Seroepidemiology of Human Metapneumovirus (hMPV) on the Basis of a Novel Enzyme-Linked Immunosorbent Assay Utilizing hMPV Fusion Protein Expressed in Recombinant Vesicular Stomatitis Virus

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Human metapneumovirus (hMPV) is a newly identified human respiratory virus now recognized as a major respiratory pathogen of infants and children. To define the seroepidemiology of hMPV, we developed a novel enzyme-linked immunosorbent assay (ELISA) based on expression of the fusion protein of hMPV (hMPV F) in recombinant vesicular stomatitis virus (VSV). Western blot analysis using an hMPV F-specific antiserum confirmed the expression of hMPV in recombinant VSV. The ELISA is specific for hMPV F; antibody specific for the most closely related human paramyxovirus, respiratory syncytial virus, does not bind to hMPV F. Overall, 216 serum specimens were tested. The percentages of seropositive individuals were 89.1% in children ≤5 months old, 55.0% in children 6 to 11 months old, 36.0% in children 12 to 23 months old, 45.0% in children 24 to 47 months old, 77.3% in children 48 to 59 months old, 91.3% in children 5 to 10 years old, and 95.5% for individuals 11 to 20 years old. This is the first seroepidemiological survey of hMPV in the United States and the first analysis to determine the prevalence of antibody to a specific hMPV protein. The data suggest that exposure to hMPV is common in childhood and that hMPV F is an antigenic determinant of hMPV.

In 2001, van den Hoogen et al. reported the discovery of a novel virus associated with respiratory tract disease in children. Genetic and phylogenetic analysis revealed that this new agent was the first human pathogen in the metapneumovirus genus of the Paramyxoviridae, and the virus was called human metapneumovirus (hMPV) (22). Since the initial report of the discovery of hMPV, the virus has been identified worldwide (2, 4, 6, 14, 25). The data from several studies suggest that hMPV is common and accounts for a significant proportion of respiratory tract disease in infants and children (7, 22, 26). hMPV has been circulating in the human population for at least 5 decades (22, 26).

The clinical manifestations associated with hMPV infection are similar to those observed with other respiratory viruses, in particular, respiratory syncytial virus (RSV). Upper respiratory tract infection and lower respiratory tract disease have been associated with hMPV infection. Coughing, wheezing, hypoxia and abnormalities on chest radiographs occur frequently in children infected with hMPV (3, 7, 24, 26). Asymptomatic infections are rare in young children (26). hMPV has also been detected in adults with clinical evidence of respiratory tract infection (8, 18).

Based on comparisons with other paramyxoviruses, the hMPV gene order and gene sequence suggest that the virus encodes three virion envelope glycoproteins, the fusion (F), attachment (G), and short hydrophobic glycoproteins. The F protein shares 33% identity with the immunodominant F protein of the most closely related human paramyxovirus RSV (21). Like the analogous protein (RSV F) in RSV, the hMPV F glycoprotein may elicit antibodies during infection. Reports from The Netherlands, United States, Canada, and elsewhere suggest that there are at least two major genotypes or clades of hMPV (1, 7, 22, 26).

Serological evidence indicates that hMPV infection is common in childhood. Reports from The Netherlands and Japan suggest that by the age of 5, nearly all individuals tested have evidence of exposure to hMPV (5, 22). The results of these studies were based on indirect immunofluorescence assays with hMPV-infected monkey kidney cells used as antigen. While the immunofluorescence assays provide some information, they have several limitations. The interpretation of immunofluorescence assays is based on subjective criteria, and these assays are prone to batch-to-batch variation. Furthermore, this assay cannot be used to study the antibody response to specific viral glycoproteins. To further study the seroepidemiology of hMPV and to determine the presence of antibody to a specific hMPV protein, we developed an enzyme-linked immunosorbent assay (ELISA) based on recombinant expression of the hMPV F glycoprotein. We chose the hMPV F protein because the corresponding protein (F) of the closely related paramyxovirus, RSV, is the immunodominant protein of the virus. To this end, we cloned the hMPV F gene from a clinical sample into recombinant vesicular stomatitis virus (VSV). This expression system was chosen for several reasons. Foreign glycoproteins are produced in relatively high amounts (~1% of total cellular protein) (17). Foreign viral glycoproteins are faithfully expressed, modified properly, and are biologically active (11, 12). This strongly suggests that these foreign viral glycoproteins maintain their antigenic configuration. This is essential for the
development of an ELISA specific for antibodies that bind hMPV F. VSV is a virus of livestock and horses; human infection is rare. The prevalence of antibodies to the VSV vector in the human population is low (20). Therefore, when recombinant VSV is used as an antigen for an ELISA, reactivity to the VSV vector is predicted to be low. Here, we describe a novel ELISA system to detect hMPV F-specific antibodies and to define the seroepidemiology of hMPV in Connecticut.

MATERIALS AND METHODS

Cloning of the hMPV F gene. The hMPV F gene was amplified from an hMPV-positive specimen identified previously (6). RNA from this hMPV-positive specimen was extracted by using a QiAamp viral RNA mini kit (QIAGEN Inc., Valencia, Calif.), and cDNA was produced by using Moloney murine leukemia virus reverse transcriptase (New England Biolabs, Beverly, Mass.) and random hexamer primers. The forward primer 5'-GGACGGCTTCTCGAGAGCCATGCTCITGGAAATGG and reverse primer 5'-CGGCTGACTCAATAATGTTGAGTATGAAAGCTCCATTGTTGTGTG were used for amplification of the hMPV F gene. These primers correspond to nucleotides 3052 to 3067 and 4641 to 4671 of the hMPV F gene (GenBank accession number AF371337). Restriction sites, C/G clamps, and an obligatory ACC (forward primer) required for VSV gene expression are underlined. The initiation codon ATG (forward primer) and termination codon (reverse codon, antisense strand) are shown in bold. Following digestion with the appropriate restriction endonucleases, the hMPV F gene was ligated into a plasmid containing the full-length genome of VSV (17). The ligation reactions were transformed into bacteria (Escherichia coli strain DH5α), and colonies containing the desired plasmid were confirmed by PCR and sequence analysis.

Recovery of recombinant VSV expressing hMPV F (VSV-hMPV F). Recovery of recombinant VSV was performed as previously described (11, 17). Briefly, individual plasmid-containing the VSV-N gene, the VSV-P gene, the VSV-L gene, and the full-length VSV genome containing the hMPV F gene were transfected into baby hamster kidney (BHK) cells (BHK-21; American Type Tissue Collection) infected with vaccinia virus (strain WR) expressing T7 RNA polymerase. Each of the VSV genes and the full-length VSV genome in the transfected plasmids were under T7 promoter control. Two days following the transfection, the infected cell supernatants were filtered (to remove vaccinia virus) and passed on BHK cells. Recovered recombinant VSV was identified by the presence of cytopathic effects. The recovery of a recombinant VSV containing the hMPV F gene was confirmed by reverse transcription-PCR (RT-PCR) and sequence analysis. The preparation of working stocks of VSV and VSV recombinants has been described previously (11).

hMPV F-specific antibody. A synthetic peptide, CQAAGSTVYPNEDKDE TRC-COOH, corresponding to amino acids 311 to 330 of the hMPV F protein (GenBank accession number AF371337), was synthesized at the W. M. Keck Biotechnology Resource Center, Yale University. To prepare antigens for animal inoculations, the synthetic peptide was first conjugated to maleimide-activated horseradish peroxidase-conjugated antibody (Jackson ImmunoResearch Laboratories, Easton, Pa.) and incubated overnight at 37°C. Following five washes with PT solution, bound secondary horseradish peroxidase-conjugated antibody (Jackson ImmunoResearch Laboratories) diluted in PGTF (1:20,000) was added to each well, and plates were incubated for 1 h at 37°C. After five washes with PT solution, bound secondary antibody was detected with 3,3′,5′,5′-tetramethyl benzidine (Pierce). Chromogenic reactions were stopped with 5 N H3PO4. Optical densities were read at 450 nm (OD450). Optical densities were normalized to the absorbance of 1.0 (OD450). Optical densities were normalized to the absorbance of 1.0 (OD450). Optical densities were normalized to the absorbance of 1.0 (OD450).

RESULTS

Cloning of the hMPV F gene into recombinant VSV. As part of an ongoing epidemiological study, hMPV-positive respiratory specimens were identified by RT-PCR and confirmed by sequence analysis. From one of these specimens, a clade B isolate (strain NH379) (7), the hMPV F gene was amplified by RT-PCR and cloned into a plasmid containing the full-length genome of VSV, and recombinant virus (VSV-hMPV F) was recovered by reverse genetics (17). The hMPV F gene was inserted into the N/P gene junction and contained the obligatory transcriptional regulatory elements required for faithful gene expression (Fig. 1A) (17). The level of gene expression in VSV is maximal at the 3′ (N) end of the genome and declines through each gene junction through the 5′ (L) end of the genome. Therefore, cloning the hMPV F gene between N and P would ensure relatively high expression in the recombinant virus (17).

The presence of the full-length, authentic copy of the hMPV F gene in the recombinant viral genome was confirmed by RT-PCR and sequence analysis of recovered virus (data not shown). To demonstrate that hMPV F was expressed in recombinant virus, a rabbit anti-hMPV F antiserum was developed. A synthetic peptide representing amino acids 311 to 330 of hMPV F was used as antigen to immunize the animal. This region of hMPV F was chosen for several reasons. This is a relatively hydrophilic domain and may be more antigenic than
hydrophobic domains. This region is not conserved among the fusion glycoproteins of paramyxoviruses, and, therefore, antibodies induced by this peptide should be specific for hMPV. Lastly, this region is conserved among isolates of the two major genotypes of hMPV. Binding of the rabbit anti-hMPV F peptide antibodies to both VSV-wt and VSV-hMPV F was assayed by Western blot analysis. Figure 1B shows the detection of two bands in the VSV-hMPV F-infected cell lysates that are not present in the VSV-wt-infected cell lysates. Based on the apparent molecular weights, these two bands likely represent the cleaved and uncleaved forms of hMPV F. The 58-kDa hMPV F protein, like the fusion protein of other paramyxoviruses, is presumably cleaved to generate a ~10-kDa N-terminal F2 and a ~46-kDa C-terminal F1 (a 2-kDa signal peptide at the N terminus is thought to be cleaved to form the mature protein). The synthetic peptide used in the rabbit immunization corresponds to a region in F1. No specific binding was detected with preimmune serum. Western blot analysis with a VSV-specific antiserum reveals that equal amounts of the viral vector were present in the lysates of both VSV-wt and VSV-hMPV F.

**hMPV F-specific ELISA based on recombinant VSV.** The VSV-hMPV F ELISA was used to measure hMPV-specific antibody titers in human serum. Although the prevalence of antibody to the VSV vector in the human population was considered to be very low, specific or nonspecific binding to the VSV antigens needed to be controlled for in the ELISA analysis. Therefore, each serum tested was assayed for binding to both VSV-wt and VSV-hMPV F lysates. For a vast majority (>95%) of human serum samples tested, the OD_{450} (at a dilution of 1:32) was <0.2 for binding to the VSV-wt lysate, and therefore a cutoff of 0.2 was used as one of the criteria for a positive ELISA for hMPV. For ≤5% of the serum samples tested, the OD_{450} was ≥0.2 for binding to the VSV-wt lysate. However, the OD_{450} for the VSV-hMPV F ELISA had to be greater than the VSV value at a fourfold-lower dilution for the sample to be considered positive for hMPV. The lysates used as antigen were standardized for protein concentration and binding of an anti-VSV antiserum. With this standardization, the assay is highly reproducible. The reproducibility of the assay was confirmed by using the anti-hMPV F peptide antibody (data not shown). ELISA results for both a representative seropositive and a representative seronegative individual are
shown in Fig. 2A and B, respectively. Assays using serum or plasma gave identical results (data not shown).

A potential shortcoming of an hMPV F ELISA is that this method may detect antibodies that bind to RSV F, thereby limiting the specificity of this assay. RSV is the most closely related human paramyxovirus, and the hMPV F and RSV F share 33% homology (21). Therefore, the potential binding of RSV-specific antibodies to the VSV-hMPV F lysates was evaluated in two ways. First, the binding of an RSV-specific antibody to VSV-hMPV F was assayed. As shown in Fig. 3A, a goat anti-RSV antiserum had a high titer of RSV-specific antibody but failed to bind VSV-hMPV F. This evidence suggested that RSV-specific antibodies do not cross-react with hMPV F. Second, serum in which RSV antibodies were subtracted was assayed for VSV-hMPV F-specific binding. The results of ELISAs in which RSV lysates were used as antigen are shown in Fig. 3B. Serum incubated with RSV lysates (Fig. 3B, sera + RSV Hep-2 lysate) had dramatically reduced binding to RSV antigens compared with serum incubated with Hep-2 lysates (Fig. 3B). However, when the same two incubated sera were assayed for VSV-hMPV F binding, no differences in the ELISA profiles were observed (Fig. 3C). These findings suggest that this ELISA is specific for hMPV antibodies.

Seroprevalence of hMPV-specific antibodies. The VSV-wt and VSV-hMPV F ELISAs were used to measure the prevalence of hMPV-specific antibodies in the pediatric population. As presented in Fig. 4, the percentage of children of ≤5 months old with hMPV-specific antibody is 89.1% (95% confidence interval [CI], 85.2 to 93.0%). The percentage of seropositive individuals drops to 55.0% (95% CI, 43.9 to 66.1%) in children 6 to 11 months of age and to 36.0% (95% CI, 26.4 to 45.6%) in children 12 to 23 months of age. The percentage of seropositive children 24 to 35 months old and 36 to 47 months old is 45.0% (95% CI, 33.9 to 56.1%) in both age groups. The percentage of seropositive individuals is 77.3% (95% CI, 68.3 to 86.2%) for children 48 to 59 months old, 91.3% (95% CI, 85.4 to 97.2%) for children 5 to 10 years old, and 95.5% (95% CI, 91.0 to 99.9%) for individuals 11 to 20 years old. Phylogenetic analysis based on the F gene of hMPV isolates from our group and others suggests that at least two distinct genotypes of hMPV circulate in the community (7, 22). Whether these genotypes represent distinct serotypes is unknown. To address this issue, we cloned the F gene from a clade A isolate (strain NH539) (7) into the VSV expression vector and tested a subset of serum samples for binding antibody by ELISA. (The ELISA results presented above were based on the F gene of a clade B isolate.) Eleven sera that tested positive by ELISA for the clade B hMPV F also tested positive for the clade A hMPV F. Likewise, nine sera that tested negative for the clade B hMPV also tested negative in the clade A hMPV ELISA (data not shown). There was no quantitative difference in the ODs for binding of antibody to the VSV-hMPV F (A) lysates and the VSV-hMPV F (B) lysates for any given serum.

In order to confirm the sensitivity of this recombinant antigen ELISA in detecting hMPV-positive sera, we developed an ELISA based on hMPV-infected (MK2-LLC) cell lysates. Uninfected cells were used as control antigens. The hMPV-infected cell ELISA detected anti-hMPV F peptide antibodies and hMPV antibodies in sera that tested positive with the recombinant antigen ELISA. Of 20 serum specimens that tested negative for hMPV F-specific antibodies (using the recombinant VSV-hMPV F antigen ELISA), all tested negative in the hMPV-infected cell ELISA, thus establishing the sensitivity of the recombinant antigen ELISA.

**DISCUSSION**

This is the first report of the seroepidemiology of hMPV in the United States and the first investigation to examine the prevalence of antibody to a specific hMPV glycoprotein. Infection with hMPV is common in our population. More than 90% of individuals of ≤5 years of age had evidence of exposure to hMPV. The dynamics of the seroepidemiology of hMPV in the pediatric population is complex. The decline in the proportion of seropositive individuals during the first year of life likely represents waning maternally acquired antibody, as observed with RSV (9). However, this decline is superimposed on an increase in antibody as results of infection during the first...
In our previous study of hMPV infection in children under 5 years old, a majority (60.4%) of hMPV-infected children were under 1 year of age (7). However, this prior study was not population based but used respiratory specimens from the hospital’s Clinical Virology Laboratory. Consequently, the true incidence of hMPV infection in children under 1 year of age remains to be determined.

There appear to be two periods of acquisition of hMPV infection in childhood. The first period occurs within the first 3 years of life. The percentage of individuals who are seropositive is essentially constant at 35 to 45% for children 12 to 47 months of age. The second period occurs in children of 48 months of age. The percentage of seropositive individuals increases to 77.3% (95% CI, 68.3 to 86.2%) in children 48 to 59 months old and to >90% in children who are >5 years old. This second peak likely reflects increased exposure to the virus, perhaps at day care or preschool environments. Our findings are similar to those reported elsewhere. In the initial report of the discovery of hMPV, van den Hoogen et al., using an immunofluorescence assay and a viral neutralization assay, found that by the age of 5 years, nearly all individuals have evidence of hMPV infection (22). Of 40 Israeli children ≥24 months old, 52% had serological evidence of hMPV infection (27). The percentage of seropositive individuals in a Japanese cohort was >90% in those ≥5 years old (5). However, only 17% (3 of 17) of children 6 months to 1 year of age had serological evidence of hMPV infection, whereas in our study the percentage of seropositive children 6 to 11 months old was 55.0% (95% CI, 43.9 to 66.1%). This difference may be explained by variation in the circulation of hMPV or by the sensitivities of the methodologies used to detect hMPV antibodies. In the Japanese study, as well as The Netherlands study, hMPV-specific antibodies were detected by assays using hMPV-infected monkey kidney cells. These methods require culture of primary monkey kidney cells and are subject to batch-to-batch variability.

The ELISA-based system has many distinct advantages over the methods used in prior studies. The amount of hMPV-specific antigen can be standardized for each assay, antibody to genotype-specific viral glycoproteins can be measured, and the results are based on defined criteria rather than subjective determinations of a positive result in an immunofluorescence study. Expression of hMPV F in recombinant VSV was validated by Western blot analysis by using a rabbit serum raised against a peptide corresponding to a hydrophilic domain of the C-terminal F1. The two bands detected by Western blotting are most likely cleaved and uncleaved forms of hMPV F. An understanding of the biochemical properties of hMPV F during infection is lacking, and therefore the relative amounts of the cleaved and uncleaved forms in hMPV-infected cells remain unknown. Our findings support the hypothesis that hMPV F is an antigenic virion glycoprotein.

The specificity and sensitivity of our ELISA were established. Antibodies to the F protein of the closely related para-
myxovirus RSV were not detected with the recombinant hMPV F ELISA. The sensitivity of our assay was confirmed by using a whole-virus ELISA, based on hMPV-infected cell lysates, as a gold standard. Assays to detect virus-specific antibody are frequently based on immunodominant glycoprotein(s). In fact, a peptide-based enzyme immunoassay for detection of antibodies to the G protein of RSV is more sensitive than conventional serological tests (13). This indicates that the VSV-hMPV F-based ELISA is an ideal tool to screen for hMPV exposure and infection.

Several groups have determined that at least two major genotypes of hMPV circulate concurrently in the community (2, 7, 22, 26), and animal studies suggest that the two genotypes may represent distinct serotypes (23). The 100% concordance between the results of the hMPV F clade A ELISA and the hMPV clade B ELISA suggests that there is significant antigenic cross-reactivity between the F glycoprotein of the two clades. This is not surprising as the hMPV F proteins of clade A and clade B are >90% identical (unpublished data). This also suggests that the two putative serotypes cannot be distinguished based on the fusion glycoprotein.

The present study also demonstrates the usefulness of the VSV expression system for the development of diagnostic reagents. An additional advantage of the VSV expression system is that, in general, the prevalence of antibodies specific for the VSV vector in the human population is low. VSV is a virus that infects animals, mainly livestock and horses, and human infection with VSV is, for the most part, rare. However, there are regions in Central and South America where VSV infection is endemic in the human population. This could potentially limit the utility of the system for individuals from those areas (19). We controlled for the possibility that some individuals may have antibodies that bind to the viral vector. Each serum specimen was screened for antibodies for hMPV F and the VSV vector. VSV is considered a controlled organism by the United States Department of Agriculture and is considered a foreign animal disease by many countries throughout the world. Any use of the virus should include the necessary precautions.

In conclusion, our results indicate that infection with hMPV is common. Most individuals are likely infected during childhood. Specific antibodies to hMPV F are transmitted transplacentally and also develop after infection with hMPV. Further studies are required to determine the immunogenic domains of hMPV F and whether hMPV F-specific antibodies are protective.

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