

PROTEIN SIGNAL TRANSDUCERS AND ACTIVATORS TRANSCRIPTION (STAT) AS GROWTH PROMOTER

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ABSTRACT

The purpose of this study was to determine the amino acid composition of proteins STAT 1 and STAT 3 existing in the liver tissue of broiler chickens that are experiencing growth due to increased growth hormone (GH). Liver tissue samples of broilers isolated from broiler chickens reared for 21 days were examined using SDS Page, followed by Western Blott and MALDI-TOP. Results of Western Blot revealed that STAT protein had molecular weight of 59.3kDa with the first amino acid composition 1, mtqwyqlqql dskfleqvqh lyddsfpmey rqlaqwlen qdwehaannv sfatlvfhdl, 61 lsnqmevggv qntmtgmldk qkeldakvka vknsvidveq diktledvqd eydfkhktfq, 121 lsqlddqfsr fliennfllq hnirkskrnl qdnfqedpih mamiihcnlk eerkilnsaq, and STAT 3 59.4kDa by arrangement amino acids 1, maqwnqlqql dtryleqlhq lysdsfpmel rqlapwies qdwayaanke shatlvfhn, 121 taaqqgggqat hptaavvtek qqmleqhlqd vrkrvqdlq knkvvenlqd dfdfnytkl, 181 sqgdmqdlng nnqsvtrqkm qqleqmtil dmqrrgivse lagllsamey vqkmladeel, 241 adwkrqqia cigppniel drlenwitsl aesqlqtrqq ikkleelqgk vsykgdpivq, 301 hrpmleriv elfnlnksa fvverqpcmp mhpdrplvik tgvqfttkvr llvkfpelny and 361 qlkikvcidk dsqdaalrg srkfnilgtm tkvmnmeesn ngslsaefkh ltlreqrcgn.

Keywords: STAT 1, STAT 3, growth, broiler

INTRODUCTION

Growth hormone has important meaning in regulating body growth and metabolism. GH metabolic effects occur when GH receptors associate with and activate tyrosine kinases. The GH bond with its receptor may activate Janus Kinase 2 (JAK 2) and further phosphorylate tyrosine in the JAK-2 GH-receptor complex. This tyrosine then forms the binding site for a number of signaling proteins, such as signal transducers and activators of transcription (STAT) to incite the effect of growth. STAT proteins that play a role in providing growth signaling are STAT 1, STAT 3, STAT 5a and STAT B. STAT proteins play a significant role in regulating metabolic and growth effects.

Increased growth in animal husbandry has great implications and appeal in the poultry field. However STAT protein signaling and its expression pattern in broilers during growth period has not been identified clearly. Therefore, knowledge on molecular weight and the arrangement of amino acids of STAT signaling protein in broilers during growth period due to increased GH can be used to make synthetic STAT protein to spur the growth of the broilers.

Until recently, the only known identification of amino acid STAT 5a protein composition was cigpppkvmnmeesn, and STAT 5b protein was datnilvspvlypdip (Anwar et al., 2013). Therefore, we need further studies to identify the composition of amino acid protein STAT 1 and STAT 3 to be used as the basis of producing synthetic STAT protein to boost the growth of the broilers.

MATERIALS AND METHODS

Chickens were placed in a battery cage with a capacity of one chicken per cage with twice daily feed at 6:00am and 6:00pm with the amount of 10% smaller than the standard. At the age of 21 days, the chickens were sacrificed to be sampled by taking their hepatic tissue for the following tests: (1) isolation of signaling proteins STAT 1 and STAT 3 from broiler liver tissue; (2) analysis of STAT 1 and STAT 3 protein signaling from broiler liver tissue; (3) identification of molecular weight of STAT 1 and STAT 3 signaling proteins by blotting method, the Western blot technique, using electrophoretically elaborated protein from polyacrylamide gel; and (4) identification of amino acids by MALDI-TOF method.

RESULTS AND DISCUSSION

SDS Page of STAT protein from broiler liver

The result of SDS-PAGE STAT protein on broiler liver tissue showed the presence of STAT protein, as in Figure 1.

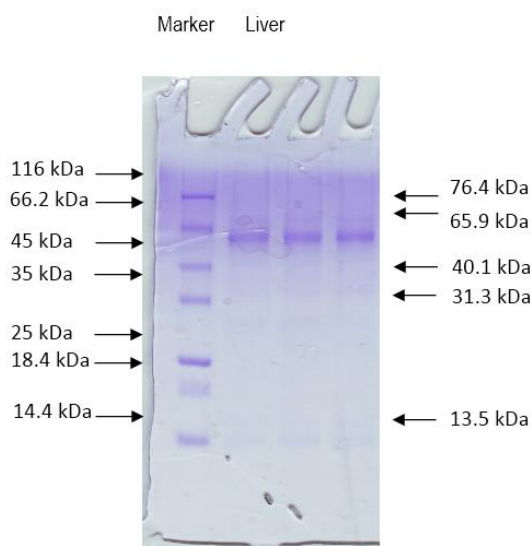


Figure 1. SDS Page STAT protein from the liver tissue

The SDS-PAGE results showed five visible protein bands between the 116kDa marker and 14.4kDa on the liver tissue.

From the calculations (Table 5.1), the protein bands formed between 116kDa markers with 14.4kDa had molecular weights of 76.4kDa, 65.9kDa, 40.1kDa, 31.3kDa and 13.5kDa. The protein bands formed on the hepatic tissue were apparently very clear at the molecular weight of 65.9kDa. Results of SDS-PAGE liver tissue protein showed the most obvious protein bands were between the 66.2kDa marker, with 45kDa, the suspected proteins STAT 1 and STAT 3. The proteins of the SDS-PAGE results could not be ascertained yet as to whether they were STAT 1 or STAT 3 proteins. To prove that the formation of protein bands were STAT 1 and STAT 3 proteins it was necessary to check with Western blot.

Western Blot of STAT 1 protein

The result of Western blot STAT protein on hepatic tissue showed the presence of STAT 1 protein with a molecular weight of 59.3kDa, as in Figure 2.

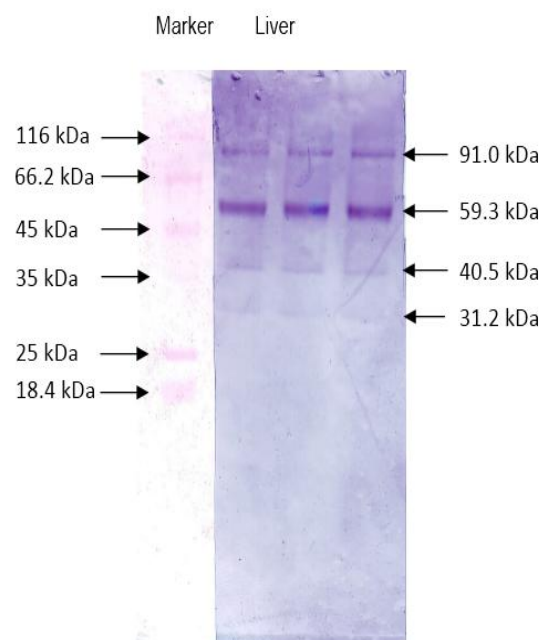


Figure 2. Western Blot of STAT-1 protein from the liver tissue

To verify that the result of protein analysis with SDS-PAGE was STAT 1 protein, then the Western blot was done using rabbit polyclonal antibody STAT 1 Ab-1 (Labvision). In Figure 3, it appears that one protein band is most apparent between the 66.2kDa marker, with 44kDa, a protein formed on the liver tissue. The protein bands formed between 116kDa markers with 66.2kDa, 45kDa with 35kDa and 35kDa with 2 kDa are not very clear.

After calculation (Table 5.2), the formation of protein bands between markers 66.2kDa with 44kDa turned out to have a molecular weight of 59.3kDa. This suggests that the SDS-PAGE-produced protein tested with Western blot was a STAT 1 protein in broiler during growth period with a molecular weight of 59.3kDa. The formation of a very clear 56.3kDa band of protein bands was due to the occurrence of a binding between the SDS-PAGE STAT 1 protein with rabbit polyclonal antibody STAT 1.

Western Blot Protein STAT 3

The result of Western blot protein STAT on hepatic tissue showed the presence of STAT 3 protein with a molecular weight of 59.4kDa, as in Figure 3.

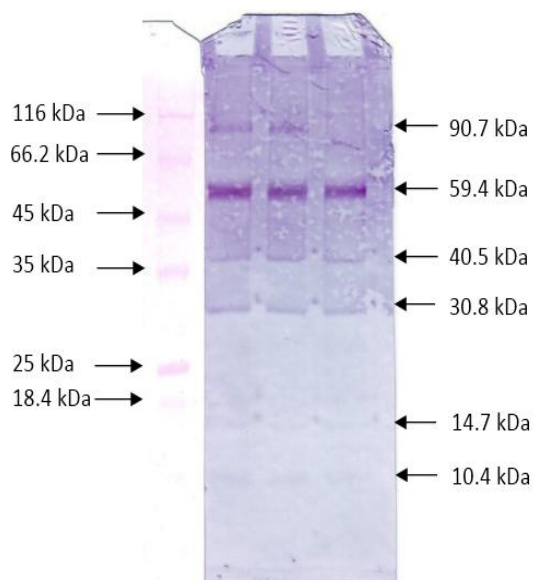


Figure 3. Western Blot of STAT-3 protein from the liver tissue

To ensure that protein analysis results with SDS-PAGE was a STAT 3 protein, the Western blot was performed using rabbit polyclonal antibody STAT 3 Ab-1 (Labvision). Figure 5.4 shows the formation of one of the clearest protein bands between the 66.2kDa marker and 44kDa; both proteins formed on adipose, muscle and liver tissue. The protein bands formed between 116kDa markers with 66.2kDa, 45kDa with 35kDa and 35kDa with 25kDa are not very clear

After calculation (Table 5.3), the protein band formed between the 66.2kDa marker and 44kDa has a molecular weight of 59.4kDa. This suggests that the SDS-PAGE protein tested with Western blot is a STAT 3 protein from broilers during growth period with molecular weight of 59.4kDa. The protein band with a molecular weight of 59.4kDa was very clearly formed due to the binding between STAT 3 protein SDS-PAGE results with rabbit polyclonal antibody STAT 3.

Amino acid of STAT 1 and 3 proteins

Results of examination using MALDI-TOP method showed that STAT 1 protein with molecular weight of 59.3kDa had amino acid 1 arrangement of mtqwyqlqql dskfleqvqh lyddsfpmel rylaqlwen qdwehaannv sfatvlfhdl, 61 lsnqmevggv qntmtgml dk qkeldakvka vknsvt dveq diktledvqd eydfkhktfq, 121 lsqlddqfsr fliennflq hnirskrnl qdnfqedpih mamiihcnlk eerkilnsaq, and STAT 3 as much as 59.4kDa with amino acid 1 arrangement maqwnqlqql dtryleqlhq lysdsfpmel rqlapwies qdwayaanke shatlvfhnl, 121 taaqqggqat hptaavvtek qqmleqlhqd vrkrvqdleq kmkvvenlqd dfdfnyktlk, 181 sqgdmqdlng nnqsvtrqkm qqleqmltal dqmrrgivse lagllsamey

vkmladeel, 241 adwkrqqia cigppncl drlenwitsl aesqlqtrqq ikkleelqqk vsykgdpivq 301 hrpml eeriv elfrnlmksa fvverqpcmp mhpdrplvik tgvqfttkvr llvkfpelny, and 361 qlkikvcidk dsgdvaalrg srkfniltgn tkvmnmeesn nglsaeafh ltlregrcn.

Growth hormone plays a role in regulating body growth and composition (Foster, 1998). Growth hormone apparently has biological effects that are influenced by insulin-like growth factor I (IGF-I) in enhancing the growth of skeletal muscle (Younken, 2000). Provision of in vivo growth factor in broilers led to an increase in growth rate and muscle mass by 15% and required 6.5% less feed than normal feed. This increase in growth has major implications and appeal in the field of poultry. However, to date, expression patterns of growth factor genes during growth period have not been clearly identified (Killefer, 2000).

STAT proteins play an important role in the regulation of gene transcription by GH and other cytokines that activate Janus Kinase (JAK). STAT proteins, originally identified in the signaling pathway of interferon (IFN) (Darnell et al., 1994), is a cytoplasmic factor containing domain SH-2. In tyrosyl phosphorylation, frequently through the cascade initiated by JAK kinases, STAT protein cytoplasm forms a complex with other STAT protein through the interaction of tyrosine which is phosphorylated on the domain of SH-2, and holds translocations leading to the nucleus, binds to DNA and subsequently activates transcription of the gene target (Ihle, 1996).

Growth hormone is known to activate STATs 1, 3, 5a and 5b. Phosphorylation of GH-dependent STATs 1, 3, 5a and 5b tyrosyl is found in 3T3-F442A fibroblasts, liver of rats with hypophysectomy, liver cell cultures and in various systems' overexpression. Phosphorylation of STATs 5a and 5b tyrosyl is also found in IM-9 human cells and muscles of the liver and skeletal muscle of normal mice (Smit et al., 1999).

STAT1, also called P91, is identified as a member of the factor 3 gene complex that is stimulated by IFN-gamma (FU, 1992). Analysis of signaling GH in JAK2 deficient cells and cells undergoing mutations in expressing GH receptor indicates that GH-dependent activation of STATs 1, 3, 5a and 5b requires activation of JAK2 (Smit et al., 1997). This is consistent with the finding that JAKs activation is required for STAT activation (Muller et al., 1993). JAK1 or JAK2 actively overexpressed in COS cells will stimulate the binding of STAT1 to DNA (Silvennoinen, 1993).

Indirect research has shown that GH stimulates the phosphorylation of STATs 1, 3 and 5 in serine or

threonine in the liver. This phosphorylation will increase DNA binding of STAT1, DNA STAT3 and substantially alter DNA binding of STAT5 (Ram et.al., 1996). STAT 1, 3, and 5a contain conserved consensus sequences for phosphorylation, the MAP kinase. Initial studies have shown that MAP kinase is responsible for serial phosphorylation of STAT1, STAT3 and STAT 5a. Whereas, in STAT 5b, since it does not contain conserved consensus sequences, the phosphorylation is performed by kinases other than MAP kinase. Proteins STAT 1, 3, 5a and 5b also contain protein kinase C and casein kinase for phosphorylation process. This suggests that double signaling pathways can converge in STAT proteins for transcriptional activation by GH.

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CONCLUSIONS OR IMPLICATIONS

1. STAT-1 protein is 1 mtqwyqlqql dskfleqvhh lyddsfpmel rylaqlwlen qdwehaannv sfatvlfhdl, 61 lsnqmevggv qntmtgml dk qkeldakvka vknsvtdeq diktledvqd eydfkhktfq, 121 lsqlddqf sr fliennflq hnirsksrnl qdnfqedpih mamiihcncl eerkilnsaq.
2. STAT-3 1 protein is maqwnqlqql dtryleqlhq lysdsfpmel rqlapwies qdwayaanke shatlvfhnl, 121 taaqgggqat hptaavvtek qqmleqhlqd vrkrvqdeq kmkvvenlqd dfdfnyktlk, 181 sqgdmqdlng nnqsvtrqkm qqleqmltal dqmrrgivse lagllsamey vqkmladeel, 241 adwkrqqia cigppnicl drlenwitsl aesqlqtrqq ikkleelqqk vsykgdpivq, 301 hrpmlleeriv elfrnlmksa fvverqpcmp mhpdrplvik tgvqfttkvr llvkfpelny, 361 qlkikvcidk dsdvaalrg srkfnilgtm tkvmnmeesn ngslsaefkh ltlreqrcgn.
3. By knowing the arrangement of amino acids, it can be employed as the basis for making STAT synthetic protein which is expected to be used to prolong the action or effect of growth hormone so that it can enhance the growth of livestock.

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