

THE DEVELOPMENT PATTERN OF IGF-1 (INSULIN-LIKE GROWTH FACTOR-1) PROTEIN EXPRESSION IN BREAST MUSCLE OF BROILER CHICKENS

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Abstract: It is proved that the protein expression of IGF-1 is detected in several tissues including liver, brain, eye, lung, pancreas and muscle. This growth factor exerts autocrine and paracrine effect during embryonic and postnatal development and is involved in muscle development. The aim of this study was to determine the level of protein expression of IGF-1 in breast muscle during embryonic and postnatal development of broiler chickens. For the experiment were used fertilized eggs and broiler chickens of heavy hybrid "Ross 308", originating from parent flock at the age of 54 weeks. Samples of breast muscle were taken from embryos during the 15th and 19th day of embryonic development and 3rd and 21st day of postnatal development from 5 birds in each age category. From all of the embryos and chickens the right side of the breast muscles were individually sampled. The muscle tissue was macerated and then homogenized in buffer with protease inhibitors. After that immunoprecipitation, electrophoresis and transfer the Western blot analysis were performed on all samples. For visualization of proteins the membranes were incubated in chemiluminescent substrate. The results indicate statistically significant increase in protein expression of IGF-1 in breast muscle on the 19th day of embryonic development and 3rd of postnatal development in compared to 15th day of embryonic development ($p < 0,05$ compared to 19th day of embryonic development and $p < 0,01$ compared to the 3rd day of postnatal development). After the 3rd day of postnatal development the protein expression was at the same level until the 21st day of postnatal development.

Key words: IGF-1, breast muscle, broiler chickens

Introduction

It has been shown that insulin-like growth factor-1 (IGF-1), in birds, is essential for normal growth and development (*Proudman et al., 1994*). The origin of IGF-1 in muscle may be twofold. IGF-1 may be derived from increased synthesis in the liver under the effect of growth hormone and, more significant,

may be of local origin, and manifest autocrine or paracrine effect (*Halevy et al., 2001*). Alternatively, the effects of IGF-1 may be, at least partially, mediated by hormones such as thyroid hormones (*Adams et al., 2000*). It has been shown that IGF-1 stimulates proliferation of satellite cells in chickens (*Duclos et al., 1991; Hodik et al., 1997*).

Growth factors in muscle define the direction of development of muscle cells in the direction of cell proliferation or the direction of cell differentiation. The same growth factor can stimulate the muscle cells in both of these processes, but through different systems of secondary messengers in the cell. The activation of phosphoinositide 3-kinase (PI3K) pathway of secondary messengers stimulates the differentiation of myoblasts, while activation of mitogen-activated protein kinase (MAPK) pathway stimulates the proliferation and inhibits differentiation of myoblasts (*Halevy and Cantley, 2004*).

IGF-1 plays an important role in the metabolism of carbohydrates, fats and protein in adipose tissue, skeletal muscle and liver. IGF-1, in skeletal muscle cells, stimulates protein synthesis and the uptake of glucose (*LeRoith and Yakar, 2007*). In this way, IGF-1 can significantly affect the chemical composition and progressive loss of moisture from the meat.

The aim of this study was to determine the level of protein expression of IGF-1 in breast muscle during embryonic and postnatal development of broiler chickens.

Materials and Methods

For this trial we have used the fertilized eggs of heavy hybrid "Ross 308". The average egg weight was 66.5 g and ranged from 63.5 g to 69 g. After the rest period the eggs were deposited in to the incubator. Temperature and relative humidity were controlled by analog device (Veb, Berlin, Germany). The humidity in all incubators was maintained between 50% and 60% until the 18th day of embryonic development and up to 75% from 19th day of embryonic development until the end of incubation. After hatching, drying and resting for 24 hours, the broilers were packed into transport boxes and transferred to the facility for fattening (the floor system with boxes). The conditions (temperature, humidity, lighting, density) were adapted in accordance with requirements of this hybrid (*Halevy et al., 2001*). Chickens were vaccinated based on a regular vaccination program against Newcastle disease and infectious bursal disease, and throughout the experiment were under constant veterinary supervision.

During the production cycle broilers were fed ad libitum in three phases. From 1st to 13th day the diet was with 22% of proteins (starter), from 14th to 34th day the diet was with 19% of proteins (grower) and from 35th day until the end of the production cycle (42nd day) the diet was with 18% of proteins (finisher).

The breast muscle samples for Western blot analysis were taken from the embryo during the 15th and 19th day of embryonic development, and during the 3rd and the 21st day of postnatal development. Samples of breast muscle were taken from 5 individuals in each age category. From all of the embryos and chickens the right side of the breast muscles were individually sampled. The samples were taken from the right side of the breast and then frozen in liquid nitrogen.

The tissue was macerated, and then homogenized in radio immune precipitation buffer with protease inhibitors (F. Hoffmann-La Roche Ltd. Basel, Swiss), three times in ten seconds intervals with ten second pauses. The sonification of samples was performed on 10 kHz, three times in fifteen seconds intervals with fifteen seconds pauses. The sample was then centrifuged in Beckham ultracentrifuge (Beckham Ultima TL, Fullerton, CA, USA), 90 minutes at 85000 g. The obtained supernatants were used for Western blot analysis.

To determine the protein expression of IGF-1, 100 mg of tissue homogenate protein was precipitated with rabbit polyclonal anti-IGF-1 antibody (Abcam, Cambridge, UK) with final dilution of 1:50. Immune complexes were precipitated adding protein A / G-Sepharose (the initial amount of 100 mg protein per 20 μ l of purified Sepharose) and were incubated for 2 hours at 4 °C. After centrifugation (5 seconds at 14000 g) pellet was rinsed three times with iced buffer. The obtained residue proteins were used for SDS-polyacrylamide electrophoresis and immunoblotting.

The proteins were separated by molecular mass in a BioRad Mini-Protean III system (BioRad, Mississauga, Ontario). Gels used to separate proteins were different depending on the molecular weight of the proteins and a gel for concentration was 5% (5% acrylamide / bisacrylamide, 0.1% SDS; 0.125 M Tris, pH 6.8). Polymerization of gels was enabled by adding 0.05% ammonium persulfate and 0.033% N,N,N',N'-tetramethyl-ethylenediamine. The buffer for reservoirs was consisted of 0.192 M glycine, 0.1% SDS and 0.025 M Tris, pH 8.3. The equal volume of sample buffer (4% SDS, 20% glycerol, 10% β -mercaptoethanol; 0.125 M Tris, pH 6.8; 0.025% brome phenol blue) was added in to the specific volume of protein solution. Samples were further denatured by boiling for 5 minutes at 95 °C. The proteins from the sample and the protein marker were poured to the gels in order to determine the molecular weight (Fermentas International, Inc., Burlington, Canada). Electrophoresis lasted for 60-90 minutes at constant voltage of 120V. After electrophoresis, the gels were used for Western blot.

Proteins were transferred from gels to polyvinylidene fluoride membrane Hybond-P (Amersham Pharmacia Biotech). The membranes were activated in methanol (10 seconds). Membranes and gels were packaged and placed in tanks with transfer buffer (0.192 M glycine, 20% methanol and 0.025 M Tris, pH 8.3). The transfer of proteins from gels to membranes was carried out at constant voltage of 100 V for 60 minutes. The success of the transfer was confirmed with 5%

Ponceau S color dissolved in glacial acetic acid. Membranes were discolored by rinsing in distilled water and then incubated for one hour at room temperature in serum to block free sites on the membrane - 5% bovine serum albumin in Tris-borate buffer supplemented with Tween detergent (TBS-T: 0.2 M Tris, 1.5 M NaCl, 0.05% Tween20, pH 7.4). After blocking the membranes were incubated with primary antibody.

Western blot was performed in a quantitative manner (*Tienrungrroj et al., 1987*). Antibody and concentration used in this experiment is: Antibody IGF-1 (ab9572), the concentration of 0.2 mg/ml. After blocking, the membranes were incubated with appropriate primary antibodies diluted in 5% bovine serum albumin, overnight at 4 °C. After incubation and washing away excess antibody, the membranes were incubated with secondary antibody obtained by immunization of goats with a rabbit immunoglobulin G (IgG) (Santa Cruz Biotechnology, Inc.), diluted 1:2000 (antibody: 5% bovine serum albumin), 2 hours at room temperature. After incubation, the membranes were rinsed with TBS solution for 30 minutes at room temperature. For visualization of proteins the membrane were incubated in luminol, with the addition of H₂O₂ for three minutes. After that, the X-ray films (Hyperfilm, Amersham Pharmacia Biotech) were subjected to membranes (0.5-2 minutes). The films were developed, scanned and used for quantification of protein bands.

Data are presented as means \pm SEM. Data were analyzed by Tukey test to separate means by using Graph Pad Software 3.03 program (GraphPad Prism 3.03, GraphPad Software Inc., San Diego, CA, USA).

Results and Discussion

It is relatively difficult to draw definitive conclusions about changes in concentration of IGF-1 during development in birds. Generally it can be concluded that concentrations of IGF-1 in plasma, during embryonic development in birds, expressed a clear single-phase profile. In chicken embryos, the blood plasma concentration of IGF-1 increases from 9th day of incubation and reaches a peak at 15th day of incubation and then gradually decreases (*Kikuchi et al., 1991; Robcis et al., 1991; Scanes et al., 1997*). In addition, there is evidence that the concentration increases again later in the development (*McMurty et al., 1994*). Ontogenetic profile of the concentration of circulating IGF-1 in chickens was given by *Hybrechts et al. (1985), McGuinness and Cogburn (1990)* and *Redecki et al. (1997)*.

Unlike mammals, where the concentration of IGF-1 in blood plasma directly related to the GH-dependent synthesis in the liver (*Kind et al., 1995*), in birds, this is only partly the case. This is based on the fact that in the liver of chicken embryos were not found the significant levels of mRNA (messenger ribonucleotid acid) for IGF-1 (*Kikuchi et al., 1991, Burnside and Cogburn, 1992*). Also, the concentration of circulating IGF-1 decreases after hypophiseectomy, but

only on the half of the initial value (*Hybrectes et al., 1985*) and a GH (growth hormone) therapy only partially establishes the initial level in the circulation (*Scanes et al., 1986*). A similar situation is found in dwarfs that lack functional GH receptors, where the level of IGF-1 in circulation decreased by about 50% (*Hybrectes et al., 1985*). In mammals, GH deficiency reduces the levels of IGF-1 in the circulation for more than 90% (*Glasscock et al., 1990*). Houston and O'Neill (*1991*) show that the liver cells are releasing IGF-1 in the presence of GH and insulin.

The explanation may be that the synthesis of IGF-1 in the liver, partially independent of GH in birds, and that the liver is not the only source of IGF-1. There is evidence that IGF-1 is synthesized in several other tissues including brain, eye, lung, pancreas, and muscles (*Serna et al., 1996, Tanaka et al., 1996*). This synthesis of IGF-1 may point to its local (autocrine or paracrine) and endocrine effects. In addition, the mere synthesis of IGF-1 does not mean, by itself, that it is entirely released into the circulation. There is indirect evidence that IGF-1, in the circulation, originates from the different tissues (*Scanes, 1997*). Because IGF-1 has local, as well as systematic effects, it is important to measure the protein expression of this growth factor in each tissue individually.

The protein expression of IGF-1 was detected in breast muscle tissue during the embryonic and postnatal development. The level of protein expression was increasing from the 15th day of embryonic development until the 3rd day of postnatal development and remained at that level until the 21st day of postnatal development. The protein expression of IGF-1 is shown in Table 1.

Table 1 Protein expression of IGF-1 in the breast musculature

Age	IGF-1 expression ¹
Prenatal	
15th day	100 ± 11
19th day	132 ± 13*
Postnatal	
3rd day	240 ± 21**
21st day	242 ± 22**

¹ % compared to the 15th day of embryonic development, * p<0,05, ** p<0,01

The results indicate statistically significant increase in protein expression of IGF-1, in breast muscle, during the late embryonic development and early postnatal development. On the 19th day of embryonic development and 3rd of postnatal development in compared to 15th day of embryonic development (p<0,05 compared to 19th day of embryonic development and p<0,01 compared to the 3rd day of postnatal development). After the 3rd day of postnatal development the

protein expression was at the same level until the 21st day of postnatal development.

Growth factors in muscle define the direction of development of muscle cells in the direction of cell proliferation or the direction of cell differentiation. The same growth factor can stimulate the muscle cells in both of these processes, but through different systems of secondary messengers in the cell (*Halevy and Cantley 2004*).

The lower levels of IGF-1 in muscle, in the earlier stages of development, may be essential for achieving better results at the end of the production cycle. The IGF-1 in muscle tissue induces differentiation process, while some factors such as FGF and HGF inhibit this process, and stimulate cell proliferation (*Barton-Davis et al., 1999; Ganong, 1995*). Lower levels of IGF-1 in muscle, in the early period of development, enable proliferation of satellite cells. The increased levels of IGF-1, in later stages of development, stimulate the differentiation and hypertrophy. This may be the basis of the mechanism through which different treatments, during the embryonic development, stimulates the body weight gain and increase of protein content in meat (*Halevy et al., 2006*).

Conclusion

The results indicate statistically significant increase in protein expression of IGF-1 in breast muscle on the 19th day of embryonic development and 3rd of postnatal development in compared to 15th day of embryonic development ($p < 0,05$ compared to 19th day of embryonic development and $p < 0,01$ compared to the 3rd day of postnatal development). After the 3rd day of postnatal development the protein expression was at the same level until the 21st day of postnatal development.

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Razvojni obrazac proteinske ekspresije IGF-1 (faktora rasta nalik insulinu – 1) u grudnim mišićima brojlerskih pilića

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Rezime

Dokazana je ekspresija IGF-1 u više tkiva uključujući jetru, mozak, oči, pluća, pankreas i miškulaturu. Ovaj faktor rasta ispoljava autokrino i parakrino delovanje tokom embrionalne i postnatalne faze razvoja i uključen je u proces razvoja mišića. Cilj ovog rada je bio da se utvrdi razvojni obrazac proteinske ekspresije IGF-1 u grudnim mišićima brojerskih pilića tokom embrionalne i postnatalne faze razvoja. Za ogled su korišćena oplodena jaja i brojerski pilići teškog hibrida "Ross 308", poreklom od roditeljskog jata u uzrastu od 54 nedelje. Uzorci grudnih mišića su uzeti od embriona tokom 15. i 19. dana embrionalnog razvoja i 3. i 21. dana postnatalnog razvoja od 5 jedinki u svakom uzrastu. Od svih embriona i pilića pojedinačno je uzet uzorak sa desne polovine grudnih mišića. Mišićno tkivo je macerirano a zatim homogenizovano u puferu sa inhibitorima proteaza. Posle imunoprecipitacije, elektroforeze i transfera na svim uzorcima je izvršena Western blot analiza. Za vizuelizaciju proteina membrane su inkubirane sa hemiluminescentnim substratom. Rezultati ukazuju na statistički značajno povećanje proteinske ekspresije IGF-1 u grudnim mišićima tokom 19. dana embrionalnog razvoja i 3. dana postnatalnog razvoja u poređenju sa 15. danom embrionalnog razvoja ($p < 0,05$ u poređenju sa 19. danom embrionalnog razvoja i $p < 0,01$ u poređenju sa 3. danom postnatalnog razvoja). Nakon 3. dana postnatalnog razvoja proteinska ekspresija je bila na istom nivou do 21. dana postnatalnog razvoja.

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