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Published in: Veterinary Immunology and Immunopathology

Link to article, DOI: 10.1016/j.vetimm.2015.01.003

Publication date: 2015

Document Version Peer reviewed version

Link back to DTU Orbit

Citation (APA):

Dalgaard, T. Ś., Skovgaard, K., Norup, L. R., Pleidrup, J., Permin, A., Schou, T. W., ... Juul-Madsen, H. R. (2015). Immune gene expression in the spleen of chickens experimentally infected with Ascaridia galli. Veterinary Immunology and Immunopathology, 164(1-2), 79-86. DOI: 10.1016/j.vetimm.2015.01.003

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16 1. Introduction

Ascaridia galli is a gastrointestinal nematode infecting chickens (Permin et al., 1999; Permin et al., 17 18 1997). Substitution of traditional cages with alternative rearing systems in modern poultry production has led to an increase in the prevalence of A. galli and recent reports from Denmark 19 20 and neighbouring countries show that the majority of chickens kept in free-range systems are 21 indeed infected with A. galli (Jansson et al., 2010; Kaufmann et al., 2011; Permin et al., 1999). Infection with A. galli may directly contribute to economic losses due to higher feed conversion 22 rates/reduced weight gain and decreased egg production (Permin and Ranvig, 2001; Skallerup et 23 al., 2005). In severe cases, A. galli infections are furthermore associated with increased mortality 24 (Das et al., 2010; Gauly et al., 2005; Kilpinen et al., 2005; Permin et al., 2006), increased 25 susceptibility to secondary infections (Dahl et al., 2002; Eigaard et al., 2006; Permin etal., 2006; 26 Saif et al., 2003), impaired vaccine responses (Pleidrup et al., 2014) and even migration of worms 27 into eggs of laying hens (Fioretti et al., 2005; Reid et al., 1973). Previously, A. galli control has been 28 29 based on synthetic anthelmintics, but concerns about parasite drug resistance and left-over residues in food products call for alternative disease control strategies (Sangster, 1999). An 30 attractive alternative is vaccination, but no successful A. galli vaccines have yet been developed. 31

Natural acquired immunity is described for avian coccidiosis, another important parasitic disease. Thus, trickle immunization may induce immunity against homologous *Eimeria* challenge (Brake et al., 1997; Joyner and Norton, 1973). Extensive *Eimeria* studies have been performed in order to understand host protective immune responses and aid vaccine development (Lillehoj et al., 2007). Natural acquired immunity against *A. galli* is less well described, but reports exist on variability in disease susceptibility. The outcome of infection may e.g. be influenced by age (Idi et al., 2004;

Tongson and McCraw, 1967) and host genetics (Herd and McNaught, 1975; Kaufmann et al., 2011; 38 39 Permin and Ranvig, 2001). Estimated heritabilities for resistance/susceptibility to A. galli infections suggest that selective breeding for disease resistance may be possible (Gauly et al., 2002; 40 41 Kaufmann et al., 2011; Schou et al., 2003). In addition, several reports describe the presence of 42 very small larvae (with so called arrested development) in the late stages of an A. galli infection 43 and acquired immunity was suggested to be related to this phenomenon (Chamanza et al., 1999a; Ferdushy et al., 2014; Herd and McNaught, 1975). Interestingly, Herd et al. (1975) reported that 44 45 the proportion of larvae with arrested development was very low in chickens treated with an immunosuppressive agent. In general, it appears that development of anti-helminthic vaccines is 46 47 far more challenging than the development of vaccines directed against viral and bacterial 48 pathogens. This is in part due to their complex life cycles and the changing host-pathogen interactions occurring during different stages of helminth infections. Thus, a detailed 49 understanding of anti-helminth immunity is essential for future disease control. 50

51 The life cycle of A .galli is direct, starting with embryonation of shedded eggs in litter or soil. After 10-20 days infective L3 stage larvae are found within the parasite eggs (Permin et al., 1997). When 52 53 ingested by chickens, the A. galli eggs hatch within the first 24 hours either in the proventiculus or the duodenum of the host (Idi et al., 2004; Saif et al., 2003). After three to nine days the larvae 54 55 enter their histotrophic phase where they move deeper into the mucosal layers of the intestine 56 (Luna-Olivares et al., 2012; Saif et al., 2003; Tugwell and Ackert, 1952). Larvae recovery from the 57 intestinal wall during the first week of infection was highest in the anterior part of the jejunum, 58 but after day 7 post infection (p.i.) larvae was also found in the posterior part of the jejunum (Ferdushy et al., 2013). A high infection dose of parasite eggs may lead to a prolonged histotrophic 59 phase, but usually young adult worms return to the intestinal lumen by day 17-30 of age during 60

which period co-existence of larvae in the intestinal wall and young worms in the intestinal 61 62 content is seen (Ferdushy et al., 2013; Herd and McNaught, 1975; Katakam et al., 2010). Recently, Luna-Olivares et al. (2012) suggested that "mucosal phase" may be a more appropriate term than 63 64 "histotrophic phase" (lamina propria invasive) as the larvae may not penetrate as deep into the 65 intestinal tissue as originally thought. They reported that most larvae were observed in the lumen (but in close contact with the epithelium) (63%) followed by "within epithelium" (32%) and only 66 few in the lamina propria (5%). However, only the very early time-point 3 days p.i. was 67 68 investigated and it is uncertain what happened later in the histotrophic/mucosal phase. However, Katakam et al. (2010) was able to recover all larvae by an EDTA method, i.e. no additional larvae 69 70 were recovered when applying additional pepsin digestion after EDTA incubation of intestinal 71 samples taken 2 weeks p.i. indicating that lamina propria associated larvae are few also at this 72 time point.

The chicken spleen works as a secondary lymphoid organ where innate and adaptive immune responses are efficiently mounted. It is hypothesized that the avian spleen plays an even more important immunological role than in mammals as avian lymphatic vessels and lymph nodes are poorly developed. The aim of this study was to investigate systemic immunological responses at different stages of an *A. galli* infection by comparing gene expression profiles in spleen tissue between infected and control chickens at week 2, 6 and 9 post infection (p.i.).

79

80 2. Materials and Methods

81 *2.1. Animals*

In the experiment, chickens of mixed gender from the Aarhus University L133 were used. Line 133 82 is of White Leghorn origin and contains only birds with the major histocompatibility complex 83 (MHC) haplotype B13. Water and commercial chicken feed were supplied ad libitum. The lighting 84 85 period was 12 h daily, and the chickens were kept at a temperature of 21°C. All experimental 86 chickens were produced from MHC-characterized parents, and the MHC haplotypes of the offspring were confirmed by genotyping the LEI0258 microsatellite locus (McConnell et al., 1999) 87 by PCR-based fragment analysis as earlier described (Dalgaard et al., 2005). Some birds in the 88 89 current experiment were shared with an already published experiment (Pleidrup et al., 2014).

90 2.2. Experimental outline

91 Experimental chickens were divided into two treatment groups; 1) negative control chickens and 2) chickens subjected to A. galli infection that were kept in separate rooms of the chicken facility. 92 93 At 4 weeks of age, chickens in group 2 were orally infected with 1750 embryonated A. galli eggs 94 recovered from female worm uteri obtained from naturally infected commercial hens and embryonated in H2SO4 as described in Permin et al. (1997). Sixteen animals from each group were 95 used for weekly blood sampling and seven other animals from each group were sacrificed at week 96 2, 6 and 9 p.i. for spleen collection. At week 6 and 9 p.i. faecal samples were collected before 97 sacrificing the chickens. Licence to conduct the animal experiment was obtained from the Danish 98 99 Ministry of Justice, Animal Experimentation Inspectorate by Helle R. Juul-Madsen. The experiment was conducted according to the ethical guidelines 100

101 2.3. A. galli-specific IgG ELISA

Blood samples from infected animals were taken at weeks 0, 6, 7, 8, 9 p.i. and from negative controls at week 0, 6, 9 p.i. and serum was used for detection of *A. galli*-specific IgG antibodies as earlier described (Norup et al., 2013).

105 2.4. Faecal A. galli egg excretion

Faecal samples were obtained from *A. galli*-infected chickens before sacrificing them for spleen sampling at weeks 6 and 9 p.i. Faeces was not sampled from chickens sacrificed 2 weeks p.i. as adult egg secreting worms are not developed until week 5-8 p.i. (Permin and Hansen, 1998). The faecal samples were examined for the presence of *A. galli* eggs using a modified McMaster counting technique (Henriksen and Aagaard, 1976; Permin et al., 1997) with a detection limit of 20 eggs per gram faeces (EPG).

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113 2.5. RNA extraction

After collection, spleens were sectioned (triangular cross-sectional slice from upper part) and identical samples from each chicken were immediately placed in RNAlater (Ambion/Life Technologies), kept overnight at 4°C and then at -20°C until further processing. Amounts of 7 to 15 mg tissue were homogenised on a TissueLyzer LT (Qiagen), and RNA isolation and DNA digestion was done using the NucleoSpin 96 RNA kit (Macherey-Nagel) according to the manufacturer's instructions. RNA quality was controlled on a 1 % agarose gel and the RNA concentration and purity were determined using a NanoDrop spectrophotometer (Saveen and Werner AB).

121

122 2.6. cDNA synthesis and pre amplification of mRNA

cDNA synthesis and preamplification was performed as described previously (Skovgaard et al., 123 124 2013). Extracted total RNA was converted into cDNA by reverse transcription of 480 ng RNA using the QuantiTECT Reverse Transcription kit (Qiagen), cDNA was diluted 1:5 in low EDTA TE-buffer 125 126 (VWR – Bie & Berntsen) prior to preamplification. Preamplification was performed using TagMan 127 PreAmp Master Mix (Applied Biosystems) and a 200 nM pooled primer mix was prepared combining each primer used in the present study. TaqMan PreAmp Master Mix (5 µl) was mixed 128 with 2.5 µl 200 nM pooled primer mix and 2.5 µl diluted cDNA, and incubated at 95°C for 10 min 129 130 and 16 cycles of 95°C for 15 sec and 60°C for 4 min. 16 U of Exonuclease I (New England BioLabs) was added to the preamplified cDNA, thermal cycling conditions were set to 37°C for 30 min 131 132 followed by 80°C for 15 min. Preamplified cDNA was diluted 1:10 in low EDTA TE-buffer (VWR -133 Bie & Berntsen) before qPCR. Primers were designed using Primer3 (http://bioinfo.ut.ee/primer3-0.4.0/) as described in (Skovgaard et al., 2010), and purchased from Sigma-Aldrich. Primer 134 sequences, efficiencies and amplicon length are shown in Table 1. 135

136

137 2.7. qPCR

Gene expression mRNA was analysed by quantitative real-time PCR (qPCR) performed in Dynamic Array Integrated Fluidic Circuits (Fluidigm) following the protocol described previously (Skovgaard et al., 2013). The following cycle parameter was used: 2 min at 50°C, 10 min at 95°C, followed by 35 cycles with denaturing for 15 sec at 95°C and annealing/ elongation for 1 min at 60°C. Melting curves were generated after each run to confirm a single PCR product (from 60°C to 95°C, increasing 1°C/ 3 sec). Reactions were performed in duplicates (cDNA replicates). Non template controls (NTC) were included to indicate potential problems with non-specific amplification or

sample contaminations. Non-reverse transcriptase controls were included to assess potential DNAcontamination.

147

Expression data (Cq values) were acquired using the Fluidigm Real-Time PCR Analysis software 148 149 3.0.2 (Fluidigm) and exported to GenEx (MultiD) for data pre-processing including interplate correction, correction for PCR efficiency for each primer assay individually, normalising to six 150 highly stable reference genes, and averaging of cDNA technical repeats. Using GeNorm (17) and 151 152 NormFinder (18), glyceraldehyde-3-phosphate dehydrogenase (GAPDH), β2 microglobulin (B2M), peptidylprolyl isomerase A (PPIA), hypoxanthine phosphoribisyl transferase I (HRPT1), TATA-box 153 154 binding protein (TBP), and tyrosine 3-monooxygenase/tryptophan 5-monooxygenase activation 155 protein, zeta polypeptide (YWHAE) were identified as the most stably expressed reference genes out of eight candidates. For each primer assay, the mean relative expression level of the control 156 group was scaled to one during data transformation log2 (Cq) to linear scale. Gene expression data 157 were log₂-transformed before testing for normal distribution, Student t test was used to analyse 158 normally distributed data, while the non-parametric test (Wilcoxon-Mann-Whitney test) was 159 160 used when data was non-normal distributed. Gene expression was considered significantly 161 different if the P value was less than 0.05 and the relative expression was greater than 2.0. Experimental practice and reporting have been performed according to the Minimum Information 162 for Publication of Quantitative Real-Time PCR Experiments (MIQE) guidelines (Bustin et al., 2009). 163

164

165 **3. Results and Discussion**

According to earlier studies, week 2 p.i. represents the mucosal phase of the *A. galli* larvae whereas at weeks 6 and 9 adult worms are present in the intestinal lumen. In the present

experiment only 43 % of the animals (data not shown) shedded A. galli eggs in faeces at week 6 168 p.i., and we hypothesise that the A. galli worms are young and have just recently started 169 producing eggs. Presumably some larvae are also still present in the mucosa at this time point as 170 171 earlier reported by Ferdushi et al. (2013). In contrast, 73 % of the chickens (data not shown) 172 shedded *A. galli* eggs in faeces at week 9 p.i. and with a higher mean EPG per animal than at week 6 (Figure 1a). Thus, this time point may represent more mature adult worms. None of the chickens 173 in the A. galli-free group tested EPG positive at any time-point during the experiment (data not 174 175 shown). Additional chickens allocated to blood sampling were sero-negative at the day of infection (data not shown). Chickens in the blood sampled A. galli-inoculated group had seroconverted by 176 177 week 6 p.i. and showed positive titres of A. galli-specific serum IgG throughout the rest of the experiment. Chickens from the blood sampled negative control group were tested at weeks 6 and 178 9 p.i. and were found to be sero-negative at both time-points (Fig. 1.b). A systemic humoral 179 immune response is reported by others as early as 2 weeks p.i., but serum titres do not appear to 180 correlate with egg excretion or worm burden (Marcos-Atxutegi et al., 2009; Norup et al., 2013; 181 Schwarz et al., 2011). 182

183

In order to understand systemic molecular response mechanisms in different stages of an *A. galli* infection we studied gene expression profiles in spleen sampled 2, 6 and 9 weeks after the experimental infection. Twelve genes (representing inflammatory cytokines, antimicrobial peptides, acute phase proteins, soluble pattern recognition receptors and T cell signature cytokines) were differentially expressed (P < 0.05) at at least one of the three analysed time points after the *A. galli* infection compared to the control group.

191 3.1. (Pro-)Inflammatory cytokines

192 Only few studies have been published concerning innate immune responses towards A. galli in chickens and focus has been on local responses in the small intestine. Thus, a single study reports 193 194 increased numbers of mast cells in the chicken jejunum 2 weeks post A. galli infection (Darmawi et 195 al., 2013). Another study reports increased numbers of presumably heterophils in the jejunum 3 days after an A. galli infection (Luna-Olivares et al., 2012). Interestingly, a genetic association 196 study indicated that chicken IFN-y gene variants may influence A. galli susceptibility (Luhken et al., 197 198 2011). In the present study, we analysed the expression of inflammatory cytokines in the spleen 199 (Table 2). Surprisingly, the expression of IFN- α , IL-1 β , IL-12 β and Il-18 was up regulated at week 6 200 p.i., but not at week 2 p.i. or week 9 p.i. The IL-8 expression was up regulated at week 2 as well as 201 week 6 p.i. in *A. galli*-infected chickens. Despite structural differences most avian cytokines display conserved functions compared to their mammalian counterparts (Staeheli et al., 2001), and roles 202 in the chicken inflammatory response have been described for IL-8, IL-1β, IL-18, IL-12β (Balu and 203 Kaiser, 2003; Barker et al., 1993; Laurent et al., 2001; Schneider et al., 2000; Weining et al., 1998; 204 Withanage et al., 2004). Also chicken IFN- α (ChIFN-I) was identified to have a similar function to 205 206 the mammalian counterpart as a potent antiviral agent (Schultz et al., 1995; Sick et al., 1996). It is 207 now accepted that IFN- α in mammals beside its antiviral properties shows additional immunomodulating effects. Although little is known of IFN- α 's role in parasite infections, 208 treatment of helminth disease in mice has been attempted with recombinant IFN- α (Godot et al., 209 210 2003).

211 *3.2. Antimicrobial peptides*

The expression of DEF_{β1} was significantly reduced at week 2 p.i. and significantly increased at 212 213 weeks 6 and 8 p.i. in spleen tissue of A. galli-infected chickens (Table 2). Antimicrobial peptides like defensins play an important role in innate immunity, and activity directed against bacteria, 214 215 fungi and viruses has been reported (Ganz, 2003). Interestingly, defensins may influence adaptive 216 immune responses as they can affect the maturation of dendritic cells as well as effector T cell recruitment (Yang et al., 2002). In the chicken genome, 14 beta-defensin/gallinacin genes exist and 217 the nomenclature AvBD1-14 was suggested (Lynn et al., 2007). Local expression of several of the 218 219 AvBD genes and their antimicrobial activity against avian enteric pathogens have been described 220 (Evans and Harmon, 1995; Harmon, 1998; Hong et al., 2012; Sugiarto and Yu, 2004). However, the 221 role of AvBD in innate immunity towards helminth infections is not clear. In humans, some beta-222 defensins are up regulated by pro inflammatory cytokines (McDermott et al., 2006; Scott and Hancock, 2000). In the present study an increased expression of DEF β 1/AvBD1 coincided with an 223 224 increase in the expression of pro-inflammatory cytokine genes at week 6 p.i., but not at week 9 p.i.

225 3.3. Acute phase proteins

Mannose binding protein (MBL) and C-reactive protein (CRP) are soluble pattern recognition 226 227 receptors. Few reports exist on chicken CRP, but it appears that infections with Eimeria spp. and Histomonas meleagridis induce high levels of CRP (Chamanza et al., 1999). In mammals, MBL binds 228 229 to microbial surface carbohydrates and mediates opsonophagocytosis directly or through activation of the lectin complement pathway. A conserved function of MBL in the chicken was 230 suggested as cMBL in a heterologous in vitro assay was shown to enhance human complement 231 factor 4 (C4) deposition in a calcium dependent way (Norup and Juul-Madsen, 2007). As in 232 233 mammals, reduced levels of serum MBL in chickens may lead to increased disease susceptibility to

viral and bacterial infections (Juul-Madsen et al., 2007; Schou et al., 2010). Chicken MBL is mainly
produced in the liver, but constitutive and inducible local expression of the gene has also been
reported (Hogenkamp et al., 2006; Laursen et al., 1998; Nielsen et al., 1998). In this study, MBL
expression was significantly increased in spleen tissue of *A. galli*-infected chickens 6 weeks p.i.
(Table 2). An *in vivo* function of MBL in intestinal helminth infections has not yet been determined,
but preliminary results suggest that faecal shedding of *A. galli* eggs is reduced in infected inbred
chickens with high MBL serum levels (unpublished, Norup).

241 3.4. Th signature cytokines

242 In mammals, Th2 polarised cells drive responses to helminth infections. Also in the chicken a Th2 243 polarised cytokine response was reported in the jejunum and spleen of A. galli-infected chickens 2 weeks p.i. (Degen et al., 2005; Kaiser, 2007; Pleidrup et al., 2014; Schwarz et al., 2011). In 244 245 agreement with former studies, we observed an increased expression of the Th2 signature 246 cytokine IL-13 at 2 weeks p.i. in the spleen of A. galli-infected chickens, but not at later stages of the infection (Table 2). This time-point corresponds to the mucosal phase of the infection which 247 co-incides with influx of both $\alpha\beta$ (including CD4+ve cells) and $\gamma\delta$ T cells in the jejunal mucosa as 248 249 reported by others (Schwarz et al., 2011). In the present study we observed a slightly decreased expression of the Th1 signature cytokine IFN-y at week 9 p.i. in spleen tissue of A. galli-infected 250 251 chickens. This observation is in contrast to earlier findings by Degen et al. (2005) who reported 252 decreased relative cytokine mRNA ratios (infected/non-infected) for IFN-y as early as 2 weeks p.i. Earlier reports do suggest that onset and length of the larvae mucosal phase depend on infection 253 dose which differed between the two experiments. 254

255 3.5. Anti-inflammatory cytokines

In human and murine infections the survival strategy of helminth parasites is largely based on 256 257 immunoregulation by excretory-secretory (ES) products through mechanisms involving regulatory T cells (Taylor et al., 2012). No Foxp3 orthologue has been identified in the chicken, but thymic 258 259 CD4+CD25+ T cells were characterised as counterparts of mammalian natural Tregs by production 260 of IL-10 and TGF-β (Shanmugasundaram and Selvaraj, 2011). In the present study, an increased expression of TGF-β4 was observed 6 weeks p.i. in spleen tissue of A. galli- infected chickens 261 (Table 2). The chicken TGF-β gene-family includes: TGF-β2, TGF-β3 and TGF-β4, of which the latter 262 263 is the chicken orthologue of mammalian TGF- β 1 acting as an anti-inflammatory cytokine (Jakowlew et al., 1997; Pan and Halper, 2003). IL-10 has a conserved function in the chicken acting 264 265 as an anti-inflammatory cytokine (Rothwell et al., 2004). No increased expression of IL-10 was 266 observed in the present study; instead the expression was lower in the spleen tissue of A. galliinfected chickens 6 and 9 weeks p.i. than in controls where expression increased by age (data not 267 shown). We have no explanation for this and further studies in other inbred chicken lines as well 268 as outbred lines need to be conducted in order to elucidate if this is a general response in A. galli 269 infections. Further studies of the expression of anti-inflammatory cytokines may also help us to 270 understand why A. galli infected chickens appear to have impaired vaccine responses towards 271 third party antigens (Pleidrup et al., 2014). 272

273 *3.6.* Conclusion

In summary, we have investigated the avian systemic immune response to *A. galli* infection by expression analyses of immune genes in spleen. Interestingly, we observed only few differentially expressed genes at week 2 p.i. which corresponds to the larvae mucosal phase. In contrast, by week 6 p.i. where the larvae expectedly have matured and migrated back into the intestinal 278 lumen, we observed increased expression of pro-inflammatory cytokines and acute phase
279 proteins. It is yet to be determined if the observed pro-inflammatory response is caused by *A. galli*

280 specific pathogen-associated molecular pattern molecules (PAMPs), host specific damage-

281 associated molecular pattern molecules (DAMPs) released by tissue damage, DAMP homologues

282 in parasite secretions of even by opportunistic secondary infections.

283

284 **Conflict of interest statement**

285 The authors declare to have no conflicts of interest.

286 Acknowledgements

The authors wish to acknowledge financial support from The Danish Council for Strategic Research, Aarhus University and the European Union Seventh Framework Network of Animal Disease Infectiology Research Facilities (NADIR; reference number FP7-228394). Pete Kaiser and Lisa Rothwell are thanked for fruitful comments and support, Karin Tarp, Lene Rosborg Dal and Helle Handll for excellent technical assistance, and Karin V. Østergaard for proof reading of the manuscript.

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