

# Respiratory infections: which pathogens should we include and when?

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ECCMID Berlin

27.04. 2013



# Aetiology of lower respiratory tract infection in the community (%)

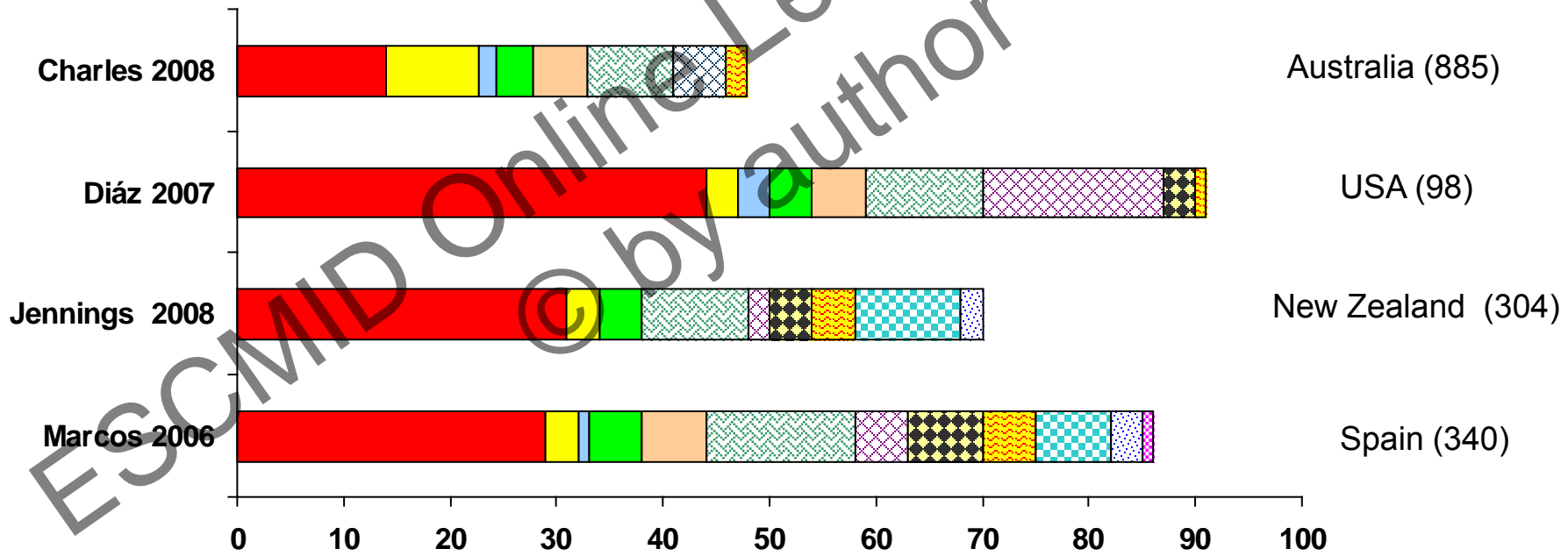
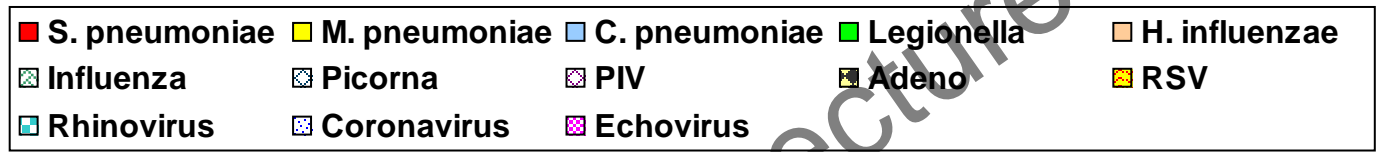


| Reference              | <i>n</i> | <i>S. pn</i> | <i>H. infl</i> | <i>M. pn</i> | <i>C. pn</i> | Virus       |
|------------------------|----------|--------------|----------------|--------------|--------------|-------------|
| Boldy et al. 1990      | 42       | 3.0          | 3.0            | 8.0          | 0            | 21.0        |
| Creer et al. 2006      | 80       | 18.8         | 6.3            | 1.2          |              | 61.3        |
| Graffelman et al. 2004 | 145      | 6.2          | 9.0            | 9.0          | 1.3          | 39.0        |
| Holm et al. 2007       | 364      | 6            | 4              | 3            | <1           | 24          |
| Hopstaken et al. 2005  | 247      | 2.9          | 13.8           |              |              |             |
| Macfarlane et al. 1993 | 206      | 30.0         | 8.0            | 0.5          |              | 8.0         |
| Macfarlane et al. 2001 | 316      | 17.1         | 9.8            | 7.3          | 17.4         | 19.3        |
| GRACE study, 2012      | 3059     | 9.1          | 14.8           | 2.9          | 2.2          | 51.1        |
| <b>Range</b>           |          | <b>3-30</b>  | <b>3-15</b>    | <b>0.5-9</b> | <b>0-17</b>  | <b>8-61</b> |

➔ ***C. pn* is in some studies reported in a large nr of cases: 0-20%**

➔ **Early data are largely based on serological analysis only**

# VD Viral Agents in CAP in Hospitalised Adult Patients



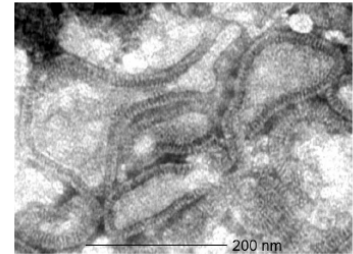


# A newly discovered human pneumovirus isolated from young children with respiratory tract disease

2001: hMPV

BERNADETTE G. VAN DEN HOOGEN<sup>1</sup>, JAN C. DE JONG<sup>1</sup>, JAN GROEN<sup>1</sup>, THijs KUIKEN<sup>1</sup>, RONALD GROOT<sup>2</sup>, RON A.M. FOUCHER<sup>1</sup> & ALBERT D.M.E. OSTERHAUS<sup>1</sup>

The NEW ENGLAND JOURNAL of MEDICINE

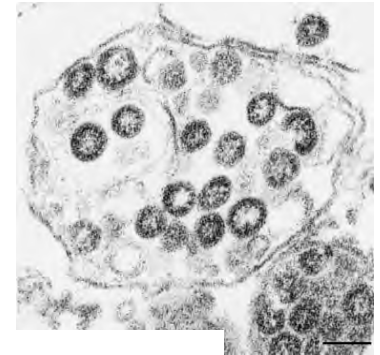


ORIGINAL ARTICLE

## Identification of a Novel Coronavirus in Patients with Severe Acute Respiratory Syndrome

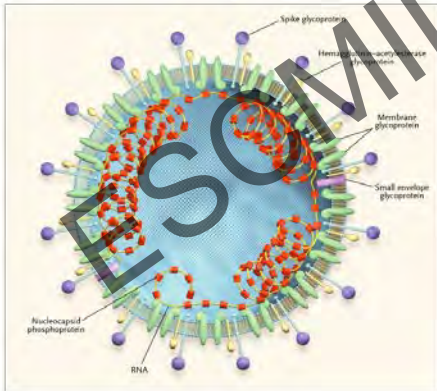
Christian Drosten, M.D., Stephan Günther, M.D., Wolfgang Preiser, M.D., Sylvie van der Werf, Ph.D., Hans-Reinhard Brodt, M.D., Stephan Becker, Ph.D., Holger Rabenau, Ph.D., Marcus Panning, M.D., Larissa Kolesnikova, Ph.D., Ron A.M. Fouchier, Ph.D., Annetmarie Berger, Ph.D., Ana-Maria Burguière, Ph.D., Jindrich Cinat, Ph.D., Markus Eickmann, Ph.D., Nicolas Escriou, Ph.D., Klaus Grywna, M.Sc., Stefanie Kramme, M.D., Jean-Claude Manuguerra, Ph.D., Stefanie Müller, M.Sc., Volker Rickerts, M.D., Martin Stürmer, Ph.D., Simon Vieth

2003: HCoV SARS



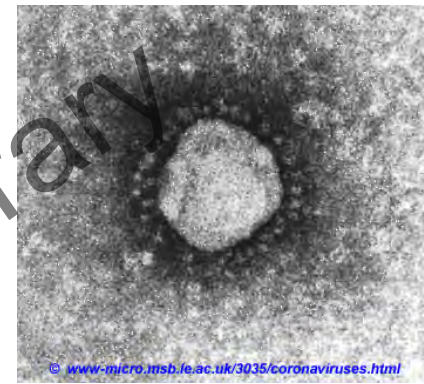
nature  
medicine

2005: HCoV NL 63



## Identification of a new human coronavirus

Lia van der Hoek<sup>1</sup>, Krzysztof Pyrc<sup>1</sup>, Maarten F Jebbink<sup>1</sup>, Wilma Vermeulen-Oost<sup>2</sup>, Ron J M Berkhout<sup>2</sup>, Katja C Wolthers<sup>1</sup>, Pauline M E Wertheim-van Dillen<sup>3</sup>, Jos Kaandorp<sup>4</sup>, Joke Spaargaren<sup>2</sup> & Ben Berkhout<sup>1</sup>



## Characterization and Complete Genome Sequence of a Novel Coronavirus, Coronavirus HKU1, from Patients with Pneumonia

Patrick C. Y. Woo,<sup>1,2†</sup> Susanna K. P. Lau,<sup>1,2†</sup> Chung-ming Chu,<sup>3</sup> Kwok-hung Chan,<sup>1</sup>  
Hoi-wah Tsoi,<sup>1</sup> Yi Huang,<sup>1</sup> Beatrice H. L. Wong,<sup>1</sup> Rosana W. S. Poon,<sup>1</sup>  
James J. Cai,<sup>1</sup> Wei-kwang Luk,<sup>4</sup> Leo L. M. Poon,<sup>1,2</sup> Samson S. Y. Wong,<sup>1,2</sup>  
Yi Guan,<sup>1,2</sup> J. S. Malik Peiris,<sup>1,2</sup> and Kwok-yung Yuen<sup>1,2†\*</sup>

2005: HCoV HKU1

## Cloning of a human parvovirus by molecular screening of respiratory tract samples

Tobias Allander<sup>\*†‡</sup>, Martti T. Tammi<sup>§1</sup>, Margareta Eriksson<sup>||</sup>, Annelie Bjerkner<sup>\*</sup>, Annika Tiveljung-Lindell<sup>\*</sup>,  
and Björn Andersson<sup>§</sup>

2005: Boca virus

**Journal of Clinical Virology**



Volume 38, Issue 3, March 2007, Pages 227–237

2007: rhinovirus C

doi:10.1016/j.jcv.2006.12.016  Cite or Link Using DOI

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## Molecular characterization of a variant rhinovirus from an outbreak associated with uncommonly high mortality

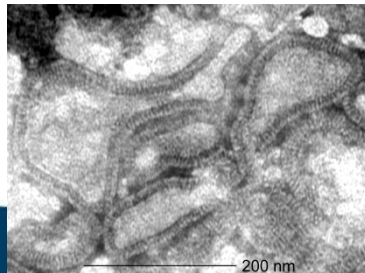
David Kiang<sup>a</sup>,  , Shigeo Yagi<sup>a</sup>, Katherine A. Kantardjieff<sup>b</sup>, Euna J. Kim<sup>b</sup>, Janice K. Louie<sup>a</sup>  
and David P. Schnurr<sup>a</sup>



# VD The main respiratory targets in molecular diagnostic tests

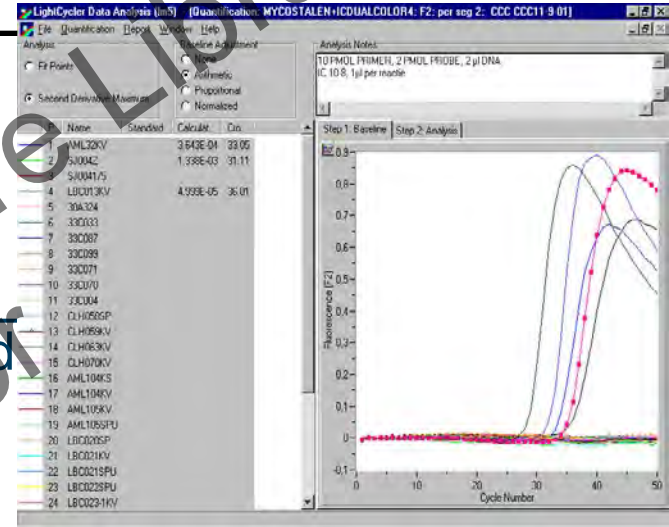
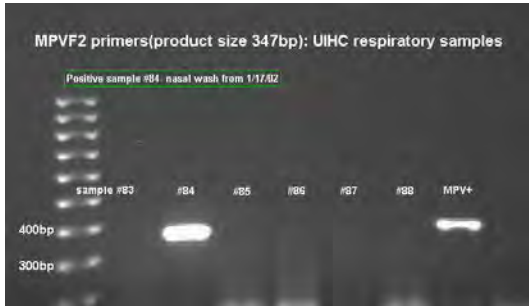


- Those in our routine panel:
  - Influenza A (IFVA)
  - Influenza B (IFVB)
  - Parainfluenza (PIV) 1-4
  - Respiratory syncytial virus (RSV)
  - Adenoviruses (ADV)
  - Metapneumovirus (hMPV)
- Those extra assays that many would consider important:
  - Rhinoviruses
  - Enteroviruses
  - Coronaviruses (OC43, 229E, NL63 and HKU1)
  - IFVA sub-typing
  - Bocavirus
  - Atypicals: *M.pn.*, *C. pn.*, *Leg. pn.*, *Bordetella pertussis*





# Conventional and Real-Time Mono- and Multiplex NAAT



| Author                 | targets | Species detected  |
|------------------------|---------|---|
| Fan, J et al. 1998     | 2       | RSVA, RSVB  |
| Scheltinga et al. 2005 | 2       | hMPN, RHI   |
| McDonough et al. 2005  | 4       | <i>M. pn.</i> , <i>C. pn.</i> , <i>L. pn.</i> , <i>B. pertussis</i>   |
| Gunson et al. 2005     | 12      | IFL A and B, PFL 1,2,3 RHI, hMPN<br>RSVA and B, COR E229, OC 43,<br>NL63 in 4 triplex reactions                                 |
| Loens et al. 2007      | 3       | <i>M. pn.</i> , <i>C. pn.</i> , <i>L. pn</i>  |
| Choi et al. 2006       | 12      | in 4 multiplex and one monoreaction   |
| Tiveljung et al. 2009  | 16      | in 13 reactions: IFL A and B, RSV A+B,<br>PFL 1+3, PFL 2+ hCoV-229E, ADE, hMPV, RHI,<br>ENT, HCoV-OC43, HCoV-NL63 and HKU, HBoV |

# Commercially available Mono- and Multiplex tests



|                               | targets | Species detected  |
|-------------------------------|---------|---|
| Xpert FluA, Cepheid           | 2       | Influenza A and subtyping   |
| RSV,ASR, Cepheid              | 2       | RSVA, RSVB  |
| ProPneumo-1, Prodesse         | 2       | <i>M. pneumoniae</i> , <i>C. pneumoniae</i>   |
| RespiFinder plus, Pathofinder | 18      | IFL A/B, PFL 1-4, RHI, hMPN,<br>RSV A/B, AV, 3 coronaviruses,<br><i>M. pn.</i> , <i>C.pn.</i> , <i>L.pn.</i> , <i>Bordetella pertussis</i>                          |
| SeeplexRV, Seegene            | 19      | <i>S. pneumoniae</i> , <i>H. influenzae</i> , <i>M. pn.</i> , <i>C.pn.</i> ,<br><i>L.pn.</i> , IFL A and B, RSV A/B, PFL 1-3,<br>RHI, 3 coronaviruses, AV, HBoV, EV |
| xTAG RVP, Luminex             | 19      | IFL A ( H1, H3, H5, non-specific ) and B,<br>PFL 1-4, RSV A/B, ADE, hMPV,<br>RHI/ENT,SARS-COR, HCoV OC43, HCoV<br>229E, HCoV NL63 and HKU1                          |





# Some commercially Multiplex tests

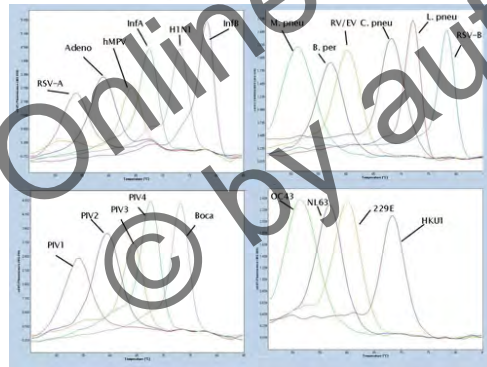


## Cepheid GeneXpert



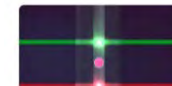
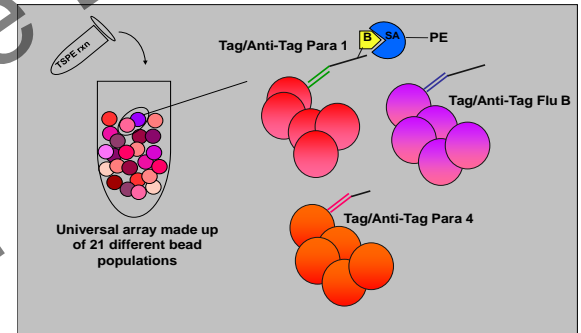
Single use  
cartridge based  
Up to 6 targets  
TAT 1.5 to 2/5 hr

## RespiFinder Mx, Pathofinder



Mx amplification  
and detection by  
melting curve  
analysis  
Up to 25 targets  
TAT 6 hr

## Luminex xTAG Universal Bead Array



Liquid microarray based  
Up to 20 targets  
RVP 10-12 hr to RVP Fast 4-5  
hr



# Comparison of some mono- and Mx methods for detection resp pathogens



| Parameter                                  | Xpert Flu,                     | RSV Respifinder     | xTAG RVP                 |
|--|--------------------------------|---------------------|--------------------------|
| Amplification platform                     | Multiplex<br>RT-PCR            | Multiplex<br>RT-PCR | Multiplex<br>RT-PCR      |
| Detection format                           | GeneXpert<br>(solid cartridge) | (real-time)         | Luminex<br>(liquid chip) |
| Nr pathogens covered                       | 5                              | 22                  | 19                       |
| No. of reactions needed                    | 2                              | 1                   | 1                        |
| Hands-on time (min) <sup>a</sup>           | 10                             | 70                  | 55                       |
| Detection throughput<br>(no. of tests/run) | 1                              | 48                  | 96                       |
| Test turnaround time (h) <sup>b</sup>      | 1.5-2                          | 6                   | 10-12                    |

<sup>a</sup> Hands-on and turnaround times include time needed for specimen processing and data analysis

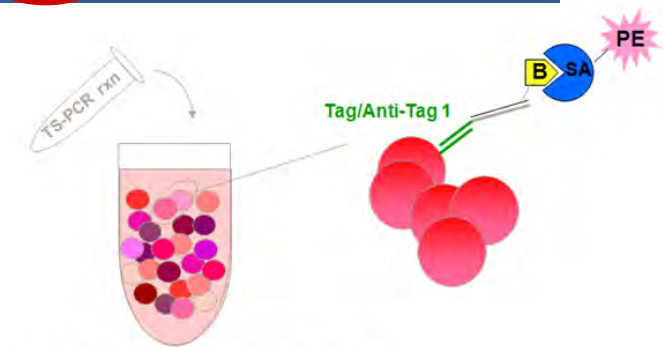
<sup>b</sup> Times are based on a full run, including specimen processing, target amplification, and product identification

# VD Sensitivity and specificity of two different commercial assays

- Evaluation of 2 CE labeled multiplex assays: RespiFinder (Pathofinder) and xTAG RVP (Luminex) on
- Compared with in-house PCRs used in GRACE

| Virus | RespiFinder |             | xTAG RVP    |             |
|-------|-------------|-------------|-------------|-------------|
|       | Sensitivity | Specificity | Sensitivity | Specificity |
| INF   | 84.8%       | 98.4        | 72.3        | 96.8        |
| HCoV  | 89.1        | 99.0        | 32.6        | 98.7        |
| hMPV  | 100         | 98.5        | 96.6        | 99.4        |
| HRV   | 95.2        | 98.3        | 88.7        | 99.6        |
| RSV   | 88.5        | 100         | 71.4        | 99.1        |

- Both commercial assays are less sensitive than in-house PCRs used in GRACE
- RespiFinder (Pathofinder) is more sensitive than Luminex assay



Universal array made up of n different bead populations with reporter (SA-PE) present



**When to use which techniques?  
Sequential approach or Mx detection?**

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# VD Some Technical, Scientific and practical hurdles

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- For number of newest technologies: limited validation and little proven clinical applications
- Few FDA cleared
- Which pathogen?
- Which clinical specimen?
- Sample preparation
- Some methodologies: low throughput tests
- Some methodologies: need for multiple instruments
- Technically demanding and labor intensive
- Costly





# Limited target detection versus multiplex detection



## Limited target detection

- Usually ↑ analytical sens.
- Lower cost
- Often lower TAT
- In outbreak situations
  - SARS Coronavirus
  - Influenza, H1N1
  - RSV, *L. pn.*, *M. pn.*
- As first approach
  - in high prevalence periods
  - if therapeutic implications
    - Influenza, *Legionella* spp, *Mycoplasma pn.*, *B. pertussis*
- Outside normal lab working hrs

## Multiplex detection

- In >90% similar results
- Expensive
- TAT usually > 4-6hours
- For epidemiological studies
  - Prevalence of respiratory etiologies
  - Role of respiratory viruses
- As add-on diagnostic test
  - In severely ill patients
  - In immunocompromised
- For virus discovery studies



# Impact of molecular methods on the diagnosis of LRTI



## Impact on Influenza H1N1 outbreak management

- Semiautomated and fully automated assays for detection of FluA (including H1N1) and FluB:

Flu A sensitivities varying between 93% - 98%,  
specificities varying between 99 -100%

Sails AD et al. J Virol Methods 2009, 162: 88-90  
Ginocchio CC et al J Clin Virol 2009; 45: 191-195  
Casalegno JS et al Clin Microbiol Infect 2009; 16: 326-329  
Beck ET, et al J Mol Diagnostics 2010; 12: 74-81

Automated **high throughput molecular system** allows clinicians and public health officials to **react quickly during outbreaks**



# Impact on patient management: Need for speed! TAT is crucial!



- Evaluation of 2 real-time RT-PCR assays ( Cepheid, Sunnyvale, USA) on SmartCycler (**TAT < 1hour**)
  - RSV Analyte Specific Reagent (ASR) bead
  - Influenza A/B ASR bead
- Comparison with “in-house” multiplex real-time PCR on +/-100 respiratory samples for Influenza and RSV
- Results: facilitates urgent testing outside batched runs or
  - RSV: sens: 98.2%, spec: 100%
  - Influenza A/B: sens: 96.5%, spec: 100%

➔ **Compared to “in house” multiplex: significant in ↓ TAT**

➔ **facilitates urgent testing outside batched runs or normal working day**

# VD RSV outbreak: an argument for real-time detection!



- Evaluation of rapid antigen testing, culture DFA and RT-PCR in symptomatic residents in a 120-bed LTCF
- In 32% (7/22) cases: RSV was detected by RT-PCR
  - None of cases was detected by Antigen testing
  - Only 2 cases (9%) were detected by culture or DFA

→ RT-PCR can provide a **more timely and accurate** diagnosis of outbreaks, which **allows for early symptomatic treatment, rational use of antibiotics, and improved infection control.**

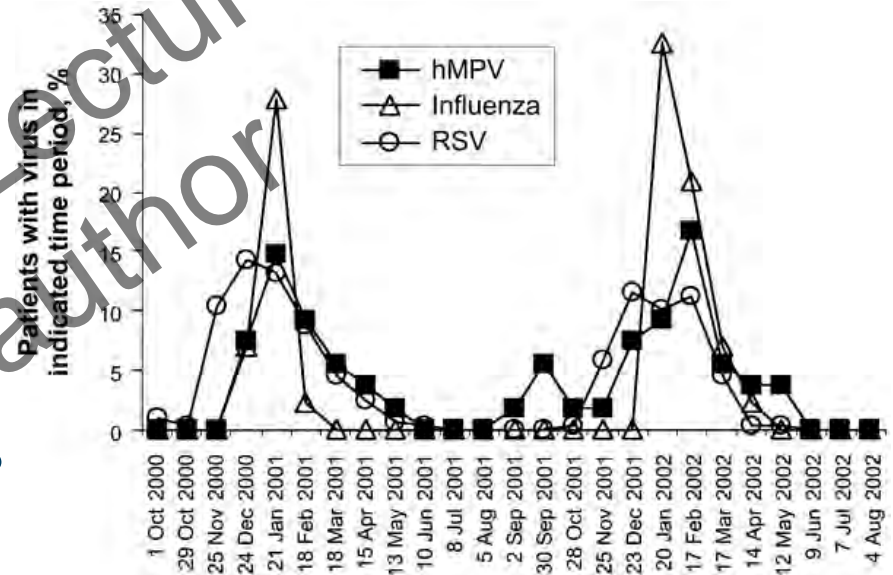
Caram BL et al. J Am Geriatr Soc 2009; 57: 482-485



# hMPV resembling RSV: Similar but Different?



- RSV: more common than hMPV in infants <6 mo.
- hMPV similar to RSV, majority of hMPV cases occur in young (<5yrs)
- Seasonality with RSV: hMPV later



- co-infection with RSV: More severe?  
Contradictory

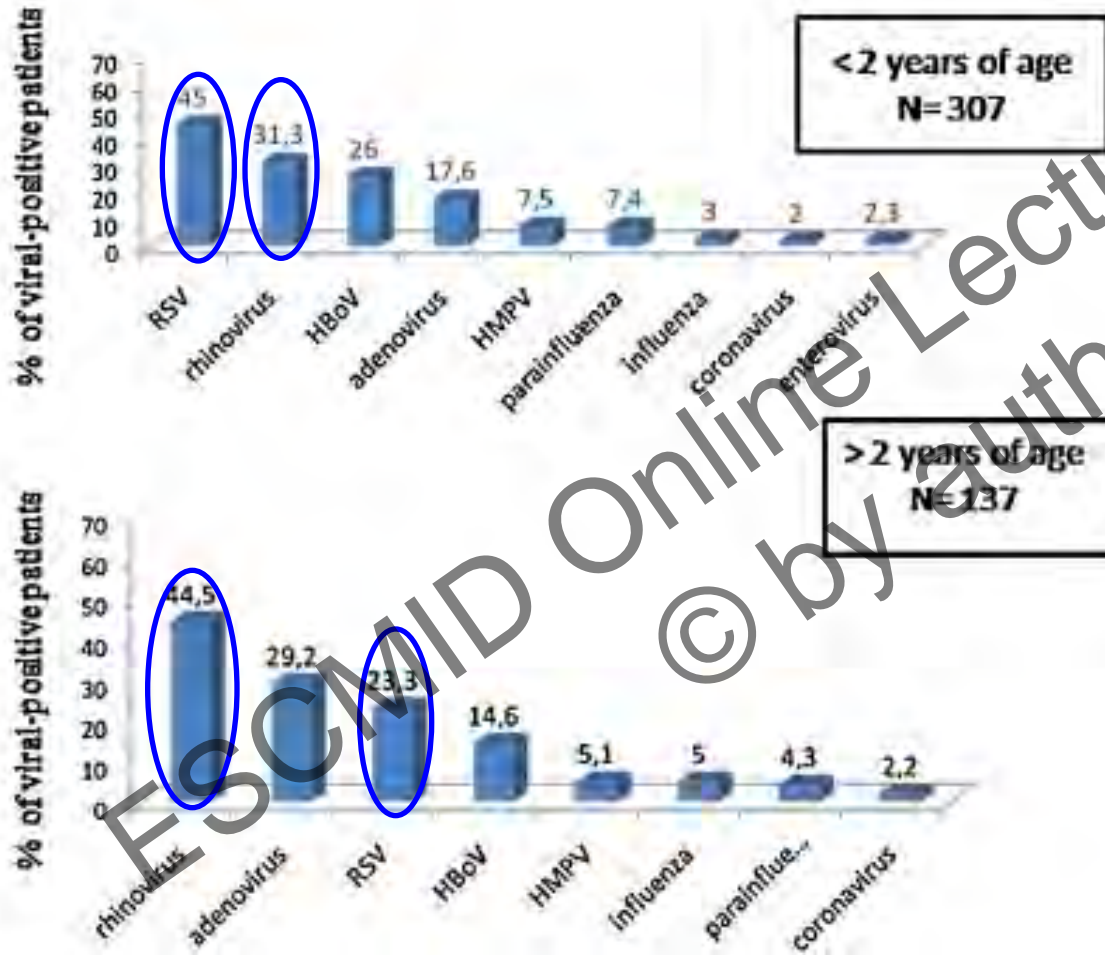
In 1 study, hMPV/RSV coinfection in 70%

Disease severity and hospitalization appears more common with RSV

Osterhaus A, et al. Lancet. 2003; 361:890-891  
 Boivin G, et al. Emerg Infect Dis. 2003;9:634-640  
 Greensill J, et al. Emerg Infect Dis. 2003; 9: 372-375.  
 McAdam AJ et al. J Infect Dis 2004; 190: 20-26  
 Esper F et al. J Infect Dis 2004; 189: 1888-96



# Emerging respiratory viruses in children with severe acute wheezing



- viruses detected in 71% of acute wheezing episodes
- RSV most commonly detected virus: 27%
- Rhinovirus in 24%
- Adenovirus 18%
- Rate of viral detection ↑ in infants (77%) than in older children (60%)
- RSV and rhino most prevalent in wheezing; emerging viruses hBoV and hMPV also important**

Garcia-Garcia ML et al. Pediatric pulmonology 2010; 45: 585-91

## Impact on epidemiological knowledge: Distribution of respiratory pathogens in LRTI



- **From 10/2007 – 12/04/2010**

- 3102 adult patients with LRTI
- 2984 controls
- 16 PCN in 12 countries

→ Presence viral etiologic agent by RT-PCR: **53.7%**

→ In total: etiology found in **> 70%** in LRTI in community



# Significance of etiologic agents?



| Target               | Patient with LRTI prevalence n/total (%) |                          |           | Matched Control subject prevalence n/total (%) |             |
|----------------------|--|--------------------------|-----------|--|-------------|
|                      | first visit (n=3059)                     | follow up visit (n=2566) | P-value   | (n=1678)                                       | P-value     |
| Parainfluenza 1-4    | 73 (2.4)                                 | 12 (0.5)                 | < 0.00001 | 8 (0.5)  | < 0.00001   |
| Rhinovirus           | 589 (19.3)                               | 112 (4.5)                | < 0.00001 | 72 (4.3)                                       | < 0.00001   |
| Human MPV            | 126 (4.1)                                | 7 (0.3)                  | < 0.00001 | 3 (0.2)  | < 0.00001   |
| Human AV             | 38 (1.2)                                 | 42 (1.6)                 | 0.26      | 24 (1.4)                                       | 0.68        |
| Bocavirus            | 13 (0.4)                                 | 12 (0.5)                 | 0.96      | 16 (1.0)                                       | <b>0.04</b> |
| RSV                  | 143 (4.7)                                | 13 (0.5)                 | < 0.00001 | 11 (0.7)                                       | < 0.00001   |
| Influenza A/B        | 300 (9.8)                                | 10 (0.4)                 | < 0.00001 | 6 (0.4)  | < 0.00001   |
| Coronaviruses        | 208 (6.8)                                | 70 (2.7)                 | < 0.00001 | 27 (1.6)                                       | < 0.00001   |
| Polyomavirus WU      | 45 (1.5)                                 | 54 (2.1)                 | 0.09      | 39 (2.3)                                       | <b>0.04</b> |
| Polyomavirus KI      | 27 (0.9)                                 | 28 (1.1)                 | 0.51      | 17 (1.0)                                       | 0.77        |
| <i>M. pneumoniae</i> | 0/809 (0)                                | 0/653 (0)                | NS        | 0/492 (0)                                      | NS          |

leven M et al on behalf of GRACE ECCMID 2010

<sup>1</sup>:Only data of first winter season



**How to implement these amplification techniques for the detection of bacterial etiologies?**

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# Individual patient care: Is bacterial quantification useful in molecular testing?



## In case of legionellosis

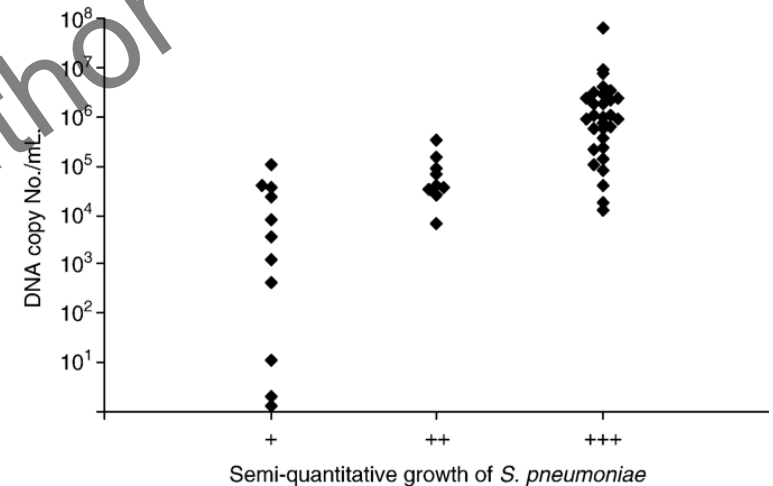
- qRT-PCR useful for predicting disease severity: high bacterial loads significantly associated with higher Fine classes, need for ICU hospitalization

## In case of pneumococcal disease

- Cut-off  $10^4$  DNA copies/ml
  - Sens: 84%
  - Spec: **94%**

Abdeldaim G et al Diagn Microbiol Infect Dis J 2008; 60: 143-50

Maurin M et al Clin Microbiol Infect 2010; 16: 379-384



➔ Q PCR enables differentiation between pathogenicity and colonization

➔ RQ-PCR particularly valuable in patients treated with AB



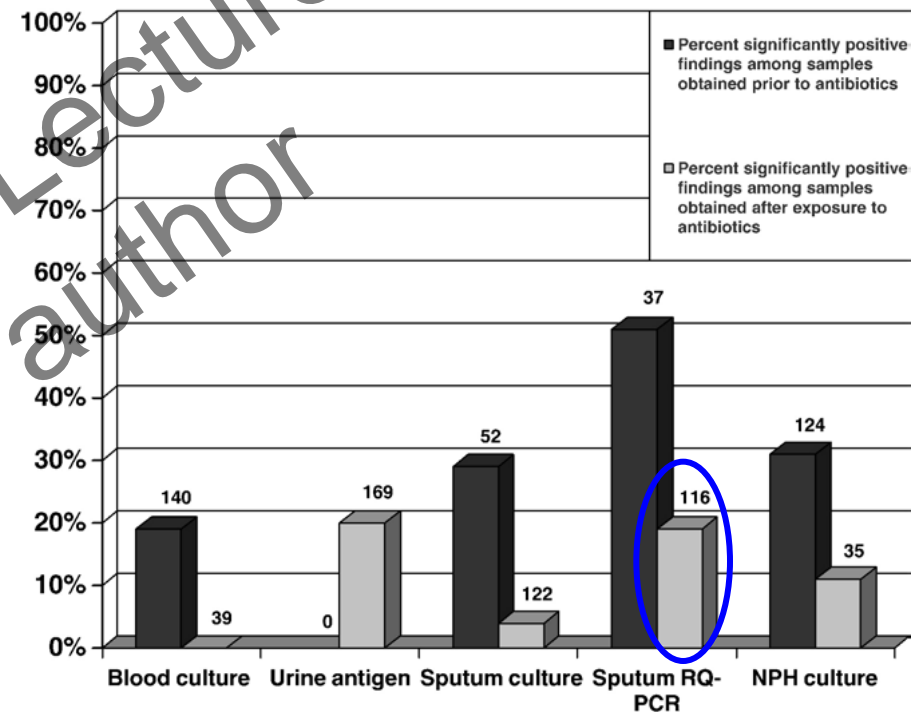


# Quantitative PCR for Diagnosis of *S. pneumoniae* Infection



70/184 (38%) patients with *S.pneumoniae*

- 15% by blood culture
- 20% by urinary Ag
- 15% culture positive sputa
- 27% by RQ-PCR
  - 82% of these also detected by other methods
  - 50% of these culture -, most of these treated with AB



⇒ RQ-PCR particularly valuable in patients treated with AB



# Individual patient care: Is viral quantification useful?



## In case of PIV, rhinovirus

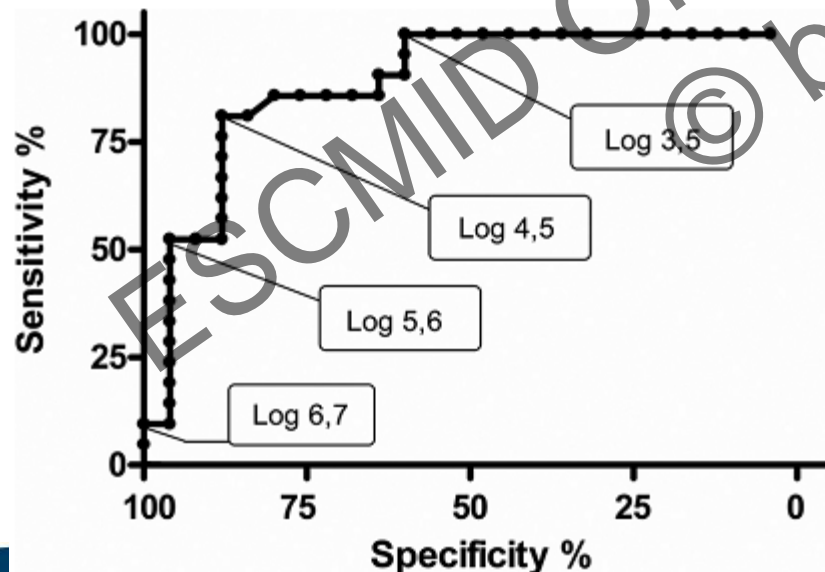
- Total viral load is related to clinical diagnosis in children presenting at emergency room

Utokaparch S et al. *Pediatr Infect Dis J* 2011; 30: e18-e23

## In case of rhinovirus

- At high viral loads ( $> 10^6$  RNA copies/ml): HRVs may cause severe LRTI
- At medium-low viral loads ( $< 10^5$  RNA copies/ml): may represent only bystander

Gerna G et al. *J Med Virol* 2009; 81:1498-1507



- At high viral loads ( $> 10^{4.5}$  RNA cps/ml): HRVs likely to be the cause of presenting LRTI
- At medium-low viral loads ( $< 10^{4.5}$  RNA copies/ml): may represent only bystander
- **Q PCR: maybe the next necessary step?**

Jansen R et al. *J Clin Microbiol* 2011; 49: 2631-36

# VD Impact of molecular detection on diagnosis of acute *M. pneumoniae* infection

- *M. pneumoniae* in 48/164 patients:
  - 45/164 (29%): PCR pos in first week
  - 44/154 (27%): significant ↑ in IgG or + IgM

## **BUT:**

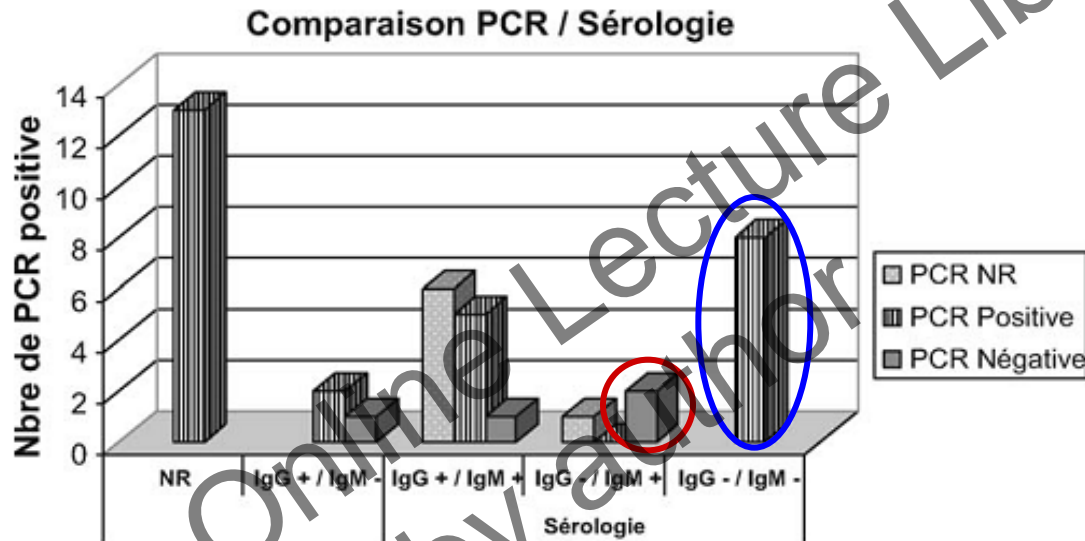
- 21% in 1st week
- 56% during second week
- 100% after 3rd week

Nilsson et al., BMC Microbiol 2008; 8: 93-100

- CAP + LRTI: study in Antwerp: 4 different tests evaluated on 224 patients (205 paired sera)
  - sensitivity of IgM: **10-31%** in first 6 days  
**20-42%** after more than 16 days

➔ **Molecular methods are superior to serology for early diagnosis of *M. pneumoniae* and impact on antibiotic management**

# Importance of PCR in the diagnosis of *Mycoplasma pneumoniae* infections



- PCR based detection: most sensitive: 28/32 (87%)
- Sensitivity of serology: 15/26 (58%)
- 7 patients only diagnosed by serology Dekeyser S et al., Pathol Biol 2011; 83-87
- *M.pn* detected in 2.9% by IgG seroconversion or significant IgG rise
- All 4/374 (1.1%) sputum PCR positive patients also IgG positive

**➔ Combination of PCR and serology detects most cases**



# Importance of dual or multiple infections?

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# Clinical relevance of infection with multiple viruses?



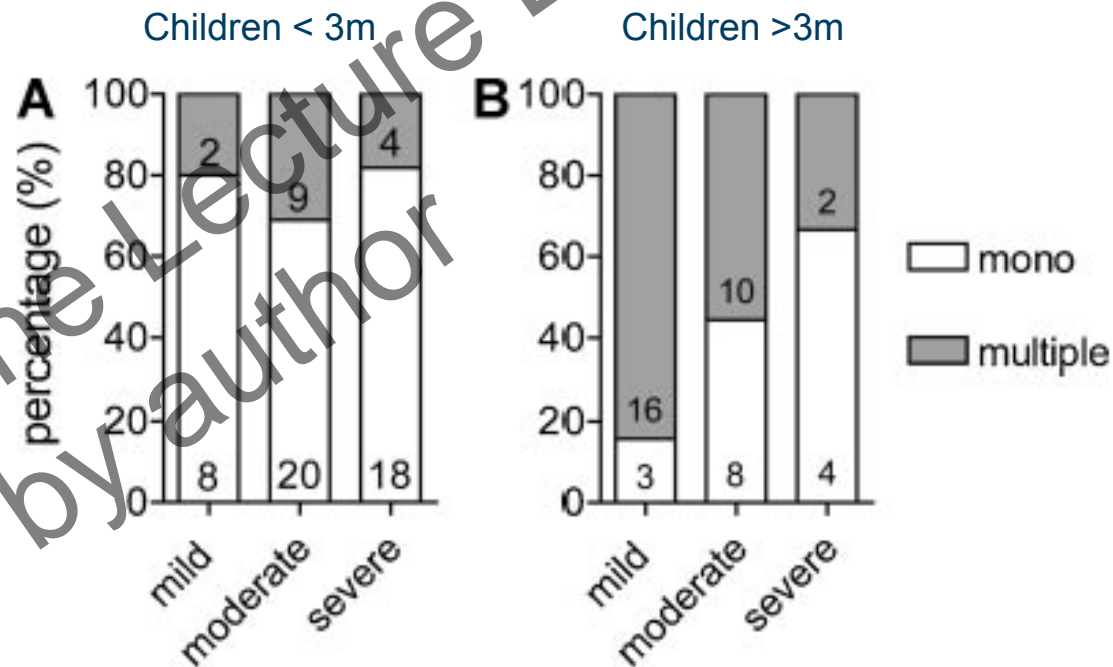
- children < 2yrs old with bronchiolitis:
  - Mild: no supportive treatment
  - Moderate: supplemental oxygen and/or nasogastric feeding
  - Severe: mechanical ventilation
- Mx PCR for 15 viruses on NPA
- Results: overall 211 viruses detected in 142 NPA
  - RSV most commonly detected virus: 73%
  - Rhinovirus in 30%
  - Other respiratory viruses in < 10% of samples

Brand HK et al. Pediatric Pulmonology 2011, 162: 88-90

# VD Importance of infection with multiple viruses?



- Children younger than 3 months: less often infected by multiple viruses compared to children older than 3 months: 25% vs 65%
- Infection with 2 or more viruses: more frequent in children with mild or moderate disease than in those with severe disease



➔ **The detection of more than one virus is not associated with increased disease severity in children with bronchiolitis**

➔ **Co-infections not associated with illness severity in acute febrile RTI**

Suryadevara M et al. Clinical Pediatrics 2011, 50: 513-51



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# When and where to implement these amplification techniques?

# VD Respiratory Virus Infections in immunocompromised Patients

- Significant viruses in URTI and LRTI in lung transplants:
  - total 27 % patients: Flu > RSV > PI; hMPV 4.3%; rhino 18%

Weinberg A et al. J Clin Virol. 2002; 25: 171-75

Dare R et al. J Clin Microbiol 2007; 45:548-52

Camps Serra M et al. Eur Resp J 2008; 31: 618-24

- Significant cause of pneumonia in hematological cancers
  - total 35 % of patients: RSV > Rhino > Influenza > PIV
  - HCoV increasingly important: 11 %

Van Elden L et al. Clin Infect Dis. 2002; 24: 177

Van Elden L et al. J Infect Dis. 2004; 189: 653

- Acute RTI in elderly and children: up to 40%:
  - mostly rhino, RSV, hMPV, and influenza

Regamey et al 2008, Renwick et al 2007, Dare et al 2007, Jartti et al 2008





Winter



Influenza epidemic in community

Influenza sporadic

Rapid influenza test with follow-up viral culture for clinically suspicious cases

⊕

Respiratory isolation x 5 days consider specific antiviral therapy

⊖

Illness severity

Severe or immunocompromised

Mild

No specific therapy

Culture = RT-PCR for RSV and other viruses

⊖

Contact isolation & consider antiviral therapy

⊕