Applied Science and Technology Annals. Vol.1, No. 1 (2020); 187–193 ISSN: 2717-5014 (Print). Available online at www.recast.tu.edu.np DOI: https://doi.org/10.3126/asta.v1i1.30305



Review Article **Coronaviruses in animals and humans, COVID-19 pandemic and one health approach**

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Received: May 23, 2020; Accepted: June 14, 2020; Published: June 25, 2020

Abstract: Coronaviruses can infect several animal species including cattle, pigs, dogs, and cats resulting in diseases related to respiratory and gastrointestinal systems. In humans, coronaviruses generally cause mild to moderate illnesses of the respiratory tract. Severe Acute Respiratory Syndrome Coronavirus (SARS-CoV) and Middle East Respiratory Syndrome Coronavirus (MERS-CoV), which emerged during 2002/03 and 2012/13 respectively, caused severe respiratory illnesses in humans. In December 2019, a novel respiratory coronavirus, SARS coronavirus 2 (SARS-CoV-2) emerged from Wuhan, China and caused coronavirus disease 2019 (COVID-19). Owing to the rapid spread of this virus, World Health Organization (WHO) declared COVID-19 outbreak as a global pandemic, which claimed over 300,000 lives by 16th May 2020. Data available so far indicate that COVID-19-associated severe illnesses, hospitalizations and deaths are more common in elderly above 65 years of age; in men; and in individuals with underlying health conditions such as cardiovascular disease, hypertension and diabetes. SARS-CoV-2 is considered to be emerged from bats and likely involved certain, yet to be identified, intermediate animal host. Prevention and control of ongoing COVID-19 pandemic and possible disease outbreaks in the future by other emerging and reemerging pathogens, requires efficient implementation of one health strategy that utilizes the expertise of human, animal and environmental health sectors.

Keywords: Coronavirus, COVID-19, One Health, Pandemic, SARS-CoV-2

सारांश: कोरोना भाइरसले गाईवस्तु, सुँगुर, कुकुर र बिरालाहरू लगायतका धेरै जनावरका प्रजातिहरूलाई संक्रमित गर्न सकछ र श्वासप्रश्वास तथा पाचन प्रणालीमा सम्बन्धित रोगहरू लगाउछ। मानवमा कोरोना भाइरसले सामान्यतया श्वासप्रश्वास प्रणालीमा हल्कादेखि मध्यम रोग निम्त्याउँछ। सन् २००२/०३ र सन् २०१२/१३ मा क्रमस सार्स र मर्स नाम गरेका कोरोना भाइरसहरु देखापरे जसले मान्छेमा श्वासप्रश्वासको गम्भीर रोगहरू निम्त्यायो। डिसेम्बर २०१९ मा, सार्स कोरोना भाइरस र नाम गरेको नया कोरोना भाइरस चीनको वुहानमा देखा पऱ्यो जसले मान्छेमा श्वासप्रश्वासको गम्भीर रोग लगायो। डिसेम्बर २०१९ मा, सार्स कोरोना भाइरस २ नाम गरेको नया कोरोना भाइरस चीनको वुहानमा देखा पऱ्यो जसले कोविड-१९ नाम गरेको गम्भीर रोग लगायो। डिसेम्बर २०१९ मा, सार्स कोरोना भाइरस २ नाम गरेको नया कोरोना भाइरस चीनको वुहानमा देखा पऱ्यो जसले कोविड-१९ नाम गरेको गम्भीर रोग लगायो। यस भाइरसको तीव्र प्रसारको कारण, विश्व स्वास्थ्य संगठनले कोभिड-१९ लाइ विश्वव्यापी महामारीको रूपमा घोषित गऱ्यो। सन् २०२० मे १६ सम्म कोभिड-१९ ले विश्वभर ३००,००० भन्दा बढीको ज्यान लिइसकेको छ। अहिलेसम्म उपलब्ध तथ्यांक अनुसार कोभिड-१९ संग सम्बन्धित अस्पताल भर्ना र मृत्यु ६५ वर्षमाथिका वृद्धहरुमा, पुरुषहरुमा, र हृदय रोग, उच्च रक्तचाप तथा मधुमेह जस्ता रोगका विरामीहरुमा बढी देखिएको छ। सार्स कोरोना भाइरस २ को उत्पति चमेराबाट भएको मानिएपनि मान्छेमा सर्नुअघि यो भाइरसले कुन जनावरको सहारा लियो त्यो पत्ता लागिसकेको छैन। अहिले संसारभर फैलिरहेको कोभिड-१९ तथा भविष्यमा फैलन सक्ने वस्तै महामारीहरुको उचित रोकथाम र नियन्त्रणको लागि एक स्वास्थ्य रणनीतिको कुशल कार्यान्वयन आवश्यक छ जसले मानव, पशु र वातावरणीय स्वास्थ्य क्षेत्रहरूको विशेषज्ञताको समुचित प्रयोग गर्दछ।

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I. Introduction

Coronaviruses are positive-sense, single-stranded RNA, and enveloped viruses belonging to the order Nidovirales, family Coronaviridae and subfamily Coronavirinae (Fehr & Perlman, 2015). They have RNA genome of 26-30 kb size, one of the largest among the RNA viruses. Coronaviruses are classified further into four genera: alpha, beta, gamma, and delta coronaviruses. They have four types of structural proteins: spike, membrane, envelope, and nucleocapsid proteins, but additional hemagglutinin-esterase protein is also present in beta coronaviruses. Spike protein is responsible for attachment of the virion to the host cell (Gallagher & Buchmeier, 2001). These viruses use different cellular receptors, such as human aminopeptidase N (CD13), angiotensin converting enzyme 2 (ACE2), 9-O-acetvlated sialic acid, and dipeptidvl peptidase 4 for attachment and cell entry (Lim et al., 2016). These receptors are regarded as the determinants of host range, tissue tropism, and pathogenicity.

Coronaviruses infect both humans and animals.

Those infecting humans are mostly clustered in alpha and beta coronaviruses whereas viruses infecting animals fall in all four genera. Human coronaviruses typically cause respiratory tract illnesses including common cold, cough, pneumonia, and bronchitis (Pene et al., 2003; Lim et al., 2016). Coronaviruses can infect several animal species including cattle, pigs, dogs, cats, and poultry resulting in diseases related to respiratory and gastrointestinal system (Saif, 2004; Fehr & Perlman, 2015). Bats are considered as a reservoir host for many coronaviruses, while zoonotic transmission to humans is possible through intermediate animal hosts (Su et al., 2016; Li et al., 2020). The current pandemic of coronavirus disease 2019 (COVID-19) is caused by a Severe Acute Respiratory Syndrome Coronavirus 2 (SARS-CoV-2). In this review, we have summarized different coronaviruses infecting animals and humans; status of COVID-19 pandemic; and the importance of one health approach to deal with outbreaks like COVID-19.

Species	Virus	Genus	System affected
Pig	Transmissible gastroenteritis virus (TGEV)	Alpha coronavirus	Gastrointestinal
	Porcine epidemic diarrhea virus (PEDV)	Alpha coronavirus	Gastrointestinal
	Porcine delta coronavirus (PdCV)	Delta coronavirus	Gastrointestinal
	Porcine respiratory coronavirus (PRCV)	Alpha coronavirus	Respiratory
Cattle	Bovine coronavirus	Beta coronavirus	Both gastrointestinal and respiratory
Chicken	Infectious bronchitis virus	Gamma coronavirus	Respiratory
Turkey	Turkey coronavirus	Gamma coronavirus	Gastrointestinal
Dog	Canine coronavirus	Alpha coronavirus	Gastrointestinal
	Canine respiratory coronavirus	Beta coronavirus	Respiratory
Cat	Feline coronavirus	Alpha coronavirus	Gastrointestinal

Table 1. Summary of coronavirus infection in different animal species

2. Coronaviruses in Animals

Historically, coronaviruses were first identified in the chickens in 1930s in the form of infectious bronchitis (IB) virus (Beach & Schalm, 1936; Estola, 1970). Since then, many coronaviruses and associated diseases are reported in livestock and pet animals (Table 1). For example, bovine respiratory coronavirus (BCoV) causes mild to severe respiratory tract and enteric infection in cattle (Saif, 2010; Amer, 2018). Interspecies transmission of BCoVs can occur between ruminants like camels, elk, deer, and water buffalo (Amer, 2018). Similarly, Transmissible Gastroenteritis Virus (TGEV) and Porcine Epidemic Diarrhea Virus (PEDV) are the alpha coronaviruses that cause severe gastroenteritis in piglets less than 2 weeks of age with almost 100% mortality (Laude et al., 1990; Jung & Saif, 2015). In pigs, porcine delta coronavirus is also responsible for acute diarrhea, vomiting, and mortality in neonatal pigs (Jung et al., 2016). The emergence of highly

virulent PEDV strain in the United States (US) in 2013, resulted in the death of around 8 million pigs leading to huge economic losses in the US swine industry (Stevenson et al., 2013). Coronaviruses have been associated with infection in pet animals as well. Canine coronaviruses cause both gastrointestinal and respiratory disease in dogs which are usually mild with severe infection in some instances (Buonavoglia et al., 2006). Coronavirus infection is also common in cat resulting in feline infectious peritonitis (Hartmann, 2005). Coronavirus infection in animals is widespread globally however, there are not many reports available on the status of coronavirus-related diseases among animals in Nepal. At least, some information is available related to avian coronavirus i.e., avian infectious bronchitis virus (IBV) infection in Nepal (Gompo et al., 2019; Office International des Epizooties [OIE], 2019a). According to an epidemiological study conducted by Gompo et al. (2019), infectious bronchitis is among the top 10 poultry diseases in Nepal and the existence of the disease in avian species is also supported by reports submitted to OIE.

3. Coronaviruses in Humans

The first human coronavirus (HCoV). B814 virus. was discovered in 1960s (Tyrrell & Bynoe, 1966; Kahn & McIntosh, 2005). This virus was isolated from the nasal wash of the individual who developed cold. Since then the other HCoVs, namely HCoV-229E, HCoV-NL63, HCoV-OC43, and HCoV-HKU1 are known to cause a significant fraction of common cold infections globally with mild to moderate diseases (Su et al., 2016). HCoV-229E, HCoV-NL63 belong to genera alpha coronavirus while HCoV-OC43 and HCoV-KHU1 belong to genera beta coronavirus. These virus infections are most common during the winter season (Gaunt et al., 2010; Uddin et al., 2018). One study conducted among infants in Nepal, between 2011-2013, showed 8% prevalence of HCoVs (Uddin et al., 2018). The four HCoVs were found to be in the infections with the highest prevalence of HCoV-OC43 followed by HKU1, NL63, and 229E in Nepal. Diseases related to human coronaviruses were considered mild before the outbreak of severely pathogenic severe acute respiratory syndrome coronavirus (SARS-CoV). The SARS outbreak which occurred in 2002/03 resulted in over 8000 cases and 774 deaths in 27 countries (de Wit et al., 2016). After a decade, Middle East Respiratory Syndrome Coronavirus (MERS-CoV) emerged. By 2019, there were 2494 confirmed cases and over 858 deaths were reported due to MERS in 27 countries (World Health Organization [WHO], 2020a). SARS-CoV was originated from bat, and palm civet is regarded as the intermediate animal

host (Wang et al., 2005; Hu et al., 2015). Likewise, for MERS, bats are the reservoir host and camels are regarded as intermediate animal host (Omrani et al., 2016). No cases of SARS and MERS were ever reported in Nepal (WHO, 2020a, 2020b). Even within less than a decade of the occurrence of MERS outbreak, another coronavirus named SARS coronavirus 2 (SARS-CoV-2) emerged that is causing the COVID-19 pandemic.

4. COVID-19 Pandemic

A novel coronavirus causing severe respiratory illness in people was first reported in December 2019 from Wuhan, China. This novel coronavirus had nearly 90% nucleotide sequence similarity to batorigin SARS-like coronaviruses and caused disease similar to earlier SARS-CoV infection in humans (Wu et al., 2020). World Health Organization (WHO) later named the virus as SARS coronavirus 2 (SARS-CoV-2) and the disease as COVID-19 (WHO, 2020b). Owing to the rapid spread of this virus worldwide, WHO declared outbreak of COVID-19 as a pandemic on March 11, 2020 (WHO, 2020b). It is likely that SARS-CoV-2 originated from bats but the information about the intermediate animal hosts involved is not available yet (Lau et al., 2020).

SARS-CoV-2 is primarily transmitted through respiratory droplets and contact with the infected people (Liu et al., 2020). Transmission may also occur through contact with contaminated fomites around the infected person (Ong et al., 2020). People infected with SARS-CoV-2 show a wide range of symptoms that range from mild to severe illnesses. The common symptoms include fever, cough, shortness of breath, headache, fatigue and muscle pain. Symptoms may develop in 2 to 14 days in the infected individuals (Centers for Disease Control and Prevention [CDC], 2020a). Most people infected with SARS-CoV-2 remain asymptomatic that could be the major reason behind rapid transmission of COVID-19 (Furukawa et al., 2020).

SARS-CoV-2 has been isolated from stool of infected person suggesting that fecal-oral transmission might be a possibility, but it needs further confirmation (WHO, 2020c; Zhang et al., 2020). Recently, multiorgan tropism of SARS-CoV-2, beyond the respiratory tract and including the kidneys, heart, liver, and brain, is also reported during autopsy of people who died with COVID-19 (Puelles et al., 2020).

As of 16th May 2020, SARS-CoV-2 has infected over 4.6 million people and claimed over 300,000 lives from 215 countries and territories in the world (Coronavirus Update Worldometer, 2020; WHO, 2020c). Host-associated factors including biological sex (being man or woman) and age (being a child or reproductive age adult or elderly) can play a significant role in the outcome of infection or immunity development (Dhakal & Klein, 2019). During COVID-19 pandemic, severe illnesses, hospitalizations, and deaths are mostly observed in elderly above 65 years of age (CDC, 2020a). Likewise, most of the countries that segregated data by biological sex (being a man or woman) have shown that death rate due to COVID-19 is greater in men than in women (Klein et al., 2020). Similarly, underlying medical conditions such as chronic respiratory disease, cardiovascular disease, cancer, diabetes, and hypertension are mostly associated with a higher severity of the disease and deaths (Jordan et al., 2020; Wu et al., 2020).

5. COVID-19 Prevention and Control Strategies

Isolation of SARS-CoV-2 infected or suspected people and restriction in their movement is highly recommended by WHO and Centers for Disease Control and Prevention (CDC) of USA in case of COVID-19 disease (WHO, 2020d; CDC, 2020b). Maintaining personal hygiene through regular handwash with soap water for at least 20 seconds and use of hand sanitizer can limit the spread of the virus infection. Additionally, other hygienic habits like avoiding touching mouth, nose, and eye with hands should be considered. Wearing nonsurgical and cloth masks for general public and N95 masks for health professionals working closely with infected or suspected people are also regarded as the best practices to prevent spread of droplets from the infected to the susceptible persons (Food and Drug Administration [FDA], 2020b).

In addition to personal hygiene, nonpharmaceutical interventions like social distancing and staving at home are also recommended and enforced in many countries (WHO, 2020c; CDC, 2020b). The effective use of the personal hygiene and nonpharmaceutical interventions combined with massive testing were helpful in getting control over COVID-19 outbreak in China and hence almost all other countries are also following the similar strategies to control the SARS-CoV-2 transmission (Adhikari et al., 2020). As of now, there is no approved treatment available for COVID-19 but several research trials at different levels using different drugs are ongoing. There are reports that drugs including remdesivir, hydoroxychloroquine, azithromycin etc. could be effective to treat COVID-19 (Gautret et al., 2020; Manli Wang et al., 2020). On May 1st, 2020, the US Food and Drug Administration (FDA) provided 'emergency use authorization' to the drug remdesivir owing to its observed benefits in animal and human studies

(FDA, 2020a). In addition to the therapeutic treatments, tremendous efforts have been applied by scientists all over the world to develop vaccines against the SARS-CoV-2 infection. So far, more than 115 vaccine candidates have been developed and are in either preclinical or early stages of clinical trials (Le et al., 2020). Under normal circumstances, development of vaccines takes years. However, owing to the devotion of each and every scientific body throughout the globe; and investment and supports from the private and public resources; it is likely that vaccine against COVID-19 will be developed in 12 to 18 month period (Amanat & Krammer, 2020).

6. One Health Approach and COVID-19 Pandemic

One health is a public health strategy to combine the efforts of multisectoral professions including human health, animal health, and environmental health to tackle the emerging and re-emerging pathogens and achieve better public health outcomes (Kelly et al., 2017; Lebov et al., 2017). About 75% of the human infectious diseases have animal origin and 60% are zoonotic. This highlights the importance of professionals working in both animal and public health to join hands together to address the heath issues at human and animal interface (OIE, 2019b). One health approach is important to address several global health issues including antimicrobial resistance and control of zoonoses (McEwen & Collignon, 2018). Prevention of zoonotic disease is only possible through coordinated efforts among veterinary and public health sectors. For example, rabies in humans can only be prevented by getting control over the animal source of the virus. Similarly, control of influenza viruses in human is only possible through control in animals to prevent the emergence of new strains and subtypes (Capua & Cattoli, 2010).

One health approach helps to identify the source of infection and track the pathogen transmission pattern which is critical to reduce the challenges of emerging and reemerging viruses (Kelly et al., 2017; Lebov et al., 2017). SARS-CoV-2 emerged from its ancestral bat host and it is likely that certain intermediate animal host was involved before its spillover to humans (Zhou et al., 2020). In the context of unknown intermediate host, one health approach can be helpful to understand the SARS-CoV-2 transmission from bat to humans. Early studies have shown that SARS-CoV-2 can infect dogs, cats, and tigers, though their roles in virus transmission to humans is not yet confirmed (American Veterinary Medical Association [AVMA], 2020). Determination of the intermediate host for SARS-CoV-2 transmission will help to investigate further on host-pathogen interaction and design effective plans to prevent the virus transmission (El Zowalaty & Järhult, 2020). The ongoing COVID-19 pandemic has once again highlighted the importance of research at humanenvironment-animal interface to ensure the timely and efficient response against such outbreaks even in the future (Amuasi et al., 2020).

Lately, the importance of one health approach has been realized in Nepal as well, but there is a long way to go before its proper implementation. The veterinary professionals in Nepal have been dealing with avian influenza virus and other outbreaks for a long time resulting in the development of better research and diagnostic expertise and facilities. In COVID-19 ongoing pandemic, multiple the veterinary laboratory facilities and resources are being utilized for the disease diagnosis in Nepal. This fact emphasizes the necessity of continued collaboration between human health sector, animal health sector, microbiologists, public health experts, and other stakeholders in the future to prevent and control any such outbreaks as COVID-19.

7. Conclusion

COVID-19 pandemic already infected millions of people and killed thousands all over the world. In the absence of approved therapeutics and vaccines, it will continuously affect a larger population in coming days leading to more cases and deaths. Despite the ongoing nonpharmaceutical mitigation efforts and apparent elimination, the detrimental effect of SARS-CoV-2 is expected to persist longer (Kissler et al., 2020). Scientists are thinking that this virus may never disappear and keep circulating as influenza viruses. Thus, there is urgent need of development of vaccines and drugs to deal with COVID-19. Moreover, the world is facing pandemics regularly at the interval of 10-50 years. Owing to the emergence and re-emergence of several zoonotic viruses, it can be predicted that another pandemic is also inevitable in the future. Proper implementation of one health approach that brings environmental, animal, and human health experts together will thus be vital for better preparedness against future disease outbreaks.

Authors Contribution: SD conceived the idea; MKC, SG, NB and SD wrote and revised the manuscript. All authors have read and approved final version of the manuscript.

Conflicts of Interest: The authors declare no conflicts of interest.

Funding: No funding was sought or obtained to conduct this study.

Ethical approval

Not required given the nature of the article.

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