

## Effects of nano-selenium on mRNA expression of markers for spermatogonial stem cells in the testis of broiler breeder males

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### Abstract

Fertility is one of the most important parameters in breeder farms and cockerels play an outstanding role in the fertility of eggs in broiler breeder farms. Today, supplementation of chicken diet with additives such as organic selenium is used to increase fertility. The aim of this study was to evaluate the effects of different levels of nano-selenium (Nano-Se) on the expression of molecular markers of spermatogonial stem cells (SSCs) in the testis of broiler breeder males. A total of 30 roosters of 40 weeks of age were randomly divided into five groups. Groups were as follows: 1) control (normal diet) group, 2) diet supplemented with 0.30 mg kg<sup>-1</sup> sodium selenite, 3) diet supplemented with 0.15 mg kg<sup>-1</sup> Nano-Se, 4) diet supplemented with 0.30 mg kg<sup>-1</sup> Nano-Se, and 5) diet supplemented with 0.60 mg kg<sup>-1</sup> Nano-Se. At the end of the experimental period (5<sup>th</sup> week), birds were autopsied and samples from testis of all birds were collected. The testis samples were used to examine the  $\beta$ 1-integrin (CD29), thy-1 (CD90) and NANOG mRNA expression by real-time PCR. The results showed that testis of the groups fed with the diets supplemented with 0.60 mg kg<sup>-1</sup> and 0.15 mg kg<sup>-1</sup> of Nano-Se had the highest and lowest mRNA expression of SSCs markers, respectively. In conclusion, the present study indicated that Nano-Se had advantages over sodium selenite. Diet supplemented with 0.60 mg kg<sup>-1</sup> of Nano-Se may contribute to optimal fertility via increasing the mRNA expression of SSCs markers of roosters' testis and could be used to delay the reduction of fertility caused by aging in broiler breeder males.

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### آثار نانو سلنیوم بر بیان مارکرهای مولکولی سلول‌های بنیادی بیضه خروس‌های گله مادر گوشتی

#### چکیده

نطفه‌داری تخم‌مرغ‌ها یکی از فاکتورهای خیلی مهم در واحدهای مرغ مادر محسوب می‌شود و خروس‌ها نقش برجسته‌ای را در باروری تخم‌مرغ‌ها ایفا می‌کنند. امروزه به منظور افزایش باروری خروس‌ها از مکمل‌های افزودنی از قبیل ترکیبات سلنیوم استفاده می‌شود. با عنایت به محاسن ترکیبات نانو ذرات، هدف از این مطالعه ارزیابی اثرات سطوح مختلف مکمل نانو سلنیوم بر بیان مارکرهای مولکولی سلول‌های بنیادی اسپرماتوگونی (SSCs) بیضه خروس‌های گله مادر گوشتی بود. در مجموع ۳۰ قطعه خروس گله مادر گوشتی با سن ۴۰ هفته، بصورت تصادفی به پنج گروه تقسیم شدند. گروه‌ها شامل: ۱) کنترل (جیره نرمال)، ۲) جیره حاوی ۰/۳۰ میلی گرم سلنیت سدیم در کیلوگرم، ۳) جیره حاوی ۰/۱۵ میلی گرم نانو سلنیوم در کیلوگرم، ۴) جیره حاوی ۰/۳۰ میلی گرم نانو سلنیوم در کیلوگرم، ۵) جیره حاوی ۰/۶۰ میلی گرم نانو سلنیوم در کیلوگرم بودند. در پایان دوره آزمایش (هفته پنجم)، کالبدگشایی و نمونه برداری از بیضه تمام پرندوها انجام گردید. نمونه‌های بیضه با روش واکنش زنجیره ای پلیمرز در زمان واقعی از نظر بیان mRNA مارکرهای مولکولی سلول‌های بنیادی اسپرماتوگونی  $\beta$ 1-integrin (CD29) و thy-1 (CD90) و NANOG مورد آزمایش قرار گرفتند. نتایج نشان داد که بیشترین میزان بیان mRNA مارکرهای مولکولی سلول‌های بنیادی اسپرماتوگونی (SSCs) به ترتیب در گروه‌های تغذیه شده با جیره‌های حاوی ۰/۶۰ و ۰/۱۵ میلی گرم نانو سلنیوم در کیلوگرم بود. نتایج این تحقیق بیانگر آن است که مکمل نانو سلنیوم نسبت به سلنیت سدیم ارجحیت داشته و مصرف ۰/۶۰ میلی گرم نانو سلنیوم در کیلوگرم جیره با افزایش بیان mRNA مارکرهای مولکولی سلول‌های بنیادی اسپرماتوگونی (SSCs) می‌تواند به بهبود باروری خروس‌ها کمک نماید و کاهش باروری وابسته به سن خروس‌های مرغ مادر گوشتی را به تأخیر بیندازد.

واژه‌های کلیدی: خروس گله مادر گوشتی، سلول‌های بنیادی بیضه، مارکرهای مولکولی، نانو سلنیوم

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## Introduction

The final product of breeder farms is one-day-old chicks. The eggs hatchability is mostly influenced by the fertility of cockerels. Nowadays, various feed additives including organic selenium are used in the diet of broiler breeder males in order to increase fertility. Recent studies indicated that nano-particles of selenium may have some advantages over organic selenium during spermatogenesis. Four types of spermatogonial cells such as spermatogonia dark type A ( $A_d$ ), pale type A ( $Ap_1$ ,  $Ap_2$ ), and type B, have been described in birds and the type  $A_d$  spermatogonia have been considered as the spermatogonial stem cells (SSCs).<sup>1-3</sup> The SSCs are the early precursor for spermatozoa and are responsible for the continuation of spermatogenesis in adult males' testis. These cells have the capability to undergo self-renewal division as well as produce daughter cells destined for differentiation into spermatozoa and transferring genetic information from an individual to the offsprings.<sup>4-6</sup> During a routine spermatogenesis process, the SSCs infrequently self-renew, but they can divide frequently in response to harmful conditions such as damage due to chemical compounds.<sup>7</sup> Therefore, the performance of SSCs during spermatogenesis influences both the number of spermatogenic cells and semen parameters including sperm concentration.<sup>8</sup> Similar to the other tissue-specific stem cells, SSCs are rare in chicken. The proportions of SSCs to testes' germ cells in young chickens and adult males are approximately 0.40% and 0.03% respectively.<sup>9,10</sup>

The SSCs reside within a specialized micro-environment called 'niche' which regulates the behavior of the stem cells and its differentiating progeny. The SSCs performance (self-renewal and differentiation) is tightly regulated by a combination of intrinsic gene expression within the SSC and the extrinsic gene signals from the niche. Communication between the niche and SSCs is crucial to maintain proper spermatogenesis.<sup>7,11,12</sup>

Aging leads to an increase in reactive oxygen (ROS) and when the generation of ROS in a system exceeds, the occurrence of oxidative stress may cause a reduction in antioxidant enzymes in spermatogonial stem cells.<sup>13-19</sup> During oxidative stress, free radical production is increased causing damages to different organs/cells. In order to decrease harmful effects of free radicals on reproductive organs and subsequently to prevent the reduction of males' fertility, dietary supplementation of an antioxidant such as selenium compounds is necessary.<sup>20,21</sup> Selenium is a major structural component of many antioxidant enzymes and has an important role in a variety of biological processes including antioxidant defense system, fertility, immune function, and muscle metabolism.<sup>22-31</sup> Selenium protects spermatozoa against oxidative damage during spermatogenesis and in fact, the optimal level of Se is required for normal development of male reproductive tissue.<sup>8,32-35</sup>

Nowadays, nanotechnology proposed several new effective forms of dietary supplementation components due to their low toxicity and high bioavailability. Among those, nano-selenium (Nano-Se) particles, because of their excellent characteristics including great surface area, effective surface activity, lots of surface active centers, high catalytic efficiency, strong absorbing ability and low toxicity have many advantages in comparison to sodium selenite.<sup>36</sup> The purpose of this study was to determine the most optimal dose of supplemental Nano-Se on the mRNA expression of molecular markers of SSCs in the testis of broiler breeder males.

## Materials and Methods

**Ethics of experimentation.** All experimental procedures were carried out according to the standard animal experimentation protocols of the Veterinary Ethics Committee of Faculty of Veterinary Medicine, Urmia University (IR-UU-AEC-266/DP3).

**Experimental design.** Thirty broiler breeder males (Arbor Acres Plus strain) at the age of 40 weeks were used in this study. The birds were randomly divided into five groups (six birds per group) and housed in the pens of identical size in a deep litter system with wood shaving floor. Each group had three replicates (two birds per pen). The first week of the experiment was designed as an adaptation period and during this period birds of all the groups were fed with standard basal diet according to the catalog of Arbor Acres Plus.<sup>37</sup> After adaptation period, group 1 (control group) was fed with basal diet, group 2 was fed with basal diet supplemented with 0.30 mg kg<sup>-1</sup> sodium selenite, groups 3, 4 and 5 were fed with basal diet supplemented with 0.15 mg kg<sup>-1</sup>, 0.30 mg kg<sup>-1</sup>, 0.60 mg kg<sup>-1</sup> of Nano-Se, respectively. All birds fed with the described diets for four weeks after the adaptation period.<sup>38</sup>

**Selenium.** Sodium selenite (Merck, Darmstadt, Germany) and Nano-Se (American Elements, Los Angeles, USA) with the size of 10-45 nm and purity of 99.99% were purchased.

**Sample collections.** At the end of the 5<sup>th</sup> week, birds of all groups were humanely euthanatized by cervical dislocation.<sup>39</sup> Then testes were collected, rinsed with normal saline, frozen immediately in liquid nitrogen and stored at -80 °C to investigate the mRNA expression levels of SSCs markers  $\beta$ 1-integrin (CD29), thy-1 (CD90) and NANOG) in testicular tissue with Real-Time PCR.<sup>8</sup>

**Primer synthesis.** The specific primers described previously based on published *Gallus gallus* mRNA sequences of the target genes in Genbank were synthesized and used in the real-time RT-PCR. The primer sequences were as follows;  $\beta$ 1-integrin (CD29) specific primers amplifying a fragment of 288 bp in size: Sense (5' GATAAGCAACGGAAGAGG 3') and antisense (5' AACCAGCAGTCATCAACG 3');<sup>8</sup> thy-1 (CD90) specific primers amplifying a fragment of 237 bp in size: Sense

(5' TCAGCCTCACCAGACAACA 3') and antisense (5' GGACGCACTTCTCCACTTT 3');<sup>8</sup> GAPDH (housekeeping gene) specific primers amplifying a fragment of 293 bp in size: Sence (5' GCCCAGAACATCATCCCA 3) and antisense (5' CCAGCACACGCATCAAAG 3');<sup>8</sup> NANOG specific primers amplifying a fragment of 195 bp in size: Sence (5' CTCCAGCAGCAGACCTCTCCTTG 3') and antisense (5' CCTTCCTTGCCACTCTCACCTT 3').<sup>40</sup>

**RNA extraction.** The total RNA of testis samples were extracted using Cinna Pure-RNA kit according to the manufacturer's instructions (CinnaGen, Tehran, Iran). The concentration and purity of the total RNA was determined by spectrophotometer (ND-2000; Thermo Fisher Scientific, Waltham, USA). Total RNA was stored at -70 °C until cDNA synthesis.

**cDNA synthesis.** First strand cDNA was synthesized from 10 µg of total RNA using random hexamers primers, dNTP Mix, M-MLV Reverse Transcriptase (Vivantis Technologies, Selangor, Malaysia), 10X Buffer M-MuLV according to the manufacturer's instructions. The reaction incubated at 42 °C for 60 min and terminated by incubating at 85 °C for 5 min followed by cooling at 4 °C. Synthesized cDNA was stored at -20 °C until processed.

**PCR assay.** The PCR thermal program for  $\beta$ 1-integrin (CD29), thy-1 (CD90), NANOG and GAPDH consisted of 95 °C for 10 sec, followed by 40 cycles of 95 °C for 10 sec, and 60 °C for 30 sec.

**Electrophoresis.** Agarose gel electrophoresis was performed with the PCR products to verify the primers specificity (Fig. 1).

**Real-Time PCR assay.** The mRNA expression levels of SSCs markers ( $\beta$ 1-integrin (CD29), thy-1 (CD90), NANOG and GAPDH) were quantified by the Real-Time RT-PCR using SinaSyber Blue HF- qPCR Mix (CinnaGen) and was performed on a Step One™ Real-Time PCR System (Applied Biosystems, Carlsbad, USA) in 25.00 µL reaction volume. PCR reactions were comprised of 12.50 µL of SinaSyber Blue HF- qPCR Mix, 0.25 µL of each primer (2.00 µM), 2.00 µL of cDNA, 9.50 µL of nuclease-free water. PCR thermal condition was 95 °C for 10 sec, followed by 40 cycles of 95 °C for 10 sec, and 60 °C for 30 sec.

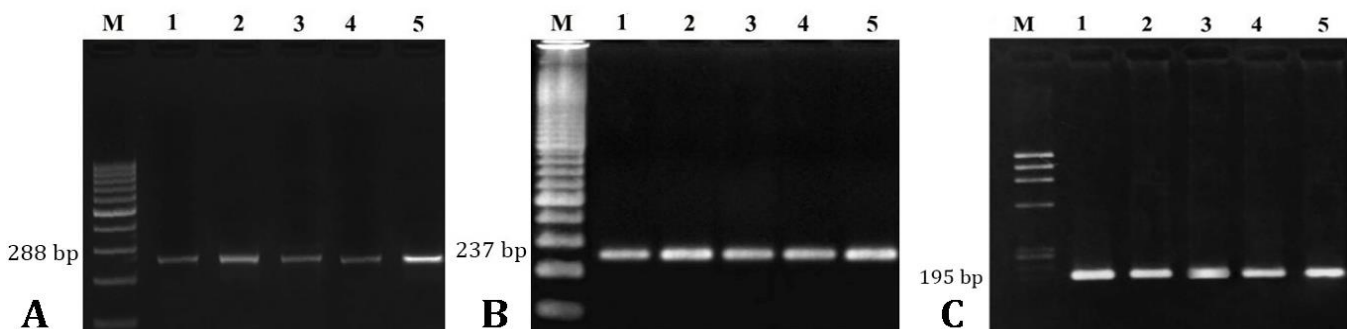
We used a standard curve from cDNA reaction mixture to determine PCR efficiencies. The efficiencies of PCR were observed between 90 and 110 percent. The values for  $R^2$  for all curves were between 0.997 to 0.999. For quantification of the product of PCR, we used the amounts of CT and expressed the relative expression level of the target gene as  $2^{-\Delta CT}$ . We measured  $\Delta CT$  after subtracting CT of housekeeping gene (GAPDH), from CT of target gene.<sup>8</sup>

**Statistical analysis.** The results were analyzed using SPSS s (version 23.0; IBM Corp, Chicago, USA) employing one-way ANOVA. The means of different treatments were compared with Tukey post-hoc test. Data are expressed as the mean  $\pm$  standard error of mean (mean  $\pm$  SE). The differences were considered to be significant at  $p \leq 0.05$ .

## Results

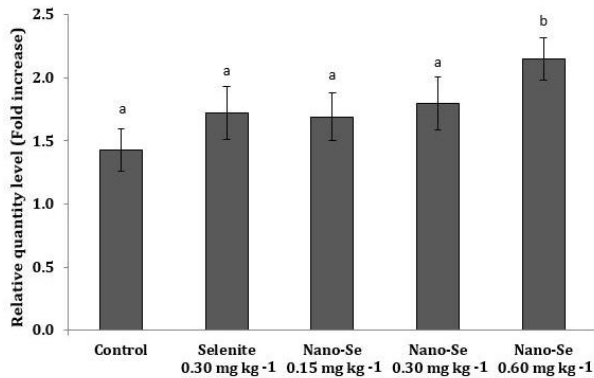
The results of mRNA expressions of the SSCs markers in the experimental groups are expressed as fold-increase in comparison to the reference gene are shown in Figures 2 to 4. As shown in Figure 2, the mRNA expressions of the  $\beta$ 1-integrin (CD29) gene of the groups 1-5 (control, 0.30 mg kg<sup>-1</sup> sodium selenite, 0.15 mg kg<sup>-1</sup> Nano-Se, 0.30 mg kg<sup>-1</sup> Nano-Se, 0.60 mg kg<sup>-1</sup> Nano-Se) were  $1.63 \pm 0.17$ ,  $1.72 \pm 0.21$ ,  $1.69 \pm 0.19$ ,  $1.80 \pm 0.21$  and  $2.15 \pm 0.17$ , respectively. The diet containing different level of Nano-Se had an increasing impact on mRNA expression of  $\beta$ 1-integrin and the group fed with 0.60 mg kg<sup>-1</sup> Nano-Se supplemented diet had the highest mRNA expressions of the  $\beta$ 1-integrin (CD29).

In regards to thy-1 (CD90) gene, the mRNA expressions of the groups 1-5 were  $1.04 \pm 0.21$ ,  $1.14 \pm 0.24$ ,  $1.07 \pm 0.14$ ,  $1.21 \pm 0.17$  and  $1.33 \pm 0.13$ , respectively (Fig. 3) and the highest mRNA expressions was seen in the group fed with diet containing 0.6 mg kg<sup>-1</sup> Nano-Se. As shown in Figure 4, the mRNA expressions of the NANOG gene of the groups 1-5 were  $0.43 \pm 0.12$ ,  $0.61 \pm 0.14$ ,  $0.50 \pm 0.11$ ,  $0.78 \pm 0.11$  and  $0.96 \pm 0.12$ , respectively and the group fed with 0.6 mg kg<sup>-1</sup> Nano-Se supplement diet had the highest level of NANOG gene expression.

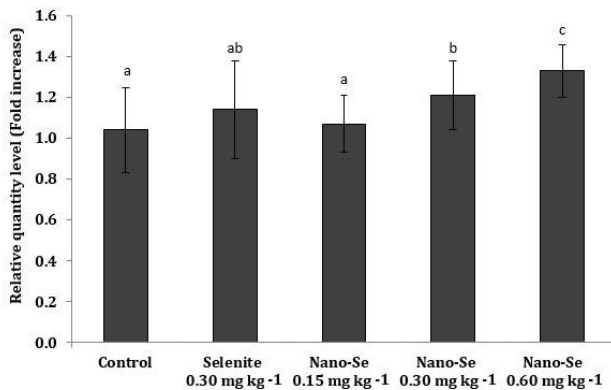


**Fig. 1.** PCR products of amplified genes visualized on agarose gel **A)**  $\beta$ 1-integrin (288 bp), **B)** thy-1 (237 bp) and **C)** NANOG (195 bp) gene primer. Lane M: DNA ladder, Lane 1: Control group; Lane 2: Selenite 0.30 mg kg<sup>-1</sup>; Lane 3: Nano-Se 0.15 mg kg<sup>-1</sup> group; Lane 4: Nano-Se 0.30 mg kg<sup>-1</sup> group; Lane 5: Nano-Se 0.60 mg kg<sup>-1</sup> group.

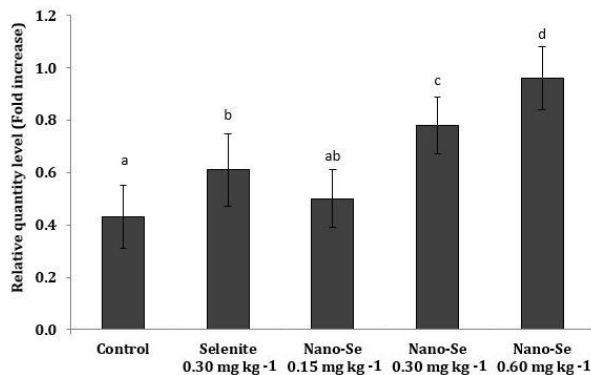
Overall, the lowest level of the mRNA expression of all three genes was in the group fed with diet containing 0.15 mg kg<sup>-1</sup> Nano-Se (Figs. 2 to 4). Primer specificity and cDNA synthesis were verified (Fig. 1).



**Fig. 2.** Level of mRNA expressions of  $\beta$ 1-integrin (CD29) in the treatment groups. Different letters indicate significant differences between the groups ( $p < 0.05$ ).



**Fig. 3.** Level of mRNA expressions of thy-1 (CD90) in the treatment groups. Different letters indicate significant differences between the groups ( $p < 0.05$ ).



**Fig. 4.** Level of mRNA expressions of NANOG genes in the treatment groups. Different letters indicate significant differences between the groups ( $p < 0.05$ ).

## Discussion

Selenium deficiency significantly decreased the population of SSCs and causing low fertility.<sup>8,32-34,41</sup> Therefore, selenium in both inorganic (selenite, selenate) and organic (selenomethionine, selenocysteine and methylselenocysteine) forms is used as a feed supplementation to improve immune responses and fertility in the poultry industry.<sup>42</sup> Recent studies have demonstrated that excess and deficiency of selenium supplementation can stimulate oxidative stress, which probably leads to the reduction of SSCs population, therefore many studies have been designed to determine the optimal dose of selenium in poultry.<sup>8,32,43,44</sup> In recent years, concurrent nano-particle science progresses, therefore Nano-Se has been paid attention as a novel prospect for nutritional supplementation.<sup>45</sup> Thus limited studies are available on Nano-Se, especially in the poultry. As Nano-Se has some advantages (lower toxicity and higher bioavailability) in comparison to inorganic forms' of selenium, therefore, investigating the effective dose of Nano-Se on poultry performances is of great interest to researchers. During this study, the effects of different doses of supplemental Nano-Se on the expression of SSCs markers in the testis of broiler breeder males were investigated by using SSCs surface antigens including  $\beta$ 1-integrin (CD29), thy-1 (CD90) and NANOG that known as the molecular markers.

As shown in Figures 2-4, the mRNA expressions of SSCs markers were elevated by the supplementation of the diet with both sodium selenite and Nano-Se when compared to the control group. The result of real-time RT-PCR indicated that similar to inorganic selenium, Nano-Se dietary supplementation could influence the mRNA expression of SSCs specific markers in testis of broiler breeder males. The results of the present study were in agreement with the previous reports which indicated that supplementation of selenium compounds increases SSCs markers.<sup>8</sup>

Our results also showed that the highest and lowest population of SSCs were in the groups of 0.60 mg kg<sup>-1</sup> and 0.15 mg kg<sup>-1</sup> Nano-Se, respectively. Although the lowest level of Nano-Se (0.15 mg kg<sup>-1</sup>) slightly increased the number of SSCs when compared with the control group. Comparison the number of SSCs markers in the group 2 (fed with 0.30 mg kg<sup>-1</sup> sodium selenite) with those of group 4 (fed 0.30 mg kg<sup>-1</sup> Nano-Se) showed that 0.30 mg kg<sup>-1</sup> Nano-Se had a higher number of SSCs markers than 0.30 mg kg<sup>-1</sup> inorganic selenium but differences were not significant. As shown in Figures 2-4, the number of SSCs markers increased when the doses of Nano-Se increased. The group fed with 0.60 mg kg<sup>-1</sup> Nano-Se, had the highest SSCs markers at the end of the experimental period and in comparisons with those of the group fed with inorganic selenium, the differences were significant ( $p < 0.05$ ).

The results obtained in the present study clearly showed that supplementation of the poultry diet with Nano-Se was more effective than sodium selenite. The dose of 0.60 mg kg<sup>-1</sup> of Nano-Se could be used in the diet of roosters in order to improve their performances and in particular their fertility without any side effects. Regarding the mechanism of action, Selenium as an essential trace element plays an important role in antioxidation<sup>46,47</sup> and protects various systems against damage caused by free radicals and oxidative stress.<sup>48</sup> The physiological functions of Selenium are considered to be mediated through selenoproteins.<sup>49,50</sup> Unlike the previous studies using inorganic selenium which showed that higher levels of sodium selenite reduces the number of SSCs markers,<sup>8</sup> the results of the present study revealed that the increasing levels of Nano-Se in the diet did not cause a detrimental effect on the population of SSCs markers (Figs. 2-4). These differences could be attributed to the differences in selenium sources.

In conclusion, the supplementation of the diet with 0.60 mg kg<sup>-1</sup> of Nano-Se can be used in roosters feeding to improve their performance in particular when fertility is declining due to aging in broiler breeder males from 40 weeks of age onward.

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#### Conflicts of interest

The authors notify that they have no conflicts of interest.

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