-injection. R in bold suggests a significant Spearman rank ordered correlation.

		IL-1 $\beta$ 1	IL-1 $\beta$ 2	TNF- $\alpha$ 1	TNF- $\alpha$ 2	TNF- $\alpha$ 3	IL-8	IL-6	IL-11	M17	IL-18	IL-21	nIL-1Fm	IL-10A
IL-1 $\beta$ 1	R	1.000	<b>0.925</b>	<b>0.846</b>	<b>0.907</b>	<b>0.949</b>	<b>0.954</b>	<b>0.818</b>	<b>0.939</b>	<b>0.837</b>	0.415	<b>0.724</b>	<b>0.905</b>	<b>0.691</b>
	p		.000	.001	.000	.000	.000	.001	.000	.001	.180	.008	.000	.013
IL-1 $\beta$ 2	R	<b>0.925</b>	1.000	<b>0.774</b>	<b>0.811</b>	<b>0.888</b>	<b>0.872</b>	<b>0.799</b>	<b>0.806</b>	<b>0.768</b>	0.439	0.544	<b>0.774</b>	<b>0.600</b>
	p	.000		.003	.001	.000	.000	.002	.002	.004	.154	.068	.003	.039
TNF- $\alpha$ 1	R	<b>0.846</b>	<b>0.774</b>	1.000	<b>0.870</b>	<b>0.809</b>	<b>0.847</b>	<b>0.882</b>	<b>0.821</b>	<b>0.747</b>	<b>0.668</b>	<b>0.789</b>	<b>0.902</b>	0.499
	p	.001	.003		.000	.001	.001	.000	.001	.005	.018	.002	.000	.099
TNF- $\alpha$ 2	R	<b>0.907</b>	<b>0.811</b>	<b>0.870</b>	1.000	<b>0.895</b>	<b>0.912</b>	<b>0.891</b>	<b>0.947</b>	<b>0.863</b>	<b>0.627</b>	<b>0.704</b>	<b>0.924</b>	<b>0.613</b>
	p	.000	.001	.000		.000	.000	.000	.000	.000	.029	.011	.000	.034
TNF- $\alpha$ 3	R	<b>0.949</b>	<b>0.888</b>	<b>0.809</b>	<b>0.895</b>	1.000	<b>0.907</b>	<b>0.855</b>	<b>0.904</b>	<b>0.853</b>	0.519	<b>0.723</b>	<b>0.865</b>	<b>0.656</b>
	p	.000	.000	.001	.000		.000	.000	.000	.000	.084	.008	.000	.020
IL-8	R	<b>0.954</b>	<b>0.872</b>	<b>0.847</b>	<b>0.912</b>	<b>0.907</b>	1.000	<b>0.823</b>	<b>0.94</b>	<b>0.896</b>	0.508	<b>0.780</b>	<b>0.949</b>	<b>0.779</b>
	p	.000	.000	.001	.000	.000		.001	.000	.000	.092	.003	.000	.003
IL-6	R	<b>0.818</b>	<b>0.799</b>	<b>0.882</b>	<b>0.891</b>	<b>0.855</b>	<b>0.823</b>	1.000	<b>0.828</b>	<b>0.787</b>	<b>0.724</b>	<b>0.747</b>	<b>0.830</b>	<b>0.599</b>
	p	.001	.002	.000	.000	.000	.001		.001	.002	.008	.005	.001	.039
IL-11	R	<b>0.939</b>	<b>0.806</b>	<b>0.821</b>	<b>0.947</b>	<b>0.904</b>	<b>0.940</b>	<b>0.828</b>	1.000	<b>.903**</b>	0.485	<b>0.743</b>	<b>0.961</b>	<b>0.745</b>
	p	.000	.002	.001	.000	.000	.000	.001		.000	.110	.006	.000	.005
M17	R	<b>0.837</b>	<b>0.768</b>	<b>0.747</b>	<b>0.863</b>	<b>0.853</b>	<b>0.896</b>	<b>0.787</b>	<b>0.903</b>	1.000	0.465	<b>0.669</b>	<b>0.907</b>	<b>0.861</b>
	p	.001	.004	.005	.000	.000	.000	.002	.000		.128	.017	.000	.000
IL-18	R	0.415	0.439	<b>0.668</b>	<b>0.627</b>	0.519	0.508	<b>0.724</b>	0.485	0.465	1.000	0.428	<b>0.589</b>	0.165
	p	0.180	.154	.018	.029	.084	.092	.008	.110	.128		.165	.044	.607
IL-21	R	<b>0.724</b>	0.544	<b>0.789</b>	<b>0.704</b>	<b>0.723</b>	<b>0.780</b>	<b>0.747</b>	<b>0.743</b>	<b>0.669</b>	0.428	1.000	<b>0.782</b>	<b>0.613</b>
	p	.008	.068	.002	.011	.008	.003	.005	.006	.017	.165		.003	.034
nIL-1Fm	R	<b>0.905</b>	<b>0.774</b>	<b>0.902</b>	<b>0.924</b>	<b>0.865</b>	<b>0.949</b>	<b>0.830</b>	<b>0.961</b>	<b>0.907</b>	<b>0.589</b>	<b>0.782</b>	1.000	<b>0.731</b>
	p	.000	.003	.000	.000	.000	.000	.001	.000	.000	.044	.003		.007
IL-10A	R	<b>0.691</b>	<b>0.600</b>	0.499	<b>0.613</b>	<b>0.656</b>	<b>0.779</b>	<b>0.599</b>	<b>0.745</b>	<b>0.861</b>	0.165	<b>0.613</b>	<b>0.731</b>	1.000
	p	.013	.039	.099	.034	.020	.003	.039	.005	.000	.607	.034	.007	

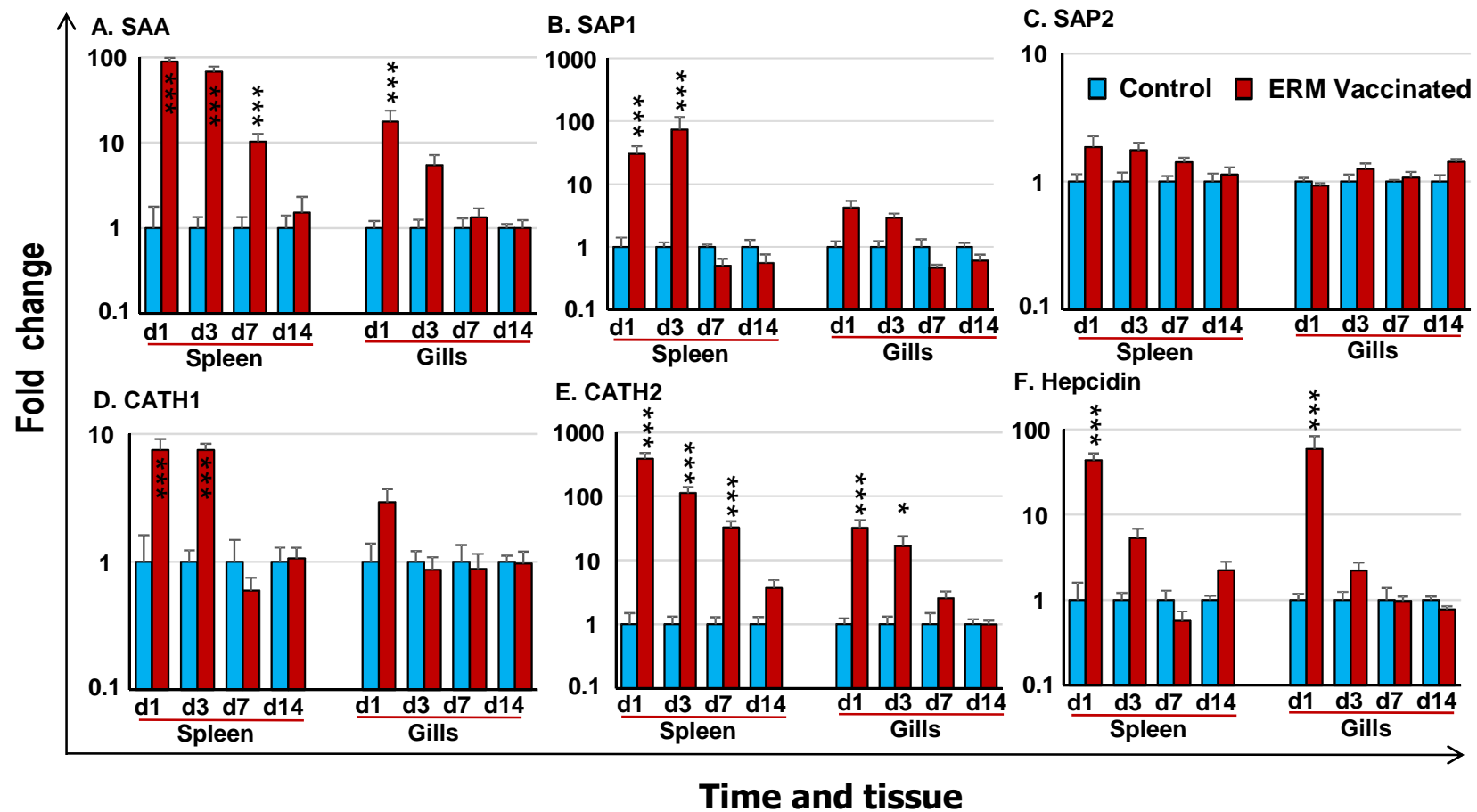
Table 4. The Spearman's rho correlation coefficient (R) and the 2-tailed significance (p) between gene expression levels of the cytokines involved in Th cell development in the spleen of vaccinated fish at day 1 and day 3 post-injection. R in bold suggests a significant Spearman rank ordered correlation.

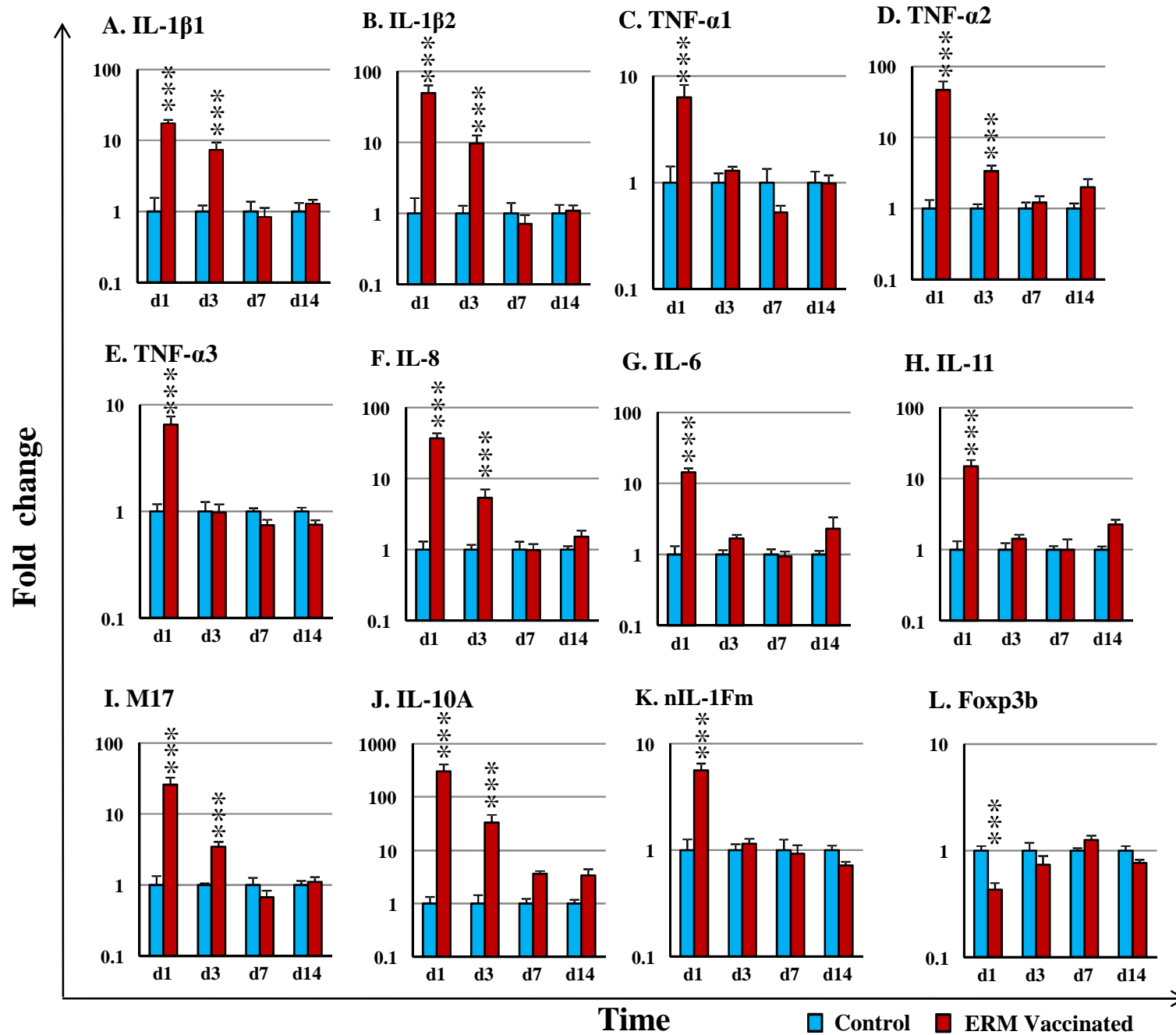
		IFN $\gamma$ 1	IFN $\gamma$ 2	IL-17A/F1A	IL-17A/F2A	IL-17A/F3	IL-4/13B1	IL-4/13B2	P35A1	P35A2	P35B1	P19	P28A	P28B	P40B2
IFN $\gamma$ 1	R	1.000	<b>0.830</b>	<b>0.904</b>	0.564	<b>0.753</b>	0.011	<b>0.652</b>	<b>0.605</b>	<b>0.840</b>	<b>0.775</b>	<b>0.789</b>	<b>0.881</b>	<b>0.793</b>	0.385
	p		.001	.000	.056	.005	.974	.022	.037	.001	.003	.002	.000	.002	.217
IFN $\gamma$ 2	R	<b>0.830</b>	1.000	<b>0.741</b>	0.394	<b>0.720</b>	-0.014	0.425	<b>0.845</b>	<b>0.839</b>	<b>0.900</b>	<b>0.646</b>	<b>0.715</b>	<b>0.620</b>	0.456
	p	.001		.006	.205	.008	.966	.169	.001	.001	.000	.023	.009	.032	.136
IL-17A/F1A	R	<b>0.904</b>	<b>0.741</b>	1.000	0.528	<b>0.755</b>	-0.210	0.386	<b>0.694</b>	<b>0.86</b>	<b>0.609</b>	<b>0.877</b>	<b>0.904</b>	<b>0.830</b>	0.186
	p	.000	.006		.078	.005	.513	.215	.012	.000	.035	.000	.000	.001	.563
IL-17A/F2A	R	0.564	0.394	0.528	1.000	0.500	0.127	0.534	0.397	<b>0.682</b>	0.473	0.530	0.490	0.215	-0.223
	p	.056	.205	.078		.098	.695	.074	.201	.015	.121	.076	.106	.502	.487
IL-17A/F3	R	<b>0.753</b>	<b>.720**</b>	<b>0.755</b>	0.500	1.000	0.168	0.498	0.528	<b>0.670</b>	0.529	0.523	<b>0.578</b>	0.420	0.102
	p	.005	.008	.005	.098		.602	.099	.078	.017	.077	.081	.049	.174	.753
IL-4/13B1	R	0.011	-0.014	-0.210	0.127	0.168	1.000	0.491	-0.412	-0.021	0.018	-0.453	-0.298	-0.347	0.207
	p	.974	.966	.513	.695	.602		.105	.183	.948	.957	.140	.347	.269	.519
IL-4/13B2	R	<b>0.652</b>	0.425	0.386	0.534	0.498	0.491	1.000	0.002	0.465	0.475	0.151	0.290	0.246	0.401
	p	.022	.169	.215	.074	.099	.105		.996	.128	.119	.639	.361	.441	.196
P35A1	R	<b>0.605</b>	<b>0.845</b>	<b>0.694</b>	0.397	0.528	-0.412	0.002	1.000	<b>0.795</b>	<b>0.762</b>	<b>0.746</b>	<b>0.691</b>	0.550	0.102
	p	.037	.001	.012	.201	.078	.183	.996		.002	.004	.005	.013	.064	.751
P35A2	R	<b>0.840</b>	<b>0.839</b>	<b>0.860</b>	<b>0.682</b>	<b>0.670</b>	-0.021	0.465	<b>0.795</b>	1.000	<b>0.830</b>	<b>0.759</b>	<b>0.779</b>	<b>0.608</b>	0.155
	p	.001	.001	.000	.015	.017	.948	.128	.002		.001	.004	.003	.036	.631
P35B1	R	<b>0.775</b>	<b>0.900</b>	<b>0.609</b>	0.473	0.529	0.018	0.475	<b>0.762</b>	<b>0.830</b>	1.000	<b>0.624</b>	<b>0.646</b>	0.460	0.290
	p	.003	.000	.035	.121	.077	.957	.119	.004	.001		.030	.023	.133	.361
P19	R	<b>0.789</b>	<b>0.646</b>	<b>0.877</b>	0.530	0.523	-0.453	0.151	<b>0.746</b>	<b>0.759</b>	<b>0.624</b>	1.000	<b>0.942</b>	<b>0.780</b>	0.030
	p	.002	.023	.000	.076	.081	.140	.639	.005	.004	.030		.000	.003	.926
P28A	R	<b>0.881</b>	<b>0.715</b>	<b>0.904</b>	0.490	<b>0.578</b>	-0.298	0.290	<b>0.691</b>	<b>0.779</b>	<b>0.646</b>	<b>0.942</b>	1.000	<b>0.896</b>	0.262
	p	.000	.009	.000	.106	.049	.347	.361	.013	.003	.023	.000		.000	.411
P28B	R	<b>0.793</b>	<b>0.620</b>	<b>0.830</b>	0.215	0.420	-0.347	0.246	0.550	<b>0.608</b>	0.460	<b>0.780</b>	<b>0.896</b>	1.000	0.513
	p	.002	.032	.001	.502	.174	.269	.441	.064	.036	.133	.003	.000		.088
p40B2	R	0.385	0.456	0.186	-0.223	0.102	0.207	0.401	0.102	0.155	0.290	0.030	0.262	0.513	1.000
	p	.217	.136	.563	.487	.753	.519	.196	.751	.631	.361	.926	.411	.088	

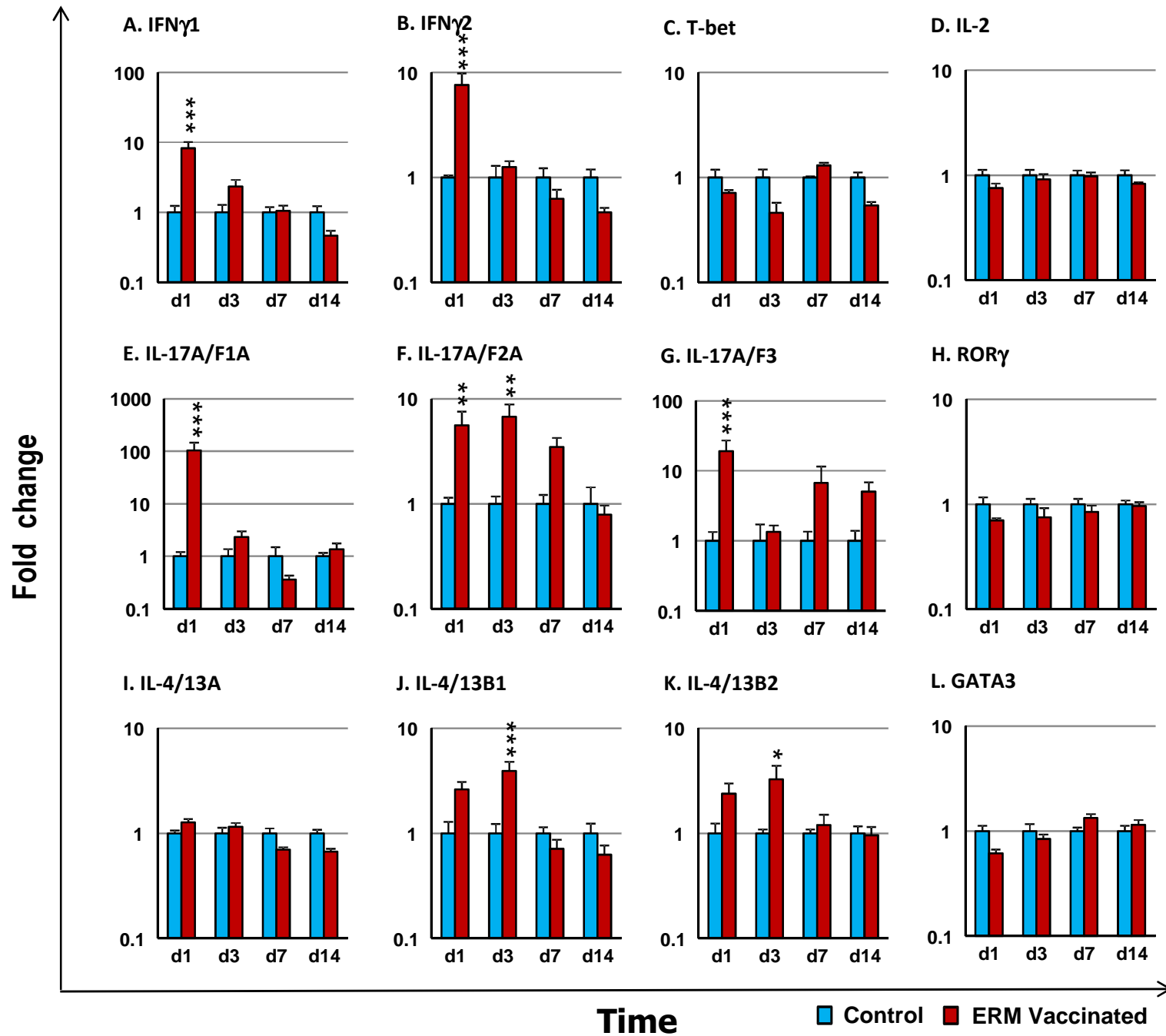
**Table 5 Comparison of gene expression modulated by ERM vaccination and *Y. ruckeri* infection in the spleen and gills of rainbow trout. “↑”= up-regulation, “↓”=down regulation and “-” = no change.**

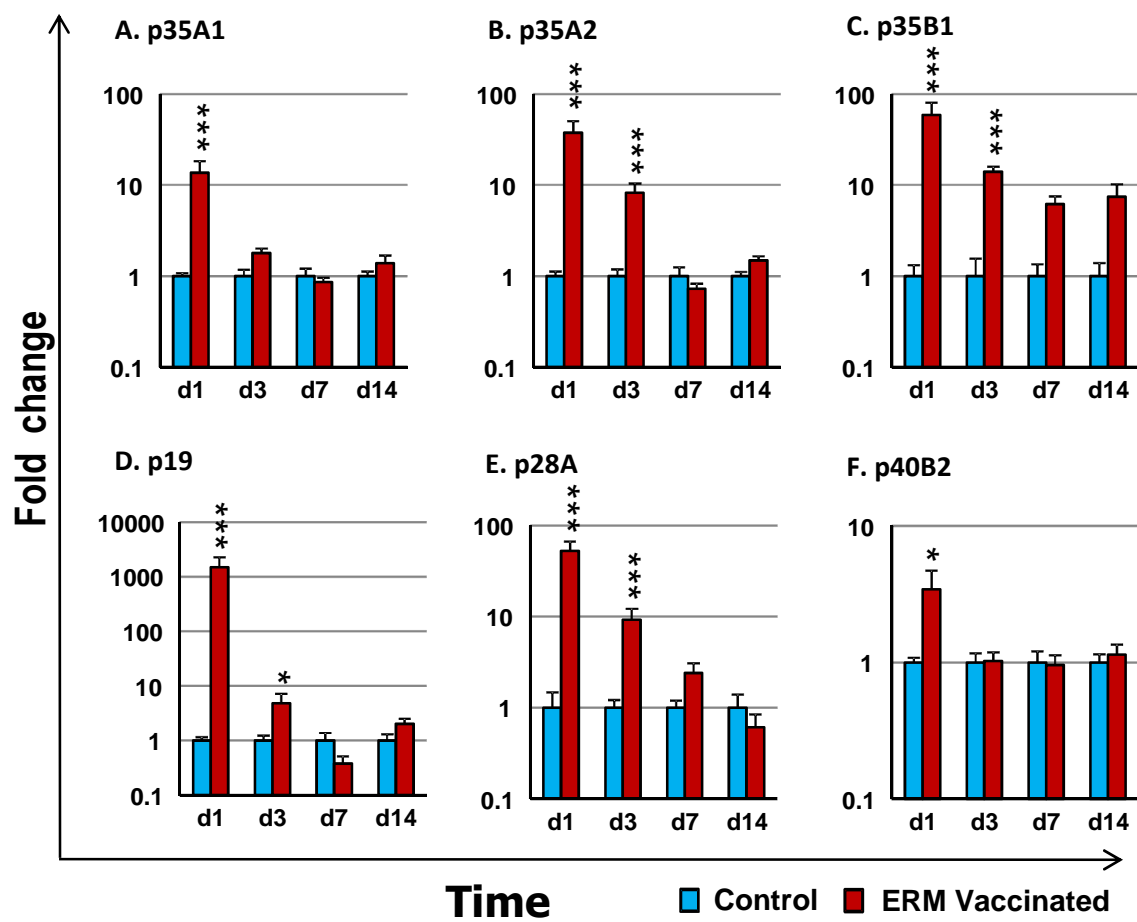
Gene	ERM vaccination		<i>Y. ruckeri</i> infection	
	Spleen	Gills	Spleen	Gills
SAA	↑	↑	↑ [9,36 ]	
SAP1	↑	-	ND	ND
SAP2	-	-	ND	ND
CATH1	↑	-	ND	ND
CATH2	↑	↑	↑ [9]	ND
Hepcidin	↑	↑	ND	ND
LEAP2A	-	-	ND	ND
β-defensin-1	-	-	ND	- [38]
β-defensin-2	-	-	ND	- [38]
β-defensin-3	-	-	ND	↑ [38]
β-defensin-4	-	-	ND	- [38]
IL-1β1	↑	-	↑ [9,17,36,45,62]	↑ [17]
IL-1β2	↑	-	↑ [45]	ND
IL-1β3	-	-	ND	ND
nIL-1Fm	↑	-	↑ [45]	ND
IL-2	-	-	↑ [17,50]	↑ [17,50]
IL-4/13A	-	-	- [9]	↓ [25]
IL-4/13B1	↑	-	ND	↓ [25]
IL-4/13B2	↑	-	ND	↓ [25]
IL-6	↑	↑	↑ [9,17,59]	↑ [17]
IL-8	↑	-	[60]	
IL-10A	↑	-	↑ [9,17,59]	↑ [17]
IL-10B	-	-	- [42]	↑ [17]
IL-11	↑	-	↑ [17,59]	↑ [17]
M17	↑	-	↑ [45]	ND
IL-12 p35A1	↑	-	↑ [26]	ND
IL-12 p35A2	↑	-	ND	ND
IL-12 p35B1	↑	-	ND	ND
IL-12 p40B1	-	-	↑ [26]	ND
IL-12 p40B2	↑	-	ND	ND
IL-12 p40C	-	-	↑ [26]	ND
IL-23 p19	↑	-	↑ [28]	ND
IL-27 p28A	↑	-	ND	ND
IL-27 p28B	↑	-	ND	ND
EBI3	-	-	ND	ND
IL-15	-	-	ND	ND
IL-17A/F1A	↑	-	↑ [24]	- [24]
IL-17A/F1B	-	-	- [24]	- [24]
IL-17A/F2A	↑	-	↑ [23-24]	- [24]
IL-17A/F2B	-	-	- [24]	- [24]
IL-17A/F3	↑	-	↑ [24]	- [24]
IL-17A/F4/N	-	-	↑ [24]	- [24]
IL-17C1	-	-	↑ [53]	ND
IL-17C2	-	-	↑ [53]	ND
IL-17D	↓	-	ND	ND
IL-18	↑	-	ND	ND
IL-20	-	-	↑ [54]	ND
IL-21	↑	-	↑ [31]	ND
IL-22	-	-	↑ [17,55]	↑ [17]
IL-34	-	-	ND	ND
IFNγ1	↑	-	↑ [9,17,36,59]	↑ [17]
IFNγ2	↑	-	ND	ND
TGF-β1A	-	-	↑ [17,36]	↑ [17]
TGF-β1B	-	-	ND	ND
TNF-α1	↑	-	↑ [9,36]	↑ [17]
TNF-α2	↑	-	↑ [17,22]	↑ [17]
TNF-α3	↑	-	↑ [22]	
T-bet	-	-	↓ [17,47], ↑ [9]	↑ [17]
GATA3	-	-	↓ [9,17,48]	↑ [17]
RORγ	-	-	↑ [49]	↑ [49]
Foxp3A	-	-	ND	ND
Foxp3B	↓	-	ND	ND











## Highlights

- ERM vaccination activates both pro- and anti-inflammatory cytokine expression in spleen.
- ERM vaccination upregulates APP and AMP expression in both spleen and gills.
- ERM vaccination induces the expression of specific IL-12 and IL-23 isoforms in spleen.
- ERM vaccination initiates a Th1/Th17 biased immune response in spleen.