

## *Molecular strategies for the differentiation and identification of local *E. coli* isolated from chicken: I. Characterization of protein profile.*

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**In this study five serotypes of *E. coli* were isolated from chickens and identified as O1, O2, O6, O78 and O126 out of 33 isolates derived from a total of 60 samples. SDS-PAGE revealed that four proteins were characteristic and shared in all these serotypes at the molecular weight of 21, 30, 55 and 74 kDa of which 55 and 74 kDa proteins were fully reacted with the antisera against *E. coli* in the western blot. Other proteins are present but varied from one serotype to another.**

*E. coli* is one of the serious problems that affect poultry industry because it is responsible for a variety of disease conditions such as colisepticemia, air sacculitis, peritonitis, perihepatitis, pericarditis, omphalitis, salpingitis, coligranuloma and enteritis (Heller *et al.*, 1972; Gross, 1991).

Most strains of *E. coli* are harmless; however some are pathogenic by virtue of plasmid. Mediated virulence factors through which disease is induced in animals and poultry (Morris and Sojka, 1985; Gyles, 1986).

*E. coli* infection in poultry is usually associated with high economical losses. This is due to high morbidity and mortality rates, decrease food conversion rate, body weight loss, and decrease in egg production, condemnation of whole affected carcass or organs and the high cost of wide range of antibacterial agents used to control. Therefore rapid and accurate mean to differentiate between the different strains of *E. coli* in poultry farms is required.

### **Materials and Methods**

**Isolation of *E. coli*.** A total of 60 suspected samples taken from colisepticaemic chickens (cloacal swabs and/or bone marrow) were used in the isolation on nutrient agar and MacConkey agar. The suspected colonies were examined for their colonial morphology, microscopical and biochemical characteristics using the API2.E system (Micro bet GNB 12 A\B/E, 24E).

**Diagnostic antisera.** Four polyvalent and 24 monovalent antisera (SEIKEN- Japan) were used for stereotyping of the *E. coli* suspected isolates using the technique described by (Edward and Ewmg, 1972). Polyvalent (1): O1, O26, O86, O111, O119, O127, O128. Polyvalent (2): O44, O55, O125, O126, O146, O166. Polyvalent (3): O6, O27, O78, O148, O159, O168. Polyvalent (4): O2, O11, O87, O127, O142.

**Harvesting of isolates.** A single bacterial colony from each isolates was streaked onto nutrient agar medium and incubated at 37°C for 24 hours then inoculated into tryptic soya broth overnight at 37°C with shaking. The cells were harvested by centrifugation at 5000 rpm/15 min using cooling centrifuge. The pellets were washed 3 times with PBS (pH 7.2) then resuspended in 5ml of ice cooled 25 mM Tris HCL (pH 7) (Chart, 1994).

**Preparation of hyperimmune sera.** Antisera to F11 and FT *E. coli* antigens were raised in chickens by intramuscular and subcutaneous injection using modified protocol described by (Condon and Owen, 1982). Isolation and concentration of immunoglobulin have been described in detail by (Owen, 1983).

**Total protein analysis.** The total protein of samples were prepared as described by (Kishor *et al.*, 1996). Total protein analysis was carried out by using SDS-PAGE and western blot technique.

**SDS-PAGE.** Extracted protein of the purified bacterial preparation was resolved on discontinuous buffer system composed of 10 % (w/v) acrylamid running gel and 4% shacking gel (Sambrook *et al.*, 1989). Electrophoresis was carried out at a constant voltage (100v) until

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**Table (1): Isolation, identification and serotyping of *E. coli* from chickens.**

Cases	No. of samples	No. of isolates	Serotypes
Apparently healthy	17	8	O1(4 isolates), O6 (2 isolates), O78 (2 isolates)
Diseased	24	14	O1 (3 isolates),O2 (2 isolates),O6 (3 isolates), O78 (4 isolates), O126 (2 isolates)
Freshly dead	19	11	O1 (2 isolates),O2 (1 isolates),O6 (2 isolates), O78 (3 isolates), O126 (3 isolates)
Total	60	33	5 serotypes

**Table (2): protein profiles of different isolated *E. coli* serotypes.**

Serogroups		O1	O2	O6	O78	O126
Protein profile	21kDa	+	+	+	+	+
	26kDa	+	-	-	+	+
	27kDa	-	+	-	+	+
	30kDa	+	+	+	+	+
	32kDa	+	+	-	-	+
	56kDa	+	+	+	+	+
	74kDa	+	+	+	+	+
	85kDa	+	-	-	+	+
	24kDa	+	-	-	-	+
	40kDa	-	-	-	+	-
	60kDa	+	+	-	-	-
	64kDa	-	-	-	+	+
	67kDa	-	-	+	-	-
	69kDa	-	+	-	-	-
78kDa	-	+	+	-	-	
95kDa	+	-	-	-	-	
100kDa	-	-	-	-	+	

tracking dye front (Bromophenol blue) moved to the bottom of the gel .Wide range protein marker (10 kDa to 200 kDa, Page Ruler protein Ladder; Fermentas) was used. The gel was stained with SDS-PAGE gel stain (Gel Code Blue Stain, PIERCE, USA.). Molecular weight of each protein band was calculated with reference to a standard curve derived from the migration patterns of the protein molecules weight marker.

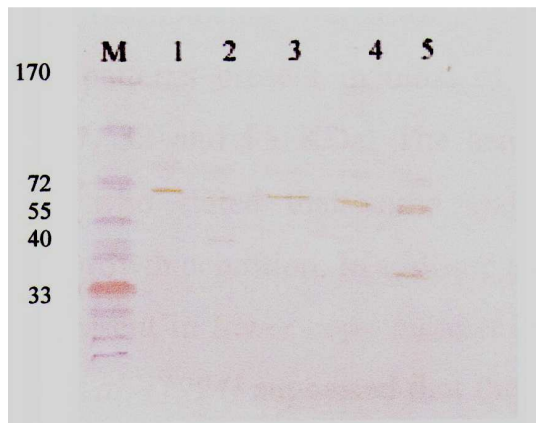
**Western blot technique.** The electrophoretic transfer of the polyacrylamide gel resolved proteins to the nitrocellulose membrane was carried out by electro blotting as described by (Towbin *et al.*, 1979) using Bio Rad Electro Transfer unit. The unoccupied sites on the membrane were blocked with blocking buffer (Tris buffer saline (TBS) pH 7.2, containing 0.1 Tween 20, 1% (w/v) western blot grade gelatin and 0.05% triton xl00). The membrane was then incubated with hyperimmune sera (1:100 in

blocking buffer) at 37°C for 1 hour followed by three washes with TBS-Tween 20. The membrane was then incubated at 37°C for 1 hour in diluted antichicken peroxidase labeled (1:5000 in TBS-Tween 20) then washed as before and incubated in freshly prepared substrate solution (10 mg aminoethyle carbazone in 50 ml PBS containing 50ul of 30% H<sub>2</sub>O<sub>2</sub>) for 3 - 4 min. for colour development and visible bands were obtained. The reaction was stopped by washing the membrane with running distilled water.

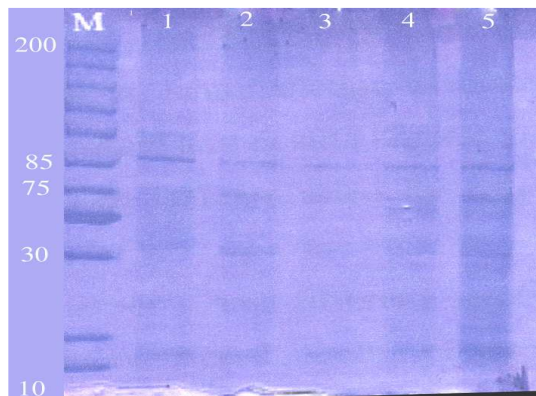
### Results

Only 5 serotypes of *E. coli* represented as O1, O2, O6, O78 and O126 were isolated biochemically identified and serogrouped out of 60 different samples as shown in Table (1)

It is clear that only O1, O6 and O78 serogroups were isolated from the apparently healthy birds while the all 5 recovered serogroups were isolated from either the



**Photo (2).** Western blot analysis of total protein of different *E. coli* serotypes. Lane (1):O1, Lane (2):O2, Lane (3):O6, Lane (4):O78, Lane (5):O126 Lane M: molecular weight protein marker.



**Photo (1).** SDS-PAGE of total protein of the five isolated *E. coli* serotypes. Lane (1):O1, Lane (2):O2, Lane (3):O6, Lane (4):O78, Lane (5):O126 Lane M: molecular weight protein marker.

colisepticaemic or diseased birds.

Regarding the protein profile as shown in the analysis of the SDS-PAGE in Table (2) and Photo (1), there is major common protein patterns of molecular weight 21, 30, 55 and 74 kDa shared in all serotypes also there are other patterns shared in the most of the isolated serotypes like that obtained at 26, 27, 32 and 85 kDa. On the other hand, there were additional protein bands related to each serotypes as shown in Table (2).

Concerning the western blot analysis against *E. coli* antisera as shown in Photo (2), it is clear that there are two major common epitopes fully reacted with the antisera prepared against *E. coli* at the molecular weight of 55 and 74 kDa other reaction appeared and varied from one serotype to another.

### Discussion

The most common clinical manifestation associated with *E. coli* infection are air sacculitis,

pericarditis, septicaemia and death (Vidotto *et al.*, 1990). Also colibacillosis due to virulent strains in chicken is manifested by respiratory disease followed by general infection (Gross, 1984).

Although chicken may harbor many different serotypes in their gastrointestinal tract, in the present study, only a restricted number of serotypes (O1, O2, O6, O78 and O126) have been most commonly found (Table 1). The most predominate serotypes were O1 and O78 where each represent nine isolates out of thirty three followed by O126 (five isolates) then O2 (four isolates). Similar serotypes (O1, O2, and O78) were obtained by Ibrahim (1997). In addition, Peighambari *et.al.* (1995) recorded that the most common serogroups of *E. coli* from avian diseases were O78, O2, and O1 which were associated with septicaemic *E. coli* infection in poultry.

As shown in Table (2) and Photo (1), there are major protein clusters shared in all isolates and appeared at the molecular weight of 21, 30, 55 and 74 kDa. Also other common patterns present in most of the isolates obtained at the molecular weight of 26, 27, 32 and 85 kDa. The same criteria were reported by Lughtenberg *et. al.* (1975) who stated that some major proteins of *E. coli* are predominate under specific growth condition. In addition to the major proteins there are minor proteins which are present in lower copy number and can be detected in SDS-PAGE. Moreover Vidotto *et. al.* (1994) suggested that the protein of approximately 74 kDa is important for the virulence of *E. coli* strains. On the other hand, Fantinatti *et al.* (1994) concluded that, the non-pathogenic strains of *E.coli* lacked two major proteins reported to be correlated either with enterocholin or with aerobactin.

As regards to the immunoreactivity in the western blot analysis, there are two major proteins fully reacted in all serotypes at the molecular weight of 55 and 74 kDa. Vidotto *et. al.* (1994) confirmed these results as they reported that the 74 kDa protein is important and predominate in the *E. coli* strains. It is clearly that the proteins of 55 and 74 kDa are products of a well-conserved gene and major cross reacting antigens within the *E. coli* serotypes. In this respect, it behaves in a manner similar to several major envelope immunogens.

In conclusion all the results of this study clearly show high discriminating potential of the protein profiles analysis in typing of *E. coli* strains isolated from chickens. The

differentiation of the isolates from each other by using protein profiles can provide a reliable additional method to aid in the characterization of the bacteria.

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### الاستراتيجيات الجزيئية لتصنيف وتفرقة العترات الحقلية للميكروب القولوني في الدجاج: (١) فصل وتحليل وتمييز البروتين الكلي للمعزولات

في هذه الدراسة تم عزل وتصنيف خمسة عترات تحت نوعية من الميكروب القولوني في الدجاج وكانت O1, O2, O6, O78, O126 من إجمالي ٣٣ عترة تم عزلها من إجمالي ٦٠ عينة تم أخذها من الدجاج ، بفصل وتحليل البروتين الكلي لهذه المعزولات وجد أنه يوجد تشابه كبير في أربعة بروتينات رئيسية ذات أوزان جزيئية ٢١ ، ٣٠ ، ٥٥ ، ٧٤ كيلو دالتون ، من هذه البروتينات ٥٥ ، ٧٤ كيلو دالتون أثبتت تفاعلي قوى وواضح في اختيار الوسترن بلوت لكل المعزولات، كذلك وجد أن هناك بعض الاختلافات في بعض الأوزان الجزيئية الأخرى والتي تختلف من عترة تحت نوعية إلى أخرى.