The Under Reported Issue of Antibiotic-Resistance in Food-Producing Animals in Pakistan

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ABSTRACT

The recent revelation of a plasmid-mediated colistin resistant superbug isolated in China from food-producing animals (FPAs) and epidemic of a pan-drug resistant human typhoid strain in Hyderabad, Pakistan proved the notion that novel mechanisms of antibiotic resistance will emerge as we continue to rely on consistent use of antibiotics. The emergence and dissemination of antibiotic resistance is accelerated up by indiscriminate and incessant usage of antibiotics due to uncontrolled infections in hospital settings, community, FPAs and aquaculture. Unfortunately, a major part of the global antimicrobial usage is utilized in FPAs either for therapeutic purpose and/or as growth promoters raising vital concerns of the emergence of novel multidrug-resistant microorganisms. Pakistan-an agriculture-based country ranking in top ten livestock producing countries is currently shifting towards modern intense farming practices whereby animals are being supplemented with antibiotics on routine basis. However, unfortunately, monitoring usage of antibiotics in FPAs is lacking. Above all, the absence of systemic surveillance data on the level, frequency and impact of antimicrobial resistance further making it more challenging to devise antimicrobial-resistance-control strategies. We are highlighting this lack of information and hope that this review will help us in getting attention of policymakers for improving infection control measures and implementing laws on restricted use of antibiotics in animals.

INTRODUCTION

Antibiotics exhibit killing or growth inhibiting activity against target microorganisms, and are naturally secreted by fungi, actinomycetes, and bacteria. It is thought that antibiotic secretion is triggered to compete for available space and nutrients and is therefore vital for the survival of host organisms. The development of resistance in a target microorganism is the acquisition of elements from the genetic pools of resistance that confer resistance against the selected antimicrobials (Levy, 1994, 2002a). Practically, drug resistance is seen when both components, the resistance conferring elements and antimicrobials, came together in the host or in an environment. Current literature indicates that there are more than 15 different classes of antibiotics (Levy, 2002b) and none has escaped a resistance mechanism (Goossens et al., 2005). Although, understanding biochemical mechanisms of drug resistance is crucial, of equal importance is elucidating the ability to disseminate these drug resistance elements.

Antibiotics are largely used in food-producing animals (FPAs) all over the world. In fact, the current modernization, intensification and efficient production of FPAs realized this partly as a result of incorporation of growth supplements, which includes mainly antibiotics (Price et al., 2005). It has been estimated that the global average annual consumption of antimicrobials per kilogram of animal produced was 45 mg·kg⁻¹, 48 mg·kg⁻¹, and 172 mg·kg⁻¹ for cattle, chicken, and pigs, respectively, for the year 2010. This usage of antibiotics in FPAs has been projected to increase by 67% in 2030 (Van Boeckel et al., 2015). This widespread use of antibiotics in FPAs contributes by means of natural selection- to the emergence of antimicrobial resistance (AMR), which has public health consequences. AMR carrying pathogens can be transmitted to human from environment (Graham et al., 2009; Mohsin et al., 2017) or food products (Price et al., 2005) and thus turns into a public health issue. In Pakistan, food-animals are becoming increasingly important for food security.
Pakistan is in the list of top 10 animal producing countries (Page and Gautier, 2012) with a recent trend of modern intense farming practices. Unfortunately, this modern animal farming in Pakistan, like in many other countries, largely relies on the consistent use of antibiotics as growth promoters and prevention against many diseases. Sadly, no proper systemic surveillance data of the antimicrobial usage (AMU) or emergence of AMR in the country is available; nonetheless, available literature suggests an overall increase in AMR. Emergence and dissemination of antibiotic-resistant pathogens from FPAs carry major public health consequences as well as constituting reservoirs for resistance elements in Pakistan (Mohsin et al., 2017; Rahman et al., 2018a (https://www.dawn.com/news/1328924)); hence, controlled and restricted use of antibiotics in livestock is imperative. In this review, we accumulated and organized the available data found in different online resources, newspaper reports and also interviewed few veterinary clinicians regarding practices, the occurrence and nature of antibiotic resistance in FPAs in Pakistan.

Development of AMR and its impact: A resistant bacterium offers drug resistance by one of the main following mechanisms (i) reducing import and increasing export of the antibiotic drugs (genetic alterations in porins, up-regulated efflux pumps). (ii) modification, mainly structural, of the drug target (mutations, enzymatic modification), (iii) over expression or increased biosynthesis of the drug target, (iv) secretion of enzymes modifying, degrading or altering targeted antimicrobials such as β-lactamase (Normark and Normark 2002; ur Rahman et al., 2018a). In addition, bacteria have evolved secretory systems for secretion of intrinsically manufactured toxins and proteins as signals and virulent factors etc. (Rahman and van Ulsen 2013; Rahman et al., 2014; van Ulsen et al., 2014; Piet et al., 2016) with potential contact-dependent killing of target bacteria.

Although, causatives due to the overuse of antibiotics in animals is hard to establish, however, evidence of close association- between livestock-associated resistant bugs in animals and human (Vieira et al., 2011; Khattak et al., 2018), level of use of antimicrobials and prevalence of resistant bacteria in animals (Aarestrup, 2005), and in humans (Schwarz et al., 2001) exist. For prevention of diseases, antimicrobials are continuously administered in drinking water or added in daily feed for a longer period at lower doses (Wegener, 2003), while growth promoters are supplemented on routine basis in feed or drinking water at low level (Stokstad and Jukes 1950), potentially encouraging growth of selected resistant bacteria (Alexander et al., 2008). There is current understanding that the addition of antibiotics in agriculture and aquaculture are the major sources, besides human use, that help in spreading antimicrobial-resistant microorganisms AMROs in the surroundings. In this context, two major determinants- the quantity and pattern of antibiotics applied in FPAs are considered crucial for the emergence and dissemination of resistant bacteria in animal reservoir. Thus, the level and pattern of resistance observed to the great extent reflects the drug usage in animals, nevertheless, other factors may also be important, such as vertical spread of clones resistant to antibacterial drugs and selective and successful adaptation of the resistant strains (Aarestrup et al., 2008).

Presence of antibiotic residues in milk of high producing animals and their spread to the environment is a matter of great concern for public health. A study analyzed marketed dairy milk for the presence of β-lactam antibiotic residues, and results indicated that 36.50% of milk samples were found positive for β-lactam antibiotics residues present at above-permissible level (Khaskheli et al., 2008). Other studies suggest a high concentration of different antibiotics in meat, milk, and eggs from various regions of Pakistan (Jabbar, 2013; Solangi et al., 2013) suggesting high usage of antibiotics in FPAs.

Resistance elements from animal pool spread through numerous ways, where the food-borne route probably is the most important route. It is particularly worrisome, when the resistance features (elements such as genes) are acquired by pathogens (Geisinger and Isberg, 2017) such as Salmonella enterica (Angulo et al., 2000; White et al., 2001), Campylobacter jejuni (Altkekruse et al., 1999; Pezzoti et al., 2003) and Yersinia enterocolitica (Frank Möller et al., 1998; Dallal et al., 2010), which are widespread worldwide mainly in industrialized countries (Aarestrup, Frank M et al., 2008). Direct contact between animals and human or human with animal environment may also be a major route of transmission, exemplified by the transmission of methicillin-resistant Staphylococcus aureus (MRSA) (Lee, 2003; Juhász-Kaszanyitzky et al., 2007). Furthermore, commensal bacteria with resistance-conferring elements (Wang et al., 2006) as well as the left-over residues of antibiotic mainly in the form of manure (Heuer et al., 2011), when it affects bacteria in soil or wild fauna, thus conveying resistance features to those living in the soil constituting reservoirs of resistance elements (Kemper 2008; Finley et al., 2013).

Besides food-borne route, the spread of zoonotic pathogens harboring AMR features is a big challenge. There are increasing reports of isolation of Salmonella and Escherichia carrying extended spectrum β-lactamase (ESBL) and quinolones resistance (Xia et al., 2010). For more detailed review readers can read referred review article (Aarestrup, et al., 2008). Altogether, there is a clear adverse effect on public-health because of resistant bacteria spread as a result of usage of antimicrobials in veterinary, agriculture, and other non-human contexts. Such consequences like infections that would not have been otherwise happened, recurrent infections due to treatment failure etc., are included.

One of the reasons that AMR became a concern is its economic impact on public, patients, ecosystem, food chain, healthcare providers and pharmaceutical industries. AMR increases hospital stay, duration of treatment, dosage of treatment and more importantly, rendering otherwise effective antibiotics ineffective. The institute of medicines estimates the annual cost of treating infections due to antibiotic resistant bacteria in the United States to about 4-5 M USD (Institute of Medicine, 1998) despite the fact that calculation of economic impact always leaves a lot of questions behind, mainly, by not incorporating all the contributing elements with its due impact. Though AMR is increasingly reported in animals and humans, however, limited data is available on the magnitude of economic impact and output on healthcare systems,
patients, livestock and agriculture. The issue is even worse in developing countries like Pakistan whereby the national action plan for the confinement of AMR has not yet been implemented, its proposed plans are formulated in line with the WHO.

Patterns of usage of antibiotics in food producing animals in Pakistan: Unfortunately, in Pakistan, AMU in food-producing animals is unregulated and there is no estimation of annual usage. AMR is one of the major health challenges in Pakistan (Levy, 2002a; Goossens et al., 2005; Ali et al., 2010; Idrees et al., 2011; Khan et al., 2013; Adnan et al., 2017; Mohsin et al., 2017; Rahman et al., 2018b). Although, it is hard to find the exact amount of antibiotics being used for treatment, prevention and as growth promoters in FPAs in Pakistan, nevertheless, practices and trends at commercial as well as domestic level indicate that major share of antibiotic usage is aimed for disease prevention and as growth promoters and is offered as supplements on routine basis [https://www.dawn.com/ news/1245840]. Approximately, 50,000 unnecessary registered products and over 600,000 quacks active in the market (http://www.who.int/medicines/areas/coordination/ pakistan.pdf) are further exacerbating the situation. A significant number of antimicrobials used in FPAs are important as human medicines (Van Boeckel et al., 2014). Interviewing different farmers and representatives of companies supplying veterinary medicines in Pakistan revealed that numerous types of antibiotics are used as growth promoters in Pakistan. These include mainly lactam antibiotics (comprising mainly penicillins), lincomamides, and macrolides including erythromycin and tetracyclines, oligosaccharide, avilamycin, flavophospholipol. Virginiamycin etc (Table 1). In European Union, monensin is widely used, which is probably one of the safest and most effective antibiotic growth-promoter with regard to human and animal health and associated bacterial-resistance problems. However, monensin is related to pristinamycin and quinupristin, both of which are applied in human medicine and so there are worries that its consistent usage may compromise human therapy (Butaye et al., 2000). Few studies reported antibiotic residues in poultry meat and dairy milk comprising oxytetracycline, gentamicin, neomycin, enrofloxacin, chloramphenicol, amoxicillin, benzylpeni-cillin, streptomycin, and tyllosin (Muhammad et al., 1997; Shahid et al., 2007a, b), suggesting overuse of these antimicrobials. In the Pakistani context, major contributing factors to the dominantly increasing AMR comprise indiscriminate and overuse of antibiotics in FPAs and all sectors of health, unsuitable selection of antibiotics, poor practices or non-existent institutional antibiotic policies, overwhelming availability of substandard antibiotics in the market, unavailability of appropriate microbiological facilities and research.

Prevalence of antibiotic resistance (AMR) in food-producing animals in Pakistan: For a strategic framework development against AMR, its extent and magnitude must be determined. Unfortunately, national surveillance AR data is lacking, however, fragmented reports have been published mainly focusing on FPAs and agriculture. The data has huge gaps, particularly the extent, quality, quantity and pattern of antibiotic usage is not known, and the overall magnitude of AR is also not determined. However, a few available reports can serve as a point of start for a strategy against AMR. Escherichia coli isolated from poultry have been reported to acquire plasmid-mediated colistin (mcr-1) resistance in recent studies in Pakistan (Azam et al., 2017; Lv et al., 2018; Aqib et al., 2019). Similarly, an estimated 90% of retail meat samples were found contaminated with multidrug-resistant Campylobacter spp. Majority of these isolates were found resistant to enrofloxacin (79.2%), followed by tyllosin (77.6%), ciprofloxacin and amoxicillin (71.2% each), colistin (69.6%), neomycin (32.8%), nalidixic acid (31.2%), gentamicin (25.6%) etc. (Nisar et al., 2017). Multi-drug resistant bacteria harboring antibiotic resistance genes were also found highly prevalent in fish and aquaculture in Pakistan (Shah et al., 2012; Phillips 2000). Environmental contamination with antibiotic residues may also contribute to drug resistance. Water samples of rivers in Punjab (the largest province of Pakistan) were found containing higher than acceptable level of antibiotic residues, and consistent to this, antibiotic resistance genes were also amplified from water samples (Khan et al., 2013; Mohsin et al., 2017). E. coli isolated from chicken and buffalo meat samples

| Antibiotic                          | Class           | Purpose                              | Type of animal     | Comments                   |
|-------------------------------------|-----------------|--------------------------------------|--------------------|----------------------------|
| Amoxicillin and ampicillin          | Peptides        | In aquaculture, treatment and prevention of bovine subclinical mastitis | Bovine and fish    | (Butaye et al., 2003)      |
| Zinc                                | Peptides        | Growth promoters for beef and poultry, prevention of poultry enteritis | Poultry and bovine | General practice in Pakistan |
| Bambermycin                         | Phosphoglycolipids | Growth promoter for poultry and cattle | Poultry and cattle | Interviewing feed mill experts |
| Colistin                            | Cyclopolyptides | Used as growth promoter mainly in poultry and rarely in cattle | Poultry and cattle | Structurally similar to vancomycin group |
| Enrofloxacin                        | Fluoroquinolone | Therapy and prevention of respiratory infections in poultry and cattle | Poultry and bovine | Structurally similar to polymyxin and share cross resistance with Polymyxin |
| Erythromycin                        | Macrolides      | Used in aquaculture, therapy for poultry and bovine infections | Bovine and poultry |                           |
| Lincomycin                          | Lincosamide     | Growth promoters in poultry           | Poultry and bovine |                           |
| Monensin                            | Ionophores      | Growth promoter in poultry, prevention of coccidiosis in poultry | Poultry            |                           |
| Neomycin                            | Aminoglycoside  | Growth promoter for poultry           | Poultry            |                           |
| Sulfonamides                        | Sulfonamides    | Growth promotes in poultry            | Poultry            |                           |
| Tetracycline                        | Tetracycline    | Growth promoter for poultry, prevention of multiple diseases in livestock and poultry | Poultry and livestock |                           |
| Virginiamycin                       | Streptogramins  | Growth promoter for poultry (broiler) | Poultry            |                           |
were found resistant to sulfamethoxazole-trimethoprim harboring sul1 and sul2 genes conferring resistance to sulfonamides (Idrees et al., 2011). A study on chicken meat samples at retail market in Hyderabad Pakistan indicated that 38% meat samples contaminated with Salmonella and all of them were found resistant to ampicillin (Soomro et al., 2011). Similarly, of 126 samples from broiler meat 71.4% were found contaminated with E. coli and majority of them were resistant to amoxicillin and tetracycline.

The percentage resistance against streptomycin and sulphonamethoxazole-trimethoprim was found to be high in backyard poultry (64.2 and 53.5%) as compared to the commercial broiler (28.5 and 35.7%) (Akhtar et al., 2016). A study carried out in Karachi reported that out of 340 samples of meat analyzed for the isolation of ARMOs 66% contained pathogenic bacteria including 33% Salmonella isolates that were all found resistant to ampicillin. Furthermore, of these pathogenic bacteria (namely E. coli, Klebsiella, Enterobacter species and Staphylococcus aureus), 16% were capable of biofilm formation (Ali et al., 2010). Similarly, a study conducted in 2014 in Lahore identified a number of pathogenic species such as E. coli, Klebsiella, Shigella, Salmonella and Enterobacter recovered from different market-food items such as dahi bhallay (a spicy appetizer), yoghurt and fruit-chat (combination of different seasonal fruits with sauce). Of these isolates, 46% were found resistant to commonly used antibiotics (Saleem et al., 2014). Similar incidence of AMROs in ready to eat raw food items has also been reported in other parts of the country (Shah et al., 2012; Capita and Alonso-Calleja 2013; Siddiqui et al., 2015 Nasim et al., 2016).

Where Pakistan stands in the perspective of global regulations and what strategy should be adopted: Pakistan has established its national body, Division of Pharmacy Services of the Drug Authority of Pakistan (DRAP) (http://www.dra.gov.pk/), in November 2012, however, unfortunately, the regulations drafted have not yet implemented strictly. Similarly, Pakistan has also established a policy of restricted use of antibiotics in animal husbandry; however, the policy is also not yet been implemented strictly Pakistan has further improved the situation by constituting a national strategic frame work for containment of AR in 2016 (https://drive.google.com/file/d/0B6zZgBqF00pF00pFV0J0a2pqeVFvS3JzR2cyd1NhR0stSTBoVzR3/view), in line with WHO global action plan to fight AMR. As a commitment to the global action plan to tackle AMR, Pakistan has recently developed its national strategic framework for containment of AMR with intention to convert this to an AMR National Action Plan (NAP) by “One-Health” approach (https://www.nih.org.pk/wp-content/uploads/2018/08/AMR-National-Action-Plan-Pakistan. pdf). As per the guidelines of national action plan for confinement of AMR, non-therapeutic use of antibiotics in FPAs and plants should be discouraged. Establishment and strengthening of AMR surveillance, initiation and encouragement of antibiotic stewardship plans, improving farm biosecurity and awareness against AMR should immediately be adopted in all relevant veterinary, public and private, stakeholders. For high producing animals, other innovative approaches like development of vaccines, alternative treatments like use of phages, and medicinal plants should be encouraged to replace the use of antibiotics. Molecular epidemiological data using One-Health approaches is required to link resistance clones or genes from different sample sources such as, from animals of different origin, human and the environment.

Maximum residual limits have been established by various countries such as Canada, China, Japan and member states of the EU in order to protect consumers from the side effects of drug residues used in production of FPAs (Reig and Toldrá 2008) and EU countries have even withdrawn the use of ionophores in feed (Casewell et al., 2003; Hunter et al., 2010). Unfortunately, despite the implementation of strict guidelines for the use of antimicrobial agents in FPAs world-wide, they are still being used illegally in many developing countries including Pakistan (Byarugaba, 2004). In Sweden, no antibiotic as growth promoters is allowed, while in the US a range of antibiotics are allowed as growth promoters. Overall, the use of antibiotics in different FPAs differ enormously. The World Health Organization (WHO) and Food and Agriculture Organization (FAO) are working together at international level for devising the AMU. The Codex Alimentarius Commission (CAC), under the WHO and FAO, has established standards and recommendations that should be implemented by all countries as a code of practice to minimize and contain AMR (Anonymous, 2005). In conclusion, challenging and daunting tasks are ahead of Pakistan to fight AMR issue in the near future. Gaps and hurdles include weak health infrastructure and relevant expertise, illiteracy, lack of awareness among the clinicians, farmers and public and lack of access to basic needs, clean water and vaccination. National level strategies including infection control and antibiotic stewardship must urgently be initiated and implemented, where it has already been drafted to improve antibiotic prescription and utilization along with nationwide vaccination to control several epidemic infections. This should further be assisted with regular surveillance programme to determine the extent and trend of AMR.

Factors affecting the spread and dissemination of resistance: To simplify the resistance phenomenon, we must keep working and concentrate on two main factors; the antibiotics itself that provide selective pressure to dominate the second important element - the resistance genes. After all, if for example, either the antibiotics or the resistance genes were not existed, we would not have the phenomenon of resistance. Shortly, a more logical way to look at this problem would be either to avoid selection and/or to stop the dissemination of resistant bacteria. Thus, dissemination of antibiotic resistance can be limited either through management of the selective pressure leading to resistance or through interventions aimed at restricting the spread of the selected resistance traits.

In the United States alone, an estimated 23×10^4 of antibiotics are used annually; half of this are provided to people for use during sickness and the rest is for agricultural use (Harrison and Lederberg 1998). About 7 x 10^5 kg of antibiotics, mainly penicillin and tetracycline, are annually used in FPAs as growth promoters (Harrison
and Lederberg 1998). On top of that, an estimated 45 x 10^3 kg of antibiotics, chiefly tetracycline and streptomycin, are sprayed over the fruits and crops annually. Most of these compounds would certainly exist in environment as residues and bacteria would confront them at a point. A pool of genes, thus, conferring resistance against these antibiotics would certainly accumulate/concentrate in the environment in response to selective pressure provided by the existing residues antibiotics in the environment, animal body and food. Such elements could also easily be spread through river water or by wild life and migratory birds to other areas as identified by Raza and his colleagues in Pakistan (Molsin et al., 2017; Raza et al., 2017). We could not identify the amount of antibiotics used in Pakistan due to unavailability of reports, however, based on the prevailing practices about how easily antibiotics are accessible in the country, it is not hard to estimate that we are currently using absolutely higher amount of antibiotics in clinical settings, agriculture and food animals. Antibiotics in Pakistan are available in the pharmacy stores, and even vendors are selling them out on the streets. This chaotic way of availability of antimicrobials and drug distribution scenario further nurtures emergence of antibiotic resistance. This blind use of antibiotics is thus encouraging new elements of AMR to emerge against antibiotics. Antibiotics are excreted in the environment, water, crops etc., where they keep exerting selective pressure resulting in “post-therapy” environmental selection phase of the antibiotics (Levy, 2001). During such phase, the antibiotic concentration would be less than the therapeutic concentration, which is ideal for selecting ARMOs. Thus, considering this scenario, it may not be the use of consistent high concentration of antibiotics in clinics for treatment (treatment period) rather the slow and persistence release of low amount of antibiotics in the environment (post therapy period) that provide ample amount of time for bacteria to develop resistance. Altogether, we should change our attitude and course of action against the use of antibiotics. First, we should emphasize to implement the shorter course of antibiotics. Secondly, cycling of the antibiotics should be considered, if new antibiotics are available in the market. Thirdly, we should alongside invest in discovering and synthesizing new antibiotics such as improvement of the current tetracycline drugs, understanding the mechanisms of resistance to currently available tetracycline drugs would help improve the efficacy and the fight against resistance. A non-chemical, non-classical approach to reversing the resistance problem would be the revival of the susceptible strain. We need to encourage the growth of the susceptible strain to take over the resistance one.

Conclusions: Despite lack of a routine nation-wide surveillance, existing literature show that AMR is escalating in Pakistan. An urgent intervention is needed to restrain emergence and spread of AMR. Pakistan should learn from other countries who observed a dramatic decrease in resistance to cephalosporin after stopping use of β-lactam and cephalosporins in poultry production. Proper planning, improved use of antimicrobials, establishment of effective surveillance systems, constituting and enforcing legislature and combined international actions should immediately be taken for the control of antibiotic resistance spread and reversal of resistance. Infection control measures and stewardship programs are vital weapons in controlling the pandemic evolution of multidrug-resistance. In conclusion, AMR is a complex multifactorial phenomenon in context of lower-middle income countries where the social behavior and knowledge of farmers and veterinarians towards AMU and AMR is imperative for the design of successful control strategies.

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REFERENCES
Aarestrup FM, 2005. Veterinary drug usage and antimicrobial resistance in bacteria of animal origin. Basic Clin Pharmacol Toxicol 96:271-81.
Aarestrup FM, Wegener HC and Collignon P, 2008. Resistance in bacteria of the food chain: epidemiology and control strategies. Expert Review Anti-infective Ther 6:673-50.
Adnan M, Khan H, Kashif J et al., 2017. Clonal expansion of sulfonamide resistant Escherichia coli isolates recovered from diarrheic calves. Pak Vet J 37:230-2.
Akhatar F, Rabbani M, Muhammad K et al., 2016. Comparative antibiotic resistance profile of the multidrug resistant E. coli isolated from commercial and backyard poultry. J Anim Poult Sci 26:1628-32.
Alexander T, Yanke L, Topp E et al., 2008. Effect of subtherapeutic administration of antibiotics on the prevalence of antibiotic-resistant Escherichia coli bacteria in feedlot cattle. Appl Environ Microb 74:4405-16.
Ali NH, Farooqui A, Khan A et al., 2010. Microbial contamination of raw meat and its environment in retail shops in Karachi, Pakistan. J Infect Dev Ctries 4:382-8.
Altekruse SF, Stern NJ, Fields PI et al., 1999. Campylobacter jejuni—an emerging foodborne pathogen. Emerg Infect Dis 5:28.
Angulo FJ, Johnson KR, Tauxe RV et al., 2000. Origins and consequences of antimicrobial-resistant nontyphoidal Salmonella: implications for the use of fluoroquinolones in food animals. Microb Drug Resist 6:77-83.
Anonymous, 2005. Code of practice to minimize and contain antimicrobial resistance. CAC/RCP:61-2005.
Aqib A, Nighat S, Ahmed R et al., 2019. Drug susceptibility profile of Staphylococcus aureus isolated from mastitic milk of goats and risk factors associated with goat mastitis in pakistan. Pak J Zool 51:307-15.
Azam M, Ehsan I, Saleemi MK et al., 2017. Detection of the colistin resistance gene mcr-I in avian pathogenic Escherichia coli in Pakistan. J Glob Antimicrob Res 11:152-3.
Butaye P, Van Damme K, Devriese L et al., 2000. In vitro susceptibility of Enterococcus faecium isolated from food to growth-promoting and therapeutic antibiotics. Int J Food Microbiol 54:181-7.
Byarugaba D, 2004. Antimicrobial resistance in developing countries and responsible risk factors. Int J Antimicrob Agents 24:105-10.
Capita R and Alonso-Callega C, 2013. Antibiotic-resistant bacteria: a challenge for the food industry. Crit Rev Food Sci Nutr 53:11-48.
Casewell M, Friis C, Marco E et al., 2003. The European ban on growth-promoting antibiotics and emerging consequences for human and animal health. J Antimicrob Chemother 52:159-61.
Dalali MMS, Doyle MP, Rezaadehbash M et al., 2010. Prevalence and antimicrobial resistance profiles of Salmonella serotypes, Campylobacter and Yersinia spp. isolated from retail chicken and beef, Tehran, Iran. Food Control 21:388-92.
Finley RL, Collignon P, Larsson D et al., 2013. The scourge of antibiotic resistance: the important role of the environment. Clin Infec Dis 57:704-10.
Geisinger E and Isberg RR, 2017. Interplay between antibiotic resistance and virulence during disease promoted by multidrug-resistant bacteria. J Infec Dis 215:59-17.
Goossens H, Ferech M, Vander Stichele R et al., 2005. Outpatient antibiotic use in Europe and association with resistance: a cross national database study. Lancet 365:579-87.
Graham JP, Evans SL, Price LB et al., 2009. Fate of antimicrobial-resistant enterococci and staphylococci and resistance determinants in stored poultry litter. Environ Res 109:682-9.
