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WEEKLY EPIDEMIOLOGICAL REPORT

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Human Metapneumovirus Infection: An Emerging Respiratory Pathogen

Introduction

Human Metapneumovirus (HMPV) is a newly emerging respiratory pathogen that can cause upper and lower respiratory disease in people of all ages, especially among young children, older adults, and people with weakened immune systems. HMPV was first discovered in 2001 and is in the Pneumoviridae family along with the respiratory syncytial virus (RSV). HMPV is a singlestranded RNA virus that is highly contagious and can spread through respiratory secretions, such as coughing and sneezing. The virus is primarily associated with bronchiolitis and pneumonia, and it poses a potential risk for bacterial pneumonia. The coexistence of HMPV with respiratory syncytial virus can lead to more severe clinical symptoms, along with complications such as asthma and exacerbations of chronic obstructive pulmonary disease. Given the potential impact on patients, enhancing our understanding of this virus is crucial to improve patient care.

Epidemiology

HMPV has a global distribution and is responsible for a substantial burden of respiratory tract infections. It exhibits a seasonal pattern, with peak activity observed during winter and spring in temperate regions. HMPV infection commonly leads to localized outbreaks in communities, healthcare settings, and long-term care facilities. Understanding the epidemiology of HMPV is crucial for implementing appropriate preventive and control measures. The COVID-19 pandemic has changed the epidemiology of respiratory viral infections, modifying the classic seasonality of respiratory syncytial virus (RSV), influenza, and other viruses. A recent study in Spain found an unusual outbreak of respiratory infections caused by hMPV in children during the sixth wave of COVID-19, associated with the Omicron variant. Patients in this outbreak were older than usual and showed more hypoxia and pneumonia, longer length of stay, and greater need for intensive care

HMPV is an important emerging respiratory pathogen that has been reported in many countries around the world. In Sri Lanka, HMPV has been identified as a significant cause of acute respiratory infections (ARI) in children under five years of age. A study conducted in 2006 found that HMPV was detected in 10.8% of children with ARI in Sri Lanka. Another study conducted in 2017 found that HMPV was detected in 13.6% of children with ARI in Sri Lanka. The increasing prevalence of HMPV infection in Sri Lanka highlights the need for increased awareness and surveillance of this emerging respiratory pathogen.

Although the description of this viral pathogen was first described in children, subsequent reports have highlighted the importance of human metapneumovirus as a cause of respiratory illness in adults of all ages, in patients with cancer, in the elderly population (as a cause of serious lower respiratory tract infection), and in adults with underlying chronic medical conditions.

Pathophysiology

HMPV is a negative-sense RNA virus that is similar to RSV in its structure and pathogenesis. The virus infects the respiratory epithelium, causing inflammation and damage to the airways. The virus can also infect immune cells, leading to a dysregulated immune response that can exacerbate the severity of the disease. The role of human metapneumovirus in causing respiratory illness in adults has been increasingly recognized in recent years.

Clinical Presentation

HMPV infection manifests as a spectrum of respiratory illnesses, ranging from mild upper respiratory tract infections to severe lower respiratory tract diseases. The clinical presentation of HMPV infection is similar to that of other respiratory viruses, making it challenging to differentiate clinically. Common symptoms include cough, nasal congestion, sore throat, fever, and malaise. In severe cases, respiratory distress may occur, necessitating hospitalization and intensive care. It can even lead to death, particularly in young children, older adults, and individuals with weakened immune systems.



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Diagnosis

Laboratory diagnosis of HMPV infection relies on the detection of viral nucleic acids in respiratory specimens. Molecular techniques such as reverse transcription-polymerase chain reaction (RT-PCR) are the gold standard for HMPV detection. Serological tests can be employed to detect HMPV-specific antibodies, aiding in retrospective diagnosis and epidemiological studies. Rapid antigen tests, although convenient, may vary in sensitivity and specificity.

Management and Prevention

There is currently no specific antiviral treatment for HMPV infection, and management primarily revolves around supportive care. Adequate hydration, fever management, and respiratory support are essential components of patient management. In severe cases, hospitalization may be required for oxygen supplementation and intensive care. Vaccines against HMPV are under development and hold promise for reducing the burden of HMPV infection.

Prevention strategies mainly focus on infection control measures to limit the spread of HMPV. Strict adherence to hand hygiene, respiratory etiquette, and appropriate isolation precautions in healthcare settings is crucial. Public health campaigns and education aimed at raising awareness among healthcare professionals and the general population are instrumental in preventing HMPV transmission.

Impact on Public Health

HMPV infection exerts a significant impact on public health globally. In children, it ranks as one of the leading causes of hospitalization for respiratory infections, second only to respiratory syncytial virus (RSV). Severe complications and increased mortality rates have been observed in older adults and immunocompromised individuals. The economic burden of HMPV infection is substantial due to healthcare costs, including hospitalizations, laboratory testing, and loss of productivity.

Future Directions

The development of effective vaccines and antiviral therapies remains a priority in the field of HMPV research. Ongoing efforts to understand the immune response and viral pathogenesis will contribute to the development of targeted interventions. Enhanced surveillance systems and increased awareness among healthcare professionals are essential for monitoring the epidemiology of HMPV and implementing appropriate preventive measures.

Conclusion

Human Metapneumovirus (HMPV) infection is an emerging respiratory pathogen that poses significant challenges to the medical community. Its clinical presentation overlaps with other respiratory viruses, necessitating accurate diagnostic techniques. Supportive care and appropriate management strategies are crucial in addressing HMPV infection. The development of effective vaccines and antiviral therapies remains a key area of research. Furthermore, enhancing surveillance systems and promoting awareness among healthcare professionals can aid in the timely diagnosis and implementation of preventive measures. As HMPV continues to emerge as a significant respiratory pathogen, collaboration among researchers, healthcare providers, and public health authorities is vital in minimizing its impact on global public health. By staying updated with the latest research and maintaining a vigilant approach, the medical community can effectively tackle the challenges posed by HMPV and improve patient outcomes.

Compiled by

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Number of microbiological water samples June 2023										
District	MOH areas	No: Expected *	No: Received							
Colombo	15	90	0							
Gampaha	15	90	NR							
Kalutara	12	72	62							
Kalutara NIHS	2	12	16							
Kandy	23	138	24							
Matale	13	78	0							
Nuwara Eliya	13	78	67							
Galle	20	120	NR							
Matara	17	102	12							
Hambantota	12	72	34							
Jaffna	12	72	NR							
Kilinochchi	4	24	2							
Manner	5	30	NR							
Vavuniya	4	24	49							
Mullatvu	5	30	31							
Batticaloa	14	84	0							
Ampara	7	42	37							
Trincomalee	11	66	0							
Kurunegala	29	174	NR							
Puttalam	13	78	3							
Anuradhapura	19	114	NR							
Polonnaruwa	7	42	0							
Badulla	16	96	NR							
Moneragala	11	66	58							
Rathnapura	18	108	NR							
Kegalle	11	66	3							
Kalmunai	13	78	0							

^{*} No of samples expected (6 / MOH area / Month)

NR = Return not received

Table 1: Selected notifiable diseases reported by Medical Officers of Health 08th-14th July 2023 (28th Week)

abl	ole 1: Selected notifiable diseases reported by Medical Officers of Health 08th-14th July 2023 (28th Week)																												
	*5	100	100	100	100	100	100	100	100	100	93	100	100	100	100	100	48	100	100	66	86	86	100	100	100	100	66	86	
WRCD	<u>*</u>	27	m	7	86	23	09	36	5 6	23	64	21	38	12	23	28	12	25	24	22	24	32	64	5 6	34	30	45	37	
Leishmania-	В	2	53	1	22	193	1	7	383	111	2	0	0	6	9	1	2	1	299	16	331	255	56	114	110	22	0	1944	
Leish	4	0	0	0	7	6	0	0	27	П	0	0	0	П	П	0	0	0	14	0	13	7	3	10	0	7	0	82	
Meningitis	В	27	47	28	18	4	8	14	16	15	6	0	7	М	0	22	22	22	101	37	34	15	32	47	110	43	22	736	
Menir	A	0	7	Н	0	0	0	0	0	0	m	0	0	0	0	Н	0	7	m	Н	4	7	2	7	П	4	0	28	
xodu	В	172	170	283	162	32	84	214	96	172	120	13	1	19	12	49	24	38	320	80	156	23	113	20	112	263	47	2855	
Chickenpox	<	m	12	20	9	П	m	10	7	7	9	1	0	0	0	m	0	7	22	m	6	Н	9	0	2	7	7	131	
<u>=</u>	В	0	0	П	П	0	0	П	0	7	П	0	0	0	0	П	0	0	7	0	0	0	0	П	7	0	0	12	
Human	<	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
	В	3	11	2	က	3	4	П	_∞	3	7	0	0	1	П	2	1	0	6	П	က	12	89	17	13	4	0	178	
Viral	<	0	0	П	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0		0	4	0	П	П	0	œ	
<u>s</u>	В	0	7	1	41	12	49	30	23	20	478	9	2	8	2	П	П	14	6	œ	28	2	31	30	21	24	1	888	
Iyphus	⋖	0	0	0	2	₩	П	н	2	0	7	0	0	0	0	0	0	0	0	Н	0	0	П	0	3	7	0	16	
Leptospirosis	В	194	341	525	168	113	9/	571	207	382	8	7	30	27	53	29	22	26	241	38	224	131	221	403	758	452	35	5326	
Lepto	⋖	4	2	19	2	m	4	13	2	12	0	0	0	2	0	3	0	Н	7	7	9	7	10	7	23	20	0	12	
Poi-	В	7	3	9	15	8	40	21	8	12	17	16	0	0	12	18	0	64	9	н	2	9	32	0	14	10	0	318	
Food	<	0	0	П	m	0	0	0	0	0	Н	0	0	0	0	0	0	0	0	0	0	0	2	0	0	7	0	12	
Dysentery Encephalit Enteric Fever Food P	В	1	m	0	7	п	m	2	П	П	6	0	1	0	m	2	0	0	0	П	П	0	0	0	2	7	0	46	
Enteri	4	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	0	
ohalit	В	10	13	7	0	0	3	12	3	9	7	0	0	1	0	7	П	П	8	7	0	2	2	2	13	7	10	111	
Ence	<	0	0	н	0	0	Н	0	0	0	0	0	0	0	0	н	0	0			0	0	0	0	0	0	0	Ŋ	
entery	В	7	12	14	27	2	98	34	7	19	24	7	9	2	6	142	1	14	31	8	9	11	56	15	53	16	4	632	
Dyse	A	0	-	0	7	0	m	7	П	0	7	0	0	0	0	7	0	0	2	0	0	-	0	0	0	7	0	76	
Fever	В	9691	10108	3312	3938	936	157	1581	1081	1183	1701	9/	73	116	104	1970	114	1912	2170	2680	581	450	730	421	1491	2061	1557	50194	
Dengue Fever	⋖	304	308	128	797	09	10	95	41	62	38	0	0	н	11	23	1	28	26	27	70	9	30	27	43	84	19	1711	
RDHS		Colombo	Gampaha	Kalutara	Kandy	Matale	NuwaraEliya	Galle	Hambantota	Matara	Jaffna	Kilinochchi	Mannar	Vavuniya	Mullaitivu	Batticaloa	Ampara	Trincomalee	Kurunegala	Puttalam	Anuradhapur	Polonnaruwa	Badulla	Monaragala	Ratnapura	Kegalle	Kalmune	SRILANKA	

Source: Weekly Returns of Communicable Diseases (esurvillance.epid.gov.Ik). T=Timeliness refers to returns received on or before 14th July, 2023 Total number of reporting units 358 Number of reporting units data provided for the current week. B = Cumulative cases for the year.

Table 2: Vaccine-Preventable Diseases & AFP

08th-14th July 2023 (28th Week)

Disease	No.	of Ca	ases	by P	rovin	ıce		Number of cases during current week in	Number of cases during same week in	Total number of cases to date in	Total num- ber of cases to date in	Difference between the number of cases to date		
	W	С	S	N	Е	NW	NC	U	Sab	2023	2022	2023	2022	in 2023 & 2022
AFP*	00	01	00	00	00	00	00	00	00	01	01	50	44	13.6 %
Diphtheria	00	00	00	00	00	00	00	00	00	00	00	00	00	0 %
Mumps	03	00	00	00	00	01	03	00	01	08	00	121	35	245.7 %
Measles	07	00	00	06	00	00	01	00	01	15	01	61	14	335.7 %
Rubella	00	00	00	00	00	00	00	00	00	00	00	01	00	0 %
CRS**	00	00	00	00	00	00	00	00	00	00	00	00	00	0 %
Tetanus	00	00	00	00	00	00	00	00	00	00	00	06	05	20 %
Neonatal Tetanus	00	00	00	00	00	00	00	00	00	00	00	00	00	0 %
Japanese Encephalitis	00	00	00	00	00	00	00	00	00	00	00	02	07	- 71.4 %
Whooping Cough	00	00	00	00	00	00	00	00	00	00	00	05	01	400 %
Tuberculosis	110	28	27	21	09	18	13	08	20	254	84	5043	3134	60.9 %

Key to Table 1 & 2

Provinces: W: Western, C: Central, S: Southern, N: North, E: East, NC: North Central, NW: North Western, U: Uva, Sab: Sabaragamuwa.

RDHS Divisions: CB: Colombo, GM: Gampaha, KL: Kalutara, KD: Kandy, ML: Matale, NE: Nuwara Eliya, GL: Galle, HB: Hambantota, MT: Matara, JF: Jaffna,

KN: Killinochchi, MN: Mannar, VA: Vavuniya, MU: Mullaitivu, BT: Batticaloa, AM: Ampara, TR: Trincomalee, KM: Kalmunai, KR: Kurunegala, PU: Puttalam,

AP: Anuradhapura, PO: Polonnaruwa, BD: Badulla, MO: Moneragala, RP: Ratnapura, KG: Kegalle.

Data Sources:

Weekly Return of Communicable Diseases: Diphtheria, Measles, Tetanus, Neonatal Tetanus, Whooping Cough, Chickenpox, Meningitis, Mumps., Rubella, CRS,

Special Surveillance: AFP* (Acute Flaccid Paralysis), Japanese Encephalitis

CRS** =Congenital Rubella Syndrome

NA = Not Available

Influenza Surveillance in Sentinel Hospitals - ILI & SARI													
Month	Human		Animal										
	No Total	No Positive	Infl A	Infl B	Pooled samples	Serum Samples	Positives						
June													
Source: Medical Research Institute & Veterinary Research Institute													

Comments and contributions for publication in the WER Sri Lanka are welcome. However, the editor reserves the right to accept or reject items for publication. All correspondence should be mailed to The Editor, WER Sri Lanka, Epidemiological Unit, P.O. Box 1567, Colombo or sent by E-mail to chepid@sltnet.lk. Prior approval should be obtained from the Epidemiology Unit before publishing data in this publication

ON STATE SERVICE

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