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Review Article

Misuse of Antibiotic during COVID 19 Outbreaks

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Abstract



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The COVID-19 outbreak, caused by the severe acute respiratory syndrome coronavirus 2, has been detected in December 2019 in Wuhan, China, and is accompanied by significant degrees of morbidity and mortality. Antibiotic resistance (AMR) threatens to wreak havoc on healthcare system and the impact of globalization, and COVID-19 is intended to keep it at bay for the time being. During the COVID-19 crisis, a wide range of antimicrobial medicines were promoted as viable treatments. While both industrialized and industrializing nations have seen a rise in antimicrobial medication usage, use and abuse have been significantly more prevalent in developing countries. Antibiotic resistance is still a concern since microorganisms that cause resistant infections develop in hospitals and medical institutions, placing all patients at risk, complicating the care of COVID-19. Improper prescriptions, a lack of care management policies and needless self-administration by the general population are examples of these. Antibiotics seem to be more motivated to abuse and misusing antibiotics than to keep them safe and take them only when prescribed. Almost all of the substantial antibiotic usage in COVID-19 patients is inevitable. Patients having lung viral infections are more likely to develop subsequent bacterial infections, which lead to higher disease severity and death. Immediately crucial components of any AMR mitigation approach are increased spending in education and increased public knowledge of AMR. More studies are needed to better understand the health risks and rate of co-infection in COVID-19 patients in order to promote a decrease in any unnecessary antimicrobial prescribing.

Keywords: COVID-19; Antibiotic use and misuse; Antimicrobial Resistance.

INTRODUCTION:

Coronaviruses (CoVs) have been linked to severe disease outbreaks in East Asia and the Middle East during the last two decades. The Middle East respiratory syndrome (MERS) and the severe acute respiratory syndrome (SARS) first appeared in 2002 and 2012, respectively. In late 2019, a new coronavirus, severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2), which causes coronavirus disease 2019 (COVID-19), appeared, posing a worldwide health hazard and generating an ongoing pandemic in numerous nations and territories¹. The World Health Organization (WHO) has named the current CoV-related disease COVID-19, which is caused by SARS-CoV-2. The main cluster of diseases has been shown to be linked to Wuhan's Huanan South Chinese seafood market². The corona virus, which belongs to the corona virus family, is primarily responsible for the illness. These members of the family are single-stranded, positively sensitive RNA viruses that cause mild respiratory infections in humans. However, in some circumstances, they can cause major difficulties in the central nervous system, gastrointestinal tract, and respiratory system^{3, 4}. This hidden illness is antimicrobial resistance (AMR), which is

aggravated by antibiotics. There is no such thing as a vacuum when it comes to the discovery of novel therapeutic antibiotics^{5,6}. Immunocompromised patients who have already been infected with a respiratory infection, such as TB, influenza, acute respiratory syndrome (SARS), or presently covid-19, are at a higher risk of getting serious illnesses⁷. The spread of SARS-CoV-2 is quite troubling because it has the potential to be very severe in low- and middle-income nations (LMICs), Antibiotics help lower morbidity and death linked with infectious illnesses in underdeveloped nations where health remains poor^{8,9}. As the globe responds to COVID-19, a major underlying threat of antimicrobial resistance (AMR) lurks. It was already killing hundreds of thousands of people worldwide at the time (approximately 700 000 deaths per year). The widespread and inappropriate use of antibiotics, among other factors, aided in the formation and dissemination of anti-pathogens¹⁰⁻¹². Unfortunately, microorganisms developed resistance to existing antibiotics within a few years after their introduction. Fungus has generated an equivalent level of resistance¹³. There has been a lack of systematic measures to minimise AMR, resulting in millions of fatalities worldwide and on a yearly basis AMR has the potential to become a

more frequent component by 2050. Death is more common than cancer¹⁴. AMR is frequently a major cause of death in developing countries due to a lack of exposure to the most economical and efficient antibiotics¹⁵. COVID-19 poses issues in all aspects of health care, including the management of chronic illnesses and non-covid-19 severe bacterial infections¹⁶. Health-care systems are agitated, Implement anti-infective policies by isolating infected COVID-19 patients effectively in order to limit the virus's spread¹⁷. Antibiotics may be prescribed to COVID-19 patients for a number of reasons. Influenza patients have a higher death rate due to bacterial superinfection¹⁸. The widespread use of antibiotics raises concerns about overuse, which can convert into Antimicrobials and the development of a plethora of medications creatures that already exist by 2050, it is anticipated to be the leading cause of mortality worldwide¹⁹. Antibiotic prophylaxis, for example azithromycin may play a role in the progression of drug-resistant strains of *Salmonella typhi*²⁰. In the setting of COVID-19 in undeveloped countries, there are also various relevant variables coming from the general population that encourage the development of antibiotic resistance.

Improper prescriptions, a lack of care management policies, and needless self-administration by the general population are examples of these. However, challenges in differentiating between Pneumonia caused by a bacterial infection and COVID-19 infection, as well as a paucity of antiviral medication with proven success, requires the use of antibiotics as part of the experience. The treatment plan for COVID-19 patients is critical ill²¹. The goal of this study was to consolidate information about the collateral effect of the COVID-19 pandemic on AMR containment based on the limited evidence that was available at the time. Our goals are to: share current understanding on the onset of the SARS-CoV-2 and COVID-19 pandemics, and emphasise the potential role of the COVID-19 pandemic to the ongoing development and spread of AMR. In this paper, we examine the influence of the present COVID-19 pandemic on the danger of AMR, focusing on the relationship of bacterial coinfections in COVID-19 patients.

INADEQUATE SUPERVISION AND IMPLEMENTATION:

Antimicrobial dispensing is controlled in developed nations, and medication is normally necessary. However, the ability to buy antimicrobials online has made them more widely available in wealthy countries^{9, 22}. Antibiotics are widely available in many underdeveloped countries, can be obtained from street vendors and are available without a prescription. In the absence of this control, antibiotics are readily available and accessible, promoting self-medication and incorrect dosing²³. Regulations are connected with inadequate drug monitoring systems, which manifest as incorrect selection, usage, and storage practices, allowing bacteria to evolve and spread resistance²⁴.

THE INFLUENCE OF COVID-19 ON ANTIMICROBIAL RESISTANCE:

Three factors influence the progression of AMR in a population: emergence, transmission, and population-level infections load. AMR is caused by selection pressures in the microbial community in people, animals, or the surroundings. Such selection forces, which are subject to the 'medicine and error' of anxiousness, promote anti-acquisition techniques such as point mutation or horizontal gene transfer. Antibiotic resistance to one or more encryption methods²⁵. Environmental factors and behaviours can allow or prevent the spread of these newly

developed antimicrobial organisms (AROs) in humans, animals and habitats. The burden of ARO-related diseases will therefore be determined by the quantity and character of infections, as well as the availability, efficacy, and safety of alternate therapies. COVID-19 has the ability to alter all three of these components directly or indirectly as a result of pandemic response. The actions implemented by the government to the general population was advised to combat COVID19 have differed, but also include a combination of internal and external immigration restrictions; school, workplace, and non-essential service cancellations, the use of telehealth increased, physical distance measures; and mask wearing Although comfortable for both patients and physicians, one of the biggest reasons of antibiotic overuse is a lack of physical examination by healthcare providers and the lack of laboratory markers for identifying the seriousness of the ailment²⁶.

ANTIMICROBIAL RESISTANCE: A THREAT HIDING IN THE SHADOWS OF THE COVID-19 PANDEMIC:

Increased antimicrobial usage is strongly linked to the establishment of worldwide AMR²⁷. In many nations, certain patients demand for antimicrobials, while others seem to be more motivated to abuse and misusing antibiotics than to keep them safe and take them only when prescribed. Patients in impoverished nations frequently self-medicate because they believe public hospitals are overloaded or would impose larger financial obligations, or they merely give antimicrobials to test whether they improve. As a result, the high prevalence of patients seeking antimicrobials directly can have a significant influence on antimicrobial consumption and resistance development²⁸. Furthermore, medicine quality suffers as a result of improper storage facilities, age, and the presence of none, too few, or too much active component. This might be deliberate or the result of bad production procedures. Inactive "excipients" in pharmaceuticals that have been adulterated can also be toxic to persons who ingest them. China and India have the unpleasant distinction of being the primary suppliers of counterfeit pharmaceuticals²⁹. Drugs in higher-level health centres in India are less prone to be outdated, but many rural clinics give out-of-date medications, particularly in areas plagued by violence³⁰. Patients' conditions worsens because the treatments contain a sub-therapeutic amount of the active component, which is inefficient in treating infections yet, the sub-optimal level leads to the establishment of tolerance³¹.

THE IMPACT OF DRUG RESISTANCE IN COVID-19:

Individualized management and global infection control programmes both prioritise observing and researching the prevalence, causes, and transmission of AMR³². In relation to COVID-19 patients, 72 percent of patients reported hospitalisation for antimicrobial therapy³³. In contrast, every time an antibiotic is taken, whether logically or irrationally, the genetic selection exerted leads to the selection of resistance microorganisms. Resistant microbes can survive in a host for minimum a year after choosing³⁴. COVID-19 poses significant problems for many areas of health care, including infection, bacterial diagnosis and management, as well as drug administration. Recognizing that patients with viral pneumonia, such as COVID-19, are at higher risk of developing serious secondary bacterial infections, there is no denying that antibiotics are empirically recommended for many COVID-19 patients, especially in frequently agitated environments. In reality, according to a

latest report, antiviral drugs and antibiotics were utilized by 93 percent and 100 percent of ICU and non-ICU patients, respectively³⁵. Aside from the increased demand on healthcare situations and hospitals, knowledge on the usage of drugs to treat COVID-19 infection is freely open to the public on the internet. The lack of adequate knowledge of the significance of antibiotics, combined with the fear of COVID-19 infection, has had a direct effect on increasing exposure to over-the-counter antibiotics, especially in low- and middle countries with impoverished antibiotic control strategies and limited access to health settings³⁶.

IN THE PRESENT COVID-19 PANDEMIC, THERE ARE SIGNS OF ANTIMICROBIAL ABUSE AND MISUSE:

Antimicrobial agent therapy is used to treat bacterial or fungal respiratory infections that are indicated or proven. This might be empirical or specific in individuals diagnosed to hospitals, or it could be used to control nosocomial infections encountered while hospitalisation, such as hospital acquired pneumonia or ventilator-associated pneumonia³⁷. Only minority patients may require antibiotic therapy, as a result, medications, particularly antibiotic, may be misused. Various investigations demonstrated that antimicrobial treatments were widely used as part of medical care plan for hospitalised COVID19 patients in some countries^{37, 38}. Whereas it implies that antibiotics or antivirals should be used in hospitalised COVID-19 patient populations with secondary bacterial infection, the chance of antibiotic prescribing in a large number of patients without confirmed secondary infection is significantly increased, potentially leading to an increase in AMR through driving selection of multidrug resistant (MDR) organisms³⁹. A retrospective study of 191 people from two local hospitals in Wuhan, China, found that 181 (95%) had received antibiotics and 41 (21%) had received antivirals³⁷. One cohort study of 37 hospitalized patients in three medical centres in Wuhan reveals that 36 (97.3%) got antiviral therapy, 29 (78.5%) got probiotics, 28 (75.7%) got managed with traditional Systems of medicine, and 22 (59.5%) got antibiotics⁴⁰. Throughout this aspect, it was revealed 68.9 percent of COVID-19 patients previously taken antibiotics (mainly azithromycin and ceftriaxone) prior to hospital admission, with a self-medication rate of 33.0 percent⁴¹. In a Wuhan investigation, 11 percent of 99 patients died, and 71 percent were given antibiotics; nevertheless, only 1 percent had co-infection with bacteria, and 4 percent had fungal co-infections⁴². Furthermore, in a single-centre research in Wuhan including 36 non-survivors with COVID-19, antimicrobial treatment was delivered to every patients, and 61.1 percent of these people got combination antibiotic medication, while 38.9 percent underwent single antibiotic³⁴. Chen et al. found similar results, showing that 71 percent of patients got preventative therapy, with 45 percent of them receiving combination medication as a preventive measure for common illnesses. Microorganisms grow in just 1% of patients, including *Klebsiella pneumoniae*, *Acinetobacter baumannii*, and *Aspergillus flavus* isolated in one case, the investigation also revealed that this *A. baumannii* had a strong resistance to antimicrobials⁴³. Antibiotics inefficacy may potentially be associated to bacterial resistance, while it is possible also that beginning and length of antibiotic therapy in each patient, as well as associated co-morbidities, had a substantial effect on the outcomes. Similarly, Clancy et al. discovered in a meta-analysis showed microbial lung superinfections were the cause of mortality in 32% of COVID-19-positive individuals worldwide. The investigators also demonstrated that 79 percent of patients were managed with antibacterial drugs, with *A. baumannii*, *P. aeruginosa*, *K.*

pneumoniae, *E. coli*, and *S. aureus* being the most common causative agents⁴⁴. Additionally, a retrospective research of 918 COVID-19 patients in Wuhan, China found that 7.1 percent of inpatients had a pathogens co-infection, including pneumonia (32.3 percent), bacteremia (24.6 percent), and urinary tract infections (21.5 percent) being the most common⁴⁵. One assessment of scientific literature on hospitalised COVID-19 patients report that while 72 percent (1450/2010) got antibiotics, only 8 percent (62/806) had recurrent microbial or fungal co-infections⁴⁶. Many of the pathogens often detected in secondary bacterial infections, such as *Staphylococcus aureus*, *Streptococcus pneumoniae*, *Neisseria meningitidis*, *Haemophilus influenzae*, *Klebsiella pneumoniae*, and members of the genera *Proteus*, *Enterobacter*, and *Citrobacter* spp., are found in hospitals⁴⁷. Previous research also reveals that drug-resistant microorganisms, such as multidrug-resistant *Escherichia coli*, *Enterococcus*, *Chlamydia pneumoniae*, *Klebsiella pneumoniae*, *Pseudomonas aeruginosa*, *Mycoplasma pneumoniae*, and extended-spectrum beta-lactamase, promote some illnesses in COVID-19 patients^{45,48,49}. Azithromycin, a broad-spectrum macrolide antibiotic used mostly to treat respiratory, gastrointestinal, and sexually contagious infectious diseases caused by a wide variety of extra- and intracellular bacteria⁵⁰, will be mass-administered for control of COVID-19 excessive usage of azithromycin, a first-line antibiotic in the treatment of pneumonia, in COVID-19 sick people will almost certainly result in prolonged selection of resistant bacteria, as well as greater resistance not only to the above antimicrobials but also to the macrolide family of drugs more broadly, and, through co selection due to the fact that resistance genes are frequently undertaken together on multidrug resistance plasmids, to other antibiotic families within the patient population⁵¹.

ANTIBIOTIC RESISTANT DEVELOPMENT IN HOSPITALS:

Species of Candida:

Species of Candida Several fungal infection were studied during the COVID-19 pandemic, however the advent of a novel multi-drug resistant Candida species has been linked to COVID-19 related candidiasis. The development of a novel Candida species is quite worrying, particularly given the prevalence of antibiotic resistance⁵². It is not understood how Candida produces co-infections among COVID-19 patients. Moreover, both Candida and SARS-CoV-2 have been identified on same hospital surfaces, and COVID-19 diagnosed ICU patients have same health risks, medicines, and associated co-morbidities of *C. Auris*. The prevalence of multi-drug resistant *C. Auris* is already higher in developing world. Chowdhary A. et al. hypothesise that this is related to a higher global impact of *C. Auris* in underdeveloped nations because to limited diagnostic techniques and insufficient funding for infection prevention and control. Overloaded medical institutions in underdeveloped countries are unable to diagnose other infections, such as *C. Auris*, because they are focusing all of their limited diagnostic skills on SARS-CoV-2. As a result, the issue of multi-drug resistant *C. Auris* and possible epidemics is far more serious in impoverished nations than in industrialised countries⁵³.

New Delhi Metallo-Beta-Lactamase-Producing Carbapenem-Resistant Enterobacterales (NDM-CRE):

New Delhi Metallo-Beta-Lactamase-Producing Carbapenem-Resistant Enterobacterales is one of the already identified possible etiological agent of healthcare-acquired infections in many countries, including developed nations such as Italy⁵⁴. The majority of first-line treatment medicines seem to be

ineffective against Enterobacterales species⁵⁵. COVID-19 individuals who had already been colonised with NDM-CRE or who developed it during their hospitalisation had an extended hospitalisation due to related comorbidities and antibiotic resistance⁵⁴.

Carbapenemase-Producing Enterobacterales (CPE):

Carbapenemase-producing enterobacterales (CPE) have resurfaced in the aftermath of the COVID-19 outbreak, owing to careless antibiotic prescribing and a paucity of drug safety. The re-emergence of CPE and superinfection in COVID-19 patients is caused by a number of reasons. The most prevalent element in the therapy of COVID-19 patients is the use of immunomodulators such as corticosteroids. Severe lung damage and subsequent mechanical ventilation are also concerns⁵⁶.

Carbapenem-Resistant Acinetobacter Baumannii (CRAB):

Hospital acquired *Acinetobacter Baumannii* strains have been associated with elevated fatality in COVID-19 patients⁵⁷. These pathogens were shown to be resistant to ciprofloxacin and gentamicin, resulting in higher morbidity and death in COVID-19 patients⁵⁶.

ANTIBIOTIC USAGE:

Almost all of the substantial antibiotic usage in COVID-19 patients is inevitable. More studies are needed to better understand the health risks and rate of co-infection in COVID-19 patients in order to promote a decrease in any unnecessary antimicrobial prescribing. More studies are needed to better understand the health risks and rate of co-infection in COVID-19 patients in order to promote a decrease in any unnecessary antimicrobial prescribing. Azithromycin is an existing medicine that has been utilized for the present pandemic while new Coronavirus therapies are discovered. Several studies, warned against taking azithromycin except in the treatment of community acquired pneumonia (CAP), stating that there is no clear evidence that azithromycin may have therapeutic benefits in COVID-19 other than antibacterial properties in bacterial superinfection⁵⁸. It is also suggested that some antibacterial, such as teicoplanin (a glycopeptide antibiotic), might be effective against COVID-19 after previously demonstrating antiviral efficacy towards coronaviruses⁵⁹. According to a group of European doctors, it can be hard to identify COVID-19 from bacterial pneumonia, which implies that antibiotics are being given to people who do not have bacterial infections³⁹. COVID-19 is becoming more common in Low Middle Income Countries (LMICs), presenting severe concerns given the poor healthcare systems, limited knowledge, and insufficient readiness for outbreaks in most of these countries. A big COVID-19 pandemic might amplify the rise in AMR in LMICs⁶⁰. Because it is hard to distinguish among bacterial and viral pneumonia, the Infectious Disease Society of America (IDSA) guideline on community acquired pneumonia (CAP) determined that complex phase cannot be used to decide whether to begin or pause medications in CAP patients⁶¹. The IDSA guideline performed wide literature reviews to evaluate the additional benefit of utilising procalcitonin, CRP, or the Updated Clinical Pulmonary Infection Index with clinical criteria for the evaluation of Hospital Acquired Pneumonia (HAP) or Ventilator Associated Pneumonia (VAP)³³.

COVID-19 ASSOCIATED WITH CO-INFECTIONS:

Patients having lung viral infections are more likely to develop subsequent bacterial infections, which lead to higher disease severity and death⁶². COVID-19 has a wide range of

illness severity, including symptomless and non-specific flu-like illness to pneumonia and life-threatening consequences such as acute respiratory distress syndrome (ARDS) and multiple organ failure⁶³. Furthermore, recurrent bacterial infections have been shown to be a substantial risk factor for poor COVID-19 results. Despite their demonstrated role in the complexity of respiratory disease symptoms, coinfections go unexplored during big pandemics of respiratory illnesses⁶². A meta-analysis and systemic review found 24 papers concentrating on microbial coinfections in hospitalized COVID-19 patients, revealing that 3.5 percent of patients suffered coinfections and 14.3 percent of COVID-19 patients had secondary infection, with total bacterial infection 6.9 percent. Another meta-analysis and systemic review that also included 30 papers assessing coinfections with COVID-19 patients found similar results, reporting that 7% of the patients had bacterial coinfections⁶⁴. COVID-19 patients had a wide range of co-infections, range from no co-infections to 100 percent coinfections in patients who died, as well as a wide range of antibiotic usage by chronic infection, ranging from 20 percent to 100 percent for antibiotics⁶⁵. The results of a retrospective mono centre investigation done in France indicated an incidence of 28 percent bacterial coinfections in individuals with severe SARS-CoV-2, and the investigators strongly recommended systemic empirical antimicrobial treatment with third generation cephalosporins. The majority of the bacterial coinfections found in their investigation were caused by methicillin-sensitive *Staphylococcus aureus*, *Haemophilus influenzae*, *Streptococcus pneumoniae*, and *Enterobacteriaceae*⁶¹.

ANTIBIOTIC RESISTANCE RESEARCH AND HEALTH-CARE SYSTEM PRIORITY:

Diagnostics:

COVID-19 has emphasised both the capabilities and confines of diagnostic technology; whereas analytical logistics with built-in redundancy have been successfully scaled up in many situations, we are still battling to identify SARS-CoV-2 infection accurately and quickly internationally. Moving forward, the AMR community should leverage on the investment in diagnostic logistics and adapt their usage for quick detection of both the causal agent of illness and any associated medication resistance, particularly in low- and middle-income countries (LMICs). In Ghana, for example, capacity building (both laboratory and staff) is required to address COVID-19 surveillance as well as guiding antibiotic dosing⁶⁶. The AMR sector could encourage the present impetus to build and make widely available a low-cost, accurate, and fast-acting diagnostic for viral infections, such as the ACT Accelerator Diagnostics Partnership. Differentiating between bacterial and viral infections and swiftly identifying resistance carriers will not only minimize excessive antibiotic treatment and, as a result, the formation of AMR⁶⁷.

Monitoring:

COVID-19 has yet again underscored the need of strengthening, open access, and continuous data gathering for infectious illnesses. This is especially crucial in the case of AMR, which frequently depends on passive surveillance: The WHO's GLASS programme is transitioning to active monitoring, although for the time being, convenience samples of isolates sent for phenotypic resistance testing are being collected. This means that they are more likely to sample people who have complicated, long-lasting diseases caused by resistant microorganisms. Global AMR observation is further complicated by significant breaks, particularly in

developing nations⁶⁸. New measures developed and deployed for COVID-19 might possibly be used to strengthen AMR environmental monitoring. For example, waste water surveillance might be utilized to identify alterations in COVID-19 and ARO occurrence as well as antibiotic concentrations early on⁶⁹.

HEALTH SYSTEM STRENGTHENING:

Antibiotic Management and Administration in the face of the CoVid-19 Pandemic:

COVID-19 has exposed several flaws in our health-care systems. To effectively address AMR, it is critical to promote appropriate medication adherence as part of antimicrobial stewardship³⁸. The World Health Organization has advocated for a larger pandemic response that ensures the continuity of vital health care services and constant supply of safety guaranteed and inexpensive antimicrobials, especially antiretroviral and tuberculosis medications, and vaccinations⁶⁷. Antimicrobial management refers to joint, synchronized programmes and interventions that promote the selection of the appropriate antimicrobial treatment regimen, comprising dose, length of treatment, and mode of administration, while minimizing undesired effects⁷⁰. The fact that the primary goal of antimicrobial stewardship is to reduce the occurrence of AMR and drug-resistant illnesses in most cases, antimicrobial stewardship is primarily a behaviour change problem for healthcare personnel and healthcare organisations in advanced healthcare facilities. Antimicrobial stewardship in LMIC healthcare institutions is exacerbated by restricted access to drugs, worries about counterfeit medicines, and insufficient hospital infection prevention programme infrastructure⁷¹. In order to promote antimicrobial management, clinicians such as infectious disease doctors, nurses, clinical pharmacists, and other care providers should be committed. Hospitals with functioning infection prevention programs would try to engage antimicrobial prescription issues even when they are not clearly observable in their absence. They must also be involved in the formation of facility guidelines^{72,73}. Before using antibiotics, a microbiological test should be done, and if the likelihood of bacterial superinfection is minimal, antibiotic treatment should be assessed and discontinued. Furthermore, as soon as it is comfortable, a transfer from intravenous to oral antibiotics should be investigated. Because of anecdotal evidence, azithromycin is being used on a regular basis antibiotics should be administered for the shortest amount of time feasible, and prophylactic antibiotics should be avoided. Despite the difficulty in distinguishing COVID-19 from bacterial pneumonia, the uncertainty about bacterial superinfections, the lack of particular antiviral therapy for the recent pandemic, and the increased rates of mortality, antibiotics as portion of the empiric treatment for the most intense suspected or confirmed COVID-19 cases (e.g., patients requiring mechanical ventilation) should not be overlooked. However, even during a pandemic, antibiotics should be taken with caution³⁹.

CONCLUSION:

The widespread utilisation of antimicrobial drugs during the COVID-19 outbreak has raised the spectre of antimicrobial resistance. Even before the outbreak, developing nations were reporting significant levels of resistance. Several variables that are particular to developing world have acquired prominence during the COVID-19 pandemic. This teaches us a lot of things, one of which is that, medical improvements, humans are still extremely vulnerable to disease with limited or no traditional therapy. Immediately crucial components of any AMR mitigation approach are

increased spending in education and increased public knowledge of AMR. Numerous studies into managing pathogens, as well as the creation of more functional antimicrobials, is required. Reserving antibiotic therapy for patients with the most severe COVID-19 illness and restricting antibiotic treatment to five days, as most recommendations, detailed observation investigation is now necessary to enhance COVID-19 patient care.

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