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# Immunostimulatory potential of Nigella sativa seeds extract on chicken macrophage function

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ARTICLE INFO	ABSTRACT
Received : 13 September 2021	In present study, the effect of Nigella sativa seeds extract on the chicken
Revised : 07 October 2021	peripheral blood mononuclear cells (PBMCs) was investigated. The chicken
Accepted : 17 October 2021	PBMCs were stimulated with two different doses (125µg/ml; 250µg/ml) of
	Nigella sativa seeds extract and the cells were harvested at different time points
Available online: 19 December 2021	till 48h post-stimulation for analysis of iNOS gene expression by quantitative
	PCR and nitric oxide (NO) estimation at 24 and 48h post-stimulation. High dose
Key Words:	(250µg/ml) of Nigella sativa seed extract showed remarkable induction of iNOS
iNOS	transcripts expression and NO production at 48h post-stimulation, which were
Macrophages	12.8040±1.03347 folds and 5.7089±.64535µM respectively. The results indicated
Nitric oxide	the immunostimulatory potential of the Nigella sativa seed extract on the
PBMCs	chicken PBMCs.

### Introduction

used in the treatment of different diseases. Nigella sativa belongs to Ranunculaceae family and known as black cumin. Nigella sativa has been widely used as food additive, food preservative or as spice in many cultures. In the field of medicine due to its various therapeutic potentials Nigella sativa broadly used in the treatment of many infectious diseases Yimer et al. (2019). The seeds of Nigella sativa showed remarkable benefits due to the presence of a variety of biochemical and pharmacological components present into it. The various active components of black cumin have been reported to show characteristics like antimicrobial. antioxidant. anti-malarial. antiinflammatory, analgesic, antihypertensive, antidiabetic, antineoplastic Darmawan et al. (2019) and Hwang et al. (2021) and immune-potentiating effects Manoharan et al. (2021). Macrophages act as scavenger cell; provide the first line of defense against microbial invasion and activate different

Natural product obtained from the plants widely types of signaling pathways Chen et al. (2017) and Ren et al. (2017). Macrophages constitute important participation in the bi-directional interaction between innate and specific immunity Carnagarin et al. (2019). The macrophage act as scavenger cell and performs many functions such as engulfment of deformed cells, attacking the invaders and providing response to infections, tumors and inflammation Wang et al. (2019). Kiboneka (2021) reported that macrophages play important role in innate and adaptive immunity and kill pathogenic microorganisms via phagocytosis process and initiate an immune response by antigen processing and presentation to T- and Blymphocytes. They also modulate adaptive immune system by producing various inflammatory and anti-inflammatory cytokines. As per Tejero et al. (2019), macrophages also release immunologically important effector biomolecules, such as reactive oxygen species (ROS) and nitric oxide (NO). Macrophages produce NO by oxidizing the guanidine nitrogen of L-arginine by an enzyme, nitric oxide synthase (NOS). This enzyme is inducible in macrophages by external stimulators like bacterial endotoxins (lipopolysaccharide) or by lymphokines and is termed as inducible NOS (iNOS) Gordon (2016). Macrophages also secrete cytokines IFN-y and IL-2 (Th1-type cytokines) and IL-13, IL-4 (a Th2-type cytokine) and aid the development of the adaptive immune responses Khan et al. (2017). Recently, Mehkri et al. (2021) the potentiation demonstrated effects of Thymoquinone (Nigella sativa) oil on mouse macrophages.

Elmowalid et al. (2013) revealed out the direct effects of Nigella sativa seed purified aqueous extract on sheep monocytes-derived- macrophages (MDM) culture. They reported that the Nigella sativa effect is not species specific and confirmed and suggested that Nigella sativa may serve as potent immunomodulator of sheep macrophage functions that could contribute to the overall host defense by preventing host cell microbial infections. AlMahmud et al. (2017) demonstrated Nigella that sativa active component Thymoquinone (TQ) modulates nitric oxide production and also inhibited the replication of intracellular *Mvcobacterium* tuberculosis in macrophages. Nigella sativa oil alone or in combination with albendazole showed antiechinoncoccosis effect by increasing IL-5 and NO production and reducing TNF-  $\alpha$  in mice model Kishik et al. (2020). Bahrami et al. (2021) reveled out that ethanolic extract of Nigella sativa significantly decreased the level of NO in peritoneal murine macrophages and RAW264.7 cells. They also demonstrated that Nigella sativa ethanolic extract significantly increased IL-10 expression, while IL-6 and TNF- $\alpha$  expression was significantly decreased in RAW264.7 cells. Nigella sativa extract was used by many workers as an immunomodulating agent and its immunostimulatory activity has been reported in sheep Elmowalid et al. (2013), in human Koshak et al. (2018) and in mice Kishik et al. (2020), but not yet explored in chicken. There is very little known about the immunostimulatory effect of Nigella sativa seed extract on chicken immune cells. Here, we demonstrated the effect of Nigella sativa extract with two doses (Low and high) on the PBMCs that helped in understanding the mode of action and

molecular mechanisms involved in the macrophage function. We analyzed the transcriptional levels of inducible nitric oxide synthase (iNOS) as well as NO production in the cell culture supernatant on stimulation with two doses; lower and higher dose of *Nigella sativa* in chicken PBMCs to understand the effect on macrophage functions.

### **Material and Methods**

### Reagents

RPMI 1640 with L-glutamine and 25 mM HEPES buffer were purchased from HiMedia Pvt. Ltd. India. Fetal bovine serum (FBS) was purchased from Hyclone (Logan, USA) and inactivated by heating at 56 °C for 30 min.

### **Seeds Extract Preparation**

Seeds of *Nigella sativa* were purchased from authenticated Local herb store available at Bareilly, UP (India). The Seeds were dried at 45 °C for removal of moisture and powdered. In 70% methanol (100ml), Ten grams of powdered seed was stirred overnight and centrifuged at 10,000 rpm at 4 °C for 10 min, the supernatant was collected. Methanol was removed by evaporation, and obtained yield was 12% w/w. The seed extract was suspended in DMSO for further *in vitro* studies. The extract was sterilized by filtration using 0.22  $\mu$ m syringe filters and stored at -20 °C till use.

### Primers

Published primers specific to chicken were synthesized from M/S Integrated DNA Technologies, Iowa, USA and used in the study (Table 1).

### **Isolation of Chicken PBMCs**

Blood was collected with heparin (20 IU/ml of blood) from the experimental birds: Chicken (n=6) maintained in the Avian Immunology laboratory, Immunology Section, ICAR-IVRI, Izzatnagar Bareilly U.P. (India). The blood was layered on to the equal amount of Ficoll Hypaque (Sigma, MO, USA) with specific gravity 1.077 g/ml and PBMCs were isoloated by density gradient centrifugation at  $500 \times g$  for 45 min. The PBMCs were washed twice with sterile PBS (pH 7.2) and the cells were re-suspended in RPMI-1640 medium containing 2 % fetal bovine serum. The viability of cells was determined by trypan blue dye exclusion method and cells were adjusted to a concentration of  $1 \times 10^7$  cells/ml and stored on ice.

Target Gene	Primer sequence (5'-3')	Product size (bp)	Reference
GAPDH	F: AGCACCCGCATCAAAGG	283	Kuo <i>et al.</i> (2017)
	R: CATCATCCCAGCGTCCA		
iNOS	F: TGGGTGGAAGCCGAAATA	241	Zhang et al. (2019)
	R:GTACCAGCCGTTGAAAGGAC		

### Table: 1 Primer used for quantitative real time PCR

# **Stimulation of macrophages for Nitric oxide** nuclease free water (20 µl). The purity of RNA was (NO) estimation checked by measuring the absorbance at 260 and

The PBMCs ( $2 \times 10^6$  cells/ml) were resuspended in RPMI 1640 medium containing L-arginine (5 mM) and stimulated with two different doses (125 and 250µg/ml) of Nigella sativa extract in triplicate. The supernatants were collected at 24 and 48h poststimulation and stored at -20 °C until further processing. NO production was determined by measuring the concentration of nitrite in the supernatant by using Griess assay. Briefly, 50 µL of standard and samples was added with 50 µL of Griess reagent (Sigma, MO, USA) followed by incubation at 37 °C for 30 min. Optical density was determined at 550 nm in a spectrophotometer. Different concentrations of sodium nitrite were used as standard and the concentration of NO was calculated from the standard curve.

# Stimulation of chicken PBMCs with Nigella sativa extract

The PBMCs (1 x  $10^6$  cells) were stimulated with two different doses of (low dose,  $125\mu$ g/ml; high dose,  $250\mu$ g/ml) of *Nigella sativa* seed extract by incubation at 40 °C, 5% CO<sub>2</sub> environment Swamy and Tan (2000) and Gholamnezhad *et al.* (2015). Then, the cells were harvested at different time points till 48h post-stimulation for analysis of iNOS gene expression by qPCR at 3, 6, 12, 24 and 48h post-stimulation.

### **Total RNA isolation from PBMCs**

Treated and untreated PBMCs were centrifuged for 5 min and supernatant was discarded. Into the pellet 750  $\mu$ l of QIAzol lysis reagent (Qiagen, CA, USA) and 250  $\mu$ l of chloroform were added and vortexed for 30 s. For separation of phases, the tubes were centrifuged at 12000 × g for 20 min at 4 °C. The supernatant aqueous phase rich in RNA were precipitated with 400  $\mu$ l of isopropanol at 12000 × g for 15 min and washed with 1 ml of 70% ethanol. Then, the tubes were air dried by inverting on to the clean filter paper for about 10 min for the removal of excess ethanol. The RNA pellet was dissolved in

nuclease free water (20  $\mu$ l). The purity of RNA was checked by measuring the absorbance at 260 and 280 nm in a Nanodrop UV spectrophotometer (Thermo Scientific, USA) Bashir *et al.* (2019).

### **Complementary DNA (cDNA) preparation**

Total RNA (2µg) was used for preparation of cDNA employing Revertaid<sup>TM</sup> First Strand cDNA Synthesis Kit (Thermo Scientific, USA), following manufacturer's instructions. For the synthesis of cDNA, 2 µg of RNA and 1 µL of random hexamer (Thermo Scientific, USA) were added to the 9.5  $\mu$ L of nuclease-free water (total volume 12.5 µL) and incubated at 65 °C for 10 min. Then, the following reagents were added to the tubes: 5 X reaction buffer (4 µL), RNase inhibitor (0.5 µL), 10mM dNTP mix (2  $\mu$ L), and reverse transcriptase (1  $\mu$ L) and mixed and then incubated at 25 °C for 10 min followed by 50 °C for 50 min for cDNA synthesis and by heating at 85 °C for 5 min, reaction was terminated. The cDNA was stored at -20 °C till further use.

### Quantitative Real-Time PCR (qPCR)

Expression levels of iNOS was analyzed by realtime PCR (Agilent Technologies AriaMx Real-Time PCR System) using the Quanti fast SYBR Green qPCR kit (Agilent) (Fig: 1). The chicken specific primers (Table 1) were used. The qPCR mixture consisted of 2 µl cDNA, 10 µl of QuantiFast® SYBR Green Master Mix (Qiagen, CA, USA), primers (0.5 µl each, 10 pmol concentration) and nuclease free water to a volume of 20 µl. Real time PCR was carried out with the following programme: 1 cycle at 95 °C for 5 min, followed by 40 cycles each of 94 °C for 30 s, 60 °C for 45 s, 70 °C for 45 s and 1 cycle of 94 °C for 30 s. All the samples were carried out in duplicates on the same plate. iNOS transcripts expression level was calculated relative to the expression of the GAPDH gene Boeglin et al. (2011) and Berzi et al. (2014) and expressed as n-fold changes relative to the control samples Pfaffl (2001) (Fig: 2). The cycle in which reporter dye concentration crossed a preset threshold was recorded as cycle threshold (Ct) value. The data of qPCR were analyzed by  $2^{-\Delta\Delta Ct}$  method of Pfaffl (2001) to derive the relative fold change in mRNA of the target gene.

### Statistical analysis

GraphPad Prism 8.0 was used for the statistical analyses. The relative fold change in the expression of immune response genes were analyzed by using one way analysis of variance (ANOVA); when the F ratio was significant, least significant difference (LSD) was used as *post hoc* test. \* (P $\leq$ 0.05) and \*\*(D $\leq$ 0.01) is lighted at the light of the light of the statistic light of the statistic light of the state of the st

\*\*(P≤0.01) indicate statistically significant.

### **Results and Discussion**

Effect of *Nigella sativa* seed extract on iNOS gene expression and NO production in the chicken PBMCs

In present study, we have analyzed the transcriptional levels of iNOS as well as NO

production in the supernatant of cell culture in the chicken PBMCs on stimulation with two doses of methanolic *Nigella sativa* seed extract to understand its effect on macrophage functions. The enzyme iNOS is involved in the production of NO, which has antimicrobial and antiviral properties. Out of three essential enzymes which produced NO from L- arginine, iNOS play very crucial role in infection, inflammation and blood pressure regulation Matsubara *et al.* (2015).

The higher dose of *Nigella sativa* extract significantly (P $\leq$ 0.01) up-regulated iNOS expression in chicken PBMCs at 12, 24 and 48h post-stimulation and a peak expression of 12.80 $\pm$ 1.033 folds was observed at 48h interval in the table 2 and figure 3. Tripathi *et al.* (2012) also studied that methanolic fraction of *Nigella sativa* on rat-peritoneal-macrophage culture

Treatments	Relative fold change in iNOS expression					
	3h	6h	12h	24h	48h	
Low dose	2.5226±1.04606	2.3591±.95981	.8843±.25474	2.3347±.68389	7.9943±2.87686 **	
High dose	1.4970±.55634	1.1401±.41510	6.3746±1.29504**	6.8396±.90284**	12.8040±1.0334 7**	

\*\* indicate statistically significant difference at P≤0.01

 Table: 3 Nitric oxide production by chicken PBMCs

 after Nigella sativa extract (Low dose and high dose)

 stimulation

Treatments	24h	48h
Low dose	1.8433±.06642*	1.7489±.42986*
High dose	3.3633±.55827**	5.7089 ±.64535**

\* indicate statistically significant difference at  $P \le 0.05$  and

\*\* indicate statistically significant difference at P≤0.01

indicated the significantly induction of iNOS expressions and NO production, which is in favor of present study conducted on macrophage of chicken PBMCs. The lower dose of the extract showed significant (P $\leq$ 0.01) up-regulation of the iNOS transcripts only at 48h post-stimulation period in the chicken PBMCs. Further, the higher dose of the extract also induced the maximum level of NO production, which was 3.36±.56 and 5.71±.65 µM at 24 and 48 h post-stimulation,

respectively as shown in table 3 and figure 4. Miliani et al. (2018) also observed enhanced production of iNOS in necrotic Jurkat T cell line lysates-pulsed macrophage co-culture system with thymoquinone treatment. In contrast to our finding, Mycobacterium Thymoquinone treatment to tuberculosis infected RAW264.7 cells inhibited expression expression of iNOS gene and reduced production of NO and eventually diminished pathogen-derived stress in host cells Al Mahmud et al. (2017). Moreover, recently, Bahrami et al. (2021) reported that ethanolic extract of Nigella sativa seeds reduced NO production and reduced IL-10 expression in RAW264.7 cells and peritoneal murine macrophages while in present study upregulation of iNOS transcripts in the chicken PBMCs along with the significant production NO by extract of Nigella sativa seeds indicated its potential for enhancing the macrophage functions in chicken.

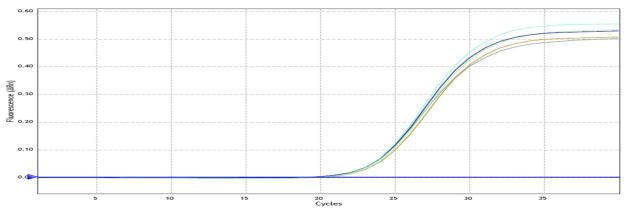


Figure 1: Amplification plot of qPCR assay for iNOS gene from Non-stimulated Control and stimulated samples

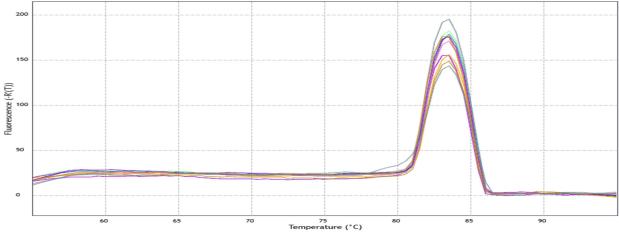
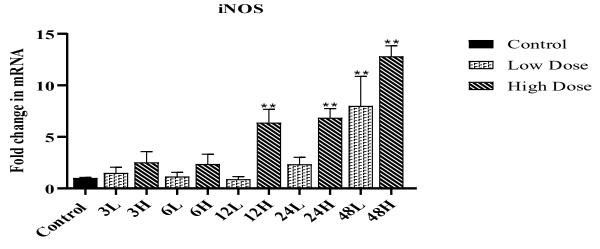


Figure 2: Melt curve of qPCR assay for iNOS gene from Non-stimulated Control and stimulated samples



Hours post-stimulation (h)

Figure 3: Relative expression of iNOS transcript in chicken PBMCs stimulated with Low dose (125µg/ml) and High dose (250 µg/ml) of *Nigella sativa* seed extract over a period of 48h

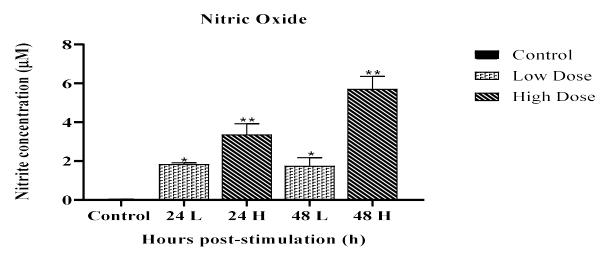


Figure 4: Production of nitric oxide (µM) in chicken PBMCs stimulated with Low dose (125µg/ml) and High dose (250µg/ml) of *Nigella sativa* seed extract stimulated at 24 and 48hr, un-stimulated PBMCs as control

#### Conclusion

The *Nigella sativa* methanolic extract induced the expression of iNOS transcripts and Nitric oxide production in the chicken PBMCs. The results of the present study also indicated the ability of the *Nigella sativa* methanolic extract to enhance macrophage functions that could regulate adaptive immunity and control infectious diseases. Hence, the *Nigella sativa* methanolic extract can be tried as

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an adjuvant with any vaccine or antigen in the chicken.

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synergize with CD40L to induce either proliferation or plasma cell differentiation of mouse B cells. *PLoS One*, *6*(10), e25542.

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