Supplementing broiler diets with bacterial selenium nanoparticles enhancing performance, carcass traits, blood indices, antioxidant status, and caecal microbiota of *Eimeria tenella*-infected broiler chickens

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ABSTRACT Nanomedicine is a critical therapeutic approach for treating most poultry illnesses, particularly parasitic infections. Coccidiosis is a severe protozoan infection affecting poultry; the emergence of drug-resistant *Eimeria* strains demands the development of new, safe therapies. Consequently, the objective of this work was to investigate the efficacy of the biosynthesized selenium nanoparticles (SeNPs) by *Paenibacillus polymyxa* (P. polymyxa) against Eimeria tenella (E. tenella) experimental infection in broiler chickens. The prepared SeNPs absorbed the UV at 270 nm were spherical with a size of 26 nm, and had a surface negative charge of -25 mV. One hundred and fifty, 1-day-old male broiler chicks were randomly allocated into 5 groups (30 birds/group with triplicates each) as follows: T1: negative control (noninfected and nontreated with SeNPs); T2: delivered SeNPs (500 μ g/kg diet) for 35 successive days, T3: E. tenella-infected (positive control birds), T4: E. *tenella*-infected and treated with SeNPs (500 $\mu g/kg$ diet) and T5: E. tenella-infected chicks and treated with anticoccidial agent (sulfadimidine, 16% solution 8 mL/Lof drinking water) for 5 successive days. At 14 d of age,

each bird in infected groups was orally treated with 3×10^3 sporulated oocvst of *E. tenella*. SeNPs considerably decreased the number of oocysts in broiler feces compared to positive control and anticoccidial drug, followed by a substantial reduction of parasite phase count in the cecum (15, 10, and 8 for meronts, gamonts, and developing oocysts) when compared with positive control birds. The *Eimeria* experimental infection lowered the activity of antioxidant enzymes, superoxide dismutase (SOD), glutathione peroxidase (GPx), and reduced glutathione (**GSH**) while increasing the stress parameters nitric oxide (**NO**) and malonaldehvde (MDA). Moreover, the production of proinflammatory (TNF- α and IL-6) and apoptotic genes (BcL2 and Cas-3) were significantly elevated. Administrating SeNPs to chicks significantly decreased oxidative stress, inflammation, and apoptotic markers in the cecum tissue. Therefore, growth performance, carcass weights, antioxidant enzymes, and blood properties of infected chicks were enhanced. The findings compared the protecting role of Se-nanoparticles against cecum damages in E. tenellainfected broilers.

Key words: anticoccidial, bacteriogenic selenium nanoparticle, broiler, Eimeria, immunity

INTRODUCTION

The poultry sector is a major economic driver worldwide, as the demand for safe and high-protein poultry products continues to grow in tandem with the world's rapidly expanding population but faces many adverse problems (Swelum et al., 2021; Abd El-Hack et al., 2022). Coccidiosis is a significant expense for broiler chicks, ranging between \$3.34 and \$3.89 per m² of broiler house if not managed (Gilbert et al., 2020; El-Maddawy et al., 2022).

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Coccidiosis induces severe depression in birds and is characterized clinically by a variable degree of diarrhea, which may reach hemorrhagic type; it reflects poorly on birds' productivity as it induces destruction in the intestinal villi and interferes with nutrients absorption and metabolism (El-Shall et al., 2022). Since 1948, different chemical anticoccidial drugs have been applied to treat coccidial infections, including ionophores antibiotics in the 1970s (Chapman, 2018).

Due to haphazard treatment resistance, coccidial endemics persist despite this (El-Shall et al., 2022). Inappropriately, no new drugs have been licensed for usage, which has heightened the desire for innovative anticoccidial options derived from nature (Noack et al., 2019).

Recently, nanotechnology has been applied in the veterinary sector for disease diagnosis, developing medicines

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and preventatives, and nanoparticle manufacturing has increased in tandem with demand therapies (El-Maddawy et al., 2022). Chemical approaches include (N, N-dimethyl formamide, sodium dodecyl sulfate, sodium borohydride, and sodium citrate) and physical (cluster beam condensation or laser ablation, sputter deposition, and thin films), they are employed for nanoparticle production (El-Ashry et al., 2022). These techniques are costly and environmentally hostile (Saad et al., 2022).

Therefore, the synthesis of green nanoparticles using natural stabilizers (plants, bacteria, macroalgae, yeast, and fungus) is highly promising; adding a biological reducing agent makes the product cheaper to produce, more accessible, and less hazardous (El-Saadony et al., 2019, 2021a; Abdel-Moneim et al., 2022). Both conventional and biologically synthesized nanoparticles are stable (El-Saadony et al., 2021b). These characteristics feature antibacterial action (Abdel-Moneim et al., 2022). This is especially crucial in light of the rising multiresistance of pathogenic and conditionally harmful bacteria, such as *E. tenella, Pseudomonas aeruginosa*, and *Escherichia coli*, which have developed a high level of antibiotic resistance (Han et al., 2021; Salem et al., 2022b).

The nanoparticles are advantageous for biological applications due to their diminutive size and distinct physicochemical characteristics; they provide regulated drug administration, release, and in vivo immunomodulation (Yousry et al., 2020). Elemental nanoparticles possess antimicrobial, antiparasitic, and free radicals scavenging activities (Bisht et al., 2022; Arafa et al., 2023).

New antiparasitic agents include nanotechnological methods and other fresh developments (Sarhan et al., 2022). Nanoparticles are interesting medicines for treating several parasitic disorders due to their unique features, different from those of solitary atoms and bulk material (Carvalho et al., 2020; Attia et al., 2022). Selenium nanoparticles were produced in this study utilizing *P. polymyxa* MS 411; *P. polymyxa* is a G+ spore-forming rhizobacterium that generates EPSs, enzymes, plant hormones, and antibiotics (Rybakova et al., 2016).

For several significant *P. polymyxa* strains, whole genome sequences have been reported that code for producing various physiologically active chemicals (Liu et al., 2017) with very diversified physiological and biotechnological roles and little toxicity (Liang and Wang, 2015). Nanoselenium provides several health advantages, such as anticoccidial, antioxidant, and antiinflammation activities (Prokisch and Zommara, 2011; Alkhudhayri et al., 2020). In addition, SeNPs have been found to have potential activities against various parasites such as murine schistosomiasis, giardiasis, and murine trichinosis (Dkhil et al., 2016; Malekifard et al., 2020; Sarhan et al., 2022).

This study's purpose was to assess the protective impact of biosynthesized SeNPs using *P. polymyxa* MS 411 against experimental infection with *E. tenella* in broiler chickens and then to examine the impact of bacterial SeNPs supplementation on the performance, carcass, and blood properties of infected chicks.

Isolation and Identification of SeNPs-Producing Bacteria; Fabrication and Characterization of Bacterial Se Nanoparticles

MATERIALS AND METHODS

Paenibacillus polymyxa MS 411 strain was used to fabricate green selenium nanoparticles (SeNPs). This strain was isolated from corn soil rhizosphere. The soil samples were collected, mixed, and transferred to the microbiology laboratory. Serial dilutions from soil samples were prepared as follows: 10 g soil was homogenized in 90 mL peptone water to obtain 10^{-1} dilution, 1 mL from the previous dilution was added to 9 mL of peptone water to get 10^{-2} dilution, and serial dilutions up to 10^{-7} . 100 μ L of each dilution was inoculated on Luria-Bertani (LB) agar plates supplemented with 1, 2, 3, 4, and 5 mM sodium selenite (Na₂SeO₃) in sterilized Petri plates (90-mm diameter) and then incubated at 30°C for 2 d. The red colonies were chosen according to their ability to reduce sodium selenite (Saad et al., 2022).

Hundred microliters of the MS 411 isolate was inoculated in 100 mL of LB broth and incubated in a shaking incubator (220 rpm, 30°C) for 2 d. The mixture was centrifuged at 6,800 $\times q$ for 25 min; The supernatant was discarded and the bacterial pellets were collected then homogenized in 100 mL of the enrichment medium (\mathbf{EM}) broth containing 4 mM sodium selenite. The mixture was then incubated in shaking incubator (30°C, 220 rpm) for 2 d. After the incubation time, the change in color of the flask (EM medium containing sodium selenite and the MS 411 isolate) from brilliant yellow to red demonstrates the reduction of sodium selenite to SeNPs by MS 411 isolate. The bacterial SeNPs were extracted from the MS 411 isolates pellets through autoclaving until the explosion of the bacterial membrane; the resulting solution containing bacterial SeNPs was centrifuged at $12,000 \times g$ for 20 min, the supernatant was subsequently obtained, and the exploded cells were discarded. The selenium-producing bacterium was initially identified by morphological and biochemical characteristics, which were then compared to the reference standards in the Bergey's manual. Further identification was conducted by MALDI-TOF spectroscopy: the selected isolate *Paenibacillus polymuxa* MS411 was identified as *Paenibacillus polymyxa* DSM 365 (El-Saadony et al., 2021b; Saad et al., 2022; Alowaiesh et al., 2023). Transmission electron microscopy (**TEM**) studies shape, and aggregation. Dynamic light scattering (**DLS**) determine the particle size distribution, and zeta potential was used to determine surface charge stability (Abdel-Moneim et al., 2022; Alowaiesh et al., 2023).

Eimeria spp., Isolation, and Identification

Oocysts utilized in this investigation were isolated from the fresh feces of naturally infected chickens; the identification of the species was based on the morphological descriptions as mentioned by Thienpont et al. (1986); the oocysts were isolated and concentrated by flotation in a salt-saturated solution. The acquired suspension of unsprouted oocysts was combined with 2.5%with an equal amount of potassium dichromate to produce the stock oocyst suspension (**OS**) (Wiedosari and Wardhana, 2018) that incubated at 27°C for 3 d. Then, the sporulated oocysts were orally inoculated in 5 healthy chickens (2-wk old) to propagate the oocysts, and at 7-days postinfection, the birds were ethically euthanized. Cecal contents were collected, then concentrated, and oocysts were sporulated and kept in a refrigerator until use (Davies et al., 1963). The McMaster technique determined the oocysts count in (mL). At 14 d of age, each bird was orally treated with 3×10^3 sporulated oocyst of E. tenella using oral gavage (Habibi et al., 2016). The shape of sporulated oocysts was used to identify isolated *Eimeria* species (Figure 1).

In Vitro Assessment of the Anticoccidial Activity of SeNPs Against E. Tenella

The anticoccidial activity of SeNPs at concentrations (100, 300, 500, 700, and 900 μ g) was in vitro evaluated compared with anticoccidial drugs (**ACD**, sulfadimidine, 16%). All treatments were tested with triplicates, each against *E. tenella* oocytes, by adding 100 μ L of tested nanoparticles to 5 mL of OS (unsporulated oocyst), then incubated at 27°C for 3 d and calculating the number of sporulated oocysts by The McMaster method and capturing microphotographs with an Olympus camera (Tokyo, Japan) (Salem et al., 2022a).

Experimental Design

One hundred and fifty, 1-day-old Cobb male broiler chicks were grown for 5 wk. The birds were reared in separate floor pens with straw bedding on the concrete floor in adjusted environmental conditions with suitable lighting programs, temperature, and humidity according to the birds' requirements. The illumination system was 23 h light:1 h dark for the first 3 d, then 20 h light:4 h dark

 Table 1. Experimental groups of infected and noninfected broiler.

Groups	Treatments
T1	Non infected chicks
T2	Non infected chicks + SeNPs 500 μ g/kg diet for 35 d
T3	E. tenella-infected chicks and non treated
T4	$E.~tenella\text{-infected}$ chicks $+$ treated with SeNPs 500 $\mu\mathrm{g/kg}$ diet for 5 d
T5	$E.\ tenella-infected\ chicks + treated\ with\ sulfadimidine, 16\%$ solution 8 mL/L of drinking water for 5 d

until the end of the experiment. Random assignment of broiler chicks to 5 treatment groups (30 birds/group), each with triplicates (10 birds per replicate), as shown in Table 1. In group 1 (T1), birds were fed commercial balanced diets [starter (1–14-days old), grower (15–28-days old), and finisher (29-35-days old)] without any anticoccidial or anticlostridial drugs and supplied with clean, clear water ad libitum. The temp was set at 35°C at the beginning of bird rearing, then decreased by 2°C to 5°C every week to adjust as 22°C at the end of the fourth week; the temperature was then maintained at 22°C for the fifth week. Based on a preliminary experiment, group 2 (T2) was continuously supplemented with SeNPs (500 μ g/kg diet) for 35 d. Group 3 (T3) E. tenellainfected chicks, Group 4 (T4), E. tenella-infected chicks and then treated with SeNPs (500 μ g/kg diet) based on in vitro study. Group 5 (T5), E. tenella-infected chicks and then treated with the anticoccidial agent (sulfadimidine, 16% solution 8 mL/L of drinking water) (Hunduma and Kebede, 2016). After 4 d from experimental infection, treatments in T4 and T5 were added (continuous 24 h) in feed and water for 5 successive days.

Fecal samples were collected from all experimental groups at 4-, 7-, 10-, and 14-days postinfection to estimate oocyst shedding using the McMaster technique to determine the oocyst count per gram of freshly voided droppings.

The experimental room was always accessible under the same environmental circumstances. At specified intervals, birds were vaccinated against common viruses



Figure 1. Morphology of *E. tenella* oocyst: (A) Unsporulated oocyst; (B) sporulated oocyst.

[avian influenza (AI), Newcastle disease (ND), infectious bronchitis (IB), infectious bursal disease (IBD)].

Growth Performance Each day, the mortality rate of the chicks was evaluated. On the first day, they were weighed separately and then weekly. The chickens' weight increase (g), feed consumption, and feed conversion ratio (**FCR**) were then measured.

FCR = Total feed consumed / Total weight of product produced

where: Total weight of product = final weight - starting weight.

Carcass Traits At the end of the experiment, 10 chicks were randomly selected from each treatment and ethically slaughtered to assess the hot carcass weight, gizzard, heart, liver, and visceral fat.

Biochemistry of Blood At the time of slaughter, blood samples were collected from 10 birds per group, centrifuged at $4,500 \times g$ for 20 min, and the plasma was stored at 20°C. Using commercial kits provided by the biodiagnostic company, El-Tahrer St., Dokki, Giza, Egypt, total protein (CAT. NO. TB2020), albumin (CAT. No. AB 10 10), AST, ALT (CAT. NO. AP 1020), GSH, MDA (CAT. No. MD 25 29), SOD, GPx, and NO (CAT. No. CA 25 17) were evaluated calorimetrically according to the instructions provided by the manufacturer (Alagawany et al., 2022).

Blood Hematology At the end of the experiment, blood samples were collected from 10 birds per group; red blood cells (**RBCs**), platelet cells (**PBCs**), and white blood cells (**WBCs**) were estimated immediately after blood was drawn and heparinized (Alagawany et al., 2021). On clean microscope slides, thin blood smears were created from each sample. They were airdried before the slides were stained with a modified form of Wright's stain. Under 100 lens power, 100 cells were tallied (Alagawany et al., 2021).

Gut Microbiota

Intestinal content from each treatment's cecum was collected separately in sterile glass flasks after slaughter. Digesta were evacuated and mixed. Flasks were kept at 4° C till the determination of microbial counts. Ten-fold serial dilutions of up to 10^{-7} of each sample were prepared. Total bacterial counts, yeasts and molds count, Escherichia coli, and lactobacilli count were estimated. A nutrient agar medium was used to enumerate aerobic bacteria (Amadi and Wami, 2023). Sabouraud dextrose agar (SDA) was used to count yeasts and molds (Alagawany et al., 2021). The eosin methylene blue (EMB) agar medium (Oxoid) was used for *Escherichia coli* counts. The deMan, Rogosaand Sharpe (MRS) agar medium was used for lactobacilli count. After incubation, colonies were counted. Numbers of colony-forming units (CFU) are expressed as log CFU per gram of cecal content.

RNA Isolation

Ten days after the experimental infection, 2 birds per replicate were ethically slaughtered, and then cecal tissues

Table 2. Primers used for the gene encoding the RNAs for some selected proteins (Salem et al., 2022).

Gene	Primer sequence $(5' \rightarrow 3')$
B cell leukemia/lymphoma 2 (Bcl-2) Caspase-3 (Casp3)	F AGCATGCGACCTCTGTTTGA R GCCACACGTTTCTTGGCAAT F GGGGAGCTTGGAACGCTAAG R CCACTGACTTGCTCCCATGT
Tumor necrosis factor- alpha (TNF- α)	<i>F</i> ACCCTCACACTCACAAACCA <i>R</i> ACCCTGAGCCATAATCCCCT
Interleukin 6 (IL-6) β -Actin (ACTB)	FCTGCAAGAGACTTCCATCCAG $R AGTGGTATAGACAGGTCTGTTGG$ $FCACCATTGGCAATGAGCGGTTC$ $R AGGTCTTTGCGGATGTCCACGT$

were hygienically dissected. The samples were collected and homogenized for 30 s at 35,000 oscillations/min in a Mini Bead-beater 24 (Bio-Spec, Bartlesville, OK) and chilled on ice, then extracted using an RNA separation kit (Ambion, Applied Biosystems, Darmstadt, Germany). Later, total RNA was isolated from preserved cecal tissues using Trizol (Invitrogentham, MA). Following the manufacturer's recommendations, RNA samples were digested for at least 1 h with DNase (Applied Biosystems, Darmstadt, Germany) before being transcribed into cDNA using the reverse transcription kit (Qiagen, Hilden, Germany). The QuantiTectTM SYBR Green PCR kit (Qiagen) performing the real-time PCR according to the manufacturer's instructions. The TaqMan 7500 system software quantifying the amplification data. Primers were used to target the genes encoding the mRNAs for caspase-3 (Casp3), B cell leukemia/lymphoma 2 (BcL2), interleukin-6 (IL-6), and tumor necrosis factor-alpha (**TNF-** α). The results are shown in Table 2. The PCR techniques were identical to those previously detailed by Dkhil et al. (2013). The Ct ($\Delta\Delta$ CT) approach outlined by Livak and Schmittgen (2001) measured the variances between the mean gene expression and the reference gene. The qRT-PCR was done 3 times on each sample. In a 40-cycle PCR amplification, the cycling conditions were denaturation at 94°C for 30 s, annealing at 60°C for 30 s, and extension at 72°C for 45 s.

Histopathological Studies

At 10-days postexperimental infection, 2 birds were ethically slaughtered per replicate, and then cecal tissues were collected. According to Adam and Caihak (1964), paraffin cecal tissue slices $5-\mu m$ thick were produced for staining with H&E for further histopathological examination. We counted meronts, gamonts, and developing oocysts in infected and infected-treated groups to distinguish between the various parasite stages in the cecum of birds (Yang et al., 2022). The values were represented in units of 10 villous crypts (VCU).

Statistical Analysis

Prior to conducting a 1-way ANOVA, pretests were administered. The assumption of normality for sample distributions were examined and found P values of 0.0001 and Levene test for homogeneity revealed P value of 0.015. The means of the triplicate data were evaluated for significant differences using 1-way ANOVA with a 95% level of confidence (Wallenstein et al., 1980), using SAS software (version 9.4, Cary, NC, 2020). The sample size was calculated from the following equation

$$n = \left(\frac{ZSD}{E}\right)^2$$

Means were compared with the least significant difference (LSD) as a post hoc test at a probability level of 5%.

RESULTS

SeNPs Characterization

The *Paenibacillus polymyxa* successfully converted the sodium selenite to selenium nanoparticles, indicating color conversion from colorless to red. The bacterial SeNPs absorbed the UV at 270 nm (Figure 2A); their shape was spherical (Figure 2B) with an average diameter of 26 nm measured by zeta sizer and a negative charge of -25 mV (Figure 2C and D) by zeta potential analysis.

Output of Developing Oocysts Under Selenium Nanoparticles Treatment

Figure 3 shows the considerable anticoccidial activity of SeNPs at concentrations (100, 300, 500, 700, and 900 μ g/mL) compared to anticoccidial drugs (**ACD**, sulfadimidine, 16%). The results showed that SeNPs reduced the *Eimeria* oocyst count by 70 to 97% based on the concentrations; the count of sporulated oocytes in control was 5,000,000; at the best SeN concentration of 500 μ g/ mL, it was reduced to 150,000 as a 97% reduction. The best concentration was 500 μ g/mL head-to-head with the anticoccidial drug; therefore, this concentration was during the in vivo experiment.

T4 and T5 revealed a significant reduction in oocysts shedding number per gram droppings in all checking points (4-, 7-, 10-, 14-days postinfection) in contrast with T3.

The life cycle of *E. tenella* was developed from meronts and gamonts to developing oocysts in the cecal epithelial cells of infected broiler chickens (Figure 4). In the infected broiler's cecal villi, these phases were recorded per 10 VCUs as 97 meronts, 25 gamonts, and 15 developing oocysts (Figure 4). Remarkably, these numbers were significantly reduced by 85, 60, and 50% in the SeNPstreated broiler (15, 10, and 8 for meronts, gamonts, and developing oocysts) compared to the anticoccidial drug



Figure 2. Physiochemical characterization of SeNPs produced by *P. polymyxa* MS 411. (A) color conversion measured by UV at 270, (B) the shape of nanoparticles by TEM, (C) the size of nanoparticles by zeta sizer, (D) the charge of nanoparticles by zeta potential.



Figure 3. Preliminary in vitro experiment for determining the best selenium nanoparticles concentration (SeN) against *E. tenella* showing that SeNPs with a concentration of $(500 \ \mu g/mL)$ as well as ACD (sulfadimidine as anticoccidial drug) significantly reduced *E. tenella* oocysts sporulation. The values are presented as mean \pm SE, lowercase letters (a–e) above columns indicate the significant differences between different treatments on the sporulated oocyst count using LSD test at P < 0.05.

(20, 11, and 9 for meronts, gamonts, and developing oocysts) the SeNPs is more effective (P < 0.001) than the anticoccidial drug.

Histological Studies on In Vivo Anticoccidial Effects of SeNPs 10 and 30-Days Postinfection

Figure 5I describes the treatment of 2 groups of infected birds with SeNPs (T4) and sulfadimidine (T5) 10-days postinfection. A) The cecum of a T1 chicken possesses typical intestinal crypts. B) Cecum of a T2 chicken with extensive coccidial infection of the intestinal crypts (black arrows). C) T4 (SeNPs 500 μ g/kg)

chicken cecum indicating restoration of normal intestinal crypts with few parasite stages (white arrow) and a considerable reduction in the number of afflicted crypts carrying macrogametes of the parasite (black arrow). D) The cecum of T5 chicken exhibits a considerable decrease in the macrogametes of *E. tenella* (black arrow).

Figure 5II depicts the treatment of 2 groups of birds with SeNPs (T4) and sulfadimidine (T5) 30-days postinfection. A) the cecum of a T1 chicken with regular duodenal crypts. B) Cecum of T2 chickens with evident coccidial schizont invasion of *E. tenella* within the intestinal crypts (arrows). C) Hyperplasia of the crypts lining the epithelial and invasion of inter mononuclear cells in the cecum of T4 (SeNPs) chickens (arrows), cecum of



Figure 4. The number of *E. tenella* developmental stages (meronts, gamonts, and oocysts) in the cecum of infested and treated groups. T3: *E. tenella*-infected (positive control birds), T4: *E. tenella*-infected and treated with SeNPs (500 μ g/kg diet), and T5: *E. tenella*-infected chicks and treated with the anticoccidial agent (sulfadimidine, 16% solution 8 mL/L of drinking water). The values are presented as mean \pm SE. Lowercase letters (a–e) above columns indicate the significant differences between different treatments on the count of oocyst stages using the LSD test at P < 0.05.



Figure 5. (I) Histopathology of cecal tissues of chicks infected with *E. tenella* 10-days postexperimental infection showing (A) normal cecal tissue (star), (B) large number of developmental stages (arrow) (schizonts) of *E. tenella* in lamina propria of villi, (C and D) reduced number of developmental stages (arrow) (schizonts) of *E. tenella* in lamina propria of villi in the treated groups. (II) Histopathology of cecal tissues of chicks infected with *E. tenella* 30-days postexperimental infection showing (A) the cecum of a T1 (control) chicken with typical intestinal crypts (H&E, 200). (B) Cecum of T2 *E. tenella*-infected chicken and non treated, chicken displaying observable invasion of coccidial schizonts of *E. tenella* inside the intestinal crypts (arrow), as well as infestation of the lamina propria with various coccidial stages of *E. tenella* that were mostly degenerated from the inside (arrow) and eosinophilic cells infiltration (arrowhead, H&E, 200, T3 (Non infected chicks + SeNPs) showed also typical intestinal crypts as control; therefore we didn't add the image in figure 5 as the 100% similarity with control). (C) cecum of T4 (*E. tenella*-infected chicken and treated with SeNPs), chicken cecal tissues displayed restoration of normal intestinal crypts (arrows, H&E, 200). Cecum of T5 (sulfadimidine) chicken displaying a substantial decrease of *E. tenella* reproducing schizonts, evident hypertrophy of the crypts lining epithelium, and modest interstitial mononuclear cell infiltration (H&E, 200).

T5 (sulfadimidine) chicken displaying a substantial decrease of *E. tenella* reproducing schizonts, apparent hypertrophy of the crypts lining epithelium, and modest interstitial mononuclear cell infiltration. This histological examination revealed the curative impact of SeNPs on *Eimeria* development and alterations in the chicken cecum.

Effect of SeNPs on RNA Expression

The gene expression of BcL-2 and caspase-3 was raised due to an infection with *E. tenella* (Figure 6A). After treatment with SeNPs, the expression of the cas-3 gene was drastically reduced from 3.5- to 2.0-fold and 2.8-fold in anticoccidial drug. BcL-2 expression was dysregulated from 7.2- to 3.2-fold, as seen in SeNPs-treated groups and 3.8-fold in anticoccidial drug-treated groups (Figure 6B). Due to *E. tenella* infection, the expression of proinflammatory genes, TNF- α and IL-6, was upregulated in Figure 6C and D. Compared to T1, the expression of these genes increased by around 8.3 and 8.5, respectively. The therapy with Se-nanoparticles (500 μ g/L) significantly decreased IL-6 and TNF- α by 2.2and 4-fold compared to the anticoccidial medication and the control.

Growth Performance

Table 3 shows the positive impact of SeNPs on the growth performance of noninfected (T2) and *E. tenella*-

infected-treated broilers (T4) compared to control-positive birds in T3 and sulfadimidine-treated birds in T5. The initial BW of chicks did not change substantially from that of the control group, nor was a significant difference across groups. The supplementation of SeNPs dramatically increased their body weight gain, whereas SeNPs supplemented group increased the BWG by 12%compared to the noninfected control; also, SeNPs had anticoccidial activity against E. tenella, where the BWG increased by 5% compared to the anticoccidial drug. Concerning the feed intake, the chicks supplemented with SeNPs had the lowest amount in noninfected groups, but in infected groups, the FI was reduced by 15% in SeNPs-treated groups. Similarly, the FCR was significantly improved in SeNPs-treated birds T4 and T2 followed by T5 when compared with T3 and even T1. Only 20% (6/30) cumulative mortalities were recorded in T3, and no mortalities have been reported in the rest of the experimental groups.

Carcass Traits

Table 4 shows a medium significant difference (P = 0.045) between chicks' groups in carcass weight. The lowest carcass weight was detected in T3 infected with *E. tenella* (75.9 mg), while the highest was in T2 (78.3 mg). Also, the other organs (liver, spleen, and heart) were maintained with SeNPs treatment; the highest weights (1.79, 0.085, and 0.56 mg) detected in SeNPs-treated groups when compared to

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Figure 6. Effect of SeNPs on the RNA expression of caspase-3 (A), BcL-2 (B), IL-6 (C), and TNF- α (D) in the cecum of *E. tenella*-infected broiler detected by RT-PCR analysis. The values are presented as mean \pm SE. Lowercase letters (a–e) above columns indicate the significant differences between different treatments on the proinflammatory markers using the LSD test at P < 0.05. T1: Non infected chicks; T2: Non infected chicks + SeNPs 500 mg/kg diet for 35 d; T3: E. tenella-infected chicks and non-treated; T4: E. tenella-infected chicks + treated with SeNPs 500 mg/kg; diet for 5 d; T5: E. tenella-infected chicks + treated with sulfadimidine, 16% solution 8 mL/L of drinking water for 5 d.

infected and noninfected control excluding spleen showed the highest weight in T3 may be contributed to *Eimeria* infection. The fat content was decreased in SeNPs, either infected or noninfected, recording the lowest values (1.20 and 1.18 mg) followed by T5; however, the breast weight was increased by SeNPs treatments with an 18 to 20% increase above controls.

Gut Microbiota

Figure 7 shows the gut microorganism's count. Total bacterial count, yeasts and molds, and *E. coli* show the highest values with the control and Eimeria-infected brolier group coefficient of 8.5, 4.9, and 7.76 Log10 CFU/g, respectively. It decreases with the augment in the dietary SeNPs level until it reaches the

Table 3. The impact of selenium nanoparticles on the growth properties of broiler chickens.

		Performance traits (g)					
Groups	IBW	FBW	BWG	FI	FCR		
T1	45.00 ± 0.0	$1960 \pm 2.1^{\rm b}$	$1915.00 \pm 2.0^{\rm b}$	$3200 \pm 1.1^{\rm b}$	$1.710 \pm 1.1^{\rm b}$		
T2	45.36 ± 0.1	$2180 \pm 1.9^{\rm a}$	$2134.64 \pm 3.0^{\rm a}$	$3150 \pm 1.3^{ m bc}$	$1.550 \pm 0.9^{\circ}$		
T3	45.10 ± 0.3	$1860 \pm 3.2^{\circ}$	$1814.91 \pm 1.5^{ m d}$	$3550 \pm 1.8^{\rm a}$	$1.840 \pm 0.7^{\rm a}$		
T4	45.22 ± 0.2	$1901 \pm 2.1^{ m bc}$	$1855.78 \pm 0.9^{\circ}$	$3050 \pm 1.5^{\circ}$	1.480 ± 0.6^{d}		
T5	45.00 ± 0.0	$1900 \pm 1.1^{\mathrm{bc}}$	$1855.12 \pm 1.1^{\rm c}$	$3080 \pm 2.0^{\rm c}$	$1.550 \pm 1.3^{\rm c}$		

The values are presented as mean \pm SE, lowercase letters (a–d) in each row indicate the significant differences between different treatments on the growth properties of broiler chickens using LSD test at P < 0.05. T1: Non infected chicks; T2: Non infected chicks + SeNPs 500 mg/kg diet for 35 d; T3: E. tenella-infected chicks and non-treated; T4: E. tenella-infected chicks + treated with SeNPs 500 mg/kg; diet for 5 d; T5: E. tenella-infected chicks + treated with sulfadimidine, 16% solution 8 mL/L of drinking water for 5 d.

Table 4. The impact of selenium nanoparticles on the carcass properties of broiler chickens.

Groups	Carcass (mg)	Liver (mg)	Spleen (g)	Heart (g)	Fat (g)	Breast muscle (g)
T1 T2 T3 T4 T5	$\begin{array}{c} 77.2 \pm 0.0^{\rm ab} \\ 78.3 \pm 0.2^{\rm a} \\ 75.9 \pm 0.2^{\rm c} \\ 76.9 \pm 0.6^{\rm b} \\ 76.2 \pm 0.3^{\rm b} \end{array}$	$\begin{array}{c} 1.78 \pm 0.1^{\rm ab} \\ 1.79 \pm 0.2^{\rm a} \\ 1.71 \pm 0.2^{\rm c} \\ 1.76 \pm 0.3^{\rm b} \\ 1.75 \pm 0.2^{\rm bc} \end{array}$	$\begin{array}{c} 0.081 \pm 0.01^{\rm c} \\ 0.085 \pm 0.02^{\rm b} \\ 0.089 \pm 0.01^{\rm a} \\ 0.079 \pm 0.03^{\rm d} \\ 0.078 \pm 0.06^{\rm d} \end{array}$	$\begin{array}{c} 0.55 \pm 0.02^{\rm ab} \\ 0.57 \pm 0.05^{\rm a} \\ 0.50 \pm 0.03^{\rm d} \\ 0.54 \pm 0.04^{\rm c} \\ 0.52 \pm 0.00^{\rm cd} \end{array}$	$\begin{array}{c} 1.35\pm0.1^{\rm a}\\ 1.20\pm0.2^{\rm cd}\\ 1.30\pm0.3^{\rm b}\\ 1.18\pm0.6^{\rm d}\\ 1.23\pm0.3^{\rm c}\end{array}$	$\begin{array}{c} 33.11 \pm 0.2^{\rm c} \\ 39.65 \pm 0.3^{\rm a} \\ 32.40 \pm 0.5^{\rm d} \\ 37.89 \pm 0.8^{\rm b} \\ 37.12 \pm 0.2^{\rm b} \end{array}$

The values are presented as mean \pm SE, lowercase letters (a–d) in each row indicate the significant differences between different treatments on the carcass properties of broiler chickens using LSD test at P < 0.05. T1: Non infected chicks; T2: Non infected chicks + SeNPs 500 mg/kg diet for 35 d; T3: E. tenella-infected chicks and non-treated; T4: E. tenella-infected chicks + treated with SeNPs 500 mg/kg; diet for 5 d; T5: E. tenella-infected chicks + treated with sulfadimidine, 16% solution 8 mL/L of drinking water for 5 d.



Figure 7. Effect of SeNPs on the gut microbiota in the cecum of *E. tenella*-infected broiler. The values are presented as mean \pm SE. Lowercase letters (a-g) above columns indicate the significant differences between different treatments on the gut microbiota using the LSD test at P < 0.05. Group 1 (T1) control, Group 2 (T2) E. tenella-infected chicks, Group 3 (T3) chicks supplemented with SeNPs (500 mg/kg diet) for 35 d. Group 4 (T4), E. tenella-infected chicks and then treated with SeNPs (500 mg/kg diet) based on *in vitro* study. Group 5 (T5), E. tenella-infected chicks and then treated with the anticoccidial agent (sulfadimidine, 16% solution 8 mL/L of drinking water).

lowest significant value with the treatment SeNPs 500 μ g/kg (6.6, 3.7, and 6.57 Log 10 CFU/g). As for lactic acid bacteria count, the lowest value appears with the control and Eimeria-infected brolier group 4 and 3.2, respectively; the values increase with the addition of bacterial SeNPs 500 μ g/kg in the diet to achieve the peak height with treatment SeNPs 500 μ g/kg (5.2 Log 10 CFU/g), consequently reduced the *Eimeria* count.

Biochemistry and Hematological Parameters

Table 5 shows Eimeria-infected broiler induced the ALT and AST enzymes (28.3 and 81.2) with a relative increase of 50 and 20% compared to the control. The

liver enzymes were downregulated to normal levels by SeNPs (500 μ g/kg diet) addition in the diet because they mitigate the *Eimeria* infection. Similarly, the activity of stress content (MDA and NO) increased by infection compared to control, while SeNPs controlled that. On the other hand, the antioxidant enzymes (GPx, GSH, and SOD) decreased by 50% in T3 (*Eimeria* infected) while supplementing the broiler diet with SeNPs upregulated the activity of these enzymes to normal levels, that is, 60, 22, and 4.6 with no significant difference compared to control.

Concerning the biochemistry of broiler blood, albumin and globulin contents were decreased by 30% in *E. tenella*-infected chicks compared with normal control;

Table 5. Effect of SeNPs on the biochemistry and hematology of broiler chickens.

Blood traits	T1	Τ2	Т3	Τ4	T5
Liver enzymes					
ALT	$17.1 \pm 0.2^{\circ}$	$9.3 \pm 0.6^{\mathrm{d}}$	$28.3\pm0.6^{\rm a}$	$17.9 \pm 0.3^{\circ}$	$20.3 \pm 0.6^{\rm b}$
AST	$59.2 \pm 0.3^{\circ}$	$38.5 \pm 0.1^{\rm d}$	$81.2 \pm 0.3^{\mathrm{a}}$	$60.2 \pm 0.1^{\circ}$	$68.1 \pm 0.5^{\rm b}$
Antioxidant enzymes					
GSH	$62.1 \pm 0.2^{\rm a}$	$60.2 \pm 0.3^{\rm b}$	$33.6 \pm 0.9^{\mathrm{d}}$	$60.9 \pm 0.1^{ m b}$	$57.5 \pm 0.2^{\circ}$
GPx	$29.8 \pm 0.1^{ m b}$	$32.1 \pm 0.5^{\rm a}$	$11.3 \pm 0.8^{\rm d}$	$22.6\pm0.5^{\rm c}$	$21.1 \pm 0.1^{\circ}$
SOD	$4.6 \pm 0.2^{\rm b}$	$4.8 \pm 0.1^{\rm a}$	$2.9 \pm 0.4^{\rm d}$	$4.6 \pm 0.4^{\mathrm{b}}$	$4.1 \pm 0.6^{\circ}$
Stress components					
MDA	$18.7\pm0.2^{ m d}$	$19.2 \pm 0.3^{\rm d}$	$45.9 \pm 0.2^{\rm a}$	$21.6 \pm 0.6^{\circ}$	$25.7 \pm 0.6^{\rm b}$
NO	$8.9\pm0.1^{ m d}$	$9.1 \pm 0.1^{\rm d}$	$22.5 \pm 0.5^{\rm a}$	$10.22 \pm 0.1^{\circ}$	$12.1 \pm 0.8^{\rm b}$
Serum biochemical					
Protein (g/dL)	$2.70 \pm 0.1^{\rm b}$	$2.9 \pm 0.9^{\mathrm{a}}$	$1.0 \pm 0.0^{\mathrm{e}}$	$2.5 \pm 0.1^{\circ}$	2.0 ± 0.1^{d}
Albumin (g/dL)	$1.36 \pm 0.2^{\rm b}$	$1.5 \pm 0.1^{\rm a}$	$1.1 \pm 0.01^{\rm d}$	$1.41 \pm 0.2^{\rm ab}$	1.32 ± 0.2^{c}
Globulin (g/dL)	$1.30 \pm 0.6^{\circ}$	$1.43 \pm 0.5^{\rm a}$	$0.95 \pm 0.02^{\rm d}$	$1.39 \pm 0.6^{\mathrm{b}}$	$1.31 \pm 0.3^{\circ}$
A/G ratio	$1.07\pm0.5^{ m b}$	$1.04 \pm 0.7^{\mathrm{b}}$	$1.15 \pm 0.01^{\rm a}$	$1.01 \pm 0.4^{ m c}$	$1.00 \pm 0.1^{\circ}$
Lipids (mg/dL)	$0.42 \pm 0.4^{\rm b}$	$0.35 \pm 0.5^{\circ}$	$0.58 \pm 0.05^{\rm a}$	$0.31 \pm 0.06^{\mathrm{d}}$	$0.34 \pm 0.04^{\circ}$
Hematological parameters					
RBCs $(10^6/\mu L)$	$2.4\pm0.2^{ m bc}$	$2.7 \pm 0.3^{\rm a}$	$1.9 \pm 0.2^{\rm d}$	$2.5 \pm 0.1^{\rm b}$	$2.3 \pm 0.2^{\circ}$
WBCs $10^2/\mu L$)	$138 \pm 0.8^{\mathrm{b}}$	130 ± 1.1^{c}	$145 \pm 1.2^{\mathrm{a}}$	131 ± 1.2^{c}	136 ± 1.2^{bc}
$\rm PLT~(10^{4}/\mu L)^{'}$	$39.2\pm0.8^{\rm b}$	$32.6\pm0.6^{\rm d}$	$48.1\pm0.2^{\rm a}$	$33.6\pm0.6^{\rm d}$	$36.6\pm0.2^{\rm c}$

The values are presented as mean \pm SE, lowercase letters (a–e) in each row indicate the significant differences between different treatments on the biochemistry and hematology of broiler chickens using LSD test at P < 0.05. T1: Non infected chicks; T2: Non infected chicks + SeNPs 500 mg/kg diet for 35 d; T3: E. tenella-infected chicks and non-treated; T4: E. tenella-infected chicks + treated with SeNPs 500 mg/kg; diet for 5 d; T5: E. tenella-infected chicks + treated with sulfadimidine, 16% solution 8 mL/L of drinking water for 5 d.

the addition of SeNPs increased the protein content by 1.5-fold than T3, while T4 and T5 recorded the lowest values in lipid profile. During *Eimeria* infection, the WBCs and PLT counts significantly increased while RBCs decreased while SeNPs corrected this content to control levels.

DISCUSSION

Eimeria infection symptoms include malnutrition, enteritis, and, in severe cases, mortality for some *Eimeria* species, endangering the health of birds and their economic production (Youssefi et al., 2023). Food security and the global agroeconomy will face significant challenges due to this pathogenic protozoan and its detrimental impact on production due to the ever-increasing human need for protein resources from poultry (Teng et al., 2023). Since the 1940s, over 30 sulfanilamide medications have been used to treat and control coccidiosis (Ekinci et al., 2023). These anticoccidial treatments affect *Eimeria* species' cofactor production, mitochondrial activity, or cell membrane function (Peek, 2010). Commercial pharmaceuticals need to develop novel anti-Eimeria therapies because of concerns about health and the environment besides parasite resistance (Wunderlich et al., 2014). In this study, we used sulfadimidine as an anticoccidial reference, and many studies have proven its efficiency (Rabo et al., 2021). This study used experimentally infected cockerels to determine the efficacy of amprolium, septrin, and sulfadimidine in controlling *Eimeria necatrix* infection. The infection was characterized by weight loss, anemia, dullness, bloody diarrhea, and death. Of the 3 drugs, septrin was found to be most effective in treating the disease, followed by sulfadimidine and amprolium, as evidenced by the amelioration of clinical signs, gross and histopathological lesions, and relative survival rates of treated birds. Also, Kuraa et al. (2021) confirmed that in vitro sporulation inhibition of garlic extract showed significant efficacy on E. magna oocysts in comparison with black seed extract and high significant efficacy of sporulation inhibition, compared to sulfadimidine and it referred to that sulfadimidine has both in vitro and in vivo an inhibitory effect against sporulation.

An experimental study was conducted to assess the efficacy of amprolium and sulfadimidine drugs in treating chickens' coccidiosis. A total of 52 chickens were purposively selected for this study. Among them, 17 chickens were used for the control group, while 35 were used for the treatment group. Of the 35 chickens used for the treatment group, 18 were treated with amprolium, and the remaining 17 were treated with sulfadimidine. The drugs were administered orally in drinking water.

The treatment response of these drugs for coccidiosis concerning the reduction in fecal scores (OPG count); the study showed that there was no statistically significant difference in OPG count between before and after treatment in amprolium-treated groups (P > 0.05), but in the sulfadimidine administered group, there was a statistically significant difference (P = 0.004). This reveals that sulfadimidine is more effective than amprolium in treating coccidial infection (Hunduma and Kebede, 2016). Furthermore, in the study of Khan et al. (2021) they found that sulfadimidine was most effective (45%) followed by amprolium (44.6%), while triquen (24.0%) showed less effectiveness against coccidiosis in pigeons.

Nanoparticles enable the development of innovative antiparasitic medicinal medicines (Alkhudhayri et al., 2020). For example, nanoselenium (SeNPs) has been used to treat different parasites (Dkhil et al., 2016; Alkhudhayri et al., 2020; Malekifard et al., 2020).

In addition, selenium at the nanoscale provides several health advantages as SeNPs possess antioxidant and anti-inflammatory effects, according to Rayman (2008). Also, antimicrobial, antiviral, and antiparasitic activities have been established (Abou Elmaaty et al., 2022; Salem et al., 2022a). SeNPs also showed anticoccidial properties (Alkhudhayri et al., 2018). This work analyses the possible anticoccidial activity of SeNPs derived from *P. polymyxa*, which is likely attributable to the composition of the bacterial pellets and the unique features of SeNPs.

Bio-SeNPs effectively lowered the number of oocysts shed by chicks by about 97.21% on the 10th day following infection. Most anticoccidial treatments are known to decrease intracellular *Eimeria* stages and reduce the oocyst shedding rate effect; some studies confirmed our results on the effect of SeNPs on *Eimeria* but were conducted on mice (Alkhudhayri et al., 2020), they found that the numbers of meronts, gamonts, and developing oocysts of *E. papillata* reduced after the infected mice were treated with the SeNPs.

Anticoccidial activities may be attributable to the high quantities of active components detected in bacterial supernatants (Oke et al., 2020). These chemicals have a significant antibacterial action by competing with the bacterial membranes, affecting their porosity for water and cations, leading to diminished function, cellular component leakage, and cell demise (Filipović et al., 2021).

Infection with *E. tenella* is followed by the inflammation of cecum cells and oxidative damage in chickens; while permitting an increase in the total count of bacteria, *E. coli*, total yeast, and molds, and reducing the lactic acid bacteria, these effects were reversed when added dietary selenium agreeing with Al-Quwaie (2023), who found that the addition of selenium nanoparticles (**BSeNPs**) and elemental selenium (**Se**) significantly reduced the levels of total bacterial count (**TBC**), total yeast mold count (**TYMC**), *E. coli*, and *Salmonella* in comparison to the control group. However, the levels of lactic acid bacteria (**LAB**) were significantly increased.

The treatment of infected animals with SeNPs significantly mitigated the oxidative injury of the cecal tissue. The powerful potential impact of SeNPs is attributable to their distinct properties, such as the antioxidant antiinflammation activity of SeNPs and stabilizing agent (Nabi et al., 2020; Qudoos et al., 2020; Abdel-Moneim et al., 2022). The coccidial infection disturbs the work of the antioxidant defense system against free radical generation (Murshed et al., 2023). Our results demonstrated that infection with *E. tenella* is linked to oxidative stress in broiler chickens' cecum, reducing the levels of GSH, GPx, and SOD enzymes, and these antioxidant defense systems are essential for preventing free radical damage to the animal body during an *Eimeria* infection. SeNPs increase these enzymes' activity, which is generally lowered during infection-induced oxidative damage (Chen et al., 2021).

Lipid oxidation leads to the formation of several carbonyl compounds, including MDA (Michalak et al., 2022). Following The SeNPs treatment, the MDA levels drop. The NO levels rose in response to pathogenic sporozoite phases that penetrated and inflamed cecal cells. In addition, the imbalance caused an infection-related rise in nitric oxide levels (Dominguez et al., 2015). Bio-SeNPs might significantly lessen the incidence of cecal infection by boosting NO levels. This demonstrates the antioxidant potential of SeNPs. According to previous research, SeNPs and plant extracts include necessary active chemicals (Seriana et al., 2021). Bacterial SeNPs and the stabilizing agent may indicate their biological activity in this work.

This study indicated how E. tenella infection reduced the expression of the MUC2 gene, as per Ciszewski et al. (2022).

A significant increase in IL-6 and TNF- α was indicative of the inflammation response to the *E. tenella* infection. Mucin synthesis is controlled by proinflammatory cytokines such as IL-6 and TNF- α (Parrish et al., 2022). In contrast to the current work, *Eimeria* parasite infection-induced cytokine influx and mucin glycosylation downregulation. These adverse effects are linked to the reduction of goblet cells and mucus thickness and accelerated transport of mucin from the Golgi to the secretory vesicles (Liu et al., 2022). The SeNPs treatment significantly decreased the inflammation response in infected chicks.

Additionally, to face the *Eimeria* stages within infected tissues, our research indicated that SeNPs act as an antiinflammation to protect host tissues. This anti-inflammatory action may be caused by active compounds in SeNPs and their precursors (Abdel-Moneim et al., 2022). In addition to immunomodulatory and anti-inflammatory properties, it can prevent the production of proinflammatory cytokines (Attia et al., 2020; Nagata et al., 2023), according to previous studies. Apoptosis may influence the chicks' response to particular intracellular parasite infections and promote the removal of damaged or contaminated cells (Al-Quraishy et al., 2019). The high levels of proapoptotic genes BCL2 and caspase-3 indicate the damage to cecum cells. Previous research revealed increased apoptotic cells in the jejunum of E. papillomainfected mice (Dkhil et al., 2016). Bio-SeNPs may ameliorate the meiosis-induced apoptotic changes in jejunal cells, as reported by many studies (Chen et al., 2022).

According to the study, SeNPs enhanced the development performance of chickens. Nano-Se improved the growth performance of treated hens, suggesting that it is more bioavailable and less harmful than inorganic forms of Se (Ibrahim et al., 2019). Our results contradict those of Yoon et al. (2007), who indicated that the Se form in the chicks' food did not affect their development performance. In contrast, the performance of chicks supplemented with 200 g/kg of organic Se or nano-Se was comparable to that of chicks given the same quantity of Se as selenite (Couloigner et al., 2015). Selenium is required for the production of hormones and also to enhance the healthy development of avian creatures (Wang et al., 2011) and has been shown to affect bird growth performance.

Using SeNPs in this study improved bird antioxidant status. Also, Visha et al. (2017) observed comparable results in birds fed nano- and organic Se; their blood and tissues showed better antioxidant capacity than those fed inorganic selenium by modulating the scavenging abilities of seleno-enzymes. Se plays a crucial role in protecting cells from oxidative damage (Sarkar et al., 2011).

In addition to serum oxidants and selenium retention in vivo, nano-Se can neutralize free radicals by boosting the activity of selenium-based enzymes and promoting growth (Sarkar et al., 2011). Unlike other selenium forms, nano-Se exhibited decreased acute toxicity but elevated selenoenzyme activity. GPXs and thioredoxin reductase are mostly responsible for the antioxidant activity of nano-Se (TR). Multiple peroxides, including H₂O₂, phospholipid hydroperoxide, fatty acid hydroperoxides, and thymine hydroperoxyl groups, can be eliminated by the GPXs (Kondaparthi et al., 2019). The study's outcome was that the inclusion of nano-Se in chickens' food can effectively increase their antioxidant potential, consequently enhancing their ability to detoxify a wide variety of peroxides, improving bird antioxidant status, performance, and blood parameters and reducing the negative impact of *E. tenella* experimental infection.

CONCLUSIONS

Remarkably, a 500 μ g/kg diet of bacteriogenic SeNPs revealed both in vitro and in vivo anticoccidial action against *E. tenella*. The dietary supplementation of 500 μ g/kg bacteriogenic SeNPs improved growth performance, gut health, carcass traits, biochemistry, and hematological indices and exhibited significant antioxidant activity in the treated chickens.

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AUTHOR CONTRIBUTIONS

Conceptualization, M. N. A., and M. T. E.-S., formal Analysis, M. N. A., and M. T. E.-S., investigation, M. N. A., and M. T. E.-S., data curation, M. N. A., and M. T. E.-S., writing original draft preparation, M. N. A., and M. T. E.-S., writing final manuscript and editing, M. N. A., and M. T. E.-S., visualization and Methodology, M. N. A., and M. T. E.-S. All authors have read and agreed to the published version of the manuscript.

DISCLOSURES

The author declares that they have no conflicts of interest.

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