















Avian pathogenic *Escherichia coli*: Epidemiology, virulence and pathogenesis, diagnosis, pathophysiology, transmission, vaccination, and control

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Abstract

Avian pathogenic *Escherichia coli* (APEC) causes colibacillosis in poultry; this type of bacteria is an extraintestinal pathogen *E. coli*. Unlike other *E. coli* pathogen groups, the characteristics of APECs cannot be identified by a single group. Serotyping and biotyping are frequently performed for isolates found in colibacillosis infections. The establishment, transmission, and persistence of this pathogenic strain in chicken populations are determined by the intricate interactions of multiple elements that make up the epidemiology of APEC. APEC employs many virulence and pathogenesis factors or mechanisms to infect chickens with colibacillosis. These factors include invasives, protectins, adhesins, iron acquisition, and toxins. In addition, the pathogenicity of APEC strains can be evaluated in 2–4 week-old chicks. The impact of unfavorable environmental conditions has also been documented, despite direct contact being demonstrated to be a significant element in transmission in APEC. Chickens are immunized against colibacillosis using a variety of vaccines. Nevertheless, commercially available vaccinations do not offer sufficient immunity to protect birds from APEC strains. Hatching egg contamination is one of the main ways that APECs spread throughout chicken flocks. Farmers also need to be mindful of storing discarded materials near the manure-watering area, removing them when necessary, and replacing wet materials with dry materials when needed. This review aimed to explain the characteristics, epidemiology, virulence, pathogenesis, diagnosis, pathophysiology, transmission, vaccination, and control of APEC.

Keywords: avian pathogenic *Escherichia coli*, colibacillosis, *Escherichia coli*, poultry, public health.

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Introduction

The well-characterized Gram-negative bacterium *Escherichia coli* is present in upper respiratory and digestive tracts of poultry and mammals [1]. Conversely, a possible risk factor for colibacillosis in poultry is the existence of pathogenic *E. coli* in the environment, digestive tract, or respiratory system [2]. Avian

pathogenic *E. coli* (APEC) causes colibacillosis in poultry; this type of bacteria is an extraintestinal pathogen *E. coli* (ExPEC) [3]. These conditions indicate colibacillosis, cellulitis, omphalitis, pericarditis, perihepatitis, and salpingitis [4]. The environment and the host's stress levels determine disease transmission; organs can still be infected, and colibacillosis can result from less pathogenic strains with fewer virulence genes [5].

APEC is common in Indonesia, and several studies have demonstrated this. The prevalence of APEC in quail was 32.26% [6]. Putri *et al.* [7] reported the prevalence of APEC in broiler chickens at 40%, but Effendi *et al.* [8] reported a higher prevalence of 73.5%. APEC causes quite high economic losses in Indonesia and results in losses of 13.10% of total poultry assets both directly (weight loss, decreased egg production, and increased total mortality) and indirectly (cleaning, disinfection, and labor compensation) if the disease occurs [9].

The numerous virulence characteristics of APEC bacterial strains enable the bacteria to leave the digestive tract and travel to different interior organs, where they can cause illness [9]. High mortality rates, weight loss, decreased egg production, and rotting of poultry carcasses during slaughter and storage are all potential financial losses associated with APEC infections in the worldwide poultry sector [10]. Human foodborne illnesses can be caused by poultry meat. Poultry producers, suppliers, consumers, and public health officials worldwide continue to be concerned about pathogenic *E. coli*, including APEC and other spoilage microbes in poultry [2]. *E. coli* is a common bacterium linked to foodborne diseases in most countries worldwide [11].

Antibiotic use for illness treatment and prevention has increased due to the prevalence of colibacillosis and other diseases in poultry [12]. It is anticipated that this will lead to the development of antibiotic resistance, also known as antimicrobial resistance (AMR), in the chicken farming industry [13]. Antibiotic resistance in APEC is a global public health concern that must be addressed by all government and social sectors. In poultry farming, uncontrolled antibiotic use can lead to antibiotic resistance [14]. Farmers believe that using antibiotics to prevent disease is a low-cost and side-effects-free endeavor, which accounts for the widespread usage of antibiotics without a prescription from a licensed veterinarian [15].

APEC transmission in chickens can be increased by several factors, including unfavorable conditions in poultry houses, high levels of fecal dust and ammonia contamination, subpar or ineffective farming practices, contaminated water and feed, stress, underlying viral diseases, contaminated eggs, insect vectors, cannibalism, and proximity to other animals in poultry farming [16]. The fecal-oral, respiratory, and vaginal pathways through the cloaca are the primary entrance points for APEC (Figure-1) [17]. The most researched forms of colibacillosis are respiratory and fecal-oral. Nevertheless, because cloacal

infection causes outbreaks of salpingitis-peritonitis syndrome, it is equally crucial to investigate this type of infection [18]. Furthermore, infection by antibiotic-resistant strains of APEC will make management extremely challenging.

Raising public awareness of the risk of APEC transmission in poultry farming and the possible harm to public health can decrease and prevent colibacillosis in poultry flocks. This review aims to comprehensively explain APEC as the main causative agent of colibacillosis, starting with its characteristics, epidemiology, virulence, pathogenesis, diagnosis, pathophysiology, transmission, vaccination, and control.

Characteristics

Unlike other *E. coli* pathogen groups, the characteristics of APECs cannot be identified by a single group. The following summarizes a few phenotypic and genotypic traits connected to this class of diseases [19].

Serotyping and biotyping are frequently performed for isolates found in colibacillosis infections. The predominant serogroups of *E. coli* recovered from sick birds are O1, O2, and O78 [19]. Thus, representative serotype strains offer a focal point for deciphering the mechanisms of APEC pathogenicity and for creating and assessing potential vaccines. Given that this dominant serogroup may also be recovered from the feces of birds that appear to be in good health, the hypothesis that the digestive tract may serve as a substantial natural reservoir for APEC and that predisposing factors may be necessary to produce the disease is supported [20]. Many investigations have revealed similarities between commensal *E. coli* and APEC, including serogroup, suggesting that APEC developed from commensal *E. coli* after gaining pathogenic qualities [21–23]. The various features observed include the outer membrane protein profile, pathogenicity profile, and multilocus enzyme electrophoresis within a serotype. To define APEC isolates, tests for motility, hemolysis, lactose fermentation, complement resistance, serotyping, aerobactin and colicin V synthesis, and embryo lethality are occasionally performed, although phenotypes can vary [24].

In APECs, plasmid-mediated horizontal gene transfer induces a great deal of variability. Several potential virulence factors are carried by APEC plasmids, yet some of these factors were discovered in isolates of *E. coli* from chickens that appeared to be in good condition [8, 25]. There is a significant variation in amount, size, and virulence traits of plasmids in both APECs and isolates from healthy chickens [26]. However, other studies did not identify colicin-related genes or APEC-specific plasmid replicons in commensal bacteria [27]. There is little question that the abundance of virulence genes carried by APEC plasmids has contributed to the present classification of APEC genotypes. APEC is more likely than avian fecal *E. coli* to carry genes encoding iron uptake and

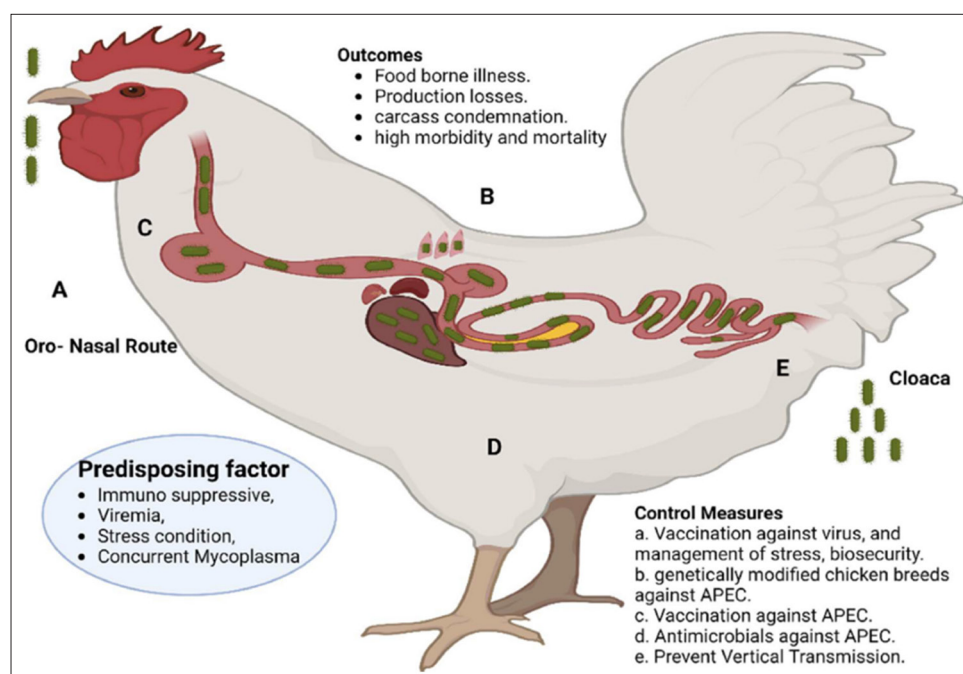


Figure-1: A schematic representation of avian pathogenic *Escherichia coli* infection in chickens after entering through oral, nasal, or cloacal routes [17].

transport mechanisms, which are located in a 94 kb area of the 180 kb ColV plasmid [28]. Numerous studies have demonstrated a connection between ColV plasmids and APEC pathogenicity [3, 9, 29].

The chicken trachea becomes more colonized when an avirulent *E. coli* strain is transformed with a recombinant plasmid (pHK11) expressing colicin V [30]. Recently, a commensal *E. coli* strain was conjugated with two APEC plasmids (large R and ColV), which improved its ability to colonize rodent kidneys, grow in human urine, and destroy chicken eggs [23]. A pair of multiplex polymerase chain reaction (PCR) methods has been recently developed to identify APECs using a set of common characteristics, such as plasmid-borne phenotypes. Strains from sick birds were classified as APEC if they contained at least four of the eight genes included in the study: Iron repressible protein (*irp2*), enteroaggregative toxin (*astA*), temperature-sensitive hemagglutinin (*tsh*), P fimbriae (*papC*), aerobactin (*iucD*), increased serum survival protein (*iss*), colicin plasmid operon genes (*cva/cvi*), and vacuolating autotransporter protein (*vat*), whereas nonpathogenic isolates had none or at most three genes [31].

Four multiplex PCRs were recently developed for virulence genotyping to determine the evolutionary links between ExPEC strains from birds and humans [32]. However, surprisingly, little is known about the gene combinations that are critical for causing systemic infections with APEC. Nonetheless, prior research has demonstrated that Tsh/Pap/Iuc pathotypes, Tsh/Pil/Iuc, and the virulence-related markers. Tsh and Iuc are crucial for APEC [33]. The relationship between APEC strain serogroup and virulence gene patterns is not very strong. The link between several

virulence genes may reflect the existence of distinct subpathotypes or pathotypes within the current APEC group, indicating that the potential subpathotypes may use distinct virulence mechanisms [34]. It is clear that more revisions to the APEC concept are required.

Epidemiology

The establishment, transmission, and persistence of this pathogenic strain in chicken populations are determined by the intricate interactions of multiple elements that constitute the epidemiology of APEC. The epidemiology of APEC includes the crucial component of avian host vulnerability [9]. Some factors affecting the risk of infection include age, immune status, and general health [2]. Although this disease affects all age groups, chicks and stressed birds are especially susceptible to this infection [35]. The environment is crucial for the transmission of APEC. In chicken farming, APEC's survivability is related to tainted water sources, overcrowding, and inadequate sanitation [36]. Thus, robust biosecurity measures are required to mitigate environmental risk factors.

Another component of its epidemiology is the genetic diversity of the APEC strains. A genomic study by Desvaux *et al.* [37] has identified numerous virulence factors and transportable genetic components that support APEC's adaptability and evolution. The significance of continuous surveillance in tracking strain fluctuation was demonstrated by the observed genetic diversity. One important aspect of APEC epidemiology is the horizontal gene transfer across bacteria [38]. A portion of the virulence and antibiotic resistance genes that are transferred among APEC strains can be attributed to cellular genetic factors such as bacteriophages and plasmids [39]. Recognizing

these pathways is essential for anticipating and managing APEC outbreaks.

Ewers *et al.* [40] have contributed to our understanding of the molecular epidemiology of APEC in chicken farming. They also discussed the significance of monitoring genetic alterations and the effects of horizontal gene transfer. Variations in the genetic makeup of the bacteria, environmental factors, and host characteristics complicate the epidemiology of APEC. Wibisono *et al.* [9] demonstrated the importance of continuing monitoring and conducting research to develop efficient management and preventative strategies that safeguard chicken health.

Virulence and pathogenesis

APEC employs many virulence and pathogenesis factors and mechanisms to infect chickens with colibacillosis. These include poisons, iron acquisition mechanisms, invasives, protectins, and adhesins. These factors include attachment to host cells, invasion of host cells, survival within phagocytic cells (macrophages), colonization of tissue, bloodstream persistence, cellular proliferation or replication, lysis, and cell damage, sequestering metals from bodily fluids for growth, resistance to oxidative and environmental stress, serum bactericidal activity, motility, and biofilm formation.

Adhesin

Adhesins are bacterial cell adhesions or surface elements that help bacteria adhere to other surfaces or cells, usually the host in which they live or infect [41]. Adherence is a crucial stage in the pathogenesis or infection of bacteria and is necessary for the organism to colonize a new host [42]. The main factors promoting adherence to APEC are type 1 fimbriae, S fimbriae, and P fimbriae [43]. Some of the genes encoding these fimbriae and additional adhesins, *fimC* (type 1 fimbriae), *fimH*, *papA*, *papEF*, *sfalsfaS* (S fimbriae), *crl*, *papGI*, *papC*, *papGII*, *flgE* (flagellar hook), *papGIII*, *felA* (P fimbriae), *focGE* (F1C fimbriae), *lpfA*, *afaIBC* (afimbriae), *lpf0154* (long polar fimbriae), *lpf0141*, *mat/ecpA* (fimbriillin), *csg* (curli), *yqiG* (putative outer membrane messenger protein), *bmaE* (M hemagglutinin), *iha* (adhesin homolog IrgA), *hra/hrlA/hek* (heat-stable agglutinin), *tsh* (temperature-sensitive hemagglutinin), and *kii* (capsule-encoding K gene) have been reported to occur in APEC [16]. These adhesins regulate macrophage APEC motility, biofilm formation, and survival [44]. In addition, colonization, adhesion, and resistance to environmental stress are facilitated by the fimbriae-encoding gene *yfcO*, whereas adhesion, intracellular survival, and motility are improved by *yadC* [45]. Adhesion, colonization, and biofilm formation are also facilitated by the adhesin autotransporter genes (*aatA*, *aatB*, and *upaB*) [46]. Several more genes (*fdeC*, *fdtA*, *yjhb*, *rldD*, and *ecpR*) were discovered to be in charge of adhesion in chicken and human cell lines by screening random transposon mutants [47].

Invasin

A class of proteins known as invasins is linked to the entry of infectious cells into host cells. Early in infection, invasiveness contributes to the admission of bacteria [48]. Several genes encoding the invasives, *tia* (toxigenic invasion locus), *ibeA* (also called *ibe10*), *ibeB* (invasion protein), and *gimB* (genetic island associated with neonatal meningitis), have been reported in APEC isolates [49]. Furthermore, invasins help APEC survive oxidative damage, colonization, biofilm formation, and proliferation in the host that is brought on by macrophages [9]. The *ibeR* operon regulator plays a role in invasion, resistance to environmental stressors and serum, and the production of virulence genes [50]. Similarly, the suspected invasive gene *ycho* contributes to biofilm formation, motility, adhesion, invasion, and the production of metabolic and membrane proteins [51].

Iron acquisition system

Once bacteria have successfully colonized or attacked their host, iron is a crucial micronutrient for proliferating and multiplying in the host [1]. To absorb iron from bodily fluids, APEC uses a distinct iron acquisition system that consists of transporters and many siderophores, including yersiniabactin, salmochelin, and aerobactin [52]. Several genes that encode for iron uptake and transport systems, *mntH* (iron and manganese transporter), *aerJ* (aerobactin), *iutA*, *sitABCD*, *fyuA* (yersiniabactin), *feoB* (iron ion transporter), *irp2* (iron repressor protein), *iroBCDEN* (salmochelin), *fepC* (ferric enterobactin transporter), *eitABCD* (putative iron transporter), *ireA* (iron-regulated virulence gene), *chuA* (outer membrane hemin receptor), *iucCD*, and *bfr* (bacterioferritin) have been reported in APEC [53]. These siderophores and transporters also mediate the expression of other virulence genes, invasion, resistance to environmental stress, colonization, and persistence of APEC in host cells [54]. Moreover, genes encoding the outer membrane efflux protein (*tolC*), enterobactin production, and transport genes (*entE* and *entS*) work together to promote invasion, colonization, and persistence [55].

Protectin

Bacteria are shielded by protectin from the host's immune system and other harmful environments [56]. In particular, protection offers defense against complement-mediated bactericidal effects in human serum and phagocytic engulfment by macrophages. The lipopolysaccharide (LPS) comprises bacterial capsules, LPS components, and outer membrane proteins [57]. Several genes that encode for protection, *traT* (complement resistance protein), *kfiC-K5* (glycosyl transferase), *iss* (increased serum survival), *kpsMT(II)*, *ompT* (outer membrane protease), *kpsMT(III)*, *kpsMT(K1)*, *neuS*, *neuC*, *neuD* (capsule), and *betA* (choline dehydrogenase), have been reported in APEC [58]. In addition to shielding APECs from host defenses, this protection facilitates

APEC adhesion, intracellular survival, colonization, invasion, and proliferation on the host [59].

In addition, the outer membrane proteins PagP and YbjX contribute to intracellular survival, invasion resistance, and resistance to environmental and serum stress [60]. In a similar vein, OmpA, another protein found on the outer membrane, also helps macrophages survive APEC [61]. LPS production genes, *wzy* (O-antigen polymerase), and *waaL* (O-antigen ligase) genes, aid in adhesion, colonization, motility, invasion, and biofilm formation, as well as intracellular survival and resistance to phagocytosis and environmental stress [62]. Similarly, the lipid A biosynthesis gene *lpxM* (myristoyl transferase) has been implicated in invasion, intracellular survival, colonization, cytokine gene expression control and nitric oxide generation [63]. In the meantime, SOD (superoxide dismutase) encourages the production of biofilms and shields APEC from host responses caused by reactive oxygen species [64].

Toxins

Toxins are biological poisons that render germs more capable of attacking and damaging tissues [65]. Several genes encode for various types of toxins: *astA*, *hlyE* (putative avian hemolysin), *hlyA*, *vat* (vacuolating autotransporter toxin), *cdtB*, *hlyF*, *stx2f* (Shiga toxin variant), *cdtS* (cytolethal distending factor), *sat* (secret autotransporter toxin), *espC* (serine protease), EAST-1 (heat-stable enterotoxin), *pic* (serine protease autotransporter), and *ace4/35* (acetylcholine esterase) have been reported in APEC [66]. In addition, this toxin promotes the development of outer membrane vesicles, agglutination, motility, colonization, biofilm formation, and vacuolization [67].

Diagnosis

Pathogenicity of one-day-old chicks

The most reliable method for identifying virulence in an *E. coli* strain is to perform a pathogenicity test on one-day-old chicks [25]. The virulence of this strain was assessed using the APEC subcutaneous inoculation method on one-day-old embryos or chicks, based on a 50% fatal dose [68]. This technique can be used to isolate APEC from infected chicks.

In addition, the pathogenicity of APEC strains can be evaluated in 2–4-week-old chicks [69]. This technique allows the illness to spread while it is in the field. This approach involves inoculating the sample into the nasopharynx or trachea following an initial challenge with a different agent [67]. This is because *E. coli* is typically caused by secondary infections caused by mycoplasma, infectious bronchitis virus, Newcastle disease, or environmental factors [70–72]. The typical clinical manifestations of colibacillosis include internal organ contamination, hemorrhage, pericarditis, perihepatitis, fibrin in the air sacs, and weight loss [18]. This model verified that the injected strain was pathogenic.

In this scenario, *E. coli* strains immediately injected into air sacs are considered high-performance

models because the triggering agent does not need to be pre-treated [73]. Compared with inoculating the upper respiratory tract, inoculating the air sac permits more uniform lesions typical of colibacillosis [74].

Isolation and biochemistry

Most microorganisms require culture media to grow *in vitro*. Some of these mediums are used to support the development of microorganisms and distinguish one sample from another according to its biochemical properties. Furthermore, because they are used to distinguish closely related organisms, selective culture medium permits the growth of certain diseases and prevents the growth of other diseases.

Levine-modified eosin-methylene blue agar, a differential medium first discovered by Holt-Harris and Teague in 1916, is used to identify and isolate Gram-negative intestinal infections. In organisms that ferment lactose and those that do not, dyes, eosin Y, and methylene blue are used as markers of differentiation in response to lactose and saccharose fermentation [75].

MacConkey agar (MCA) is a selective medium containing bile salts and crystal violet, which prevent the growth of Gram-positive bacteria, particularly enterococci and staphylococci [76]. Gram-negative bacteria grow more effectively with MCA. Furthermore, MCA is frequently used to separate *Salmonella* Typhi from the coliform group as well as to identify and isolate all strains of typhoid, paratyphoid, and dysentery [77].

The isolation of *E. coli* in Sorbitol MCA (SMCA) and SMCA plus cefixime and tellurite media is widely used [78]. This medium can potentially provide false-positive results for bacteria that do not ferment sorbitol, including *Hafnia alvei*, *Morganella morganii*, *Proteus* spp., *Providencia* spp., and *Aeromonas* spp. [79].

E. coli's primary biochemical trait is its ability to ferment maltose, rhamnose, mannitol, xylose, glycerol, glucose, arabinose, mannose, and sorbitol to produce acids and gases. There are differences in the applications of additional sugars such as dulcitol, ornithine, saccharose, adonitol, salicin, raffinose, and arginine [80]. Positive results were observed for the synthesis of lysine, indole, and motility. Tests for oxidase, citrate utilization, urea hydrolysis, gelatin melting, and H₂S (hydrogen sulfide) production are likely to yield negative results [81]. Other tests like Voges Proskauer and methyl red are anticipated to yield positive and negative results, respectively.

The ability of *E. coli* to ferment glucose by releasing gas and acid is its primary trait [82]. The β-galactose enzyme converts lactose into glucose and galactose [83]. This enzyme is used to distinguish *E. coli* from *Shigella* and *Salmonella* species.

Serotype

Within the serogroup, APEC strains are categorized as serotype O: K:H [84]. Somatic antigens

belong to serogroup O, capsular antigen to K1, flagella antigens to serogroup H, and antigens type 1 (F1A), P (F11), and curli fimbriae related to fimbriae [9]. APEC strains include 177 “O” antigens, 100 “K” antigens, and 56 “H” flagellar antigens [59]. In contrast, a different study found that APEC has 167 “O,” 74 “K,” 53 “H,” and 17 fimbriae antigens [23].

Somatic “O” antigen is an endotoxin secreted during the lysis of *E. coli* and is largely composed of polysaccharides [82]. The O serogroups involved in avian colibacillosis are O1, O2, O3, O4, O6, O8, O11, O15, O18, O21, O35, O36, O50, O64, O71, O74, O75, O78, O87, O88, O95, O100, O103, O109, O115, O119, O132, O141, and O152 [85]. APEC’s most prevalent serogroups are O1, O2, O8, O35, and O78 [12].

The adherence of *E. coli* to the cell surface is linked to fimbrial antigen, also known as the “F” antigen [1]. Fimbriae, also known as pili, are associated with the presence of this carbohydrate in cells and exhibit sensitivity or resistance to mannose agglutination [86, 87]. Most APECs have type 1 and curli fimbriae [88]. Curli, type 1 and P fimbriae APEC may contain F1C, which clings to buccal epithelial cells; Dr fimbriae, which persist in kidney tissue; and type 1-like fimbriae [89]. Avian *E. coli* 1 fimbriae (AC/1 fimbriae) are a novel fimbrial group of the fimbrial adhesin family [22].

Another antigen that codes for *E. coli* is flagellin, which is sometimes known as the “H” antigen [90]. The majority of APECs include flagellar antigens, which indicate APEC motility. Flagella offers intestinal mucus penetration, invasion, colonization, and persistence in a specific-pathogen-free chicken model [91]. The virulence of *E. coli* is not linked to this antigen.

The major APEC serotype contains the polysaccharide polysialic acid, which comprises the K capsule antigen [92]. This is related to virulence. There is a connection between this antigen and extraintestinal infections. In the avian isolate O2:K1, K antigen was found to be a virulence factor [93]. Colibacillosis frequently involves serogroups O1:K1, O2:K1, and O78:K80 [94]. The serotype O86:K61 has not yet been isolated from commercial poultry. This illness has only been linked to diarrheal sickness in horses, pigs, and calves [95]. It has also been linked to cellulitis in broiler chickens. The human *E. coli* clonal group that was recovered from cases of newborn meningitis, urinary tract infections, and septicemia included the APEC serotypes O1:K1, O2:K1, and O18:K1, which are potentially zoonotic strains [96]. Humans and animals can contract the UroPathogenic *E. coli* strain serogroup O25b: H4-ST131 (sequencing type 131), which was recently found in chickens [97].

Enzyme-linked immunosorbent assay (ELISA)

ELISA is widely used in veterinary medicine as a diagnostic tool and quality control measure to detect

specific pathogens like *E. coli* in food products [98]. For research, this technique is also employed as an official veterinary diagnostic test to identify particular antigens or antibodies in production animals (such as those associated with Avian Influenza, Newcastle disease, Aujeszky’s disease, and classical swine fever) [99].

Many commercial ELISA kits are capable of detecting *E. coli*. Some of them detect it in food products: SafePath® *E. coli* O:157 Microwell ELISA® (SafePath®, USA), REAGENT™ *E. coli* O157:H7 Elisa Test Kit (REAGENT™, USA), and 3M™ TECRA™ *E. coli* O:157 (3M™ TECRA™, Germany) [85, 87]. This particular anti-*E. coli* O:157 antibody is used in a sandwich ELISA test [96]. Abnova® Laboratories has a double antibody (sandwich) ELISA that uses the anti-*E. coli* O:157 antibody; BIO K 345 *E. coli* F5 ELISA Kit® from Bio-X Diagnostics® detects F5 pilus antigen (K99) (Bio-X Diagnostics, Belgium) antibodies from *E. coli*; and DAI® *E. coli* O:157 captures the antigen from stool samples [85]. The sandwich enzyme immunoassay method known as the *E. coli* protein (*E. coli* P) ELISA Kit® from MyBiosource® (MyBiosource®, USA) is used to measure the amount of *E. coli* P in serum, plasma, and tissue homogenates [100].

Molecular detection

The expansion of molecular biology tools to evaluate the genetic variability of many strains of bacteria, such as *E. coli*, occurred due to the discovery that prokaryotic genomes contain repetitive sequences: Repetitive extragenic palindrome, palindromic unit sequences, and enterobacterial repetitive intergenic consensus sequences [101]. The PCR reaction produces an amplified band pattern unique to each strain; subsequent use of particular primers that are homologous to this region [102].

The *chuA*, *yjaA*, and an unidentified DNA fragment known as TSPE4.C2 were detected using a quick triplex PCR technique, which was used to identify the phylogenetic group of *E. coli* strains [103]. This technique yielded 230 strains previously collected using other reference methods.

APEC research has involved the use of numerous molecular typing techniques; however, none of these techniques have identified specific genotypes. A multiplex PCR panel that targets the plasmid-borne genes *iutA*, *hlyF*, *iss*, *iroN*, and *ompT*, which are linked to extremely dangerous APEC strains [104]. Multiplex PCR was used to amplify 994 avian *E. coli* samples for these five genes [104]. These five genes allowed PCR to distinguish APEC strains from *E. coli* isolates from bird feces [105].

Pathophysiology

Omphalitis and yolk sac infection

One of the most frequent reasons why chickens die in their first few days of life is omphalitis, or inflammation of the navel [106]. The bacteria most

commonly linked to these deaths are *E. coli*. Eggs are more vulnerable to fecal contamination. Nevertheless, germs can enter the bloodstream through the intestines. For example, *E. coli* infections can enter through an unhealed navel and contaminate the egg yolk sac [107]. Clinical indicators of omphalitis include edema, erythema, hardness of the skin around the navel, yolk sac, and swelling [108].

Cellulitis

This illness is known as cellulitis and is caused by subcutaneous inflammation in the thighs and lower abdomen [109]. *E. coli* is the pathogen responsible for this condition. Cellulitis in chickens does not show any overt symptoms, and it is only observed in slaughterhouses, where it spreads to chickens kept in cages [110]. The condition is typified by superficial skin lesions arising from bird and bedding that come into close contact [111]. Cellulitis-related lesions in hens are believed to be caused by a compromised innate immune response, especially in heterophils [112].

Swollen head syndrome (SHS)

SHS is characterized by acute or subacute cellulitis affecting the head's periorbital region and surrounding subcutaneous tissue [83]. The first documented case of SHS was linked to coronavirus and *E. coli* infection and reported in South Africa [113]. An inflammatory exudate produced by bacteria under the skin, generally *E. coli*, causes swelling in the head, which is followed by viral infections, such as infectious bronchitis virus and avian pneumovirus [114].

Acute vaginitis

This condition is more common in turkey and, shortly after insemination, results in acute and deadly vaginitis [115]. The most common cause of *E. coli* infection is hymen perforation, which can lead to internal laying, protrusion of the intestines and digestive tract, peritonitis, vaginitis, and egg binding [116].

Salpingitis/peritonitis

Salpingitis affects laying hens and broiler chickens and is characterized by *E. coli*-induced oviduct inflammation [117]. Salpingitis may be linked to infections in the fallopian tubes, vents, or entire body [118]. In hatcheries, asymptomatic infections can cause decreased egg production and higher embryo mortality, even when not noticeable [119]. Peritonitis is characterized by caseous discharge and significant inflammation in the body cavity [120].

Orchitis and epididymitis

Genital tract infections in male chickens can result in orchitis, similar to salpingitis in hens [121]. The frequency of *E. coli* infections is increasing, resulting in bigger, harder, irritated, and irregular testicles [122].

Colisepticemia

High morbidity and mortality rates are associated with colisepticemia in laying hens and broiler chickens [35]. The prevalence of additional secondary

infectious agents, such as viral infections and mycoplasma, is correlated with mortality rates [84, 123]. Approximately 80% of colibacillosis cases are associated with serotypes O2, O78, and O1, the most prevalent serotypes worldwide [124]. The primary cause of colibacillosis is high AMR, which renders antibiotic treatment useless [125]. Nevertheless, laying hens, broilers, and turkeys are susceptible to this disease because the vaccinations currently on the market do not offer sufficient immunity to protect poultry against it.

Coligranuloma (Hjarre's disease)

An uncommon type of systemic colibacillosis that can affect turkeys, laying hens, and broiler chickens is called coligranuloma [39]. Normally, this disease affects only some birds, but if it spreads to all birds, it can result in a significant fatality rate of up to 75% [74]. Granulomas are the hallmark of this condition and can be found in the liver, duodenum, cecum, and mesentery, among other organs [121]. Common lesions of coligranulomas resemble leukosis tumors [126]. Hepatic coagulation, some heterophils, and large cell counts were noted [85]. Pyogranuloma typhlitis and hepatitis in turkey are linked to coligranuloma [127].

Transmission

Housing conditions, ventilation, and stress

The impact of unfavorable environmental conditions has also been documented despite direct contact being demonstrated to be a significant element in transmission in APEC [128, 129]. The respiratory systems of birds are harmed by inadequate ventilation, high dust concentrations, or other chemical vapors in chicken houses [130]. Scratches or wounds in the injured respiratory tract can allow APEC to enter, leading to the development of airsacculitis, polyserositis, and potential septicemia [88]. Excessive ammonia concentrations can weaken the cilia lining the epithelium, impairing the bird's ability to filter dust and dangerous pathogens from its respiratory system [131]. High ammonia levels are typical because ventilation is decreased in colder climates to reduce heating expenses. Reduced ventilation leads to moist air inside the poultry house, which raises the water content in the droppings and provides a perfect environment for bacteria to release large amounts of ammonia and break down uric acid [132]. This can damage the bird's respiratory system and increase the possibility of APEC transmission.

Cleaning the poultry house while the birds are in their cages is not recommended because it can release feces-contaminated dust. Inhalation of contaminated fecal dust is a confirmed method of APEC transmission in poultry flocks [133]. Dust serves as a place for *E. coli* to grow. For this reason, *E. coli* has been identified in livestock dust that has been kept for up to 35 years at 4°C [134]. Furthermore, a preliminary investigation on poultry farms found that most poultry

houses had a lot of dust on the walls and windows, which could raise the risk of APEC transmission in the bird house [16]. According to additional research, dust in and around chicken buildings is believed to be a significant factor in the spread of APEC [135]. Therefore, well-ventilated poultry barns are believed to prevent the spread of APEC and other diseases in chickens.

Poor husbandry practices, high chicken densities, and the advent of sexual maturity have all been demonstrated to stress chickens, which raises the risk of infection [132]. Increased housing density has also been linked to increased rates of germs, including *Salmonella* Enteritidis, in the intestines of chickens [136]. The same might also hold true for APEC transmission, although more investigation is necessary because various bacteria might have distinct growth needs.

Contaminated water, feed, and eggs

Water could play a significant role in APEC's dissemination to poultry. Pathogenic *E. coli* serotypes can be introduced into chicken flocks through contaminated well water, leading to APEC transmission [2]. The spread of APEC can be assisted by urban chicken farms that use recycled wastewater [137]. *E. coli* bacteria isolated from final effluent released from two wastewater treatment plants in the Eastern Cape Province, South Africa, were multidrug-resistant [138]. If water is used for chicken rearing, the final effluent discharge may represent an equally significant risk [139]. There is a possibility that contamination of feed and feed ingredients could result in the emergence of novel disease strains. In addition, there is evidence that the type of food fed to hens affects their intestinal microbiota [140]. It has been demonstrated that some food ingredients encourage the growth of specific gastrointestinal bacteria while inhibiting the growth of other bacteria.

Feces-contaminated eggs can cause *E. coli* infection in the yolk sac during hatching, which is typically linked to high rates of chick mortality [141]. Poultry can transmit typhus and pullorum from one generation to another through contaminated eggs [142]. Although there may be differences in the risk factors for APEC transmission between chickens and typhoid and pullorum, it is believed that tainted eggs may increase the likelihood of APEC transmission [143]. Reduced incidence of avian colibacillosis has been demonstrated by fumigating eggs within 2 h after laying and discarding eggs with cracked shells or stained by feces [74].

Underlying chicken disease

Colibacillosis frequently coexists with other illnesses, such as *Mycoplasma* and respiratory virus infections, making diagnosis and treatment challenging for farmers [1]. Conversely, healthy hens have a strong immune system and are naturally resistant to *E. coli*, which is present naturally [144]. Livestock can be more vulnerable to APEC infections if they have immunity abnormalities brought on by acute

infections, specifically infectious bursal disease, adenovirus Newcastle disease, reovirus, Marek's disease, and infectious bronchitis [145]. Furthermore, the role of underlying illnesses and serum antibodies against viruses such as Newcastle disease and infectious bronchitis virus in the spread of APEC are currently being studied.

Vectors of disease and cannibalism

Both domesticated and farmed poultry can transmit APECs to other animals. Surveys have revealed that prey birds frequently enter chicken homes, raising the possibility of infection transmission, including APEC [12]. Rodents, flies, chicken mites, foreign objects, and rodents can also serve as APEC vectors in addition to humans [146]. It has been shown that bacteria identical to those found in animal waste and multidrug-resistant clonal lineages can be transferred to different substrates by flies and other insects [147]. According to reports, flies gathered in poultry barns contained *E. coli* that produce extended-spectrum beta-lactamase, which raises the possibility of APEC transmission even more [148].

In addition, rodent droppings may be the primary cause of APEC. Antibiotic-resistance genes can be transferred between bacterial strains by rodent droppings [146]. Furthermore, it has been demonstrated that flea infestations in poultry flocks stress chickens. Therefore, it has been hypothesized that flies can transfer APEC from lesions in sick hens to healthy chickens and that fleas or other parasites feed on the blood of chickens [16]. In this study, insect control was suggested as a crucial herd management strategy to lower the spread of APEC. In addition, it was discovered that cannibalism and perking wounds are two ways in which APEC spreads among chickens [26].

Proximity to other animals, poultry farming, and poultry density

The risk factor for the transmission of poultry diseases is the distance between poultry farms. Reducing the number of chicken farms in a region and the number of chickens on each farm is one of the most effective ways to prevent colibacillosis [142]. If biosecurity regulations are not strictly implemented, backyard flocks of chickens that usually coexist with wild birds should be separated from commercial chickens [149]. It was reported that between 2002 and 2003, private flocks in the United States experienced an outbreak of exotic Newcastle disease, which later spread to commercial chicken flocks [150]. Backyard flocks can spread zoonoses and other highly contagious infectious diseases to commercial poultry and are frequently exposed to avian influenza. The possibility of APEC transmission in poultry is thought to be increased by interactions between maintained birds and other chicken species, particularly backyard flocks. However, APEC transmission differs from that of avian influenza and Newcastle disease viruses [151].

Vertical transmission

E. coli oviduct infection is a common cause of mortality in laying hens and broilers and egg production. Vertical transmission has previously been demonstrated as a means by which bacteria with AMR genes can spread [152]. Vertical transmission has also been demonstrated in other bacterial species, such as *Salmonella enterica* and *Enterococcus faecalis* [153]. For the 1st time, Giovanardi *et al.* [154] documented vertical transmission of APECs from parent to offspring; previously, *E. coli* studies only examined outbreaks.

Peterson *et al.* [155] and Bortolaia *et al.* [152] later reported the transmission of *E. coli* resistant to fluoroquinolones, nalidixic acid, and ampicillin from parent to broiler. Vertical transmission of APEC has been documented by isolating *E. coli* clones from salpingitis-peritonitis lesions in broiler broodstock [156]. The transmission of *E. coli* to day-old chicks is linked to increased risk [157].

Vaccination

Chickens are immunized with various vaccines to prevent colibacillosis. Vaccines are available in subunit, recombinant, inactivated, and live attenuated forms. Nevertheless, commercially available vaccinations do not provide sufficient immunity to protect birds from APEC strains [158].

Numerous efforts are being made to identify the best and most immunogenic *E. coli* vaccination strategies for chickens. A previous by Qiu *et al.* [159] employed bacterins; currently, recombinant or subunit vaccinations are popular. The primary issue here is that this vaccine can immunize birds in three different ways, regardless of whether it is a recombinant, attenuated, inactivated, or subunit vaccine. In summary, the vaccine should provide cross-immunity against several serotypes present in APEC. This can be achieved using various delivery methods, including food, drink, *in ovo*, and spraying [160]. This technique enables the widespread use of vaccination vaccines in chicken houses. Not less significant than the previously listed elements, the vaccine for APEC must immunize broiler chickens against APEC strains by the time they are 21 days old, as this is the critical period at which the birds can become infected [161].

Studies have demonstrated the benefits of vaccine use [158, 159]. Live vaccines in broilers reduce antibiotic use. Spray vaccination of day-old broilers against *E. coli* reduced the number of *E. coli* isolates from internal organs in the 6th week of life. *E. coli* isolates from vaccinated birds were more susceptible to antimicrobials. Vaccination of broilers against *E. coli* should be considered in routine immunoprophylaxis [162, 163]. Autogenous *E. coli* vaccines are widely used in the field to prevent *E. coli* peritonitis syndrome in laying hens. Based on the results of the study, groups of laying hens that had been vaccinated intramuscularly at 14–18 weeks of age with an

inactivated vaccine formulated either as an aqueous suspension or as a water-in-oil emulsion were challenged homologously or heterologously by aerosol at 30 weeks of age. Vaccination has been shown to have no effect on body growth, and both types of vaccines induced (almost) complete protection against homologous challenge [164].

Control

Hatching egg contamination is one of the main ways that APECs spread throughout chicken flocks [165]. Frequent precautions should be taken to prevent contamination. To maintain the cleanliness of nest materials, it is necessary to regularly gather eggs, dispose of eggs that are contaminated with dust and excrement on the ground, and disinfect or fumigate eggs as soon as they are laid, ideally within 2 h [36]. These actions assist in lowering APEC transmission. APEC can be decreased or eliminated by sanitizing shell surfaces [158]. Cleaning agents work better when applied by electrostatic spraying [166]. Irradiation with UV light is another efficient strategy that can minimize or remove APEC and other infections while not interfering with incubation or affecting the hatchability of eggs [158]. Care and handling of the contaminated eggs must be performed as closely as possible during the incubation and hatching phases, as broken eggs could spread the infection to other chicks.

There is also a possibility that the hatchery's equipment and handling staff could contaminate other chicks [2]. The vulnerable stage of the egg is until it hatches. During incubation and hatching, several recommendations for prevention and dissemination strategies exist. By opening the incubator to the outside air and, if possible, using various configurations, cross-contamination and losses can be reduced [167]. Hatcheries can contaminate people who are exposed to APEC on farms or in other hazardous areas [20]. The chicks that could be exposed to APEC should be fed frequently and kept warm.

A balanced meal high in selenium, protein, and Vitamins A and E generally increases the probability of survival in poultry [168]. However, since selenium inhibits antibody formation and causes cellulitis and colibacillosis, excessive consumption of this mineral in food can also be harmful [169]. The nutritional value of chickens is directly correlated with the severity of colibacillosis. It has been demonstrated that a feeding regimen based on alternating days is more effective for curing APEC infections in birds than regular feeding [170].

Various aspects need to be taken into account to lower the rate of APEC transmission in the digestive tract and feces: Rat droppings are a source of APEC pathogen transmission, contaminated water might contain APEC, and pelleted feed has less APEC than ground feed [3]. APEC is removed by heat during the pelletization process. Another attempt to eradicate

APEC infections involves supplementing feed with 5%–10% egg yolk powder [85]. Contamination can come from water sources. It is recommended that chlorination systems and nipple irrigation be employed to lessen the transmission of APEC through water [9]. These steps lower the incidence of airsacculitis and colibacillosis. Another strategy for eliminating APEC strains from chick intestines is competitive exclusion. It has been reported that several competitive exclusion techniques, such as the use of *Bacillus subtilis* spores or commercial competitive exclusion products, provide resistance to chick microflora [171].

Maintaining adequate air quality and bedding is recommended to prevent colibacillosis from infecting the flock [172]. Proper cage ventilation reduces the amount of bacteria exposed while maintaining low levels of dust and ammonia [36]. Due to the high concentrations of dust and ammonia in the poultry environment, APEC can attach itself to these particles and be swallowed by chickens, leading to respiratory system infections [173].

APECs can endure and proliferate in moist waste. To minimize the moisture content of litter, approximately 100 feet/min of proper air velocity is required. At this rate, the garbage stays dry, lessening APEC growth [16]. Farmers should use caution when irrigation because water leaks can dampen litter and alter the conditions in which APEC breeds [2]. Farmers also need to be mindful of storing discarded materials near the manure-watering area, removing them when necessary, and replacing wet materials with dry materials when necessary.

APEC infections in poultry are frequently managed with antibiotics [174]. Numerous antibiotics from various classes, including tetracyclines (tetracycline, oxytetracycline, and chlortetracycline), sulfonamides (sulfadimethoxine, trimethoprim, sulfadiazine, sulfamethazine, sulfaquinoxaline, and ormetoprim), aminoglycosides (apramycin, gentamicin, neomycin, and spectinomycin), penicillins (amoxicillin and ampicillin), cephalosporins (ceftiofur), quinolones (danofloxacin, sarafloxacin, and enrofloxacin), polymyxins (colistin), chloramphenicols (florfenicol), macrolides (erythromycin), and lincosamide (lincomycin), have all been used in the poultry industry globally to control APEC infections. However, APEC is resistant to several antibiotics, suggesting that their use will be difficult in the future [16, 175–179].

Conclusion

The most prevalent bacterial infection of chickens, APEC, is causing the global poultry sector to suffer significant financial losses. Animal health can be improved by efficient APEC control. Poultry systemic infection is caused by the coordinated action of several APEC virulence and pathogenesis factors. Given the serious problem of antibiotic resistance and the increasing potential for human infection by bacteria and genes resistant to antibiotics; it is necessary

to develop antibacterials. The creation of antibacterials exclusively for veterinary use that do not exhibit cross-resistance to existing antibiotics may offer a future solution to the serious issue of antibiotic resistance and the high risk of the spread of bacteria and genes resistant to these antibiotics. In addition, an APEC vaccine that offers cross-protection against several APEC serotypes is required. The pathophysiology and virulence mechanisms of APEC should be studied to identify potential novel vaccines.

Authors' Contributions

ARK, IBM, SW, and RR: Drafted the manuscript. IBM, SMY, and AW: Revised and edited the manuscript. DAA, KHPR, IF, and MKJK: Prepared and critically checked the manuscript. SCR, KAF, and AHF: Edited the references. All authors have read and approved the final manuscript.

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Competing Interests

The authors declare that they have no competing interests.

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References

- Pokharel, P., Dhakal, S. and Dozois, C.M. (2023) The diversity of *Escherichia coli* pathotypes and vaccination strategies against this versatile bacterial pathogen. *Microorganisms*, 11(2): 344.
- Joseph, J., Zhang, L., Adhikari, P., Evans, J.D. and Ramachandran, R. (2023) Avian pathogenic *Escherichia coli* (APEC) in broiler breeders: An overview. *Pathogens*, 12(11): 1280.
- Newman, D.M., Barbieri, N.L., de Oliveira, A.L., Willis, D., Nolan, L.K. and Logue, C.M. (2021) Characterizing avian pathogenic *Escherichia coli* (APEC) from colibacillosis cases, 2018. *PeerJ*, 9(1): e11025.
- Sharma, V., Jakhar, K.K., Nehra, V. and Kumar, S. (2015) Biochemical studies in experimentally *Escherichia coli* infected broiler chicken supplemented with neem (*Azadirachta indica*) leaf extract. *Vet. World*, 8(11): 1340–1345.
- Meguenni, N., Chanteloup, N., Tourtereau, A., Ahmed, C.A., Bounar-Kechih, S. and Schouler, C. (2019) Virulence and antibiotic resistance profile of avian *Escherichia coli* strains isolated from colibacillosis lesions in central of Algeria. *Vet. World*, 12(11): 1840–1848.
- Prihtiyantoro, W., Khusnan, K., Slipranata, M. and Rosyidi, I. (2019) Prevalence of avian pathogenic *Escherichia coli* (APEC) strains causing colibacillosis in quail [Prevalensi strain avian pathogenic *Escherichia coli* (APEC) penyebab kolibasilosis pada burung puyuh]. *J. Sains. Vet.*, 37(1): 69.
- Putri, M.F.R., Kendek, I.A., Wibisono, F.J., Effendi, M.H.,

- Rahardjo, D., Tyasningsih, W. and Ugbo, E.N. (2023) Molecular detection of iron gene on multidrug-resistant avian fecal *Escherichia coli* isolated from broiler on traditional markets, Surabaya, Indonesia. *Biodiversitas*, 24(12): 6454–6460.
8. Effendi, M.H., Faridah, H.D., Wibisono, F.M., Wibisono, F.J., Nisa, N. and Fatimah dan Ugbo, E.N. (2022) Detection of virulence factor encoding genes on *Escherichia coli* isolated from broiler chicken in Blitar District, Indonesia. *Biodiversitas*, 23(7): 3437–3442.
 9. Wibisono, F.J., Effendi, M.H. and Wibisono, F.M. (2022) Occurrence, antimicrobial resistance, and potential zoonosis risk of avian pathogenic *Escherichia coli* in Indonesia: A review. *Int. J. One Health*, 8(2): 76–85.
 10. Landman, W.J.M. and van Eck, J.H.H. (2015) The incidence and economic impact of the *Escherichia coli* peritonitis syndrome in Dutch poultry farming. *Avian Pathol.*, 44(5): 370–378.
 11. Syahrul, F., Wahyuni, C.U., Notobroto, H.B., Wasito, E.B., Adi, A.C. and Dwirahmadi, F. (2020) Transmission media of foodborne diseases as an index prediction of diarrheagenic *Escherichia coli*: Study at elementary school, Surabaya, Indonesia. *Int. J. Environ. Res. Public Health*, 17(21): 8227.
 12. Jeong, J., Lee, J.Y., Kang, M.S., Lee, H.J., Kang, S.I., Lee, O.M., Kwon, Y.K. and Kim, J.H. (2021) Comparative characteristics and zoonotic potential of avian pathogenic *Escherichia coli* (APEC) isolates from chicken and duck in South Korea. *Microorganisms*, 9(5): 946.
 13. Hedman, H.D., Vasco, K.A. and Zhang, L. (2020) A review of antimicrobial resistance in poultry farming within low-resource settings. *Animals*, 10(8): 1264.
 14. Habiba, U.E., Khan, A., Mmbaga, E.J., Green, I.R. and Asaduzzaman, M. (2023) Use of antibiotics in poultry and poultry farmers- a cross-sectional survey in Pakistan. *Front. Public Health*, 11(1): 1154668.
 15. Ozturk, Y., Celik, S., Sahin, E., Acik, M.N. and Cetinkaya, B. (2019) Assessment of farmers' knowledge, attitudes and practices on antibiotics and antimicrobial resistance. *Animals*, 9(9): 653.
 16. Kathayat, D., Lokesh, D., Ranjit, S. and Rajashekara, G. (2021) Avian pathogenic *Escherichia coli* (APEC): An overview of virulence and pathogenesis factors, zoonotic potential, and control strategies. *Pathogens*, 10(4): 467.
 17. Nawaz, S., Wang, Z., Zhang, Y., Jia, Y., Jiang, W., Chen, Z., Yin, H., Huang, C. and Han, X. (2024) Avian pathogenic *Escherichia coli* (APEC): Current insights and future challenges. *Poult. Sci.*, 103(12): 104359.
 18. Apostolakos, I., Laconi, A., Mughini-Gras, L., Yapicier, Ö.Ş. and Piccirillo, A. (2021) Occurrence of colibacillosis in broilers and its relationship with avian pathogenic *Escherichia coli* (APEC) population structure and molecular characteristics. *Front. Vet. Sci.*, 8(1): 737720.
 19. Schouler, C., Schaeffer, B., Brée, A., Mora, A., Dahbi, G., Biet, F., Oswald, E., Mainil, J., Blanco, J. and Moulin-Schouleur, M. (2012) Diagnostic strategy for identifying avian pathogenic *Escherichia coli* based on four patterns of virulence genes. *J. Clin. Microbiol.*, 50(5): 1673–1678.
 20. Levy, S., Islam, M.S., Sobur, M.A., Talukder, M., Rahman, M.B., Khan, M.F.R. and Rahman, M.T. (2020) Molecular detection of avian pathogenic *Escherichia coli* (APEC) for the first time in layer farms in Bangladesh and their antibiotic resistance patterns. *Microorganisms*, 8(7): 1021.
 21. Kazibwe, G., Katami, P., Alinaitwe, R., Alafi, S., Nanteza, A. and Nakavuma, J.L. (2020) Bacteriophage activity against and characterisation of avian pathogenic *Escherichia coli* isolated from colibacillosis cases in Uganda. *PLoS One*, 15(12): e0239107.
 22. Lymberopoulos, M.H., Houle, S., Daigle, F., Léveillé, S., Brée, A., Moulin-Schouleur, M., Johnson, J.R. and Dozois, C.M. (2006) Characterization of Stg fimbriae from an avian pathogenic *Escherichia coli* O78:K80 strain and assessment of their contribution to colonization of the chicken respiratory tract. *J. Bacteriol.*, 188(18): 6449–6459.
 23. Solà-Ginés, M., Cameron-Veas, K., Badiola, I., Dolz, R., Majó, N., Dahbi, G., Viso, S., Mora, A., Blanco, J., Piedra-Carrasco, N., González-López, J.J. and Migura-García, L. (2015) Diversity of Multi-drug resistant avian pathogenic *Escherichia coli* (APEC) causing outbreaks of colibacillosis in broilers during 2012 in Spain. *PLoS One*, 10(11): e0143191.
 24. Wang, Z., Zheng, X., Guo, G., Hu, Z., Miao, J., Dong, Y., Xu, Z., Zhou, Q., Wei, X., Han, X., Liu, Y. and Zhang, W. (2022) O145 may be emerging as a predominant serogroup of Avian pathogenic *Escherichia coli* (APEC) in China. *Vet. Microbiol.*, 266(1): 109358.
 25. Widodo, A., Effendi, M.H. and Khairullah, A.R. (2020) Extended-spectrum beta-lactamase (ESBL)-producing *Escherichia coli* from livestock. *Sys. Rev. Pharm.*, 11(7): 382–392.
 26. Mellata, M., Ameiss, K., Mo, H. and Curtiss, R. (2010) Characterization of the contribution to virulence of three large plasmids of avian pathogenic *Escherichia coli* chi7122 (O78:K80:H9). *Infect. Immun.*, 78(4): 1528–1541.
 27. Cummins, M.L., Li, D., Ahmad, A., Bushell, R., Noormohammadi, A.H., Wijesurendra, D.S., Stent, A., Marenda, M.S. and Djordjevic, S.P. (2023) Whole genome sequencing of avian pathogenic *Escherichia coli* causing bacterial chondronecrosis and osteomyelitis in Australian poultry. *Microorganisms*, 11(6): 1513.
 28. Barbieri, N.L., Pimenta, R.L., de Melo, D.A., Nolan, L.K., de Souza, M.M.S. and Logue, C.M. (2021) mcr-1 identified in fecal *Escherichia coli* and avian pathogenic *E. coli* (APEC) from Brazil. *Front. Microbiol.*, 12(1): 659613.
 29. Reid, C.J., Cummins, M.L., Börjesson, S., Brouwer, M.S.M., Hasman, H., Hammerum, A.M., Roer, L., Hess, S., Berendonk, T., Nešporová, K., Haenni, M., Madec, J.Y., Bethe, A., Michael, G.B., Schink, A.K., Schwarz, S., Dolejska, M. and Djordjevic, S.P. (2022) A role for ColV plasmids in the evolution of pathogenic *Escherichia coli* ST58. *Nat. Commun.*, 13(1): 683.
 30. Roland, K., Karaca, K. and Sizemore, D. (2004) Expression of *Escherichia coli* antigens in *Salmonella* Typhimurium as a vaccine to prevent airsacculitis in chickens. *Avian Dis.*, 48(3): 595–605.
 31. Amer, M.M., Mekky, H.M., Fedawy, H.S., El-Shemy, A., Bosila, M.A. and Elbayoumi, K.M. (2020) Molecular identification, genotyping of virulence-associated genes, and pathogenicity of cellulitis-derived *Escherichia coli*. *Vet. World*, 13(12): 2703–2712.
 32. Xia, F., Cheng, J., Jiang, M., Wang, Z., Wen, Z., Wang, M., Ren, J. and Zhuge, X. (2022) Genomics analysis to identify multiple genetic determinants that drive the global transmission of the pandemic ST95 lineage of extraintestinal pathogenic *Escherichia coli* (ExPEC). *Pathogens*, 11(12): 1489.
 33. Fujimoto, Y., Inoue, H., Kanda, T., Ijiri, M. and Uemura, R. (2021) Virulence-associated gene profiles of *Escherichia coli* isolated from chickens with colibacillosis in Japan and their correlation with pathogenicity in chicken embryos. *Avian Dis.*, 65(3): 401–405.
 34. Cordoni, G., Woodward, M.J., Wu, H., Alanazi, M., Wallis, T. and La Ragione, R.M. (2016) Comparative genomics of European avian pathogenic *E. coli* (APEC). *BMC Genomics*, 17(1): 960.
 35. Abdelhamid, M.K., Hess, C., Bilic, I., Glösmann, M., Rehman, H.U., Liebhart, D., Hess, M. and Paudel, S. (2024) A comprehensive study of colisepticaemia progression in layer chickens applying novel tools elucidates pathogenesis and transmission of *Escherichia coli* into eggs. *Sci. Rep.*, 14(1): 8111.
 36. Gržinić, G., Piotrowicz-Cieślak, A., Klimkowicz-Pawlas, A., Górny, R.L., Ławniczek-Walczyk, A., Piechowicz, L.,

- Olkowska, E., Potrykus, M., Tankiewicz, M., Krupka, M., Siebielec, G. and Wolska, L. (2023) Intensive poultry farming: A review of the impact on the environment and human health. *Sci. Total Environ.*, 858(Pt 3): 160014.
37. Desvaux, M., Dalmasso, G., Beyrouthy, R., Barnich, N., Delmas, J. and Bonnet, R. (2020) Pathogenicity factors of genomic islands in intestinal and extraintestinal *Escherichia coli*. *Front. Microbiol.*, 11(1): 2065.
 38. Young, M.M., de Oliveira, A.L., Nolan, L.K., Barbieri, N.L. and Logue, C.M. (2022) Identification of novel genes involved in the biofilm formation process of avian pathogenic *Escherichia coli* (APEC). *PLoS One*, 17(12): e0279206.
 39. Feng, A., Akter, S., Leigh, S.A., Wang, H., Pharr, G.T., Evans, J., Branton, S.L., Landinez, M.P., Pace, L. and Wan, X.F. (2023) Genomic diversity, pathogenicity and antimicrobial resistance of *Escherichia coli* isolated from poultry in the southern United States. *BMC Microbiol.*, 23(1): 15.
 40. Ewers, C., Janssen, T., Kiessling, S., Philipp, H.C. and Wieler, L.H. (2004) Molecular epidemiology of avian pathogenic *Escherichia coli* (APEC) isolated from colisepticemia in poultry. *Vet. Microbiol.*, 104(1–2): 91–101.
 41. Di Martino, P. (2018) Bacterial adherence: Much more than a bond. *AIMS Microbiol.*, 4(3): 563–566.
 42. Stones, D.H. and Krachler, A.M. (2016) Against the tide: The role of bacterial adhesion in host colonization. *Biochem. Soc. Trans.*, 44(6): 1571–1580.
 43. Mellata, M., Dho-Moulin, M., Dozois, C.M., Curtiss, R., Lehoux, B. and Fairbrother, J.M. (2003) Role of avian pathogenic *Escherichia coli* virulence factors in bacterial interaction with chicken heterophils and macrophages. *Infect. Immun.*, 71(1): 494–503.
 44. de Paiva, J.B., da Silva, L.P., Casas, M.R., Conceição, R.A., Nakazato, G., de Pace, F., Sperandio, V. and da Silveira, W.D. (2016) *In vivo* influence of *in vitro* up-regulated genes in the virulence of an APEC strain associated with swollen head syndrome. *Avian Pathol.*, 45(1): 94–105.
 45. Li, Y., Wang, H., Ren, J., Chen, L., Zhuge, X., Hu, L., Li, D., Tang, F. and Dai, J. (2016) The YfcO fimbriae gene enhances adherence and colonization abilities of avian pathogenic *Escherichia coli* *in vivo* and *in vitro*. *Microb. Pathog.*, 100(1): 56–61.
 46. Paxman, J.J., Lo, A.W., Sullivan, M.J., Panjikar, S., Kuiper, M., Whitten, A.E., Wang, G., Luan, C.H., Moriel, D.G., Tan, L., Peters, K.M., Phan, M.D., Gee, C.L., Ulett, G.C., Schembri, M.A. and Heras, B. (2019) Unique structural features of a bacterial autotransporter adhesin suggest mechanisms for interaction with host macromolecules. *Nat. Commun.*, 10(1): 1967.
 47. Ali, A., Kolenda, R., Khan, M.M., Weinreich, J., Li, G., Wieler, L.H., Tedin, K., Roggenbuck, D. and Schierack, P. (2020) Novel avian pathogenic *Escherichia coli* genes responsible for adhesion to chicken and human cell lines. *Appl. Environ. Microbiol.*, 86(20): e01068-20.
 48. Calvo, M., Stefani, S. and Migliorisi, G. (2024) Bacterial infections in intensive care units: Epidemiological and microbiological aspects. *Antibiotics*, 13(3): 238.
 49. Maciel, J.F., Matter, L.B., Trindade, M.M., Camillo, G., Lovato, M., de Ávila Botton, S. and de Vargas, A.C. (2017) Virulence factors and antimicrobial susceptibility profile of extraintestinal *Escherichia coli* isolated from an avian colisepticemia outbreak. *Microb. Pathog.*, 103(1): 119–122.
 50. Wang, S., Bao, Y., Meng, Q., Xia, Y., Zhao, Y., Wang, Y., Tang, F., ZhuGe, X., Yu, S., Han, X., Dai, J. and Lu, C. (2015) *IbeR* facilitates stress-resistance, invasion and pathogenicity of avian pathogenic *Escherichia coli*. *PLoS One*, 10(3): e0119698.
 51. Pilatti, L., de Paiva, J.B., Rojas, T.C., Leite, J.L., Conceição, R.A., Nakazato, G. and da Silveira, W.D. (2016) The virulence factor ychO has a pleiotropic action in an Avian Pathogenic *Escherichia coli* (APEC) strain. *BMC Microbiol.*, 16(1): 35.
 52. Gao, Q., Wang, X., Xu, H., Xu, Y., Ling, J., Zhang, D., Gao, S. and Liu, X. (2012) Roles of iron acquisition systems in virulence of extraintestinal pathogenic *Escherichia coli*: Salmochelin and aerobactin contribute more to virulence than heme in a chicken infection model. *BMC Microbiol.*, 12(1): 143.
 53. Varga, C., Brash, M.L., Slavic, D., Boerlin, P., Ouckama, R., Weis, A., Petrik, M., Philippe, C., Barham, M. and Guerin, M.T. (2018) Evaluating virulence-associated genes and antimicrobial resistance of avian pathogenic *Escherichia coli* isolates from broiler and broiler breeder chickens in Ontario, Canada. *Avian Dis.*, 62(3): 291–299.
 54. Dziva, F. and Stevens, M.P. (2008) Colibacillosis in poultry: Unravelling the molecular basis of virulence of avian pathogenic *Escherichia coli* in their natural hosts, *Avian Pathol.*, 37(4): 355–366.
 55. Alav, I., Kobylka, J., Kuth, M.S., Pos, K.M., Picard, M., Blair, J.M.A. and Bavro, V.N. (2021) Structure, assembly, and function of tripartite efflux and type 1 secretion systems in gram-negative bacteria. *Chem. Rev.*, 121(9): 5479–5596.
 56. Soni, J., Sinha, S. and Pandey, R. (2024) Understanding bacterial pathogenicity: A closer look at the journey of harmful microbes. *Front. Microbiol.*, 15(1): 1370818.
 57. Sun, J., Rutherford, S.T., Silhavy, T.J. and Huang, K.C. (2022) Physical properties of the bacterial outer membrane. *Nat. Rev. Microbiol.*, 20(4): 236–248.
 58. Silveira, F., Maluta, R.P., Tiba, M.R., de Paiva, J.B., Guastalli, E.A. and da Silveira, W.D. (2016) Comparison between avian pathogenic (APEC) and avian faecal (AFEC) *Escherichia coli* isolated from different regions in Brazil. *Vet. J.*, 217(1): 65–67.
 59. Sora, V.M., Meroni, G., Martino, P.A., Soggiu, A., Bonizzi, L. and Zecconi, A. (2021) Extraintestinal pathogenic *Escherichia coli*: Virulence factors and antibiotic resistance. *Pathogens*, 10(11): 1355.
 60. Song, X., Hou, M., Tu, J., Xue, M., Shao, Y., Jiang, H., Liu, H., Xue, T., Wang, G. and Qi, K. (2019) Outer membrane proteins YbjX and PagP co-regulate motility in *Escherichia coli* via the bacterial chemotaxis pathway. *Res. Vet. Sci.*, 125(1): 279–284.
 61. Nielsen, D.W., Ricker, N., Barbieri, N.L., Allen, H.K., Nolan, L.K. and Logue, C.M. (2020) Outer membrane protein A (OmpA) of extraintestinal pathogenic *Escherichia coli*. *BMC Res. Notes*, 13(1): 51.
 62. Zuo, J., Tu, C., Wang, Y., Qi, K., Hu, J., Wang, Z., Mi, R., Huang, Y., Chen, Z. and Han, X. (2019) The role of the *wzy* gene in lipopolysaccharide biosynthesis and pathogenesis of avian pathogenic *Escherichia coli*. *Microb. Pathog.*, 127(1): 296–303.
 63. Dovala, D., Rath, C.M., Hu, Q., Sawyer, W.S., Shia, S., Elling, R.A., Knapp, M.S. and Metzger, L.E. (2016) Structure-guided enzymology of the lipid A acyltransferase LpxM reveals a dual activity mechanism. *Proc. Natl. Acad. Sci. U S A*, 113(41): E6064–E6071.
 64. Afonso, V., Champy, R., Mitrovic, D., Collin, P. and Lomri, A. (2007) Reactive oxygen species and superoxide dismutases: Role in joint diseases. *Joint Bone Spine*, 74(4): 324–329.
 65. Chiloeches, M.L., Bergonzini, A. and Frisan, T. (2021) Bacterial toxins are a never-ending source of surprises: From natural born killers to negotiators. *Toxins*, 13(6): 426.
 66. Zhao, S., Wang, C.L., Chang, S.K., Tsai, Y.L. and Chou, C.H. (2019) Characterization of *Escherichia coli* isolated from day-old chicken fluff in Taiwanese hatcheries. *Avian Dis.*, 63(1): 9–16.
 67. Murase, K., Martin, P., Porcheron, G., Houle, S., Helloin, E., Pénary, M., Nougayrède, J.P., Dozois, C.M., Hayashi, T. and Oswald, E. (2016) HlyF produced by extraintestinal pathogenic *Escherichia coli* is a virulence factor that regulates outer membrane vesicle biogenesis. *J. Infect. Dis.*, 213(5): 856–865.
 68. Nicolas, M., Faurie, A., Girault, M., Lavillatte, S.,

- Menanteau, P., Chaumeil, T., Riou, M., Velge, P. and Schouler, C. (2023) *In ovo* administration of a phage cocktail partially prevents colibacillosis in chicks. *Poult. Sci.*, 102(11): 102967.
69. Tuntufye, H.N., Lebeer, S., Gwakisa, P.S. and Goddeeris, B.M. (2012) Identification of Avian pathogenic *Escherichia coli* genes that are induced *in vivo* during infection in chickens. *Appl. Environ. Microbiol.*, 78(9): 3343–3351.
70. Bhuiyan, M.S.A., Sarker, S., Amin, Z., Rodrigues, K.F., Bakar, A.M.S.A., Saallah, S., Shaarani, S. and Siddiquee, S. (2023) Seroprevalence and molecular characterisation of infectious bronchitis virus (IBV) in broiler farms in Sabah, Malaysia. *Vet. Med. Sci.*, 10(2): e1153.
71. Wu, Z., Ding, L., Bao, J., Liu, Y., Zhang, Q., Wang, J., Li, R., Ishfaq, M. and Li, J. (2019) Co-infection of *Mycoplasma gallisepticum* and *Escherichia coli* triggers inflammatory injury involving the IL-17 signaling pathway. *Front. Microbiol.*, 10(1): 2615.
72. El Tayeb, A.B. and Hanson, R.P. (2002) Interactions between *Escherichia coli* and Newcastle disease virus in chickens. *Avian Dis.*, 46(3): 660–667.
73. Santos, A.C.M., Silva, R.M., Valiatti, T.B., Santos, F.F., Santos-Neto, J.F., Cayó, R., Streling, A.P., Nodari, C.S., Gales, A.C., Nishiyama-Jr, M.Y., Carvalho, E. and Gomes, T.A.T. (2020) Virulence potential of a multidrug-resistant *Escherichia coli* strain belonging to the emerging clonal group ST101-B1 isolated from blood-stream infection. *Microorganisms*, 8(6): 827.
74. Kabir, S.M.L. (2010) Avian colibacillosis and salmonellosis: A closer look at epidemiology, pathogenesis, diagnosis, control and public health concerns. *Int. J. Environ. Res. Public Health*, 7(1): 89–114.
75. Leininger, D.J., Roberson, J.R. and Elvinger, F. (2001) Use of eosin methylene blue agar to differentiate *Escherichia coli* from other gram-negative mastitis pathogens. *J. Vet. Diagn. Invest.*, 13(3): 273–275.
76. Al-Blooshi, S.Y., Latif, M.A.A., Sabaneh, N.K., Mgaogao, M. and Hossain, A. (2021) Development of a novel selective medium for culture of Gram-negative bacteria. *BMC Res. Notes*, 14(1): 211.
77. Paniel, N. and Noguer, T. (2019) Detection of *Salmonella* in food matrices, from conventional methods to recent aptamer-sensing technologies. *Foods*, 8(9): 371.
78. Fujisawa, T., Sata, S., Aikawa, K., Takahashi, T., Yamai, S. and Shimada, T. (2000) Modification of sorbitol MacConkey medium containing cefixime and tellurite for isolation of *Escherichia coli* O157:H7 from radish sprouts. *Appl. Environ. Microbiol.*, 66(7): 3117–3118.
79. Park, S.H., Ryu, S. and Kang, D.H. (2011) Improved selective and differential medium for isolation of *Escherichia coli* O157:H7. *J. Clin. Microbiol.*, 49(1): 405–408.
80. Carreón-Rodríguez, O.E., Gosset, G., Escalante, A. and Bolívar, F. (2023) Glucose transport in *Escherichia coli*: From basics to transport engineering. *Microorganisms*, 11(6): 1588.
81. Mazumder, R., Hussain, A., Rahman, M.M., Phelan, J.E., Campino, S., Abdullah, A., Clark, T.G. and Mondal, D. (2023) Genomic and functional portrait of multidrug-resistant, hydrogen sulfide (H₂S)-producing variants of *Escherichia coli*. *Front. Microbiol.*, 14(1): 1206757.
82. Dien, B.S., Nichols, N.N. and Bothast, R.J. (2002) Fermentation of sugar mixtures using *Escherichia coli* catabolite repression mutants engineered for production of L-lactic acid. *J. Ind. Microbiol. Biotechnol.*, 29(5): 221–227.
83. Saqib, S., Akram, A., Halim, S.A. and Tassaduq, R. (2017) Sources of β -galactosidase and its applications in food industry. *3 Biotech*, 7(1): 79.
84. Koutsianos, D., Athanasiou, L.V., Mossialos, D., Franzo, G., Cecchinato, M. and Koutoulis, K.C. (2022) Investigation of serotype prevalence of *Escherichia coli* strains isolated from layer poultry in Greece and interactions with other infectious agents. *Vet. Sci.*, 9(4): 152.
85. Filho, H.C.K., Carvalho, D., Grassotti, T.T., Soares, B.D., Rossato, J.M., Cunha, A.C., Brito, K.C.T., Cavalli, L.S. and Brito, B.G. (2015) Avian pathogenic *Escherichia coli*-methods for improved diagnosis. *Worlds Poult. Sci. J.*, 71(2): 249–258.
86. Müller, C.M., Aberg, A., Strasevičienė, J., Emody, L., Uhlin, B.E. and Balsalobre, C. (2019) Type 1 fimbriae, a colonization factor of uropathogenic *Escherichia coli*, are controlled by the metabolic sensor CRP-cAMP. *PLoS Pathog.*, 5(2): e1000303.
87. Effendi, M.H., Harijani, N., Yanestria, S.M. and Hastutiek, P. (2018). Identification of Shiga toxin-producing *Escherichia coli* in raw milk samples from dairy cows in Surabaya, Indonesia. *Philipp. J. Vet. Med.*, 55(S1): 109–114.
88. Ghunaim, H., Abu-Madi, M.A. and Kariyawasam, S. (2014) Advances in vaccination against avian pathogenic *Escherichia coli* respiratory disease: Potentials and limitations. *Vet. Microbiol.*, 172(1–2): 13–22.
89. Vega-Hernández, R., Ochoa, S.A., Valle-Rios, R., Jaimes-Ortega, G.A., Arellano-Galindo, J., Aparicio-Ozores, G., Ibarra, J.A., Hernández-Castro, R., Cruz-Córdova, A. and Xicohtencatl-Cortes, J. (2021) Flagella, type I fimbriae and curlin of uropathogenic *Escherichia coli* promote the release of proinflammatory cytokines in a coculture system. *Microorganisms*, 9(11): 2233.
90. Wolfson, E.B., Elvidge, J., Tahoun, A., Gillespie, T., Mantell, J., McAteer, S.P., Rossez, Y., Paxton, E., Lane, F., Shaw, D.J., Gill, A.C., Stevens, J., Verkade, P., Blocker, A., Mahajan, A. and Gally, D.L. (2020) The interaction of *Escherichia coli* O157: H7 and *Salmonella* Typhimurium flagella with host cell membranes and cytoskeletal components. *Microbiology (Reading)*, 166(10): 947–965.
91. Best, A., La Ragione, R.M., Sayers, A.R. and Woodward, M.J. (2005) Role for flagella but not intimin in the persistent infection of the gastrointestinal tissues of specific-pathogen-free chicks by shiga toxin-negative *Escherichia coli* O157:H7. *Infect. Immun.*, 73(3): 1836–1846.
92. Azurmendi, H.F., Veeramachineni, V., Freese, S., Lichaa, F., Freedberg, D.I. and Vann, W.F. (2020) Chemical structure and genetic organization of the *E. coli* O6:K15 capsular polysaccharide. *Sci. Rep.*, 10(1): 12608.
93. La Ragione, R.M. and Woodward, M.J. (2002) Virulence factors of *Escherichia coli* serotypes associated with avian colisepticaemia. *Res. Vet. Sci.*, 73(1): 27–35.
94. Vandekerchove, D., De Herdt, P., Laevens, H. and Pasmans, F. (2004) Colibacillosis in caged layer hens: Characteristics of the disease and the aetiological agent. *Avian Pathol.*, 33(2): 117–125.
95. Mohamed, M.A., Shehata, M.A. and Rafeek, E. (2014) Virulence genes content and antimicrobial resistance in *Escherichia coli* from broiler chickens. *Vet. Med. Int.*, 2014(1): 195189.
96. Tivendale, K.A., Logue, C.M., Kariyawasam, S., Jordan, D., Hussein, A., Li, G., Wannemuehler, Y. and Nolan, L.K. (2010) Avian-pathogenic *Escherichia coli* strains are similar to neonatal meningitis *E. coli* strains and are able to cause meningitis in the rat model of human disease. *Infect. Immun.*, 78(8): 3412–3419.
97. Alqasim, A., Jaffal, A.A. and Alyousef, A.A. (2020) Prevalence and molecular characteristics of sequence type 131 clone among clinical uropathogenic *Escherichia coli* isolates in Riyadh, Saudi Arabia. *Saudi J. Biol. Sci.*, 27(1): 296–302.
98. Law, J.W., Ab Mutalib, N.S., Chan, K.G. and Lee, L.H. (2015) Rapid methods for the detection of foodborne bacterial pathogens: Principles, applications, advantages and limitations. *Front. Microbiol.*, 5(1): 770.
99. Mason, P.S., Holder, T., Robinson, N., Smith, B., Hameed, R.T., Al Dulayymi, J.R., Hughes, V., Stevenson, K., Jones, G.J., Vordermeier, H.M., Mc Kenna, S. and Baird, M.S. (2024) An ELISA using synthetic

- mycolic acid-based antigens with DIVA potential for diagnosing Johne's disease in cattle. *Animals*, 14(6): 848.
100. Salhi, I., Rabti, A., Dhehibi, A. and Raouafi, N. (2022) Sandwich-based immunosensor for dual-mode detection of pathogenic F17-positive *Escherichia coli* strains. *Int. J. Mol. Sci.*, 23(11): 6028.
 101. Brocchi, M., Ferreira, A., Lancellotti, M., Stehling, E.G., Campos, T.A., Nakazato, G., de Castro, A.F.P. and Silveira, W.D. (2006) Typing of avian pathogenic *Escherichia coli* strains by REP-PCR. *Pesq. Vet. Bras.*, 26(2): 69–73.
 102. Álvarez-Fernández, R. (2013) Explanatory chapter: PCR primer design. *Methods Enzymol.*, 529(1): 1–21.
 103. Mahmoud, A.T., Salim, M.T., Ibrahim, R.A., Gabr, A. and Halby, H.M. (2020) Multiple drug resistance patterns in various phylogenetic groups of hospital-acquired uropathogenic *E. coli* isolated from cancer patients. *Antibiotics*, 9(3): 108.
 104. Johnson, T.J., Wannemuehler, Y., Doetkott, C., Johnson, S.J., Rosenberger, S.C. and Nolan, L.K. (2008) Identification of minimal predictors of avian pathogenic *Escherichia coli* virulence for use as a rapid diagnostic tool. *J. Clin. Microbiol.*, 46(12): 3987–3996.
 105. Borzi, M.M., Cardozo, M.V., Oliveira, E.S., Pollo, A.S., Guastalli, E.A.L., Santos, L.F.D. and Ávila, F.A. (2018) Characterization of avian pathogenic *Escherichia coli* isolated from free-range helmeted guineafowl. *Braz. J. Microbiol.*, 49(Suppl 1): 107–112.
 106. Shahjada, Z., Hussain, K., Islam, M.M., Majumder, S., Hasan, I., Rahman, M. and Saha, S. (2017) Bacteria causing omphalitis in newly hatched chicks from broiler and layer flocks and their antibiotic profiles. *Int. J. Nat. Soc. Sci.*, 4(2): 73–81.
 107. Oliveira, G.D.S., Pires, P.G.D.S., McManus, C., de Jesus, L.M., Santos, P.H.G.S. and Dos Santos, V.M. (2024) Plant extract in the control of poultry omphalitis. *Pathogens*, 13(6): 438.
 108. Tawk, A., Abdallah, A., Meouchy, P., Salameh, J., Khoury, S., Kyriakos, M., Abboud, G., Dagher, M., Semaan, P., Metri, A. and Ashou, R. (2021) Omphalitis with umbilical abscess in an adult with a urachal remnant. *Case Rep. Gastroenterol.*, 15(3): 966–971.
 109. Amer, M.M., Mekky, H.M., Fedawy, H.S., Amer, A.M. and Elbayoumi, K.M. (2020) Cellulitis in broiler chickens. *Korean J. Food Health Converg.*, 6(5): 1–10.
 110. Bernd, K.S., Kump, A.W.S., Rohn, K., Reich, F. and Kehrenberg, C. (2020) Management factors influencing the occurrence of cellulitis in broiler chickens. *Prev. Vet. Med.*, 183(1): 105146.
 111. Sullivan, T. and de Barra, E. (2018) Diagnosis and management of cellulitis. *Clin. Med. (Lond)*, 18(2): 160–163.
 112. Wlazlak, S., Pietrzak, E., Biesek, J. and Dunislawska, A. (2023) Modulation of the immune system of chickens a key factor in maintaining poultry production-a review. *Poult. Sci.*, 102(8): 102785.
 113. Yehia, N., Salem, H.M., Mahmmud, Y., Said, D., Samir, M., Mawgod, S.A., Sorour, H.K., AbdelRahman, M.A.A., Selim, S., Saad, A.M., El-Saadony, M.T., El-Meihy, R.M., Abd El-Hack, M.E., El-Tarabily, K.A. and Zanaty, A.M. (2023) Common viral and bacterial avian respiratory infections: an updated review. *Poult. Sci.*, 102(5): 102553.
 114. Lu, Y.S., Shien, Y.S., Tsai, H.J., Tseng, C.S., Lee, S.H. and Lin, D.F. (1994) Swollen head syndrome in Taiwan-isolation of an avian pneumovirus and serological survey. *Avian Pathol.*, 23(1): 169–174.
 115. Gazdzinski, P. and Barnes, J. (2004) Venereal colibacillosis (acute vaginitis) in turkey breeder hens. *Avian Dis.*, 48(3): 681–685.
 116. Landman, W.J.M., Heuvelink, A. and van Eck, J.H.H. (2013) Reproduction of the *Escherichia coli* peritonitis syndrome in laying hens. *Avian Pathol.*, 42(2): 157–162.
 117. Li, L.L., Xu, P.T., Liu, Z.P., Liu, C.A., Dong, X.Y., Zhang, Z.F., Guo, S.S. and Ding, B.Y. (2023) Effects of salpingitis simulation on the morphology and expression of inflammatory-related genes of oviduct in laying hens. *Poult. Sci.*, 102(1): 102246.
 118. van der Putten, M.E., Engel, M. and van Well, G.T. (2008) Salpingitis. A rare cause of acute abdomen in a sexually inactive girl: A case report. *Cases J.*, 1(1): 326.
 119. Yousef, H.M.Y., Hashad, M.E., Osman, K.M., Alatfeehy, N.M., Hassan, W.M.M., Elebeedy, L.A., Salem, H.M., Shami, A., Al-Saeed, F.A., El-Saadony, M.T., El-Tarabily, K.A. and Marouf, S. (2023) Surveillance of *Escherichia coli* in different types of chicken and duck hatcheries: one health outlook. *Poult. Sci.*, 102(12): 103108.
 120. Shibuki, S., Saida, T., Hoshiai, S., Ishiguro, T., Sakai, M., Amano, T., Abe, T., Yoshida, M., Mori, K. and Nakajima, T. (2024) Imaging findings in inflammatory disease of the genital organs. *Jpn. J. Radiol.*, 42(4): 331–346.
 121. Monleon, R., Martin, M.P. and Barnes, H.J. (2008) Bacterial orchitis and epididymo-orchitis in broiler breeders. *Avian Pathol.*, 37(6): 613–617.
 122. Lu, Y., Liu, M., Tursi, N.J., Yan, B., Cao, X., Che, Q., Yang, N. and Dong, X. (2021) Uropathogenic *Escherichia coli* infection compromises the blood-testis barrier by disturbing mTORC1-mTORC2 balance. *Front. Immunol.*, 12(1): 582858.
 123. Kendek, I.A., Putri, MFR, Wibisono, F.J., Effendi, M.H., Tyasningsih, W., Ugbo, E.N. and Agumah, N.B. (2024) Molecular detection of *hlyF* gene on multidrug resistance of avian pathogenic *Escherichia coli* isolated from ducks on wet markets of Surabaya, Indonesia. *Biodiversitas*, 25: 1246–1252.
 124. Wilczyński, J., Stepien-Pyśniak, D., Wystalska, D. and Wernicki, A. (2022) Molecular and serological characteristics of avian pathogenic *Escherichia coli* isolated from various clinical cases of poultry colibacillosis in Poland. *Animals*, 12(9): 1090.
 125. Hu, J., Afayibo, D.J.A., Zhang, B., Zhu, H., Yao, L., Guo, W., Wang, X., Wang, Z., Wang, D., Peng, H., Tian, M., Qi, J. and Wang, S. (2022) Characteristics, pathogenic mechanism, zoonotic potential, drug resistance, and prevention of avian pathogenic *Escherichia coli* (APEC). *Front. Microbiol.*, 13(1): 1049391.
 126. Ateş, M.B., Zeynep Çelik, Z. and Çiftçi, M.K. (2020) Pathological and immunohistochemical examinations on co-infection with coligranulomatosis and Marek's disease in a turkey flock. *J. Poult. Res.*, 17(2): 87–95.
 127. Landman, W.J.M. and van Eck, J.H.H. (2017) Coligranulomatosis (Hjärre and Wramby's disease) reconsidered. *Avian Pathol.*, 46(3): 237–241.
 128. Zhou, Z., Sharif, A., Inglesi-Lotz, R. and Bashir, M.F. (2024) Analysing the interplay between energy transition, resource consumption, deforestation, and environmental factors on agricultural productivity: Insights from APEC countries. *J. Clean. Prod.*, 446(1): 141408.
 129. Yanestria, S.M., Dameanti, F.N.A.E.P., Musayannah, B.G., Pratama, J.W.A., Witaningrum, A.M., Effendi, M.H. and Ugbo, E.N. (2022) Antibiotic resistance pattern of extended-spectrum β -lactamase (ESBL) producing *Escherichia coli* isolated from broiler farm environment in Pasuruan district, Indonesia. *Biodiversitas*, 23(9): 4460–4465.
 130. Wang, K., Shen, D., Dai, P. and Li, C. (2023) Particulate matter in poultry house on poultry respiratory disease: A systematic review. *Poult. Sci.*, 102(4): 102556.
 131. Wang, G., Liu, Q., Zhou, Y., Feng, J. and Zhang, M. (2022) Effects of different ammonia concentrations on pulmonary microbial flora, lung tissue mucosal morphology, inflammatory cytokines, and neurotransmitters of broilers. *Animals*, 12(3): 261.
 132. Van Limbergen, T., Sarrazin, S., Chantziaras, I., Dewulf, J., Ducatelle, R., Kyriazakis, I., McMullin, P., Méndez, J., Niemi, J.K., Papisolomontos, S., Szeleszczuk, P., Van Erum, J. and Maes, D. (2020) Risk factors for poor

- health and performance in European broiler production systems. *BMC Vet. Res.*, 16(1): 287.
133. Stromberg, Z.R., Johnson, J.R., Fairbrother, J.M., Kilbourne, J., Van Goor, A., Curtiss, R.R. and Mellata, M. (2017) Evaluation of *Escherichia coli* isolates from healthy chickens to determine their potential risk to poultry and human health. *PLoS One*, 12(7): e0180599.
 134. Nguyen, X.D., Zhao, Y., Evans, J.D., Lin, J. and Purswell, J.L. (2022) Survival of *Escherichia coli* in airborne and settled poultry litter particles. *Animals*, 12(3): 284.
 135. Mehat, J.W., van Vliet, A.H.M. and La Ragione, R.M. (2021) The avian pathogenic *Escherichia coli* (APEC) pathotype is comprised of multiple distinct, independent genotypes. *Avian Pathol.*, 50(5): 402–416.
 136. Ahmed, M.F.E., Abd El-Wahab, A., Kriewitz, J.P., Hankel, J., Chuppava, B., Ratert, C., Taube, V., Visscher, C. and Kamphues, J. (2021) Mitigating the spread and translocation of *Salmonella Enteritidis* in experimentally infected broilers under the influence of different flooring housing systems and feed particle sizes. *Microorganisms*, 9(4): 874.
 137. Walker, G.K., Suyemoto, M.M., Gall, S., Chen, L., Thakur, S. and Borst, L.B. (2020) The role of *Enterococcus faecalis* during co-infection with avian pathogenic *Escherichia coli* in avian colibacillosis. *Avian Pathol.*, 49(6): 589–599.
 138. Adefisoye, M.A. and Okoh, A.I. (2016) Identification and antimicrobial resistance prevalence of pathogenic *Escherichia coli* strains from treated wastewater effluents in Eastern Cape, South Africa. *Microbiologyopen*, 5(1): 143–151.
 139. Koutsianos, D., Athanasiou, L., Mossialos, D. and Koutoulis, K. (2021) Colibacillosis in poultry: A disease overview and the new perspectives for its control and prevention. *J. Hellenic Vet. Med. Soc.*, 71(4): 2425–2436.
 140. Artdita, C.A., Zhuang, Y.R., Liu, T.Y., Cheng, C.Y., Hsiao, F.S. and Lin, Y.Y. (2021) The effect of feeding restriction on the microbiota and metabolome response in late-phase laying hens. *Animals*, 11(11): 3043.
 141. Swelum, A.A., Elbestawy, A.R., El-Saadony, M.T., Hussein, E.O.S., Alhotan, R., Suliman, G.M., Taha, A.E., Ba-Awadh, H., El-Tarabily, K.A. and Abd El-Hack, M.E. (2021) Ways to minimize bacterial infections, with special reference to *Escherichia coli*, to cope with the first-week mortality in chicks: An updated overview. *Poult. Sci.*, 100(5): 101039.
 142. El-Saadony, M.T., Salem, H.M., El-Tahan, A.M., Abd El-Mageed, T.A., Soliman, S.M., Khafaga, A.F., Swelum, A.A., Ahmed, A.E., Alshammari, F.A. and Abd El-Hack, M.E. (2022) The control of poultry salmonellosis using organic agents: An updated overview. *Poult. Sci.*, 101(4): 101716.
 143. Waktole, H., Ayele, Y., Ayalkibet, Y., Teshome, T., Muluneh, T., Ayane, S., Borena, B.M., Abayneh, T., Deresse, G., Asefa, Z., Eguale, T., Amenu, K., Ashenafi, H. and Antonissen, G. (2024) Prevalence, molecular detection, and antimicrobial resistance of *Salmonella* isolates from poultry farms across central Ethiopia: A cross-sectional study in Urban and Peri-Urban Areas. *Microorganisms*, 12(4): 767.
 144. Hess, C., Troxler, S., Jandreski-Cvetkovic, D., Zloch, A. and Hess, M. (2022) *Escherichia coli* isolated from organic laying hens reveal a high level of antimicrobial resistance despite no antimicrobial treatments. *Antibiotics*, 11(4): 467.
 145. Umar, S., Munir, M.T., Ahsan, U., Raza, I., Chowdhury, M.R., Ahmed, Z. and Shah, M.A.A. (2017) Immunosuppressive interactions of viral diseases in poultry. *Worlds Poult. Sci. J.*, 73(1): 121–135.
 146. Domanska-Blicharz, K., Opolska, J., Lisowska, A. and Szczotka-Bochniarz, A. (2023) Bacterial and viral rodent-borne infections on poultry farms. An attempt at a systematic review. *J. Vet. Res.*, 67(1): 1–10.
 147. Zurek, L. and Ghosh, A. (2014) Insects represent a link between food animal farms and the urban environment for antibiotic resistance traits. *Appl. Environ. Microbiol.*, 80(12): 3562–3567.
 148. Blaak, H., Hamidjaja, R.A., van Hoek, A.H., de Heer, L., de Roda Husman, A.M. and Schets, F.M. (2014) Detection of extended-spectrum beta-lactamase (ESBL)-producing *Escherichia coli* on flies at poultry farms. *Appl. Environ. Microbiol.*, 80(1): 239–246.
 149. Mace, J.L. and Knight, A. (2024) From the backyard to our beds: The spectrum of care, attitudes, relationship types, and welfare in non-commercial chicken care. *Animals*, 14(2): 288.
 150. Chakrabarti, S., King, D.J., Afonso, C., Swayne, D., Cardona, C.J., Kuney, D.R. and Gerry, A.C. (2007) Detection and isolation of exotic Newcastle disease virus from field-collected flies. *J. Med. Entomol.*, 44(5): 840–844.
 151. Subhashree, A.V.S., Krishna, G.S., Kumari, G.D., Reddy, T.N. and Kumar, P.A. (2015) Pathogenicity study on extra-intestinal avian pathogenic *Escherichia coli* isolated from broiler chickens. *Indian J. Agric. Bus.*, 1(2): 85–90.
 152. Bortolaia, V., Bisgaard, M. and Bojesen, A.M. (2010) Distribution and possible transmission of ampicillin and nalidixic acid resistant *Escherichia coli* within the broiler industry. *Vet. Microbiol.*, 142: 379–386.
 153. Liljebjelke, K.A., Hofacre, C.L., Liu, T., White, D.G., Ayers, S., Young, S. and Maurer, J.J. (2005) Vertical and horizontal transmission of *Salmonella* within integrated broiler production system. *Foodborne Pathog. Dis.*, 2(1): 90–102.
 154. Giovanardi, D., Campagnari, E., Ruffoni, L.S., Pesente, P., Ortali, G. and Furlattini, V. (2005) Avian pathogenic *Escherichia coli* transmission from broiler breeders to their progeny in an integrated poultry production chain. *Avian Pathol.*, 34(4): 313–318.
 155. Peterson, A., Christensen, J.P., Kuhnert, P., Bisgaard, M. and Olsen, J.E. (2006) Vertical transmission of a fluoroquinolone-resistant *Escherichia coli* within an integrated broiler operation. *Vet. Microbiol.*, 116(1–3): 120–128.
 156. Poulsen, L.L., Thofner, I., Bisgaard, M., Christensen, J.P., Olsen, R.H. and Christensen, H. (2017) Longitudinal study of transmission of *Escherichia coli* from broiler breeders to broilers. *Vet. Microbiol.*, 207: 13–18.
 157. Christensen, H., Bachmeier, J. and Bisgaard, M. (2021) New strategies to prevent and control avian pathogenic *Escherichia coli* (APEC). *Avian Pathol.*, 50(5): 370–381.
 158. Lindsey, L.L., Elliott, K.E.C., Fatemi, S.A., Evans, J.D., Mousstaaid, A., Gerard, P.D. and Peebles, E.D. (2022) Variable effects of the *in ovo* administration of an *Escherichia coli* vaccine in the amnion or air cell on commercial layer embryo and hatchling development. *Poultry*, 1(4): 278–290.
 159. Qiu, L., Chirman, D., Clark, J.R., Xing, Y., Santos, H.H., Vaughan, E.E. and Maresso, A.W. (2024) Vaccines against extraintestinal pathogenic *Escherichia coli* (ExPEC): Progress and challenges. *Gut Microbes*, 16(1): 2359691.
 160. Hu, R., Li, J., Zhao, Y., Lin, H., Liang, L., Wang, M., Liu, H., Min, Y., Gao, Y. and Yang, M. (2020) Exploiting bacterial outer membrane vesicles as a cross-protective vaccine candidate against avian pathogenic *Escherichia coli* (APEC). *Microb. Cell. Fact.*, 19(1): 119.
 161. Paudel, S., Apostolakos, I., Ngom, R.V., Tilli, G., de Carvalho Ferreira, H.C. and Piccirillo, A. (2024) A systematic review and meta-analysis on the efficacy of vaccination against colibacillosis in broiler production. *PLoS One*, 19(3): e0301029.
 162. Wibisono, F.J., Sumiarti, B., Untari, T., Effendi, M.H., Permatasari, D.A. and Witaningrum, A.M. (2021) Molecular identification of ctx gene of extended-spectrum beta-lactamase (ESBL) producing *Escherichia coli* on layer chicken in Blitar, Indonesia. *J. Anim. Plant. Sci.*, 31(4): 954–959.
 163. Smialek, M., Kowalczyk, J. and Koncicki, A. (2020) Influence of vaccination of broiler chicken against *Escherichia coli* with live attenuated vaccine on general properties of *E. coli* population, IBV vaccination efficiency,

- and production parameters a field experiment. *Poult. Sci.*, 99(11): 5452–5460.
164. Landman, W.J.M. and van Eck, J.H.H. (2017) The efficacy of inactivated *Escherichia coli* autogenous vaccines against the *E. coli* peritonitis syndrome in layers. *Avian Pathol.*, 46(6): 658–665.
 165. Fatemi, S.A., Lindsey, L.L., Evans, J.D., Elliott, K.E.C., Leigh, S.A., Robinson, K.J., Mousstaaid, A., Gerard, P.D. and Peebles, E.D. (2023) Effects of the *in ovo* injection of an *Escherichia coli* vaccine on the hatchability and quality characteristics of commercial layer hatchlings. *Poult. Sci.*, 102(11): 103057.
 166. Chai, M., Lu, M., Keener, T., Khang, S.J., Chaiwatpongsakorn, C. and Tisch, J. (2009) Using an improved electrostatic precipitator for poultry dust removal. *J. Electrostat.*, 67(6): 870–875.
 167. Bae, D., Song, K.Y., Macoy, D.M., Kim, M.G., Lee, C.K. and Kim, Y.S. (2022) Inactivation of airborne avian pathogenic *E. coli* (APEC) via application of a novel high-pressure spraying system. *Microorganisms*, 10(11): 2201.
 168. Marangoni, F., Corsello, G., Cricelli, C., Ferrara, N., Ghiselli, A., Lucchin, L. and Poli, A. (2015) Role of poultry meat in a balanced diet aimed at maintaining health and wellbeing: An Italian consensus document. *Food Nutr. Res.*, 59(1): 27606.
 169. Weyh, C., Krüger, K., Peeling, P. and Castell, L. (2022) The role of minerals in the optimal functioning of the immune system. *Nutrients*, 14(3): 644.
 170. Monson, M.S. and Lamont, S.J. (2021) Genetic resistance to avian pathogenic *Escherichia coli* (APEC): Current status and opportunities. *Avian Pathol.*, 50(5): 392–401.
 171. Khalid, A., Khalid, F., Mahreen, N., Hussain, S.M., Shahzad, M.M., Khan, S. and Wang, Z. (2022) Effect of spore-forming probiotics on the poultry production: A review. *Food Sci. Anim. Resour.*, 42(6): 968–980.
 172. Dhaka, P., Chantziaras, I., Vijay, D., Bedi, J.S., Makovska, I., Biebau, E. and Dewulf, J. (2023) Can improved farm biosecurity reduce the need for antimicrobials in food animals? A scoping review. *Antibiotics*, 12(5): 893.
 173. Guabiraba, R. and Schouler, C. (2015) Avian colibacillosis: Still many black holes. *FEMS Microbiol. Lett.*, 362(15): fnv118.
 174. Agunos, A., Leger, D. and Carson, C. (2012) Review of antimicrobial therapy of selected bacterial disease in broiler chickens in Canada. *Can. Vet. J.*, 53: 1289–1300.
 175. Tyasningsih, W., Ramandinianto, S.C., Ansharieta, R., Witaningrum, A.M., Permatasari, D.A., Wardhana, D.K., Effendi, M.H. and Ugbo, E.N. (2022) Prevalence and antibiotic resistance of *Staphylococcus aureus* and *Escherichia coli* isolated from raw milk in East Java, Indonesia. *Vet. World*, 15(8): 2021–2028.
 176. Hussein, A.H., Ghanem, I.A., Eid, A.A., Ali, M.A., Sherwood, J.S., Li, G., Nolan, L.K. and Logue, C.M. (2013) Molecular and phenotypic characterization of *Escherichia coli* isolated from broiler chicken flocks in Egypt. *Avian Dis.*, 57(3): 602–611.
 177. Sondhi, P., Maruf, M.H.U. and Stine, K.J. (2019) Nanomaterials for biosensing lipopolysaccharide. *Biosensors*, 10(1): 2.
 178. Fernández, A., Hernández, M., Moreno, Y. and García-Hernández, J. (2024) Specific and simultaneous detection of *E. coli* O157:H7 and Shiga-like toxins using a label-free photonic immunosensor. *Photonics*, 11(4): 374.
 179. Meconi, S., Vercellone, A., Levillain, F., Payré, B., Al Saati, T., Capilla, F., Desreumaux, P., Darfeuille-Michaud, A. and Altare, F. (2007) Adherent-invasive *Escherichia coli* isolated from Crohn's disease patients induce granulomas *in vitro*. *Cell Microbiol.*, 9(5): 1252–1261.
