International Journal of Current Advanced Research

ISSN: O: 2319-6475, ISSN: P: 2319-6505, Impact Factor: 6.614 Available Online at www.journalijcar.org Volume 10; Issue 11 (B); November 2021; Page No.25521-25527 DOI: http://dx.doi.org/10.24327/ijcar.2021. 25527.5096



Research Article

CO-AFFECT OF SARS-COV-2 WITH OTHER RESPIRATORY VIRUSES

Raghavendra Rao¹., Mubasheer Alii M.V²., Chennamchetty Vijay Kumar³., Mahendra Kumar Verma⁴., Dilip Mathai⁵., Nikita Chitrapu⁶., Ritika Chitrapu⁷., Gil C Apacible⁸.,Maddineni Sai Aditya⁹ and H.N.Raghu¹⁰

¹Department of Medicine, Apollo Institute of Medical Sciences and Research, Jubilee Hills, Hyderabad, Telangana, India

²Consultant, MD Internal Medicine, Apollo Hospitals and Apollo TeleHealth Services, Associate Professor Department of General Medicine, Shadan Medical College, India

³Department of Pulmonary Medicine, Apollo Institute of Medical Science and Research, Hyderabad, TS, India, ⁴American University School of Medicine Aruba, Caribbean islands

⁵Department of Medicine, & Dean, Apollo Institute of Medical Sciences and Research, Hyderabad, TS, India ^{6,7}Shadan Degree and PG College for Women, Khiratabad, Hyderabad, TS, India

⁸Anatomical & Developmental Sciences, Neuroscience, Behavioral Science, and Preventive Medicine Epidemiology

Epidemiology

⁹Medical student finishing up electives in Beckley, WV USA,

¹⁰Medical student finishing up electives at Loretto hospital ,645 S Central Ave, Chicago, IL 60644,USA

ARTICLE INFO

Article History: Received 12th August, 2021 Received in revised form 23rd September, 2021 Accepted 7th October, 2021 Published online 28th November, 2021

Key words:

Coronavirus, Influenza virus, Dengue virus, Rhinovirus, 229E (alpha coronavirus), NL63 (alpha coronavirus), OC43 (beta coronavirus), HKU1, MERS-CoV, SARS-CoV, SARS-CoV-2

ABSTRACT

The earth is brimming with viruses. Co-infection with respiratory viruses and SARS-CoV-2 raising danger bells around the world. The world is living on the brink.COVID-19, an exciting outstanding pandemic co-infected with respiratory viruses, which demands crucial public health Intervention. The extent of co-infections of SARS-CoV-2 with other respiratory viruses remains unknown. Data on the prevalence and outcomes of co-infection and super infection are limited. The most significant challenge raised by co infection is perhaps the unsuitable propagation of recombination viruses through the interchange of genetic material amidst different strains. Influenza virus is a habitual pathogen that causes pneumonia in winter. It was found clinically that very few patients were diagnosed with both COVID-19 and influenza virus Rhinoviruses and coronaviruses are perceived as the significant source of the common cold syndrome.

Copyright©2021 **Raghavendra Rao et al.** This is an open access article distributed under the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.

INTRODUCTION

Identifying co-infections of more than one respiratory virus can help with understanding the various clinical symptoms, long-lasting effects on health, and appropriate methods of prevention (1)

Hospitalization in children under 3 years of age, following simultaneous infection with RSV and Human metapneumovirus (hMPV) (2) Furthermore, patients with viral respiratory tract coinfections are more likely to be admitted to the hospital than individuals with a single respiratory virus infection (3) Moreover, due to the similarity of common clinical symptoms of COVID-19 with that of the diseases caused by other respiratory viruses, it has become quite challenging to precisely distinguish SARS-CoV-2 infection from other viral infections (4) It is thought that co-infection with common respiratory viruses can still occur in individuals infected with other respiratory-specific strains (5)

Human infections with zoonotic corona viruses (CoVs), including severe acute respiratory syndrome (SARS)-CoV and Middle East respiratory syndrome (MERS)-CoV, have raised great public health concern globally.(6)

The virus, referred to as severe acute respiratory syndrome corona virus-2 (SARS-CoV-2), is transmitted through

^{*}Corresponding author: Raghavendra Rao

Department of Medicine, Apollo Institute of Medical Sciences and Research, Jubilee Hills, Hyderabad, Telangana, India

respiratory tract and could induce pneumonia (7)

Influenza virus has similar transmission characteristics with COVID-19, including direct contact (human-to-human transmission) and transmission via airborne droplets (8,9)

COVID-19 and influenza co-infection is limited. Coinfections with viral respiratory pathogens, including corona virus-influenza co-infections (10) and influenza-Middle East respiratory syndrome coronavirus infection, have been reported in a substantial number of patients.(11) CMV can predispose individuals to severe cases of COVID-19 through disruption of peripheral blood T-cell differentiation, and upregulation of inflammatory cytokines, including IL-6, especially in the elderly(12)

DENV,116 which is more prevalent in rainy seasons, causing conditions varying from mild dengue fever (DF) to severe hemorrhagic DF and dengue shock syndrome (13,14)

An investigation conducted by Yan et al. suggested the possibility of DENV and SARS-CoV-2 co infection, characterized by prolonged fever (15)

History

Coronavirus disease 2019 (COVID-19) is a contagious disease caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2).

The first known case was identified in Wuhan, China, in December 2019 (16) The disease has since spread worldwide, leading to an ongoing pandemic (17)

The first confirmed human infections were in Wuhan, Hubei, China. A study of the first 41 cases of confirmed COVID-19, published in January 2020 in The Lancet. (18,19)

Official publications from the WHO reported the earliest onset of symptoms as 8 December 2019 (20)

Human-to-human transmission was confirmed by the WHO and Chinese authorities by 20 January 2020 (21,22)

According to official Chinese sources, these were mostly linked to the Huanan Seafood Wholesale Market. (23)

In May 2020 George Gao, the director of the CDC, said animal samples collected from the seafood market had tested negative for the virus, indicating that the market was the site of an early super spreading event, but that it was not the site of the initial outbreak (24) Possibly the first influenza epidemic occurred around 6,000 BC in China, and possible descriptions of influenza exist in Greek writings from the 5th century BC (25) In both 1173–1174 and 1387, epidemics occurred across Europe that were named "influenza".

Whether these epidemics and others were caused by influenza is unclear since there was no consistent naming pattern for epidemic respiratory diseases at that time. (26)

Influenza may have been brought to the Americas as early as 1493, when an epidemic disease resembling influenza killed most of the population of the Antilles (27,28) The flu pandemic in 1557 was potentially the first time influenza was connected to miscarriage and death of pregnant women (29) The first record of a case of probable dengue fever is in a Chinese medical encyclopedia from the Jin Dynasty (266–420) which referred to a "water poison" associated with flying insects (30)

There have been descriptions of epidemics in the 17th century, but the most plausible early reports of dengue epidemics are from 1779 and 1780.(31) In 1953, when a cluster of nurses developed a mild respiratory illness, Winston Price, from the Johns Hopkins University, took nasal passage samples and isolated the first rhinovirus, which he called the JH virus, named after Johns Hopkins.His findings were published in 1956 (32)

Different types of human corona viruses

229E (alpha coronavirus) NL63 (alpha coronavirus) OC43 (beta coronavirus) HKU1 (beta coronavirus)

Influenza amidst Covid-19

Influenza, commonly called "the flu", and caused by influenza viruses. Fever, runny nose, sore throat, muscle pain, headache, coughing, and fatigue are the common symptoms. These symptoms begin 1–4 days after exposure to the virus and last for about 2–8 days..

Influenza and Corona viruses have similar transmission and clinical manifestations. The details of COVID-19 and influenza co infection is limited (33) laboratory findings Skiagraphy and other imaging studies, show similar clinical characteristics both in Influenza and COVID-19 patients Nasal tampon and pharyngalgia may be more prone to appear for those coinfection patients (34) The influenza season and the COVID-19 pandemic has created a similar and strenuous challenge. COVID-19 and influenza are both contagious respiratory illnesses With the ongoing winter season in the northern hemisphere, it is expected that the seasonal influenza virus will be in circulation along with SARS-CoV-2 (35)

Dengue amidst COVID-19

The mystery of plummeting cases Classical dengue fever is characterized by the onset of fever, malaise, headache, facial flushing, retrobulbar pain that worsens on eye movements, conjunctival suffusion and severe back ace, which is a prominent symptom.

Lymphadenopathy, petechiae on the soft palate and transient morbilliform skin rashes may also appear on the limbs with subsequent spread to involve the trunk Desquamation occurs. Cough is uncommon. Dengue fever syndrome is characterized by capillary leak syndrome, thrombocytopenia, hemorrhage, hypotension, and shock. It is characteristically a disease of children occurring most commonly in South-East Asia. Warning Signs of Dengue are stomach pain, tenderness, Vomiting (at least 3 times in 24 hours),Bleeding from the nose or gums, Blood in vomit or stool, Feeling tired, restless, or irritable but Trouble breathing, Persistent pain or pressure in the chest, are the signs of Covid-19 (36)

Rhinovirus amidst Covid-19

How the common cold can boot out Covid The viruses of this group are responsible for the most frequent of all human infections, the common cold. Most people suffer from 2-4 colds every year and, although the primary infection is not a severe one, secondary bacterial infection often follows with temporary incapacity. These viruses cause the loss of many million man hours of work. When rhinoviruses are instilled into nose of human volunteers a mild sore throat and cough are the pulmonary signs which precede the profuse nasal discharge of typical common cold. Chimpanzees are the only other creatures known to be susceptible to these viruses. The viruses can be recovered from the secretions of the nose and throat but only rarely from the faeces.

The virus settles on the ciliated nasal epithelial cells, enters the cells and spreads from cell to cell the epithelium. The cilia become immobilized and both cilia and cell degenerate as the virus replicates. Although it is often suggested that bacterial invasion of the damaged tissue occurs and is responsible for the purulent nasal discharge which is a feature, the engulfing of dead or virus damaged epithelial cells by phagocytes of various sorts may be primary cause of this discharge. The cells lining the throat and lower airways are exposed to the environment making them a prime target for co-infection by respiratory viruses.

Corona viruses and Acute Respiratory Syndromes MERS, and SARS

Fever, cough, headache, fatigue, breathing difficulties, and loss of smell and taste are the common symptoms of Covid-19 (37) 30%-40% of people who are infected do not develop noticeable symptoms (38) Of those people who develop symptoms noticeable enough to be classed as patients, most (81%) develop mild to moderate symptoms (up to mild pneumonia), while 14% develop severe symptoms (dyspnea, hypoxia, or more than 50% lung involvement on imaging), and 5% suffer critical symptoms (respiratory failure, shock, or multiorgan dysfunction (39) MERS-CoV is a coronavirus believed to be originally from bats.Typical symptoms include fever, cough, diarrhea, and shortness of breath. The disease is typically more severe in those with other health problems (40) Middle East respirator syndrome coronavirus, or MERS-CoV) was first diagnosed in Saudi Arabia in 2012.

Middle East respiratory syndrome (MERS) is a viral respiratory disease caused by novel coronaviruses. Coronaviruses can cause diseases fluctuating from common cold to Severe Acute Respiratory Syndrome (SARS). 35% of patients died with MERS-CoV infection. Middle East respiratory syndrome is caused by the MERS coronavirus (MERS-CoV), a species with single-stranded RNA belongs to the genus beta corona virus which is distinct from SARS coronavirus and the common-cold corona virus (41).

COVID-19 is correlated with an acute respiratory distress syndrome (ARDS) and severe pneumonia (42)

COVID-19 was the first severe disease to make an appearance in the 21st century and spread all over the globe. The receptor binding domain (RBD) in S protein of this virus gets attached with the ACE2 receptors present on human cells.

MERS CoV was first reported in 2012 in Middle East, originated from bat and transmitted to humans through camels (43)

COVID-19 virus produces mild to moderate respiratory illness. The COVID-19 virus spreads primarily through droplets of saliva or when an infected person coughs or sneezes.

Bird flu outbreak amidst COVID-19

Pandemic South Africa has testimony for several outbreaks of avian influenza (H5N1) in dissimilar poultry farms.

It is a shock that was already suffering from the deadly COVID-19 pandemic. South Africa is facing the economic COVID-19 crisis, the bird flu causing a significant incision in the economy, as well as severely impacting people's day-to-day life. Bird flu-also called avian influenza-is a viral infection that can infect not only birds but also humans and other animals

Unavailability of suitable vaccines, correct treatment, and deficiency of public health measures are major setbacks.(44)

COVID-19 and Zika association

Zika virus was first identified in Uganda in 1947. WHO announced Zika virus disease as a "Public Health Emergency of International Concern".

Till last year it was a concern of Latin America and Africa but now it sporadically spread in India too, where pregnant mothers usually have no history of traveling to an affected area of foreign nations. Zika virus disease is caused by Zika virus which belongs to the Flavivirus. Zika virus is transmitted through the bite of an infected Aedes mosquito. Ades mosquito which is known to transmit infections like dengue and chikungunya. It is associated with severe neurological complications, particularly congenital Zika virus syndrome and Guillain-Barre syndrome. It is transmitted from an infected pregnant mother to her baby during pregnancy Transmitted from sexual contact with an infected person.

Difference between corona virus and Viral fever

COVID-19 and Influenza both cause respiratory sickness, and are caused by different viruses. COVID-19 is caused by infection with a coronavirus, and flu is caused by infection with influenza viruses. COVID-19 spreads more easily than flu. A person experiences flu symptoms from 1 to 4 days after infection, But Covid-19, symptoms can appear 2 to 14 days after infection.

Virology

Dengue virus (DEVN) RNA virus belongs to the family of Flavivirus genus of the Flaviviridae family. Other viruses that are related to Dengue are Japanese B encephalitis and Yellow fever.

There are four serotypes DEVN 1, DEVN 2, DEVN 3 and DEVN 4. A fifth serotype DEVN 5 has been described recently. Infection from one virus produces lifelong immunity to that strain of the virus. The viruses contain a single strand of RNA. It is enclosed by a lipoprotein and has a nucleocapsid. It is composed of three structural proteins encoding the nucleocapsid protein(C), a membrane associated protein (M) and an envelope protein (E). There are seven nonstructural proteins known as NS proteins. The NS proteins are involved in viral replication and packaging.

These processes are closely linked to host endoplasmic reticulum and secretory pathway function.

Coronaviruses belong to the family Coronaviridae. They are enveloped viruses with a positive-sense singlestranded RNA genome. The genome size of coronaviruses is one of the largest among RNA viruses. They have clubshaped spikes that project from their surface, Middle East respiratory syndrome caused by the MERS is has single-stranded coronavirus (MERS-CoV), It RNA belonging to the genus betacoronavirus. Its genomes are phylogenetically classified into two clades, A and B. Influenza viruses have a negative-sense, single-stranded RNA genome. The negative sense of the genome means it can be used as a template to synthesize messenger RNA (mRNA). IAV and IBV have eight genome segments that encode 10 major proteins. ICV and IDV have seven genome segments that encode nine major proteins. Rhinoviruses belong to the genus Enterovirus in the family Picornaviridae. The three species of rhinovirus (A, B, and C) include around 160 recognized types of human rhinovirus that differ according to their surface proteins, and are the smallest viruses.

Research on viral infection of the lung: Host Response and Sequelae

The recognition of pathogen-associated molecular patterns by the pattern recognition receptors such as the stromal cells (CD45-) and cells in the lung (CD45+) is the key to initiate the host immune response to the microbial invasion. As well as the recognition of damage-associated danger signals are important to activate the intracellular innate protein complex, the inflammasome, which are responsible in the organization of both the innate and adaptive immune responses.

In response to a viral respiratory tract infection, interferon (type I and III) are produced by the recognition of: The melanoma differentiation-associated protein 5 and the retinoids acid-inductively gene I, both cytotoxic receptors, will recognize the double-stranded RNA. Unknown cytosolic DNA receptors or DAI will recognize the double-stranded DNA. Retinoids acid-inductively gene I is also triggered by 5'pppRNA from the transcription of double-stranded DNA, using an RNA polymerase III. After the recognition has been made, the interferon regulatory transcription factors will be activated through the kinase TANK-binding kinase; which then leads to the production of interferon type I and III in the site of the viral infection.

Viral molecules arrive at their respective TLR3, TLR4, TLR7, and TLR9 which will then stimulate through the adaptor protein molecules for the production of interferon type I; leading to the transcription of interferon genes and proinflammatory cytokines. Type I interferons (IFN-a, IFN-ß) and type III (IFN-?) share many functions, including induction by viral infection, activation of shared signaling pathways, and transcriptional programs. Interferon type I binds to the receptor IFNAR2 with high-affinity, which then activates a low-affinity IFNAR1 to form a signaling complex.

How the Corona virus Attacks Lungs

The genetic material of viruses enveloped by protein and fat.

They are connecting links between life and death. As long as the virus is present in the cell, they are living, but when present outside the cell they are non-living.

Lung macrophages challenged with respiratory viruses

Most of the cells of the immune system derived from the hematopoietic system. Phagocytic cells are found in the circulation (Monocytes and granulocytes) and reside in the tissues (macrophages).Each cell type expresses characteristic surface molecules (CD3,CD4,CD8.) Alveolar or pulmonary macrophages present freely on the outer surface of the lung.These cells scavenging the dust particles, microorganisms and other debris.The primary function of macrophage is phagocytosis. The macrophages, by their property of amoeboid movement, put forth pseudopodia which help in engulfing any solid particle such as the invading microorganisms. The macrophages have lysosomal granules, containing acid hydrolases and degradative enzymes with which it destroys the phagocytosed substances. The attachment of antigens to macrophage is specific.

All the macrophages have surface receptors for the C3 component of complement as well as for Fc component of antibody. The macrophages are involved in processing of antigens before they are presented to the T and B cells. (45) Lower respiratory tract infection due to RSV is a complicated process and several cell types are implicated in the disease progression (46) Disease severity due to extensive lung tissue damage correlates with enhanced pro-inflammatory cytokine secretion and inflammation. (47).

Pulmonary macrophage cells may behave differently with respect to RSV infection (48)

Laboratory Findings

Early detection may help to use appropriate antiviral agents. 1. Complete blood count may reveal leukocytosis with mild left shift. 2. Gram stains of the sputum may show few too many polymorphonuclear leukocytes.

Radiographic Findings

Chest X ray shows perihilar pulmonary edema, or dense opacities.

Diagnosis

Culture- Influenza virus can be isolated from upper and lower respiratory samples. However, because of the long turnaround time of reports, they are not used routinely. 2. Serology-Complement fixation and hemagglutinin inhibition during the acute phase of the illness and repeat test at 2-3 weeks during the convalescent phase to demonstrate a fourfold rise in antibody titer may be used. 3. Rapid antigen assays- less sensitive in detecting Influenza A infections as compared to RT-PCR assays. 4. Molecular diagnosis- Molecular methods like multiplex PCR / RT PCR are increasingly useful for the detection of H1N1, H3N2, and novel swine H1N1v.

The retrospective study included 461 patients who had been hospitalized with acute Covid-19 within four days of development of symptoms and positive RT PCR test for SARS-CoV-2 using a nasopharyngeal sample. (49)

Treatment and prevention

Influenza patients should be kept in bed until the fever has subsided. A mild analgesic usually relieves the head and backache. The vaccine is administered in cold countries to protect elderly people with preexisting cardiopulmonary or nasal disease and to other high-risk individuals with influenza outbreak is imminent.

RSV is worldwide in distribution and causes a winter epidemic in infants and children below one year of age Both formalininactivated crude whole-virus vaccine as well as a live vaccine have been tried but none is found satisfactory. Attempts to prepare vaccines with purified F and G surface glycoproteins of RSV are in progress.

COVID-19 Vaccine

The specific immune response against the pathogen is the key element stimulated by vaccine administration, Nanomaterialbased vaccines have proven prodigious prophylaxis of various infectious and non-infectious diseases of human and animal concern. The control of the spread of the virus and preventing severe fatal illness are the primary goals of the COVID-19 vaccine. It is widely accepted that neutralizing antibodies are key for the protection, which prevents the viruses from attacking the cells by blocking the SARS-CoV-2 spike protein interaction domain from binding with the ACE To control the pandemic, a constructive vaccine candidate should achieve immunity around 60–70% of the population to the infection to accomplish the concept of "herd immunity". The prominent genomic homology of the SARS-CoV2 virus with other beta coronaviruses like SARS-CoV and MERS-CoV supported the vaccine researchers to immensely accelerate the development of efficacious vaccine formulations to the status of a global healthcare issue. On January 11, 2020, the sequence of the SARS-CoV-2 RNA genome was published, which boosted in designing various varieties of innovative vaccine technologies, including modified or recombinant proteins, peptide fragments, particle mimicking the virus, viral vectors, nucleic acids (RNA and DNA), and complete virus vaccines (inactivated or live attenuated virus). Development of vaccines typically takes decades, but overwhelming demand of vaccination, knowledge on years of previous studies on relevant diseases which made significant achievements, sufficient funding for vaccine developers to run multiple trials simultaneously, and regulators decision making at unprecedented speed. Attenuated vaccine pathogens are known for their complexities. Even if considering tremendous advancements in rDNA technology in recent times empower broad exploration of antigen (protein or peptide) encoded recombinant vaccines, subunit vaccines, and nucleic acidbased vaccines.

Research programs for the next generation world

The main goal of most vaccination strategies is to be able to induce a strong and healthy virus-specific neutralizing antibody response in the host.

By doing so through the induction of virus-specific CD8 T-cell response and its humoral immunity. This dual approach may allow optimal viral control. Nowadays with viruses such as Respiratory Syncytial Virus, Parainfluenza Virus, Influenza Virus, Human Metapneumovirus, Rhinovirus, Parainfluenza Virus and Coronavirus. Researchers are exploring different technologies to tackle these respiratory infections. Virus vaccines: by using the virus in a weakened form or inactivated. Viral-vector vaccines: by genetically engineering the measles of adenovirus (weakened) to produce coronavirus proteins in the body.

Two types are those that can still replicate within the cells without producing a disease and the other one that cannot reproduce in the cells by disabling key genes. Nucleic-acid vaccines: by using genetic instructions, in the form of DNA or RNA, for a coronavirus protein that will encourage in the host the immune response. The nucleic acids are inserted into human cells that will churn out copies of the virus protein. Protein-based vaccine: injection of the virus proteins, fragments or protein shells that mimics the virus outer coat, directly into the host. (50)

CONCLUSION

Advances in immunology, immunopathology, and immunopharmacology have already opened a way to a clearer understanding of the nature of certain respiratory diseases. Immunology has a further contribution to make in the field of respiratory infection Coronaviruses are known to cause respiratory infections ranging from the common cold to more severe diseases.

References

- 1. Ozaras R, Cirpin R, Duran A, et al. Influenza and COVID-19 coinfection: report of six cases and review of the literature. J Med Virol. 2020; 92(11): 2657-2665.
- König B, König W, Arnold R, Werchau H, Ihorst G, Forster J. Prospective study of human metapneumovirus infection in children less than 3 years of age. J Clin Microbiol. 2004; 42(10): 4632- 4635.
- Drews AL, Atmar RL, Glezen WP, Baxter BD, Piedra PA, Greenberg SB. Dual respiratory virus infections. Clin Infect Dis. 1997; 25(6): 1421-1429
- 4. .4. Huang C, Wang Y, Li X, et al. Clinical features of patients infected with 2019 novel coronavirus in Wuhan, China. The Lancet. 2020; 395(10223): 497-506.
- 5. Lee N, Chan PK, Yu IT, et al. Co-circulation of human metapneumovirus and SARS-associated coronavirus during a major nosocomial SARS outbreak in Hong Kong. J Clin Virol. 2007; 40(4): 333- 337.
- Li-Li Ren 1, Ye-Ming Wang 2 3, Zhi-Qiang Wu 4, Zi-Chun Xiang 1, Li Guo 1, Teng Xu 5, Yong-Zhong Jiang 6, Yan Xiong 7, Yong-Jun Li et al,
- 7. Identification of a novel coronavirus causing severe pneumonia in human: a descriptive study, Chin Med J (Engl), 2020 May 5;133(9):1015-1024.
- 8. Chen N, Zhou M, Dong X, et al. Epidemiological and clinical characteristics of 99 cases of 2019 novel coronavirus pneumonia in Wuhan, China: a descriptive study. Lancet. 2020; 395(10223): 507- 513
- 9. Zhang N, Wang L, Deng X, et al. Recent advances in the detection of respiratory virus infection in humans. J Med Virol. 2020; 92(4): 408- 417.
- 10. Jiang C, Yao X, Zhao Y, et al. Comparative review of respiratory diseases caused by coronaviruses and influenza A viruses during epidemic season. Microbes Infect. 2020.
- Stefanska I, Romanowska M, Donevski S, Gawryluk D, Brydak LB. Co-infections with influenza and other respiratory viruses. Adv Exp Med Biol. 2013; 756: 291-301.
- 12. Alfaraj SH, Al-Tawfiq JA, Alzahrani NA, Altwaijri TA, Memish ZA. The impact of co-infection of influenza A virus on the severity of Middle East respiratory syndrome coronavirus. J Infect. 2017; 74(5): 521-533
- Kadambari S, Klenerman P, Pollard AJ. Why the elderly appear to be more severely affected by COVID-19: the potential role of immunosenescence and CMV. Rev Med Virol. 2020;30(5):e2144.
- 14. 13, Halstead SB. Dengue. The Lancet. 2007; 370(9599): 1644-1652.
- 15. Setiati TE, Wagenaar JFP, de Kruif M, Mairuhu A. Changing epidemiology of dengue haemorrhagic fever in Indonesia. Dengue Bull. 2006; 30: 1- 14.

- 16. Yan G, Lee CK, Lam L, et al. Covert COVID-19 and false-positive dengue serology in Singapore. Lancet Infect Dis. 2020; 20(5): 536.
- 17. Page J, Hinshaw D, McKay B (26 February 2021). "In Hunt for Covid-19 Origin, Patient Zero Points to Second Wuhan Market – The man with the first confirmed infection of the new coronavirus told the WHO team that his parents had shopped there". The Wall Street Journal. Retrieved 27 February 2021.
- Zimmer C (26 February 2021). "The Secret Life of a Coronavirus – An oily, 100-nanometer-wide bubble of genes has killed more than two million people and reshaped the world. Scientists don't quite know what to make of it". Retrieved 28 February 2021.
- 19. Wu YC, Chen CS, Chan YJ (March 2020). "The outbreak of COVID-19: An overview". Journal of the Chinese Medical Association. 83 (3): 217–220.
- 20. Wang C, Horby PW, Hayden FG, Gao GF (February 2020). "A novel coronavirus outbreak of global health concern". Lancet. 395 (10223): 470–473.
- 21. "Novel Coronavirus China". World Health Organization (WHO). 12 January 2020.
- Kessler G (17 April 2020). "Trump's false claim that the WHO said the coronavirus was 'not communicable'". The Washington Post. Archived from the original on 17 April 2020. Retrieved 17 April 2020.
- 23. Kuo L (21 January 2020). "China confirms human-tohuman transmission of coronavirus". The Guardian. Retrieved 18 April 2020.
- 24. Epidemiology Working Group For Ncip Epidemic Response, Chinese Center for Disease Control Prevention (February 2020). "[The epidemiological characteristics of an outbreak of 2019 novel coronavirus diseases (COVID-19) in China]". Zhonghua Liu Xing Bing Xue Za Zhi = Zhonghua Liuxingbingxue Zazhi (in Chinese). 41 (2): 145–151.
- 25. Areddy JT (26 May 2020). "China Rules Out Animal Market and Lab as Coronavirus Origin". The Wall Street Journal. Retrieved 29 May 2020.
- 26. Martin PM, Martin-Granel E (June 2006). "2,500-year evolution of the term epidemic". Emerging Infectious Diseases. 12 (6): 976–980.
- 27. Morens DM, North M, Taubenberger JK (4 December 2010). "Eyewitness accounts of the 1510 influenza pandemic in Europe". Lancet. 376 (9756): 1894–1895.
- 28. Guerra F (1988). "The earliest American epidemic. The influenza of 1493". Soc Sci Hist. 12 (3): 305–325.
- 29. Guerra F (1993). "The European-American exchange". Hist Philos Life Sci. 15 (3): 313–327.
- Institute of Medicine (US) Forum on Microbial Threats (2005). "1: The Story of Influenza". In Knobler S, Mack A, Mahmoud A, Lemon S (eds.). The Threat of Pandemic Influenza: Are We Ready? Workshop Summary (2005). Washington, DC: The National Academies Press. pp. 60–61
- Anonymous (2006). "Etymologia: dengue" (PDF). Emerg. Infect. Dis. 12 (6): 893.from the original on 3 December 2013
- 32. Kuno G (November 2015). "A re-examination of the history of etiologic confusion between dengue and chikungunya". PLOS Neglected Tropical Diseases. 9 (11): e0004101. 1779–1780: 'Knokkel-koorts' in Batavia (now Jakarta, Indonesia) and 'break bone fever'

in Philadelphia

- 33. Kennedy, Joshua L; Turner, Ronald B.; Braciale, Thomas; Heymann, Peter W.; Borish, Larry (June 2012). "Pathogenesis of Rhinovirus Infection". Current Opinion in Virology. 2(3): 287–293
- 34. Resat Ozaras, Rasim Cirpin, Arif Duran, Habibe Duman, Ozgur Arslan, Yasin Bakcan, Metin Kaya, Huseyin Mutlu, Leyla Isayeva, Fatih Kebanlı, Bekir A Deger, Eldar Bekeshev, Fatma Kaya, Suat Bilir, Influenza and COVID-19 coinfection: Report of six cases and review of the literature, J Med Virol2020 Nov;92(11):2657-2665
- 35. The clinical characteristics of pneumonia patients coinfected with 2019 novel coronavirus and influenza virus in Wuhan, China,J Med Virol2020 Sep;92(9):1549-1555.
- 36. A. Agarwal, M. Agarwal, A. Sharma, R. Jakhar, Impact of Influenza A Co-Infection With Covid-19,2021, A Centuury of Leadership In Health
- Saniasiaya J, Islam MA (November 2020). "Prevalence and Characteristics of Taste Disorders in Cases of COVID-19: A Meta-analysis of 29,349 Patients". Otolaryngology–Head and Neck Surgery. 165 (1): 33– 42.
- Agyeman AA, Chin KL, Landersdorfer CB, Liew D, Ofori-Asenso R (August 2020). "Smell and Taste Dysfunction in Patients With COVID-19: A Systematic Review and Meta-analysis". Mayo Clin Proc. 95(8). 95 (8): 1621–1631.
- Oran DP, Topol EJ (January 2021). "The Proportion of SARS-CoV-2 Infections That Are Asymptomatic: A Systematic Review". Annals of Internal Medicine. 174 (5): M20-6976.
- Interim Clinical Guidance for Management of Patients with Confirmed Coronavirus Disease (COVID-19)".
 U.S. Centers for Disease Control and Prevention (CDC). 6 April 2020. Archived from the original on 2 March 2020. Retrieved 19 April 2020.
- 41. "Middle East respiratory syndrome coronavirus (MERS-CoV)". www.who.int. Retrieved 15 April 2020.
- 42. Saey, Tina Hesman (2013). "Story one: Scientists race to understand deadly new virus: SARS-like infection causes severe illness, but may not spread quickly among people". Science News. 183 (6): 5–6
- 43. N Petrosillo, G Viceconte, O Ergonul, G Ippolito, E Petersen,, COVID-19, SARS and MERS: are they closely related? Clin Microbiol Infect,2020 Jun; 26(6):729-734.
- 44. Nosheen Masood, Saima Shakil Malik, Muhammad Naqqi Raja, Sumaira Mubarik, Chuanhua Yu, Unraveling the Epidemiology, Geographical Distribution, and Genomic Evolution of Potentially Lethal Coronaviruses (SARS, MERS, and SARS CoV-2),Front Cell Infect Microbiol,2020 Aug 27;10:49
- 45. Chandrakant Lahariya, A K Sharma, S K Pradhan, Avian flu and possible human pandemic, Indian Pediatr,2006 Apr;43(4):317-25.
- Dulsy Fatima and Arumugam,2002.Saras publication 35G, ARP Camp road, Periavillai, Kottar Post, Nagarkoil, Kanyakumari, Tamilnadu, India
- Collins PL, Graham BS: Viral and host factors in human respiratory syncytial virus pathogenesis. J Virol. 2008, 82 (5): 2040-2055. 10.1128/JVI.01625-07

- 48. Johnson JE, Gonzales RA, Olson SJ, Wright PF, Graham BS: The histopathology of fatal untreated human respiratory syncytial virus infection. Mod Path. 2007, 20 (1): 108-119. 10.1038/modpathol.3800725.
- 49. Haeberle HA, Takizawa R, Casola A, Brasier AR, Dieterich HJ, Van Rooijen N, Gatalica Z, Garofalo RP: Respiratory syncytial virus-induced activation of nuclear factor-kappaB in the lung involves alveolar macrophages and toll-like receptor 4-dependent pathways. J Infect Dis. 2002, 186 (9): 1199-1206. 10.1086/344644.
- 50. https://www.nature.com/articles/d41586-020-01221-y
- 51. Veena Aggarwal, Consultant Womens' Health, CMD and Editor-in-Chief, IJCP Group & Medtalks Trustee, Dr KK's Heart Care Foundation of India, 29 September 2021

How to cite this article:

Raghavendra Rao et al (2021) 'Co-Affect Of Sars-Cov-2 With Other Respiratory Viruses', International Journal of Current Advanced Research, 10(11), pp. 25521-25527. DOI: http://dx.doi.org/10.24327/ijcar.2021. 25527.5096
