



Human Metapneumovirus and Influenza Viruses in Children with Severe Acute Respiratory Infections in Iran

Jila Yavarian,¹ Nazanin Zahra Shafiei-Jandaghi,¹ Abbas Rahimi-Foroushani,² Azadeh Shadab,¹ Masoumeh Ghamarchehreh,¹ and Talat Mokhtari-Azad^{1,*}

¹Virology Department, School of Public Health, Tehran University of Medical Sciences, Tehran, IR Iran

²Epidemiology and Biostatistics Department, School of Public Health, Tehran University of Medical Sciences, Tehran, Iran

*Corresponding author: Talat Mokhtari-Azad, Virology Department, School of Public Health, Tehran University of Medical Sciences, Poursina St, POB: 14155-6446, Tehran, IR Iran. Tel/Fax: +98-2188962343, E-mail: mokhtari@tums.ac.ir

Received 2017 August 27; Revised 2018 January 07; Accepted 2018 January 24.

Abstract

Background: Influenza viruses and human metapneumovirus (HMPV) are among the common causes of acute respiratory tract infections in children worldwide.

Objectives: The aim of this study was to determine the frequency of HMPV and influenza viruses in children less than 5 years old hospitalized with severe acute respiratory infections (SARI) during 2013 - 2016.

Methods: For this study, 770 nasopharyngeal swabs from children with SARIs were tested for detection of influenza viruses by One step real time RT-PCR method in National Influenza Center, Tehran, Iran. Influenza virus negative samples were tested for HMPV using One step RT-PCR kit.

Results: Influenza viruses were detected in 263 of 770 specimens (34%) with influenza A/H1N1 being the prominent subtype. All 507 influenza negative samples were tested for HMPV. HMPV was detected in 7 (1.4%) cases.

Conclusion: In this research two different viruses were detected in 35% of cases, the remaining 65% of the patients might have been infected by other viruses or bacteria. Active surveillance studies with large sample size in different geographical regions and population based studies besides hospitalized patients with SARIs are needed to obtain more data about respiratory infections epidemiology.

Keywords: Influenza Viruses, Human Metapneumovirus, Severe Acute Respiratory Infections, Children, Iran

1. Background

Acute respiratory tract infections (ARTIs) are the most common illnesses worldwide. They account for considerable morbidity and mortality especially in young children and elderly. Viruses causing ARTIs include influenza viruses, parainfluenza viruses, respiratory syncytial virus (RSV), adenoviruses, coronaviruses, enteroviruses, rhinoviruses and human metapneumovirus (HMPV). Among these, influenza viruses within *Orthomyxoviridae* family are major pathogens of the lower respiratory tract infections (LRTIs) in all ages with significant rates of hospitalization (1). HMPV, a recently discovered virus, was identified in the Netherlands by van den Hoogen et al in 2001. HMPV is responsible for 5% - 8% of ARTIs in hospitalized children (2). It had been classified within *Paramyxoviridae* family, but according to the new taxonomy of the international committee on taxonomy of viruses (ICTV), now it is a member of *Pneumoviridae* family (3). In Iran, the influenza season starts in late November and lasts until late April, peak-

ing in January and February. The national influenza center (NIC) in Iran, located in Virology Department, School of Public Health, Tehran University of Medical Sciences, examines clinical samples from patients with severe acute respiratory infections (SARI) for influenza virus surveillance throughout the year. During influenza season many other respiratory viruses can cause ARTIs which usually share a common set of symptoms. HMPV is an emerging respiratory pathogen which can cause mild upper respiratory tract infections (URTIs) and severe LRTIs (2). Although there are many reports of HMPV prevalence worldwide, data from Iran is limited. The aim of this study was to describe the contribution of HMPV and influenza virus infections in hospitalized children less than five years of age with SARI during four consecutive years.

2. Methods

2.1. Clinical Samples

From January 2013 until December 2016 a total of 770 nasopharyngeal swabs from children less than five years of age with SARIs were sent in a cold chain system by Ministry of Health to NIC Iran. This study has been approved by Tehran University of Medical Sciences ethics committee.

SARI is an acute respiratory infection with history of fever $\geq 38^{\circ}\text{C}$ and cough with onset within the last 10 days which requires hospitalization. Of these specimens 161, 137, 299 and 173 were collected in 2013, 2014, 2015 and 2016 respectively (Table 1).

2.2. Molecular Analysis

Viral nucleic acids were extracted using high pure viral nucleic acid kit (Roche, Germany) according to the manufacturer's instructions. Influenza virus detection and typing/subtyping were carried out with real time RT-PCR kit, SuperScript III Platinum (Invitrogen) using CDC protocol (Table 2). After influenza screening all negative samples for influenza viruses, were tested for HMPV RNA using One step RT-PCR kit (Qiagen, Germany) with primers amplifying a conserved region of M gene. The sense and anti-sense primers are as follows: 5'-TGARTCAGCYACTGTTGAAGC-3' and 5'-TTTGGATTGTTTCATGGTCATRATC-3'. Negative and positive controls were used in each run. Plasmid containing a fragment of HMPV M gene was kindly denoted by Dr B. Khansari Nejad to use as a positive control.

3. Results

Of 770 children enrolled in this research, influenza viruses were detected in 263 cases (34%). Influenza A/H1N1pdm09, A/H3N2 and B viruses were detected in 120 (45.6%), 89 (33.9%) and 54 (20.5%) of positive samples respectively. Of 507 negative samples for influenza viruses, HMPV was positive in 7 (1.4%).

As shown in Table 1, in the year 2013, 44 (27.3%) of the specimens were positive for influenza viruses and the prominent type was influenza B and HMPV was not detected. During 2014, 36 (26.2%) and 1 (1%) of the specimens were positive for influenza and HMPV infections and the prominent influenza subtype was A/H3N2. In 2015, 101 (33.8%) and 1 (0.5%) influenza and HMPV viruses were detected respectively with A/H1N1 being prominent. In 2016, 82 of 173 (47.4%) specimens were positive for influenza viruses and A/H1N1 was the prominent subtype (54.9%). Five (5.5%) specimens were identified as positive for HMPV.

Of 770 patients, 459 (60%) were male and 311 (40%) female. Among positive children 97 (36%) were female and

173 (64%) male. Boys were more frequently infected by influenza viruses than the girls. HMPV was detected in four girls and three boys.

Incidence of influenza virus infections was higher in age group 0 - 1 year (46.8%). HMPV was detected in age group 3-5 years.

The peak prevalence of influenza was in December (n = 63, 24%), January (n = 63, 24%) and February (n = 56, 21.3%). The majority of HMPV detections were in January 2016 (5 out of 7), one positive sample was detected in June 2015 and one in December 2014.

4. Discussion

This paper showed the results of study of HMPV and influenza virus infections among hospitalized children less than 5 years of age with SARI during 2013 - 2016. In many studies detection rate of respiratory viruses are different which can be the result of different age groups, method of sampling and different detection methods. In our study distribution of HMPV, influenza A/H1N1, A/H3N2 and influenza B viruses was 1.4%, 15.6%, 11.5% and 7% respectively. Influenza A/H1N1 had the highest distribution. The reported incidence of influenza viruses and HMPV ranges from 1% to 40% in different studies as follows:

In a survey on 9274 respiratory samples for detection of HMPV and other viruses in west of Scotland, HMPV, influenza A and B viruses were positive in 2.2%, 4.5% and 3.7% of patients respectively (4). In a study in Pakistan, 169 children with WHO-defined severe pneumonia were tested for HMPV and influenza viruses' detection. HMPV and influenza A viruses were detected in 24 (14.2%) and 9 (5.3%) cases respectively (5). In a study in San Francisco in 2002, in 266 cases with ARTIs, 54 were positive for influenza A and B viruses and 4 were positive for HMPV infection (6). Of 380 specimens from influenza illness-like (ILI) patients in Greece, influenza viruses were detected in 151 (39.7%) patients and HMPV in 23 (6.05%) (7). Of 94 specimens from children less than 2 years of age hospitalized with bronchiolitis, 6 (6.4%) were positive for HMPV and 1 (1.5%) for influenza A viruses in a French study (8). In a study in the United States, of 668 specimens taken from inpatient children, 26 (3.9%) and 23 (3.4%) were positive for HMPV and influenza viruses respectively (9). In a study in Hong Kong, of 587 children with ARTIs, 5.5% were HMPV positive and 8% influenza positive (10). Of 7091 respiratory specimens collected in Scotland, 2% were HMPV positive and 2.5% and 1.3% were influenza A and B positive respectively (11). In a study in Netherlands, of 685 samples obtained from patients with ARTIs, 48, 11 and 7 were positive for HMPV, influenza A and B respectively (12). Of 4989 nasopharyngeal samples in Stockholm, 2.9% were positive for HMPV and

Table 1. Detection of Influenza Viruses and HMPV in Children Less Than Five Years with SARI in National Influenza Center, Tehran, Iran, 2013 - 2016^a

Year	SARI	Influenza Positive	A/H1N1	A/H3N2	Influenza B	HMPV
2013	161 (21)	44 (16.7)	9 (7.5)	14 (15.8)	21 (38.9)	0 (0)
2014	137 (17.8)	36 (13.7)	5 (4.1)	19 (21.3)	12 (22.2)	1 (14.3)
2015	299 (38.8)	101 (38.4)	61 (50.9)	20 (22.5)	20 (37)	1 (14.3)
2016	173 (22.4)	82 (31.2)	45 (37.5)	36 (40.4)	1 (1.9)	5 (71.4)
Total	770 (100)	263 (100)	120 (100)	89 (100)	54 (100)	7 (100)

Abbreviation: SARI, severe acute respiratory infection.

^aValues are expressed as No. (%).**Table 2.** Primers and Probes for Detection of Influenza Viruses According to CDC REF. #1-007-05^a

Primers and Probes	Sequence 5' - 3'
Influenza A/B detection	
InfA Forward	GAC CRA TCC TGT CAC CTC TGA C
InfA Reverse	AGG GCA TTY TGG ACA AAK CGT CTA
InfA Probe	FAM-TGC AGT CCT CGC TCA CTG GGC ACG-BHQ1
Influenza A/H1N1	
SW H1 Forward	GTG CTA TAA ACA CCA GCC TYC CA
SW H1 Reverse	CGG GAT ATT CCT TAA TCC TGT RGC
SW H1 Probe	FAM-CA GAA TAT ACA TaCC RGT CAC AAT TGG ARA A-BHQ1
Influenza A/H3N2	
AH3 Forward	AAG CAT TCC YAA TGA CAA ACC
AH3 Reverse	ATT GCR CCR AAT ATG CCT CTA GT
AH3 Probe	FAM- CAG GAT CAC ATA TGG GSC CTG TCC CAG-BHQ1

Abbreviation: SW, Swine.

^aTaqman® probes are labeled at the 5'-end with the reporter molecule 6-carboxyfluorescein (FAM) and quenched internally at a modified "T" residue with BHQ1.

14.8% for influenza A and 2.3% for influenza B viruses (13). In a Brazilian study, of 240 nasal washes of ILI patients, 49, 35 and 23 were positive for influenza A, B and HMPV respectively (14). In a study in Italy, of 1505 children younger than 15 years, HMPV was detected in 42 (2.8%) and influenza viruses in 230 (15.3%) (15). In a study by Boivin G et al. in 208 children less than 3 years old hospitalized with ARTIs, HMPV and influenza viruses were detected in 12 (6%) and 49 (24%) respectively (16). As shown above, incidence percentage of 34% for influenza viruses in this study was in agreement with previously reported studies; however, 1.4% incidence of HMPV was lower than the above and following studies.

In a similar study in Japan during 2002 - 2003 and 2003

- 2004 influenza seasons HMPV was positive in 5.7% and 5.2% of ILI patients respectively (17). In a study in Finland, HMPV was found in 7% of children with respiratory infections from 2000 to 2002 (18). In a survey on 1132 respiratory specimens in a pediatric center, 41 (3.6%) were HMPV positive especially during winter (19). In a study in Southern Brazil on 156 samples collected during the winter months from the patients less than 4 years, HMPV was detected in 10 (6.4%) samples (20). Of 220 children tested in Thailand between March 2001 and September 2003, 12 (5.4%) specimens were positive for HMPV (21). In a study of 296 specimens from children less than 5 years which were negative for other respiratory viruses, HMPV was detected in 19 (6.4%) in USA (22). In a study of 146 patients with respiratory illness, HMPV was detected in 5 (3.4%) in Rochester in 2004 (23).

However in our study the relatively low detection rate of HMPV is not unusual. They are different similar studies with low detection rate. In 200 nasopharyngeal aspirates specimens from children with RTIs which were negative for influenza viruses, 3 (1.5%) HMPV were detected in an Australian study (24). In a study in Germany, frequency of HMPV positivity was determined < 1% among children less than 3 years of age with ARTIs but 18% among those admitted in intensive care units (25). In a study by Scheltinga SA et al. on 239 patients, HMPV was detected in 6 (2.5%) during one year period from December 2001 - 2002 (26). In a multiyear study at US Midwestern Medical Referral Center, 34 (2.6%) of 1294 clinical specimens were positive for HMPV (27). Of 171 children with ARTIs, 4 (2.3%) were positive for HMPV in a UK study (28). In a study by Stockton et al, of 408 samples negative for influenza and RSV in patients with ILI, 9 (2.2%) were positive for HMPV (29). In a study in the Republic of Ireland on 171 bronchoalveolar lavage samples from adults and 122 specimens from children less than 5 years of age with RTIs, 2.4% of adults were positive for HMPV but no HMPV RNA was detected in samples from children (30). Of 400 samples collected in Finnish population, 4 (1.1%) were positive for HMPV during November 2007 to

June 2008 (31). In a ten year study of patients with ILI of different age groups, HMPV was detected in 3% (138/4549) of specimens in Germany (32). In an Iranian study during 2008 to 2009, of 202 specimens collected from children less than 6 years old, one (0.49%) HMPV was detected (33).

HMPV infection had temporal variation in our study with 5.5% in 2016, 0.0%, 1.0% and 0.5% during 2013 - 2015 respectively, the same as some other studies. In Italy during a 3 year study, HMPV positivity rate was 7% in 2001 in comparison to 40% in 2000 and 2002 (34). Another Italian study showed variation of HMPV infection from 1.5% to 7% in 1999 - 2000 and 2000 - 2001 respectively (35).

Some potential limitations in our research need to be addressed. First, we missed detection of coinfections with working only on influenza-negative specimens. Second, other groups of viruses such as RSV were not included by us because of limited sample amounts. Third, we did not have enough clinical and demographic information of the patients.

Finally, in our research two different viruses were detected in 35% of cases, the remaining 65% of the patients might have been infected by other viruses or bacteria, so more active surveillance studies with large sample size in different geographical regions and population based studies besides hospitalized patients with SARIs are needed to obtain more data about respiratory infections epidemiology.

Acknowledgments

We express our thanks to our colleagues in Virology Department, School of Public Health, Tehran University of Medical Sciences. We also say thanks to Dr B. Khansari Nejad for HMPV positive control. This study was supported by Tehran University of Medical Sciences, grant No.28370.

References

- Teodosieva A, Angelova S, Korsun N. Infections with influenza viruses, respiratory-syncytial virus and human metapneumovirus among hospitalized children aged ≤ 3 years in Bulgaria. *Trakia J Sci*. 2014;**12**(1):226-32.
- van den Hoogen BG, de Jong JC, Groen J, Kuiken T, de Groot R, Fouchier RA, et al. A newly discovered human pneumovirus isolated from young children with respiratory tract disease. *Nat Med*. 2001;**7**(6):719-24. doi: [10.1038/89098](https://doi.org/10.1038/89098). [PubMed: [11385510](https://pubmed.ncbi.nlm.nih.gov/11385510/)].
- Afonso CL, Amarasinghe GK, Banyai K, Bao Y, Basler CF, Bavari S, et al. Taxonomy of the order Mononegavirales: update 2016. *Arch Virol*. 2016;**161**(8):2351-60. doi: [10.1007/s00705-016-2880-1](https://doi.org/10.1007/s00705-016-2880-1). [PubMed: [27216929](https://pubmed.ncbi.nlm.nih.gov/27216929/)].
- Sivaprakasam V, Collins TC, Aitken C, Carman WF. Life-threatening human metapneumovirus infections in West of Scotland. *J Clin Virol*. 2007;**39**(3):234-7. doi: [10.1016/j.jcv.2007.03.011](https://doi.org/10.1016/j.jcv.2007.03.011). [PubMed: [17512777](https://pubmed.ncbi.nlm.nih.gov/17512777/)].
- Ali A, Khowaja AR, Bashir MZ, Aziz F, Mustafa S, Zaidi A. Role of human metapneumovirus, influenza A virus and respiratory syncytial virus in causing WHO-defined severe pneumonia in children in a developing country. *PLoS One*. 2013;**8**(9). e74756. doi: [10.1371/journal.pone.0074756](https://doi.org/10.1371/journal.pone.0074756). [PubMed: [24058625](https://pubmed.ncbi.nlm.nih.gov/24058625/)].
- Louie JK, Hacker JK, Gonzales R, Mark J, Maselli JH, Yagi S, et al. Characterization of viral agents causing acute respiratory infection in a San Francisco University Medical Center Clinic during the influenza season. *Clin Infect Dis*. 2005;**41**(6):822-8. doi: [10.1086/432800](https://doi.org/10.1086/432800). [PubMed: [16107980](https://pubmed.ncbi.nlm.nih.gov/16107980/)].
- Gioula G, Chatzidimitriou D, Melidou A, Exindari M, Kyriazopoulou-Dalaina V. Contribution of human metapneumovirus to influenza-like infections in North Greece, 2005-2008. *Euro Surveill*. 2010;**15**(9). [PubMed: [20214868](https://pubmed.ncbi.nlm.nih.gov/20214868/)].
- Bouscambert-Duchamp M, Lina B, Trompette A, Moret H, Motte J, Andreoletti L. Detection of human metapneumovirus RNA sequences in nasopharyngeal aspirates of young French children with acute bronchiolitis by real-time reverse transcriptase PCR and phylogenetic analysis. *J Clin Microbiol*. 2005;**43**(3):1411-4. doi: [10.1128/JCM.43.3.1411-1414.2005](https://doi.org/10.1128/JCM.43.3.1411-1414.2005). [PubMed: [15750120](https://pubmed.ncbi.nlm.nih.gov/15750120/)].
- Mullins JA, Erdman DD, Weinberg GA, Edwards K, Hall CB, Walker FJ, et al. Human metapneumovirus infection among children hospitalized with acute respiratory illness. *Emerg Infect Dis*. 2004;**10**(4):700-5. doi: [10.3201/eid1004.030555](https://doi.org/10.3201/eid1004.030555). [PubMed: [15200863](https://pubmed.ncbi.nlm.nih.gov/15200863/)].
- Peiris JS, Tang WH, Chan KH, Khong PL, Guan Y, Lau YL, et al. Children with respiratory disease associated with metapneumovirus in Hong Kong. *Emerg Infect Dis*. 2003;**9**(6):628-33. doi: [10.3201/eid0906.030009](https://doi.org/10.3201/eid0906.030009). [PubMed: [12781000](https://pubmed.ncbi.nlm.nih.gov/12781000/)].
- Gaunt E, McWilliam-Leitch EC, Templeton K, Simmonds P. Incidence, molecular epidemiology and clinical presentations of human metapneumovirus; assessment of its importance as a diagnostic screening target. *J Clin Virol*. 2009;**46**(4):318-24. doi: [10.1016/j.jcv.2009.09.016](https://doi.org/10.1016/j.jcv.2009.09.016). [PubMed: [19819755](https://pubmed.ncbi.nlm.nih.gov/19819755/)].
- van den Hoogen BG, van Doornum GJ, Fockens JC, Cornelissen JJ, Beyer WE, de Groot R, et al. Prevalence and clinical symptoms of human metapneumovirus infection in hospitalized patients. *J Infect Dis*. 2003;**188**(10):1571-7. doi: [10.1086/379200](https://doi.org/10.1086/379200). [PubMed: [14624384](https://pubmed.ncbi.nlm.nih.gov/14624384/)].
- Rafiefard F, Yun Z, Orvell C. Epidemiologic characteristics and seasonal distribution of human metapneumovirus infections in five epidemic seasons in Stockholm, Sweden, 2002-2006. *J Med Virol*. 2008;**80**(9):1631-8. doi: [10.1002/jmv.21242](https://doi.org/10.1002/jmv.21242). [PubMed: [18649344](https://pubmed.ncbi.nlm.nih.gov/18649344/)].
- Bellei N, Carraro E, Perosa A, Watanabe A, Arruda E, Granato C. Acute respiratory infection and influenza-like illness viral etiologies in Brazilian adults. *J Med Virol*. 2008;**80**(10):1824-7. doi: [10.1002/jmv.21295](https://doi.org/10.1002/jmv.21295). [PubMed: [18712837](https://pubmed.ncbi.nlm.nih.gov/18712837/)].
- Bosis S, Esposito S, Niesters HG, Crovari P, Osterhaus AD, Principi N. Impact of human metapneumovirus in childhood: comparison with respiratory syncytial virus and influenza viruses. *J Med Virol*. 2005;**75**(1):101-4. doi: [10.1002/jmv.20243](https://doi.org/10.1002/jmv.20243). [PubMed: [1543589](https://pubmed.ncbi.nlm.nih.gov/1543589/)].
- Boivin G, De Serres G, Cote S, Gilca R, Abed Y, Rochette L, et al. Human metapneumovirus infections in hospitalized children. *Emerg Infect Dis*. 2003;**9**(6):634-40. doi: [10.3201/eid0906.030017](https://doi.org/10.3201/eid0906.030017). [PubMed: [12781001](https://pubmed.ncbi.nlm.nih.gov/12781001/)].
- Sasaki A, Suzuki H, Saito R, Sato M, Sato I, Sano Y, et al. Prevalence of human metapneumovirus and influenza virus infections among Japanese children during two successive winters. *Pediatr Infect Dis J*. 2005;**24**(10):905-8. [PubMed: [16220090](https://pubmed.ncbi.nlm.nih.gov/16220090/)].
- Heikkinen T, Osterback R, Peltola V, Jartti T, Vainionpaa R. Human metapneumovirus infections in children. *Emerg Infect Dis*. 2008;**14**(1):101-6. doi: [10.3201/eid1401.070251](https://doi.org/10.3201/eid1401.070251). [PubMed: [18258088](https://pubmed.ncbi.nlm.nih.gov/18258088/)].
- Chano F, Rousseau C, Laferriere C, Couillard M, Charest H. Epidemiological survey of human metapneumovirus infection in a large pediatric tertiary care center. *J Clin Microbiol*. 2005;**43**(11):5520-5. doi: [10.1128/JCM.43.11.5520-5525.2005](https://doi.org/10.1128/JCM.43.11.5520-5525.2005). [PubMed: [16272480](https://pubmed.ncbi.nlm.nih.gov/16272480/)].
- do Carmo Debur M, Bordignon J, Duarte dos Santos CN, Vidal LR, Nogueira MB, de Almeida SM, et al. Acute respiratory infection by human metapneumovirus in children in southern Brazil. *J Clin Virol*. 2007;**39**(1):59-62. doi: [10.1016/j.jcv.2007.01.012](https://doi.org/10.1016/j.jcv.2007.01.012). [PubMed: [17350886](https://pubmed.ncbi.nlm.nih.gov/17350886/)].

21. Samransamruajkit R, Thanasugarn W, Prapphal N, Theamboonlers A, Poovorawan Y. Human metapneumovirus in infants and young children in Thailand with lower respiratory tract infections; molecular characteristics and clinical presentations. *J Infect.* 2006;**52**(4):254–63. doi: [10.1016/j.jinf.2005.07.001](https://doi.org/10.1016/j.jinf.2005.07.001). [PubMed: [16183133](https://pubmed.ncbi.nlm.nih.gov/16183133/)].
22. Esper F, Boucher D, Weibel C, Martinello RA, Kahn JS. Human metapneumovirus infection in the United States: clinical manifestations associated with a newly emerging respiratory infection in children. *Pediatrics.* 2003;**111**(6 Pt 1):1407–10. [PubMed: [12777560](https://pubmed.ncbi.nlm.nih.gov/12777560/)].
23. Falsey AR, Criddle MC, Walsh EE. Detection of respiratory syncytial virus and human metapneumovirus by reverse transcription polymerase chain reaction in adults with and without respiratory illness. *J Clin Virol.* 2006;**35**(1):46–50. doi: [10.1016/j.jcv.2005.04.004](https://doi.org/10.1016/j.jcv.2005.04.004). [PubMed: [15919236](https://pubmed.ncbi.nlm.nih.gov/15919236/)].
24. Nissen MD, Siebert DJ, Mackay IM, Sloots TP, Withers SJ. Evidence of human metapneumovirus in Australian children. *Med J Aust.* 2002;**176**(4):188. [PubMed: [11913922](https://pubmed.ncbi.nlm.nih.gov/11913922/)].
25. 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):4632–5. doi: [10.1128/JCM.42.10.4632-4635.2004](https://doi.org/10.1128/JCM.42.10.4632-4635.2004). [PubMed: [15472321](https://pubmed.ncbi.nlm.nih.gov/15472321/)].
26. Scheltinga SA, Templeton KE, Beersma MF, Claas EC. Diagnosis of human metapneumovirus and rhinovirus in patients with respiratory tract infections by an internally controlled multiplex real-time RNA PCR. *J Clin Virol.* 2005;**33**(4):306–11. doi: [10.1016/j.jcv.2004.08.021](https://doi.org/10.1016/j.jcv.2004.08.021). [PubMed: [15994117](https://pubmed.ncbi.nlm.nih.gov/15994117/)].
27. Gray GC, Capuano AW, Setterquist SF, Erdman DD, Nobbs ND, Abed Y, et al. Multi-year study of human metapneumovirus infection at a large US Midwestern Medical Referral Center. *J Clin Virol.* 2006;**37**(4):269–76. doi: [10.1016/j.jcv.2006.08.016](https://doi.org/10.1016/j.jcv.2006.08.016). [PubMed: [17008122](https://pubmed.ncbi.nlm.nih.gov/17008122/)].
28. Hopkins MJ, Redmond C, Shaw JM, Hart IJ, Hart CA, Smyth RL, et al. Detection and characterisation of human metapneumovirus from children with acute respiratory symptoms in north-west England, UK. *J Clin Virol.* 2008;**42**(3):273–9. doi: [10.1016/j.jcv.2008.03.011](https://doi.org/10.1016/j.jcv.2008.03.011). [PubMed: [18455957](https://pubmed.ncbi.nlm.nih.gov/18455957/)].
29. Stockton J, Stephenson I, Fleming D, Zambon M. Human metapneumovirus as a cause of community-acquired respiratory illness. *Emerg Infect Dis.* 2002;**8**(9):897–901. doi: [10.3201/eid0809.020084](https://doi.org/10.3201/eid0809.020084). [PubMed: [12194763](https://pubmed.ncbi.nlm.nih.gov/12194763/)].
30. Carr MJ, McCormack GP, Crowley B. Human metapneumovirus-associated respiratory tract infections in the Republic of Ireland during the influenza season of 2003–2004. *Clin Microbiol Infect.* 2005;**11**(5):366–71. doi: [10.1111/j.1469-0691.2005.01129.x](https://doi.org/10.1111/j.1469-0691.2005.01129.x). [PubMed: [15819862](https://pubmed.ncbi.nlm.nih.gov/15819862/)].
31. Jokela P, Piiparinen H, Luiro K, Lappalainen M. Detection of human metapneumovirus and respiratory syncytial virus by duplex real-time RT-PCR assay in comparison with direct fluorescent assay. *Clin Microbiol Infect.* 2010;**16**(10):1568–73. doi: [10.1111/j.1469-0691.2010.03191.x](https://doi.org/10.1111/j.1469-0691.2010.03191.x). [PubMed: [20156219](https://pubmed.ncbi.nlm.nih.gov/20156219/)].
32. Reiche J, Jacobsen S, Neubauer K, Hafemann S, Nitsche A, Milde J, et al. Human metapneumovirus: insights from a ten-year molecular and epidemiological analysis in Germany. *PLoS One.* 2014;**9**(2). e88342. doi: [10.1371/journal.pone.0088342](https://doi.org/10.1371/journal.pone.0088342). [PubMed: [24505479](https://pubmed.ncbi.nlm.nih.gov/24505479/)].
33. Shatizadeh Malekshahi S, Mokhtari Azad T, Yavarian J, Shahmahmoodi S, Naseri M, Rezaei F. Molecular Detection of Respiratory Viruses in Clinical Specimens From Children With Acute Respiratory Disease in Iran. *Pediatr Infect Dis J.* 2010;**29**(10):931–3. doi: [10.1097/INF.0b013e3181e2062e](https://doi.org/10.1097/INF.0b013e3181e2062e).
34. Falsey AR, Erdman D, Anderson LJ, Walsh EE. Human metapneumovirus infections in young and elderly adults. *J Infect Dis.* 2003;**187**(5):785–90. doi: [10.1086/367901](https://doi.org/10.1086/367901). [PubMed: [12599052](https://pubmed.ncbi.nlm.nih.gov/12599052/)].
35. Maggi F, Pifferi M, Vatteroni M, Fornai C, Tempestini E, Anzilotti S, et al. Human metapneumovirus associated with respiratory tract infections in a 3-year study of nasal swabs from infants in Italy. *J Clin Microbiol.* 2003;**41**(7):2987–91. [PubMed: [12843031](https://pubmed.ncbi.nlm.nih.gov/12843031/)].