History and Spread of Viruses (COVID-19) and Associated Demographic and Clinical Parameters – A systematic review and meta-analysis
Swambhavi Awasthi¹, Sunil Sharma², Saurav Attri³, Sakshi Malik Attri⁴, Rajesh Sharawat⁵, Gayatri Vishwakarma⁶

¹BSc in Molecular Biology and Biotechnology, Independent Research Scholar, Sector-16, Indira Nagar, Lucknow, 226016; ²Dean, Indian Spinal Injuries Centre, New Delhi; ³MS, Sr. Consultant, Civil Hospital, Gurgaon, India; ⁴Consultant, Civil Hospital, Gurgaon, India; ⁵Indian Spinal Injuries Centre, New Delhi; ⁶Indian Spinal Injuries Centre, New Delhi

Abstract
COVID-19 made a huge impact on the world due to its rapid transmission and no treatments being available for it. The virus affected more people and spread to various countries than what was predicted when COVID-19 initially began spreading. There have been numerous pandemics and epidemics in the 21st century yet COVID-19 has affected more people and spread widely. The primary objective of the study was to explore history, spread and associated parameters of existing viruses especially COVID-19. Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) guideline was followed for a systematic search to identify eligible published articles. Clinical data, regarding COVID-19 patients, was obtained from previously published articles. The main cause of COVID-19 spreading rapidly was noted to be due to a high percentage of asymptomatic patients, transmission being air-borne, and the lack of knowledge and preventative measures being implemented when the virus began spreading. The common co-morbidity that found in patients was Diabetes Mellitus, Hypertension, and Coronary Heart Disease. The common symptoms, found through the Meta-analysis, that the patients faced included cough (55.4%), fever (68.4%), fatigue (20.3%), and shortness of breath (18.1%). The proportion of asymptotic positive cases was measured 58.3% (95%CI: 24.7% – 87.9%) while mortality proportion was found to be 6.7% (fixed-effect model) and 13.4% (random-effect model). The Meta-analysis indicated that a higher percentage of males were affected by COVID-19 than females and more patients are found to be asymptomatic. Moreover, the mortality rate of patients that have had COVID-19 was found to be low.

Keywords
Covid-19; Spread; Transmission; Asymtomatic; Co-Morbidity; Symptoms; Mortality

Introduction
The novel coronavirus pandemic, COVID-19, has become a concerning health emergency for the world due to its rapid and prevalent spread. As of 19 July 2020, there have been over 14 million confirmed cases reported accompanied by 597,583 deaths from over 200 countries. The virus was first identified as numerous patients from Wuhan, China were...
diagnosed with pneumonia in December 2019, and then in March, the World Health Organization (WHO) declared COVID-19 a pandemic. (2) The virus primarily spreads from human to human transmission and numerous studies have proposed that the virus has the capability of spreading during the incubation period of 14 days. (3) The transmission of the virus takes place through respiratory droplets that are released when coughing and sneezing. The virus enters the body through either the mouth, eyes, or nose and has caused severe pneumonia and led to death in several patients. (4) The COVID-19 pandemic has been hard to contain due to several factors like the presence of asymptomatic patients, chances of transmission during the incubation period, a relatively high reproduction number (R0), and the lack of education regarding social distancing norms. The most effective method of controlling the outbreak is preventing human to human transmission which can be achieved via implementing social distancing and hygiene norms. (5) This has led to multiple countries and cities following lockdown or curfew conditions to flatten the curve formed by recording the spread of the virus.

**Aims & Objectives**

Multiple viruses have caused either a pandemic, epidemic, or an outbreak in the 21st century, yet, COVID-19 remains to be one of the most contagious and widespread pandemics that the world would have witnessed. COVID-19 may not be as fatal as MERS or SARS, but it has impacted more areas and a higher population across the globe. A question that arises from COVID-19’s vast and speedy spread is the reason that the virus has had a greater impact on a larger proportion of the population in comparison to the outbreaks that came before COVID-19. This review aims at summarizing different factors of COVID-19 that have allowed to become a widespread pandemic in comparison to previous outbreaks like SARS, MERS, Zika, H1N1 (Swine flu), H1N1 (Spanish flu), and Ebola. Furthermore, the review aims at analyzing the clinical characteristics of COVID-19 through a systematic review and meta-analysis.

**Material & Methods**

A rapid review for a broad range of COVID-19 following a standardized guideline i.e. Preferred Reporting Items for Systematic Reviews and Meta-Analyses (PRISMA) was conducted. (6) A systematic literature search of electronic databases (PubMed, Scopus, Cochrane, and Google scholar) was done using keywords COVID-19 (including all mesh terms), symptoms, co-morbidity, and death (mortality). A Systematic literature search in electronic databases was done for Dec 2019 to July 2020 using Boolean operation with combinations of the search terms: “COVID-19 OR COVID19 OR COVID-19 pandemic OR SARS-CoV-2 infection OR COVID-19 virus disease OR 2019 novel coronavirus infection OR 2019-nCoV infection OR coronavirus disease 2019 OR coronavirus disease-19 OR 2019-nCoV disease OR COVID-19 virus infection AND Symptoms AND Comorbidity OR Comorbidity AND Death OR Mortality AND Asymptomatic”.

In addition, the reference lists of each eligible publication were carefully checked in order to identify additional records. Two-three authors independently read the full manuscript and judged on the basis of inclusion and exclusion criteria to include it in the present systematic review. In case of any disagreement, Delphi Method was used to resolve the issue. Quantitative data synthesis was used for included studies that are sufficiently homogenous in nature. The aim of the initial broad search was to identify all COVID-19 scientific and medical literature to answer a range of research questions related to demographic and clinical outcomes. We followed a strict criteria (Dec 2019 onwards) to select references which contain demographic and clinical outcomes related to COVID-19 and removed various articles from the search due to the proliferation of COVID-19 literature.

MedCalc Statistical Software version 19.2.6 (MedCalc Software bv, Ostend, Belgium; https://www.medcalc.org; 2020) was used for meta-analysis. Random effects model was considered more appropriate, in which both the random variation within the studies and the variation between the different studies are incorporated. This assumption was tested by the "Heterogeneity test". If this test yields a low P-value (P<0.05), then the fixed effects model may be considered invalid. I2 statistics with 95% Confidence Interval (CI) was used to show the percentage of inconsistency.

**Results**

A total of 25 studied were included and a selection of studies was presented in [Figure 1](#). The total sample size was observed 55420 subjects with a
pooled average age of 52.7±3.1 years ranging from 46.7 – 58.8 years (95%CI) (Figure 2). (Figure 3) shows the forest plots for proportions of gender. It is observed that 52.6% male and 47.4% female were diagnosed with COVID-19 and the difference in proportion was found statistically significant (P<0.001) as the sample size was large.

17 studies out of 25 discussed about co-morbidities (Figure 4): 53.8% (95%CI: 41.1% – 66.3%) subjects in these studies were having co-morbidities. 11 studies showed number of events about some common co-morbidities such as Hypertension (HT), Diabetes Mellitus (DM), and Coronary Heart Disease (CAD). (Figure 5) reveals pooled proportion of HT (p=29.2%; 95%CI: 10.8 – 52.1), DM (p=15.3%; 95%CI: 10.3 – 21.0), and CAD (p=11.0%; 95%CI: 6.8 – 16.2). (Figure 6) depicts forest plots of the proportion of asymptotic subjects in eight studies. 58.3% (95%CI: 24.7% – 87.9%) of the study subjects were asymptotic or not having any symptoms at the time of diagnosis of COVID-19. Furthermore, 17 studies mentioned mortality due to COVID-19. (Figure 7) reveals that 6.7% (fixed-effect model) and 13.4% (random-effect model) was the pooled mortality percentage. However, none of the studies were found to be discussing the survival rate as almost most of the studies were cross-sectional studies.

Funnel plot for assessing publication bias based on study proportions is shown in (Figure 8). Circles indicate study point estimates. There was no clear asymmetry found and therefore, funnel plot demonstrates no significant publication bias.

**Discussion**

(Table 1) depicts the comparison of origin, spread, transmission, and WHO recommendation for pandemic viruses such as H1N1 (Spanish Flu), Zika, Ebola, SARS, H1N1 (Swine Flu), and MERS with the COVID-19 virus. No treatment or vaccine is still available till date to fight against these viruses.

Multiple viruses have caused either a pandemic, epidemic, or an outbreak in the 21st century, yet, COVID-19 remains to be one of the most contagious and widespread pandemics that the world would have witnessed. COVID-19 may not be as fatal as MERS or SARS, but it has impacted more areas and a higher population across the globe. A question that arises from COVID-19’s vast and speedy spread is the reason that the virus has had a greater impact on a larger proportion of the population in comparison to the outbreaks that came before COVID-19. This review aims at summarizing different factors of COVID-19 that have allowed to become a widespread pandemic in comparison to previous outbreaks like SARS, MERS, Zika, H1N1 (Swine flu), H1N1 (Spanish flu), and Ebola.

**Spread and Transmission:** A systematic review adequately explained that Corona-viruses are large, lipid-enveloped, single-stranded RNA viruses found in avian and mammalian species. The review made the authors question why COVID-19 transmitted globally, when compared to other viruses. It is the fact that COVID-19 spread around the world far quicker than originally predicted. The article published in Lancet medical journal confirmed that Wuhan’s first known COVID-19 patient began developing symptoms in December 2019. The World Health Organization (WHO) declared the COVID-19 epidemic a global health emergency in January. In Japan, more than 3,700 passengers form 48 countries were quarantined on a cruise, called Diamond Princess after a passenger who departed in Hong Kong was found to be a confirmed and asymptotic patient of COVID-19(7). COVID-19 appears to have a fatality rate of 4.4%, which is much lower than the 10% fatality rate for SARS and approximately 30% for MERS-CoV. Nevertheless, COVID-19 is found to be more contagious than SARS or MERS-CoV, and critically it spreads undetected.(8)

At the beginning of 2020, it was unclear whether or not asymptomatic COVID-19 patients had the ability to transmit the virus. Numerous studies indicate that the virus particles are present in the saliva of COVID-19 patients and thus asymptomatic patients have the capability of transmitting the virus. Though the possibility of transmission before symptoms develop seems to be infrequent, there are suggestions that asymptomatic individuals could transmit the virus.(9) Limited studies and data are available for asymptomatic infections.(10)

In a recent study,(11) an immense number of total infections were estimated out of which undocumented (80% to 90%) cases and undetected infections were the sources for 79% of documented cases in China.

However, a study reveals that asymptomatic cases are assumed half as infectious as symptomatic ones (relative infectivity fa=0.5).(12) COVID-19 infections are often not accompanied by symptoms. The most reported symptoms include mild flu, cough, fatigue, and headaches.(13) A review study shows that there...
is some evidence that patients may not be infectious for the whole period that they are SARS-CoV-2 (COVID-19) positive and that infectivity may be related to the viral load and time since symptom onset.(14)

Prevention and treatment: To date, no drugs are currently approved for COVID-19 although some have been tried such as Chloroquine and hydroxychloroquine. Two small human studies have been conducted with both these drugs in COVID-19, and have shown significant improvement in some parameters in patients with COVID-19.(15,16,17,18,19)

The term ‘household isolation’ refers to households going into isolation for a fixed duration of time. This approach reduces the risk that other household members would transmit in the community when patients are pre-symptomatic or if they are asymptomatic but still infectious.

Social distancing is the policy widely adopted in many countries to limit transmission of SARS-CoV-2 (COVID-19), at a great economic and social cost. Social-distancing policies may have an important role in mitigating the spread of pandemics; for many infectious diseases, such as influenza.(20) Multi-site screening including nasal and pharyngeal swabs, stool, and sputum specimens could be considered to improve the diagnosis, treatment, and infection control in patients with COVID-19.(21)

The purpose of the study has been to understand why COVID-19 has proven to be a widely spread virus across the globe. COVID-19 has been established to be less deadly than the viruses that have caused other pandemics, yet it is more contagious.(3) The COVID-19 virus can easily transmit from one person to another through the air, respiratory droplets, or contaminated surfaces, and the virus can go unnoticed in asymptomatic patients making detection very difficult.(4,13,14)

Another important aspect that leads to the virus spreading so rapidly was the lack of knowledge that was present when the virus first crossed over from animals to humans.(8,22) This meant that at the early stages there was not adequate medical protection, social distancing measures, protective measure, and awareness in the general public which permitted the virus to infect numerous people.(23)

The Meta-analysis permits for the evaluation of clinical characteristics of various COVID-19 patients. Through the results it can be derived that 58.3% of COVID-19 patients were asymptomatic which was found using eight studies that had clinical data regarding asymptomatic patients and confirmed by the published studies.(10) A high percentage of asymptomatic patients along with risks of transmission allows for the virus to spread in a larger percentage of the population, going unnoticed.(24,25,26,27,28,29,30,31,32)

There was a slight difference between the proportion of male and female patient’s data, indicating that there may be more males that are affected by the virus than females. Certain studies suggest that women are less susceptible to the virus due to the presence of the X chromosome and certain sex hormones which allow for the development and strengthening of adaptive immunity. (32) Further exploration of immunity between males and females who get impacted by COVID-19 needs to be carried out to establish the difference. (7,10,20,22-31,33-45) The common symptoms, found through the Meta-analysis, that the patients faced included cough (55.4%), fever (68.4%), fatigue (20.3%), and shortness of breath (18.1%). (7,10,20,22-28,30,31,33-45) Co-morbidities weaken the immune system of a person, which then allows the COVID-19 virus to have a severe impact and cause further complications to the patient. (22,25,26,28,29,30,40,44,45) The common co-morbidity that was found in patients was Diabetes Mellitus, Hypertension, and Coronary Heart Disease (7, 10,20,22–29,30,31,33–45). The death rate for a patient that has co-morbidities and gets infected by COVID-19 are higher than a healthy patient. (7,10,22,24,26,28,29,31,36,37,38,40,42,44,45)

**Conclusion**

This systematic review and meta-analysis found that the evidence suggests that the viral load of COVID-19 is more on males as compared to females. However, the clinical significance of this finding is indecisive. There is some evidence that more patients are found asymptomatic in nature. The study revealed a low mortality percentage due to COVID-19. However further studies are required to measure pure percentage of death with or without co-morbidity to understand the seriousness of the disease. It is noted that due to the highly fluid, dynamic, asymptomatic patients, the evolving nature of the pandemic, and the describing natural history of the disease process it spreads globally.
Limitation of the study

The current study has a few limitations. In our study, a high degree of heterogeneity was observed; however, we did not find the sources of heterogeneity in our analysis. The observed heterogeneity might be due to the population characteristics, most of the studies were published on the Chinese population. However, heterogeneity was minimized by using a random effect model of analysis. Further, the quantity of the number of studies available on COVID-19 was large enough to assess the quality of study hence we focused only on studies discussing demographic and clinical parameters. In addition, we tried to reduce selection bias by using stringent searching strategy.

Relevance of the study

This review allows for a better understanding of the spread of COVID-19 when compared to other pandemics and epidemics that have occurred. It lays out a comparison and analyzes clinical characteristics of COVID-19. Understanding the clinical characteristics can permit for a better understanding of the pandemic and its reaction on different patients.

Authors Contribution

SA: Literature search, data extraction, manuscript writing, SS: Critical review of included full articles and manuscript writing and reviewing, SA: Critical review of included full articles and manuscript writing, SMA: Critical review of included full articles and manuscript writing, RS: Data extraction, submission and coordination with authors, GV: Conceptualization, Literature search, data extraction, Methodology, manuscript writing

Acknowledgement

We would like to acknowledge Dr. H.S. Chhabra, Chief of Spine Services and Medical Director, Indian Spinal Injuries Centre, New Delhi for allowing us to conduct this study and for providing all his support during the study.

References

33. Antibody tests for identification of current and past infection with SARS-CoV-2 - Deeks, JJ - 2020 | Cochrane Library. Available from:


## Tables

### TABLE 1 HISTORY, SPREAD AND TRANSMISSION OF THE VIRUSES

<table>
<thead>
<tr>
<th>Virus</th>
<th>Place of Origin</th>
<th>Year</th>
<th>Spread</th>
<th>Transmission</th>
<th>Reasons that virus did not spread as much as COVID-19</th>
<th>WHO initiative/ comment for controlling the spread</th>
<th>Reference #</th>
</tr>
</thead>
</table>
| H1N1, Pandemic, Spanish Flu | Not Known believed to be from China, France, Britain, or USA | 1918 | • After world-war-1, the H1N1 Spanish Flu was the deadliest pandemic that the world faced.  
• Estimates suggest that the pandemic resulted in the death of almost one-third of the world’s population with approximately 40 million deaths(46).  
• The pandemic affected US, Europe, and parts of Asia. Studies also indicate that the pandemic resulted in deaths amongst young adults, which is unusual as most strains of influenza do not have a severe impact on their age group (46).  
• In US the pandemic came to an end by the summer, as the weather did not favor the virus, and by this time people either developed immunity or had died(47).  | • In 1918, due to the lack of scientific and medical advances, very little was known about the H1N1 Spanish Flu virus. The genome of the virus has been sequenced in modern-day, using the fixed and frozen lung tissue from a 1918 influenza victims(48). The host of the H1N1 1918 influenza has not been identified(49).  
• The transmission of the virus has not completely been understood and a few research studies suggest that the influenza virus transmits in a similar way to the common cold virus.  
• The human-to-human transmission is believed to occur via the following:  
  o Respiratory droplets from coughing and sneezing  
  o A patient contaminating inanimate object and surfaces  
  o Inadequate protection at the healthcare setting | Since very little is known about the H1N1 virus that caused a pandemic in 1918, comparing it to the current COVID-19 virus is difficult however with suggestions found from research the following can be suggested:  
1. People began to develop immunity to the H1N1 virus  
   o Studies suggest that the reason that the 1918 virus had very little impact on the elderly population was that they had some immunity to the virus from the 1847 pandemic(49).  
   o As more people got infected with the virus and recovered, they had immunity from the virus(49).  
   o With COVID-19, WHO states that there are currently no studies that verify that patients who have recovered from COVID-19 have immunity from a second infection(50).  
2. The H1N1 virus was very lethal and resulted in numerous deaths.  
   o The death of a patient meant a decrease in the population and decreased the chance of the virus spreading further  
   o It is suggested that one third of the population died from the H1N1 pandemic in 1918(51).  
3. International Travel was not as advanced in 1918.  
   o Globalization has connected the world and advances in aviation have interconnected the world. The movement of people from one country to another allows for imported cases.  
   o In 1918, travel was limited and that prevented the H1N1 virus | N/A | [46], [47], [48], [49], [50], [51] |
<table>
<thead>
<tr>
<th>Virus</th>
<th>Countries</th>
<th>Year</th>
<th>Notes</th>
</tr>
</thead>
<tbody>
<tr>
<td>ZIKA</td>
<td>Uganda and The Republic of Tanzania</td>
<td>1952</td>
<td>There have been numerous spreads and outbreaks of the ZIKA virus. The largest outbreak occurred in 2007 on the island of Yap. During which approximately 5000 people were infected and it lasted for three months. In 2013, there was a big outbreak that occurred in the French Polynesia. The virus spread eastwards toward Oceania and then to South America. There were approximately 20,000 people infected. In 2015, there was an outbreak that began in Brazil and it spread to nearly 50 countries. This outbreak was declared an epidemic. The ZIKA virus is transmitted when the Aedes genus mosquito infected with the ZIKA virus bites a human. Human-to-human transmission can occur via the following: Pregnant women infected with the ZIKA virus can pass the virus to the fetus and it has been noticed that the babies that get infected during pregnancy can have birth defects like microcephaly. Transmitted through sexual contact with an infected person. Blood transfusion of an infected patient. The blood of an infected person comes in contact with another person. The incubation period of the virus is 3-14 days. Most people who get infected with the ZIKA virus generally do not develop major symptoms and the symptoms are generally mild. These symptoms last for 2-7 days.</td>
</tr>
<tr>
<td>Ebola</td>
<td>Near the Ebola River in Democratic Republic of Congo (formerly known as Zaire)</td>
<td>1976</td>
<td>There have been 26 outbreaks of the Ebola Virus Disease to date. The first two were consecutive outbreaks that occurred in Central Africa. This was the deadliest outbreak with 218 deaths out of the 318 patients that were infected. The virus is transmitted from human to human through blood, bodily fluids, and being in contact with people who are infected or have died from the Ebola Virus Disease. The Ebola Virus Disease made the spillover to humans through a non-human primate via direct contact of secretions, organs, bodily fluids, or blood. There is no evidence that suggests that Ebola virus can be transmitted through skin-to-skin contact or through small 1. Ebola Virus cannot be transmitted during the incubation period, whereas research shows that COVID-19 can be transmitted during the incubation period. Ebola virus disease can only be transmitted through bodily fluids or blood whereas COVID-19 spreads through the small droplets that a person lets out during coughing or sneezing. This makes COVID-19 more contagious.</td>
</tr>
</tbody>
</table>

1. The mode of transmission for the ZIKA Virus is the mosquito (55). Using mosquito repellents and staying in places that do not have mosquitoes can prevent a person from getting the virus. The COVID-19 virus mainly spreads from human-to-human transmission, and hence social distancing is the most important aspect in preventing it from spreading. 2. The human to human transmission in ZIKA Virus does not occur easily. When a person has the ZIKA virus they can only transmit it to another person when their blood comes in contact with another person. With COVID-19, respiratory droplets are the main mode of transmission. This makes the COVID-19 virus more contagious and allows it to spread easily. COVID-19 has asymptomatic patients and it can spread during the incubation period. 4. ZIKA virus can either be transmitted from a mosquito or an infected person’s body fluids. COVID-19 can go undetected in an asymptomatic patient and it can be transmitted to others. |

There are no vaccines or treatments available for the ZIKA virus and hence supportive care is recommended. Protection against mosquito bites during the day and the evening by using mosquito repellents that contain DEET, IR3535, or icaridin. Wearing clothing that covers most of the body and wearing light colored clothes to prevent mosquito bites. Physical barriers should be used like closed windows and doors. Sleep under the mosquito nets. Avoid sexual contact with people who are infected with the ZIKA virus. |

Reducing the contamination of wildlife-to-human transmission. It is advised to minimize contact with non-human primates as well as avoid the consumption of their raw meat. Reduction of human-to-human transmission. The patients that are infected and have died need to be buried properly to prevent the spread. |
| SARS       | Guangdong, China | 2002 | • The SARS outbreak began in November 2002 in the Guangdong province in China (61) 26 countries were impacted by the outbreak and there were over 8000 cases reported. Due to its large impact and spread it was announced to be a pandemic by WHO. The spread of the virus from China to other countries was found to be due to imported cases. SARS caused the death of nearly 1800 people in China. • The first outbreak that began in 2002, ended in 2004 and no more outbreaks of SARS have occurred after that. WHO had declared Guangdong to be a zone where re-emergence of the virus can occur.
|           |                  |      | • The spillover of the SARS virus is believed to be from an animal reservoir, though the animal species has not been identified yet but believed to be a bat (62). • Most transmission of the SARS virus occurred through person-to-person contact.  
  ○ Respiratory droplets of infected people produced when coughing and sneezing.  
  ○ The virus can spread through objects that can get contaminated by an infected person.  
  ○ Stool of an infected person.  
  ○ Inadequate prevention and control in the healthcare setting.  
|           |                  |      | • Certain studies regarding COVID-19 suggest that airborne transmission is possible (60).  
  1. Studies suggest that SARS does not spread during the incubation period (63).  
  • COVID-19 spreads during the incubation period which makes the virus more contagious and makes prevention harder.  
  • There are many patients of COVID-19 that are asymptomatic, which allows them to go undetected and spreads the virus further (64).  
  2. The reproduction number, R0, for SARS is found to be under 1, in isolation, but the median R0 is found to be between 2-4 (65).  
  • COVID-19 has an R0 that ranged between 2-2.5.  
  3. COVID-19 has asymptomatic and the virus spread during the incubation period, which overall makes it harder to track and allows the virus to spread from one person to another even before the original patient can be diagnosed with the virus (64).  
  4. Besides SARS being a big pandemic of the 21st century there are no treatments available to help patients.
|           |                  |      | At present there are no vaccines or FDA approved treatments that are available for SARS-CoV and supportive care is recommended.  
  • Provide healthcare workers with adequate protective gear to prevent the transmission from patients to healthcare workers  
  • Wash hand and maintain proper hygiene  
  • Promote testing and isolation of infected patients  
  • Avoid contact with animals and do not consume undercooked or uncooked meat.  

- The other significant breakout took place between December 2013 to June 2016 and it was the largest spread. The affected areas included Guinea, Liberia, and Sierra Leone (58). There were over 26,000 cases detected.
- droplets that are released when coughing and sneezing (58).
- The incubation period of the virus is 2 to 21 days and the virus cannot spread during the incubation period (59).
- transmission of the virus from the dead body.
- Sexual transmission is possible and hence awareness needs to be spread. Recovered patients need to be advised to practice safe sex for 12 months after the appearance of their symptoms as studies suggest that the virus can be present in the semen (56).
- Pregnant women infected with Ebola Virus can have the virus present in their milk even after they have recovered.
- Inadequate prevention and control in healthcare setting.
- Certain studies regarding COVID-19 suggest that airborne transmission is possible (60).  
  
1. Studies suggest that SARS does not spread during the incubation period (63).  
   • COVID-19 spreads during the incubation period which makes the virus more contagious and makes prevention harder.  
   • There are many patients of COVID-19 that are asymptomatic, which allows them to go undetected and spreads the virus further (64).  
   2. The reproduction number, R0, for SARS is found to be under 1, in isolation, but the median R0 is found to be between 2-4 (65).  
   • COVID-19 has an R0 that ranged between 2-2.5.  
   3. COVID-19 has asymptomatic and the virus spread during the incubation period, which overall makes it harder to track and allows the virus to spread from one person to another even before the original patient can be diagnosed with the virus (64).  
   4. Besides SARS being a big pandemic of the 21st century there are no treatments available to help patients.

- The spread of the virus can occur.  
- The emergence of the virus can be due to imported cases from China to other countries was found to be imported cases.
- The death of nearly 1800 people in China.
- The first outbreak that began in 2002, ended in 2004 and no more outbreaks of SARS have occurred after that. WHO had declared Guangdong to be a zone where re-emergence of the virus can occur.
- The SARS outbreak began in November 2002 in the Guangdong province in China (61) 26 countries were impacted by the outbreak and there were over 8000 cases reported. Due to its large impact and spread it was announced to be a pandemic by WHO. The spread of the virus from China to other countries was found to be due to imported cases. SARS caused the death of nearly 1800 people in China. • The first outbreak that began in 2002, ended in 2004 and no more outbreaks of SARS have occurred after that. WHO had declared Guangdong to be a zone where re-emergence of the virus can occur.
- The spillover of the SARS virus is believed to be from an animal reservoir, though the animal species has not been identified yet but believed to be a bat (62). • Most transmission of the SARS virus occurred through person-to-person contact.  
  ○ Respiratory droplets of infected people produced when coughing and sneezing.  
  ○ The virus can spread through objects that can get contaminated by an infected person.  
  ○ Stool of an infected person.  
  ○ Inadequate prevention and control in the healthcare setting.
- transmission of the virus from the dead body.
- Sexual transmission is possible and hence awareness needs to be spread. Recovered patients need to be advised to practice safe sex for 12 months after the appearance of their symptoms as studies suggest that the virus can be present in the semen (56).
- Pregnant women infected with Ebola Virus can have the virus present in their milk even after they have recovered.
- Inadequate prevention and control in healthcare setting.
- Certain studies regarding COVID-19 suggest that airborne transmission is possible (60).  
  
1. Studies suggest that SARS does not spread during the incubation period (63).  
   • COVID-19 spreads during the incubation period which makes the virus more contagious and makes prevention harder.  
   • There are many patients of COVID-19 that are asymptomatic, which allows them to go undetected and spreads the virus further (64).  
   2. The reproduction number, R0, for SARS is found to be under 1, in isolation, but the median R0 is found to be between 2-4 (65).  
   • COVID-19 has an R0 that ranged between 2-2.5.  
   3. COVID-19 has asymptomatic and the virus spread during the incubation period, which overall makes it harder to track and allows the virus to spread from one person to another even before the original patient can be diagnosed with the virus (64).  
   4. Besides SARS being a big pandemic of the 21st century there are no treatments available to help patients.

- The spread of the virus can occur.  
- The emergence of the virus can be due to imported cases from China to other countries was found to be imported cases.
| H1N1, Swine Flu | Mexico | 2009 | • The first case of the novel H1N1 appeared in Mexico in 2009 (48). By June of 2009, WHO had declared the outbreak of H1N1 virus that caused the swine flu a pandemic.  
• It impacted approximately 74 countries and territories. By September of 2009, there were nearly 200,000 confirmed cases from over 100 countries and there were over 17,000 deaths (66). The pandemic was declared to be over by the WHO in 2010. | It was found that the genes of the H1N1 virus are very similar to the genes of many influenza viruses that occur in pigs. H1N1 Swine Flu is believed to be passed from an animal, like a pig, to humans. Another mode of transmission is through human-to-human contact (67). The transmission can occur through the following:  
○ Respiratory droplets that are produced when sneezing and coughing.  
○ Infected patients contaminating inanimate objects or areas.  
○ The virus can enter through the eyes, or the nose. | 1. The H1N1 swine flu virus has a lower reproduction number, R0 value, than COVID-19. The 2009 swine flu virus had the reproduction number of 1.46 (68).  
○ COVID-19 has a reproduction number in between 2-2.5.  
○ This means that COVID-19 is more contagious than H1N1 and hence it can spread more easily.  
2. Difference in Viral Load across age (68)  
The viral load in adults and children is found to be very similar this shows that the virus impacts all patients similarly as the amount of virus found is similar (69)  
○ Studies suggest that H1N1, swine flu affects children more and the viral load is higher in children (70)  
○ Asymptomatic patients with COVID-19 make detection and isolation harder (65).  
○ H1N1, swine Flu, asymptomatic patients are very rare and this means that the patient can be diagnosed and isolated in the early stages preventing the virus from spreading further (71).  
WHO recommendations (72):  
• Washing hands regularly.  
• Maintaining a good respiratory hygiene which would include covering the mouth and nose when coughing and sneezing and using disposable tissues.  
• Self-isolation upon feeling unwell or feeling any of the symptoms.  
• Avoid touching the nose and the mouth.  
• Using proper protective gear when near sick patients. |
| MERS | Jeddha, Saudi Arabia(28) | 2012 | • The first outbreak of (Middle East Respiratory Syndrome) MERS occurred in 2012. The first patient that was diagnosed with MERS had a lower respiratory disease caused by the virus.  
• The countries that were majorly affected included the countries in Middle East. In total, 27 countries were affected by the Virus (73). | There is very little known about the MERS virus. It is a novel coronavirus that is believed to infect adults. The mortality rate of the patients infected with MERS virus was recorded to be 35%.  
• The transmission of the virus to humans is believed to have occurred from an animal. Many studies suggest that this spillover occurred from a camel (75).  
• There is very little known about the human-to-human transmission of the virus, as WHO has reported that most of the cases in Middle East have occurred through unprotected contact with infected animals like the camel. | 1. There is still very little known about the MERS virus, with the information that is gathered it can be established that MERS-CoV did not spread as much Ebola due to these possible reasons:  
1. Transmission of the MERS virus from one person to another was found to be rare and it was only noted in people who were close to the infected patient like healthcare workers and family members.  
○ The R0, reproduction number which determines how contagious a virus is, of MERS is reported to be less than 1. In a healthcare setting it has crossed (76). This means that if one There are no vaccines or treatments that have been FDA approved for the prevention and treatment of MERS-CoV(28).  
WHO advises the following:  
• Precaution while visiting places like the barn, farm, or any other places where dromedary camels and other animals are kept.  
• Ensuring that basic hygiene is kept in places where dromedary camels and other animals are kept.  
• Wash hands after touching the animals.  
[48], [66], [67], [68], [69], [70], [71], [72], [73], [74], [75], [76], [77]. |
Since the first outbreak there have been 2220 laboratory confirmed cases out of which 790 deaths occurred.

In 2015, there was a MERS outbreak in the Republic of Korea. It was the most impacted country outside of Middle East with the death of 36 people. The outbreak was linked to a traveler that travelled from the Middle East to The Republic of Korea (74).

Research also suggests that human-to-human transmission is really rare and in most cases the transmission has occurred in healthcare facility centers due to inadequate protection and prevention.

Avoid consumption of uncooked and undercooked meat. Be precautions while consuming milk.

People who have are immunocompromised or have diabetes, renal failure, and chronic lung disease are considered to be highly compromised and they should practice extra precaution.

- The R0 of COVID-19 is believed to be 2-2.5. This shows that COVID-19 is more contagious.
- As MERS-CoV was less contagious than COVID-19 it was easier to contain and prevent its spread (77).

By following the WHO guideline and restricting travel to the affected countries it was easier to contain MERS-CoV, especially as it was noted that the human to human transmission was rare.
FIGURE 3 FOREST PLOT FOR GENDER

<table>
<thead>
<tr>
<th>Study</th>
<th>Male</th>
<th>Proportion (%)</th>
<th>95% CI</th>
<th>Female</th>
<th>Proportion (%)</th>
<th>95% CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Zhang et al 2000</td>
<td>60.00</td>
<td>56.47 to 63.53</td>
<td>60.00</td>
<td>56.47 to 63.53</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Rhee et al 2000</td>
<td>45.00</td>
<td>35.55 to 54.45</td>
<td>45.00</td>
<td>35.55 to 54.45</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Zhou et al 2000</td>
<td>35.00</td>
<td>25.55 to 44.45</td>
<td>35.00</td>
<td>25.55 to 44.45</td>
<td></td>
<td></td>
</tr>
<tr>
<td>Hu et al 2000</td>
<td>50.00</td>
<td>40.00 to 60.00</td>
<td>50.00</td>
<td>40.00 to 60.00</td>
<td></td>
<td></td>
</tr>
</tbody>
</table>

FIGURE 4 FOREST PLOT FOR CO-MORBIDITY

<table>
<thead>
<tr>
<th>Study</th>
<th>Sample Size</th>
<th>Proportion (%)</th>
<th>95% CI</th>
</tr>
</thead>
<tbody>
<tr>
<td>Deng et al 2020</td>
<td>61</td>
<td>40.00</td>
<td>28.56 to 51.43</td>
</tr>
<tr>
<td>Yang et al 2020</td>
<td>200</td>
<td>32.00</td>
<td>25.96 to 38.04</td>
</tr>
<tr>
<td>RH et al 2020</td>
<td>179</td>
<td>84.38</td>
<td>78.19 to 89.37</td>
</tr>
<tr>
<td>Zhou et al 2020</td>
<td>191</td>
<td>47.84</td>
<td>40.38 to 55.34</td>
</tr>
<tr>
<td>Bogan et al 2020</td>
<td>19</td>
<td>100.00</td>
<td>82.53 to 100.00</td>
</tr>
<tr>
<td>Hu et al 2020</td>
<td>24</td>
<td>8.33</td>
<td>1.02 to 26.97</td>
</tr>
<tr>
<td>Shahintrai et al 2020</td>
<td>113</td>
<td>69.91</td>
<td>60.58 to 79.18</td>
</tr>
<tr>
<td>Jin et al 2020</td>
<td>34</td>
<td>61.76</td>
<td>53.56 to 70.07</td>
</tr>
<tr>
<td>Le et al 2020</td>
<td>5</td>
<td>40.00</td>
<td>10.27 to 85.33</td>
</tr>
<tr>
<td>Bhagat et al 2020</td>
<td>24</td>
<td>58.67</td>
<td>36.63 to 79.80</td>
</tr>
<tr>
<td>Kung et al 2020</td>
<td>83</td>
<td>18.07</td>
<td>10.48 to 25.00</td>
</tr>
<tr>
<td>Zhang et al 2020</td>
<td>28</td>
<td>100.00</td>
<td>87.05 to 100.00</td>
</tr>
<tr>
<td>Wan et al 2020</td>
<td>135</td>
<td>31.85</td>
<td>24.10 to 40.41</td>
</tr>
<tr>
<td>Yang et al 2020</td>
<td>143</td>
<td>43.48</td>
<td>37.65 to 52.96</td>
</tr>
</tbody>
</table>

Test for heterogeneity

| Q | 377.2354 |
| DF | 16 |
| Significance level | P < 0.0001 |
| I² (Inconsistency) | 95.76% |
| 95% CI for I² | 84.37 to 96.80 |
FIGURE 5 FOREST PLOT FOR HYPERTENSION, DIABETES MELLITUS AND CORONARY HEART DISEASE

[Diagram showing forest plots for various studies with effect sizes and 95% CIs for each study.]

Test for heterogeneity
- Q: 67.22
- DF: 10
- P: < 0.0001
- I²: 96.37%
- R²: 99.77%

FIGURE 6 FOREST PLOT FOR PROPORTIONS OF ASYMPTOMATIC SUBJECTS

[Diagram showing forest plots for various studies with effect sizes and 95% CIs for each study.]

Test for heterogeneity
- Q: 153.78
- DF: 9
- P: < 0.0001
- I²: 98.41%
- R²: 99.89%
Funnel plot for assessing publication bias based on study proportions is shown in Figure 8. Circles indicate study point estimates. There was no clear asymmetry found and therefore, funnel plot demonstrates no significant publication bias.