

Multiplex PCR to detect respiratory viral infections

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ESCMID Venice
25.10.2010



Molecular methods for diagnosis of respiratory tract infections



- Virus discovery → new molecular diagnostic tests
- Will revolutionize infectious disease diagnosis and patient care
- Early detection of infection impacts outbreak management
- Impact on epidemiological knowledge
 - Role Distribution of respiratory viruses in LRTI
 - In +/- 75% of outbreaks of RTI in the community never have an etiological agent identified
- Medical impact on individual patient care
 - Stop inappropriate AB use: reduces overall costs, resistance:
 - Although viruses account for 80% of all human respiratory tract infections, 60% of patients will receive antibiotics
 - Decrease unnecessary diagnostic studies



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

2001: hMPV

BERNADETTE G. VAN DEN HOOGEN¹, JAN C. DE JONG¹, JAN GROEN¹, THUIS KUIKEN¹, RONALD GROOT², RON A.M. FOUCHIER¹ & ALBERT D.M.E. OSTERHAUS¹

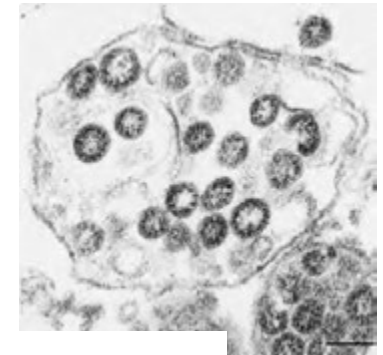
The NEW ENGLAND JOURNAL of MEDICINE

ORIGINAL ARTICLE

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

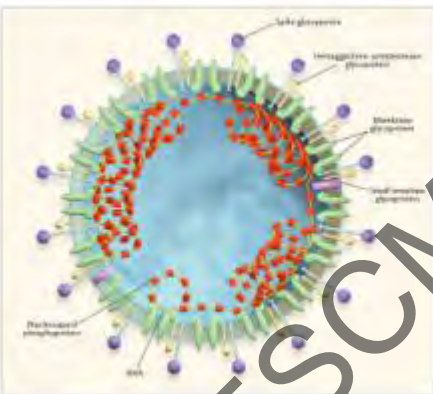
Christian Drosten, M.D., Stephan Günther, M.D., Wolfgang Preiser, M.D., Sylvie van der Werf, Ph.D., Hans-Reinhard Brodt, M.D., Stephan Becker, Ph.D., Holger Rabenau, Ph.D., Marcus Panning, M.D., Larissa Kolesnikova, Ph.D., Ron A.M. Fouchier, Ph.D., Annemarie Berger, Ph.D., Ana-Maria Burguière, Ph.D., Jindrich Cinatl, 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 Riekerts, M.D., Martin Stürmer, Ph.D., Simon Vieth

2003: HCoV SARS



nature
medicine

2005: HCoV NL 63

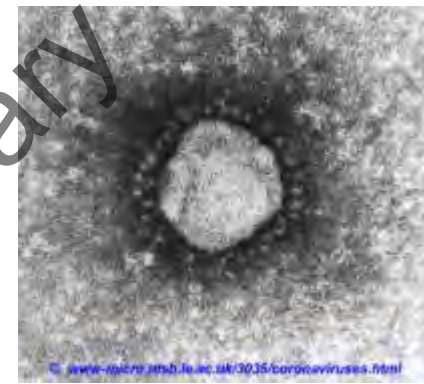


Identification of a new human coronavirus

Lia van der Hoek¹, Krzysztof Pyrc¹, Maarten F Jebbink¹, Wilma Vermeulen-Oost², Ron J M Berkhout², Katja C Wolthers¹, Pauline M E Wertheim-van Dillen³, Jos Kaandorp⁴, Joke Spaargaren² & Ben Berkhout¹

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

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



2005: HCoV HKU1

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

Tobias Allander^{*†‡}, Martti T. Tammi^{§¶}, Margareta Eriksson^{||}, Annelie Bjerkner^{*}, Annika Tiveljung-Lindell^{*},
and Björn Andersson[§]

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

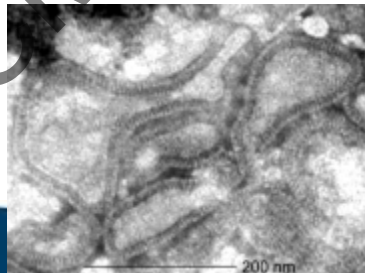
David Kiang^{a,✉}, Shigeo Yagi^a, Katherine A. Kantardjieff^b, Euna J. Kim^b, Janice K. Louie^a
and David P. Schnurr^a



The main respiratory targets in molecular diagnostic tests

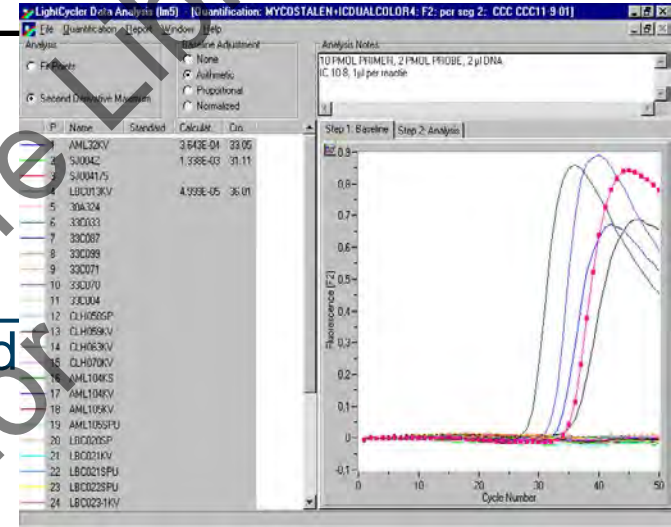


- Those in our routine panel:
 - Influenza A (IFVA)
 - Influenza B (IFVB)
 - Parainfluenza (PIV) 1-4
 - Respiratory syncytial virus (RSV)
 - Adenoviruses (ADV)
 - Metapneumovirus (hMPV)
- Those extra assays that many would consider important:
 - Rhinoviruses
 - Enteroviruses
 - Coronaviruses (OC43, 229E, 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 RSVA and B, COR E229, OC 43, 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, PFL 1+3, PFL 2+ hCoV-229E, ADE, hMPV, RHI, 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, RSV A/B, AV, 3 coronaviruses, <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> , <i>L.pn.</i> , IFL A and B, RSV A/B, PFL 1-3, RHI, 3 coronaviruses, AV, HBoV, EV
xTAG RVP, Luminex	19	IFL A (H1, H3, H5, non-specific) and B, PFL 1-4, RSV A/B, ADE, hMPV, RHI/ENT,SARS-COR, HCoV OC43, HCoV 229E, HCoV NL63 and HKU1



- Rapid, real-time PCR/TMA/NASBA/LAMP
- Fast cyclers
- New innovative detection systems
- Multiplex capabilities expanding
- Automation for NA extraction
- Automated amplification and detection platforms
- Full integration of MDx steps
- Molecular “Point of Care” testing
- Next gen sequencing



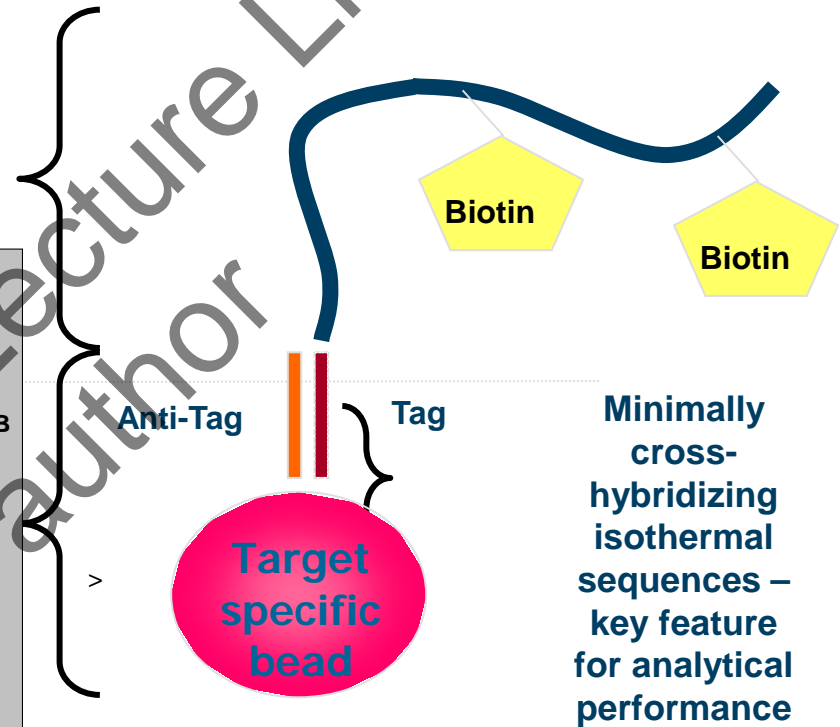
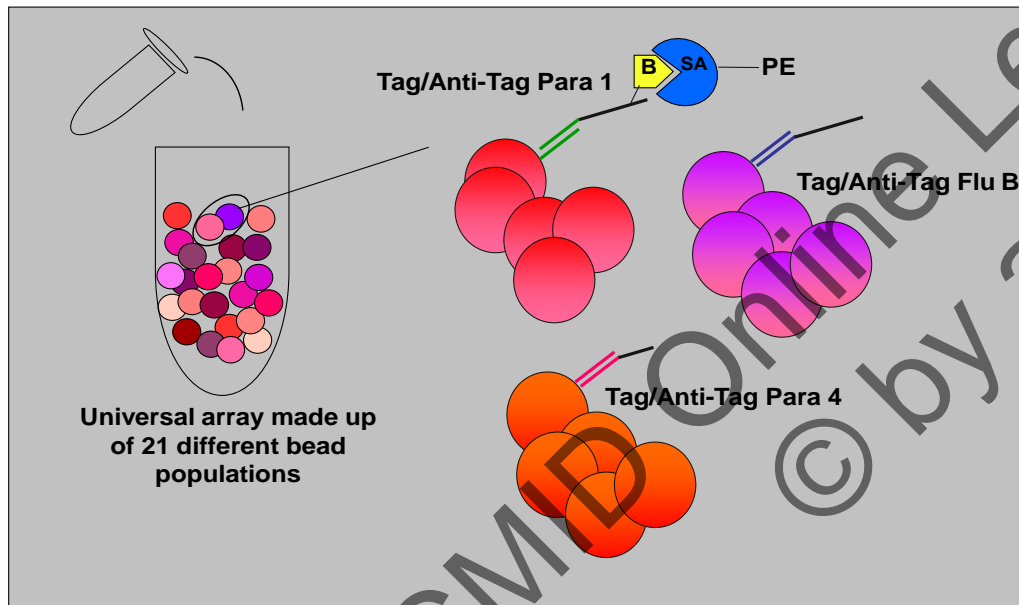
- Rapid, real-time PCR/TMA/NASBA/LAMP
- Fast cyclers
- **New innovative detection systems**
- **Multiplex capabilities expanding**
- Automation for NA extraction
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- Full integration of MDx steps
- **Molecular “Point of Care” testing**
- Next gen sequencing

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VD Luminex xTAG Universal Bead Array

Universal Array Sorting

Oligo-coupled beads used with Tag-It products.
Use MFI for every target-specific primer to determine presence or absence of each virus



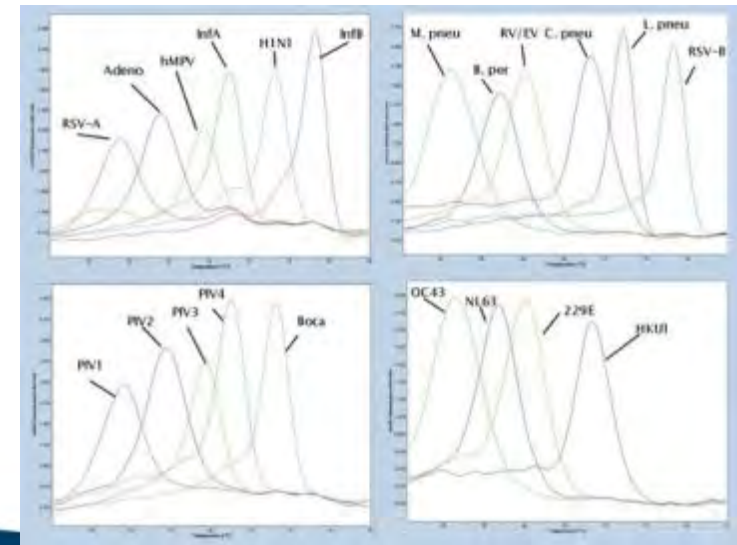
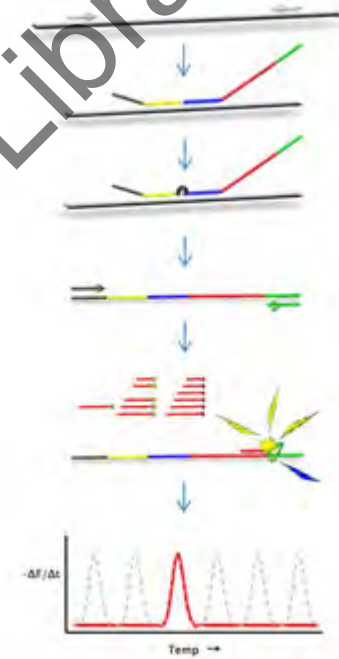
RVP 10-12 hr to RVP Fast 4-5 hr



VD RespiFinder Mx assay, Pathofinder



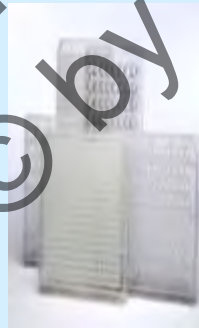
- Rapid, conventional & real-time PCR
- Multiplex capabilities expanding till up to 25 targets
- 18 viral + 4 bacterial pathogens in 1 RT assay
- Gene-specific Mx reverse transcription step
- 2 pathogen specific probes hybridized by ligation, amplified and detected by melting curve analysis
- Contains a competitive internal amplification control
- Diagnosis within 6 hours
- Validated on QCMD panels
 - CE-IVD labelled





The future

- Major advances in microfluidics and -electronics:
"Biochips", "Lab-on-a-chip", "nanotechnology" or POCT



Change of scale

Miniaturization

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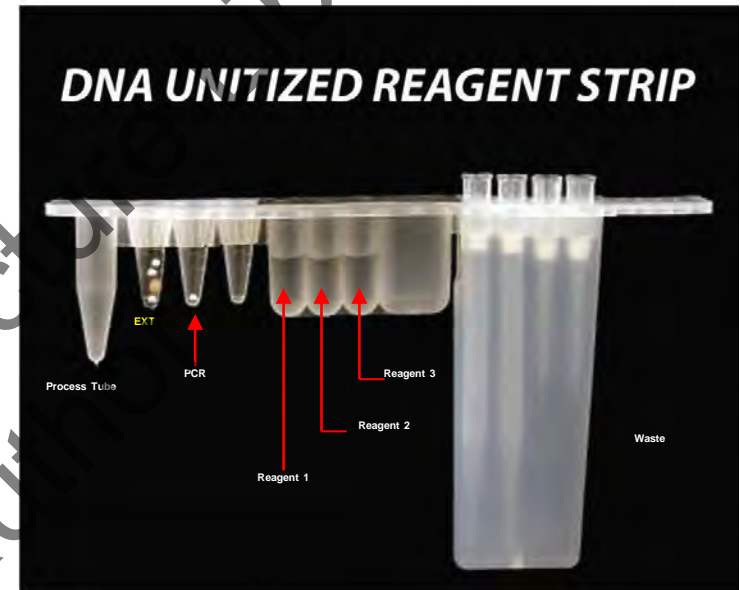


Cepheid GeneXpert System



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

- Complete System: extraction, amplification and real time detection
- Reagent strip and PCR cartridge
- 1st generation: 2 colors
- 2nd generation: 6 colors
- BD developed assays
- Open system for LDTs



PCR Cartridge





One manual pipetting step



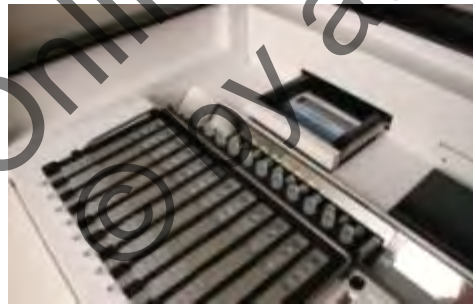
Load reagents and specimens



Place rack in Max



Load PCR cartridge



Place on Max



Create worklist and Close door to initiate run

2 positions with 12 slots per position



- System integrates sample preparation, amplification, detection and analysis
- All reagents are freeze dried in 1 pouch
- Closed system prevents cross-contamination
- Internal controls for each step
- Advanced software runs the system and automatically analyzes and reports results
- Multiplexed testing analyzes up to 120 tests per sample
- Rapid results in 1 hr from sample injection

Viral: Adeno, Boca, Corona (229E, HKU1, OC43, NL63), Flu A, Flu A H1, Flu A H1 2009, Flu A H3, Flu B, hMPV, Para (1, 2, 3, 4), RSV, Rhino
Bacterial: *B. pertussis*, *C. pneumoniae*, *M. pneumoniae*



Some Technical, Scientific and practical hurdles



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

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V_iD_i EQA panel to evaluate the performance of different real-time PCRs in GRACE



GRACE Sample Number	Sample Content	Conc. / Dilution (Ct)	Centre 2 Results	Centre 3 Results	Centre 4 Results
GRACE-01	HMPV-I	1x10 ⁻⁶ (36)	hMPV Positive	hMPV Positive	Negative
GRACE-02	Influenza A virus Type H3	1x10 ⁻⁷	Negative	PIV 2/4 Positive	Negative
GRACE-03	Parainfluenza virus 2	-34	PIV 2 Positive	PIV 2/4 Positive	PIV 2 positive
GRACE-04	Negative Transport Medium	0	Negative	Negative	Negative
GRACE-05	Negative Transport Medium	0	Negative	Negative	Negative
GRACE-06	Influenza B virus	1x10 ⁻⁶	INF B Positive	INF Positive	Negative
GRACE-07	HMPV-II	1x10 ⁻⁵ (35)	hMPV Positive	hMPV Positive	Negative

Sample content	Conc. / Dilution	Centre 2	Centre 3	Centre 4
Influenza A	1x10 ⁻⁶	Negative	INF positive	Negative
Influenza B	1x10 ⁻⁶	INF B positive	INF positive	Negative
Influenza A	1x10 ⁻⁷	INF A positive	Negative	Negative
hMPV-II	1x10 ⁻⁵	hMPV positive	hMPV positive	Negative
RSV A	1x10 ⁻⁵	RSV positive	RSV positive	RhV positive

GRACE-21	RSVA	1x10 ⁻⁶ (24)	RSV Positive	RSV Positive	ADV + RSV positive
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VDI EQA panel to evaluate the performance of different real-time PCRs in GRACE




GRACE Sample Number	Sample Content	Conc. / Dilution (Ct)	Centre 2 Results	Centre 3 Results	Centre 4 Results
GRACE-22	Coronavirus NL63(2x10 ⁻⁶)	(35-37)	Negative	hCoV Positive	Negative

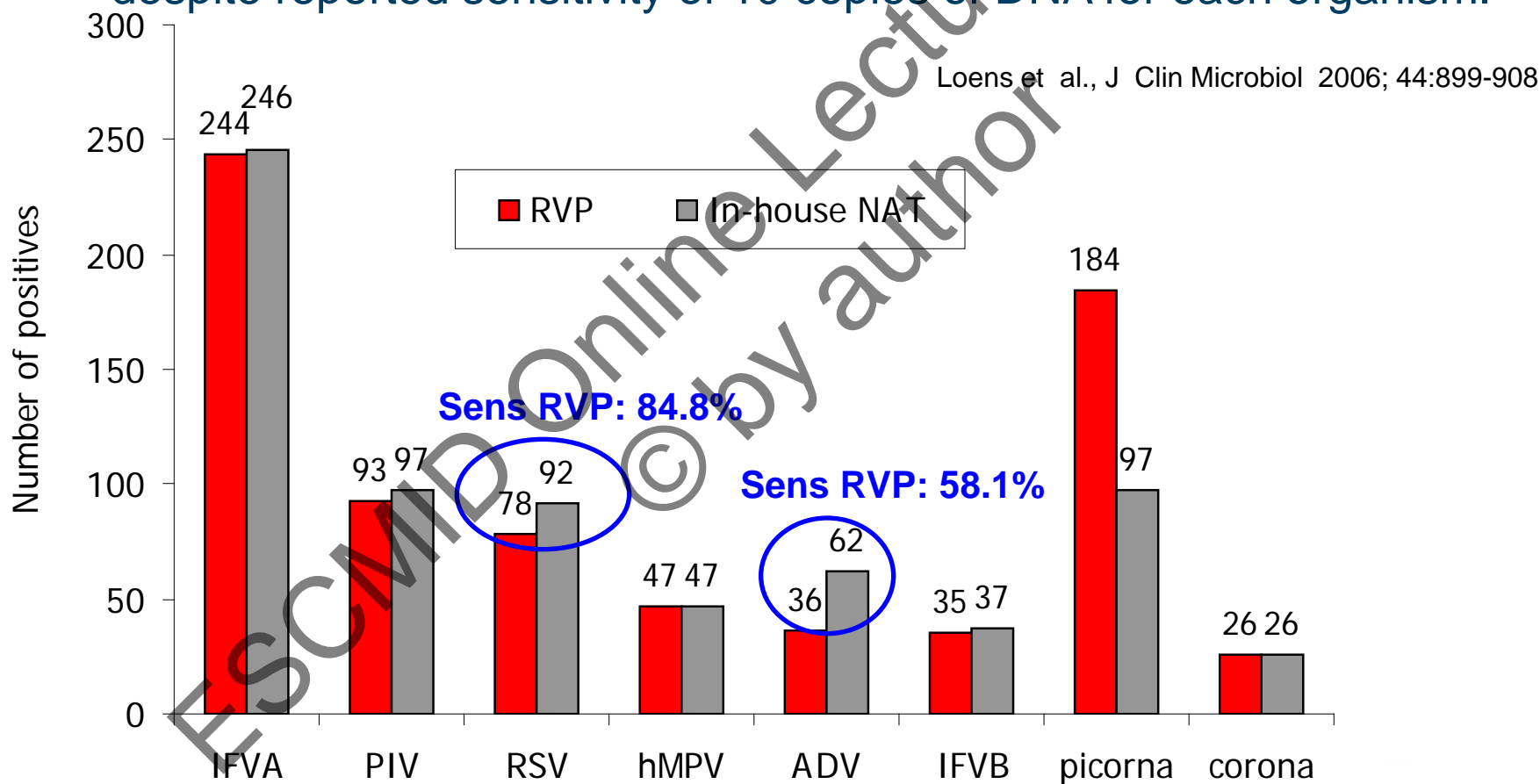
Sample content	Conc. / Dilution	Centre 2	Centre 3	Centre 4
hCoV NL63	2x10 ⁻⁶	Negative	hCoV positive	Negative
hCoV OC43	1x10 ⁻⁴	hCoV positive	hCoV positive	Negative
RhV 90	1x10 ⁻⁵	RhV positive	RhV positive	Negative
ADV 1	1x10 ⁻²	ADV positive	ADV positive	Negative



Single detection versus Mx testing?



- Both multiplex PCR and NASBA formats scored a smaller number of samples positive for *M. pn.*, *C.pn.*, *L. pn.* than monoplex reactions .
- One commercial Mx kit scored only 1/3 of all + samples positive despite reported sensitivity of 10 copies of DNA for each organism.





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



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

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**



GRACE: Etiologic diagnosis of LRTI in primary care: impact of RT-PCR



- EU FP6 Network of excellence
16 PCN in 11 countries
- **From 10/2007 – 12/04/2010**
 - 3102 adult patients with LRTI
 - 2984 controls
 - 16 PCN in 12 countries
- Blood & respiratory samples taken, transported to Antwerp

➔ 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)	0.04
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)	0.04
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

1 :Only data of first winter season



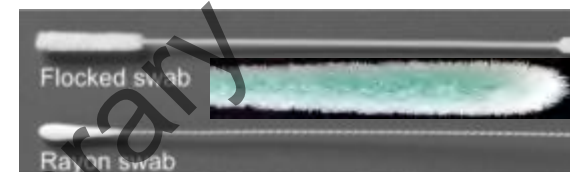
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CA-LRTI: comparison of samples/organism



Pathogen	Sample ranking	Method	Age (yr)	Total no. of specimens/ no. of patients	Reference
<i>M. pneumoniae</i>	Sputum > TW > NPS > OPS	PCR	20-93	552/144	(1)
	OPS > NPS	PCR	NSp	132/66	(2)
	OPS > BAL > Sputum	PCR	NSp	325/197	(3)
	Sputum > OPS	Gene-probe test	>18	160	(4)
	Sputum > NPA	Ag-EIA	>18	102/51	(5)
	Sputum > OPS	Culture, PCR, NASBA	NSp	302/180	(6),(7)
	NPS = OPS	PCR	NSp	63	(8)
	Sputum > NPA = OPS	PCR	22-29	96/32	(9)
	OPS > NPA	PCR	NSp	102	(10)

(1) J. Clin. Microbiol. 2001, 39: 1184-6

(2) Scan. J. Infect. Dis. Suppl. 1997, 104: 11-2

(3) J. Clin. Microbiol. 2000, 38: 1382-4

(4) J. Infect. Dis. 1990, 162: 70-5

(5) Eur. J. Clin. Microbiol. Infect. Dis. 1993, 12: 872-5

(6) J. Microbiol. Methods 2008, 73: 257-62

(7) J. Clin. Microbiol. 2008, 46: 185-91

(8) J. Clin. Microbiol. 2004, 42: 3339-41

(9) J. Med. Microbiol. 2005, 54: 287-91

(10) Eur. J. Clin. Microbiol. Infect. Dis. 1995, 14: 58-61



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TheraEDGE Project



An Integrated Platform Enabling Theranostic Applications at the Point of Primary Care

- 4 year project
- Budget: 11M€
- 9 different countries

Theranostics & Personalized Medicine



www.theraedge.org



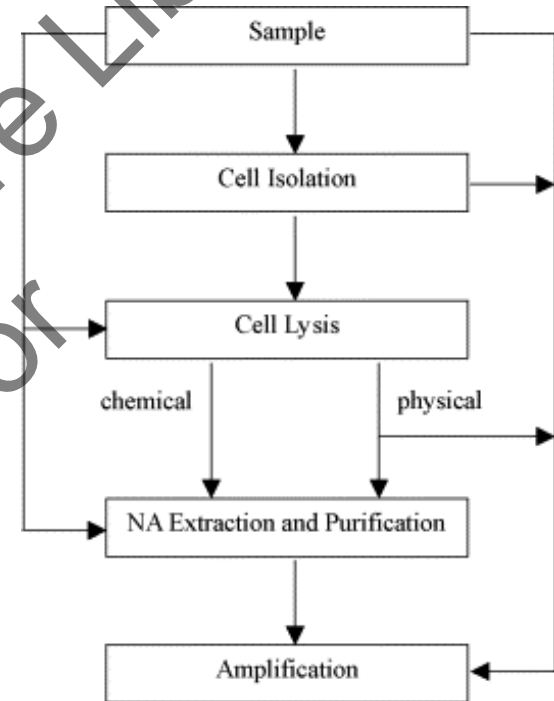
UNIVERSITAT DE BARCELONA



VD A Big Bottleneck in Developing POCTs: Sample Preparation



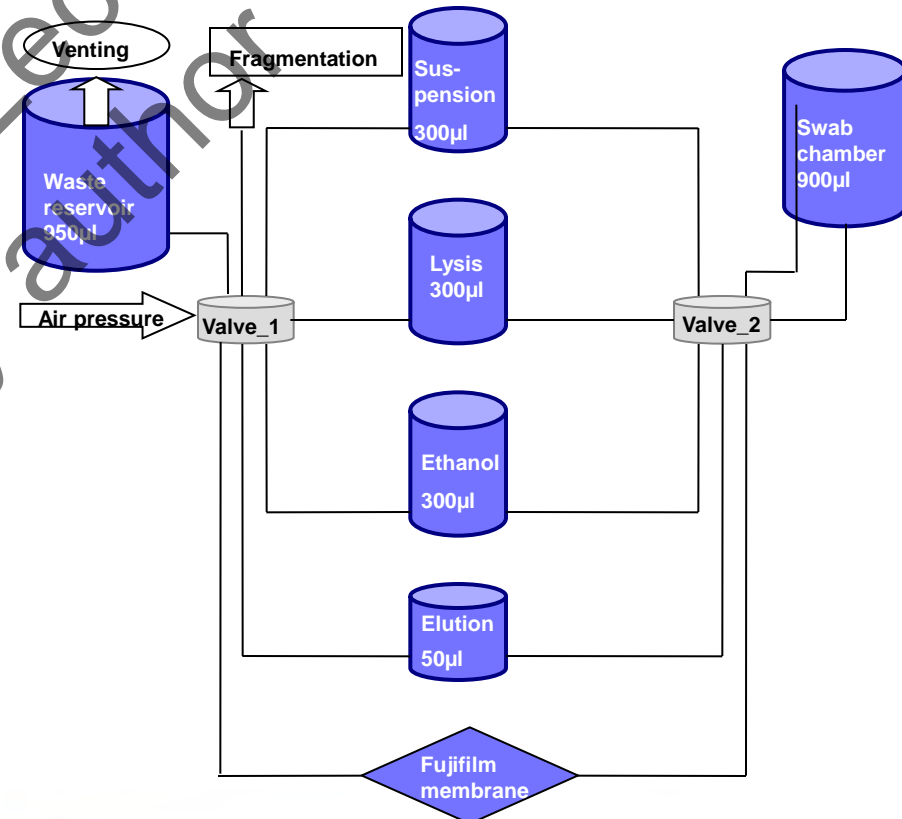
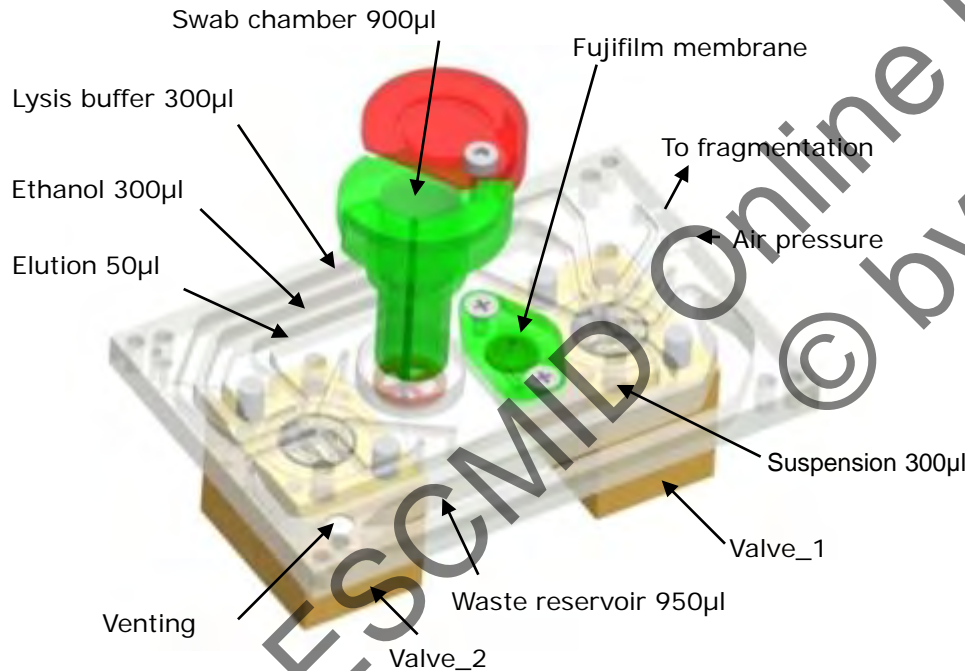
- Off-chip (macroscale) sample prep
 - Laborious
 - Refrigerated/frozen reagents
 - Large sample volumes
 - Requires centrifuges, bead beaters, several machines
 - Few hours
- On-chip sample prep
 - Room temperature stable reagents (disposable chips with on-chip storage)
 - Microliter volumes
 - Few minutes!!



Development of a proprietary bacterial lysis and DNA purification protocol and its successful application on a prototypal microfluidic chip for a CA-LRTI assay

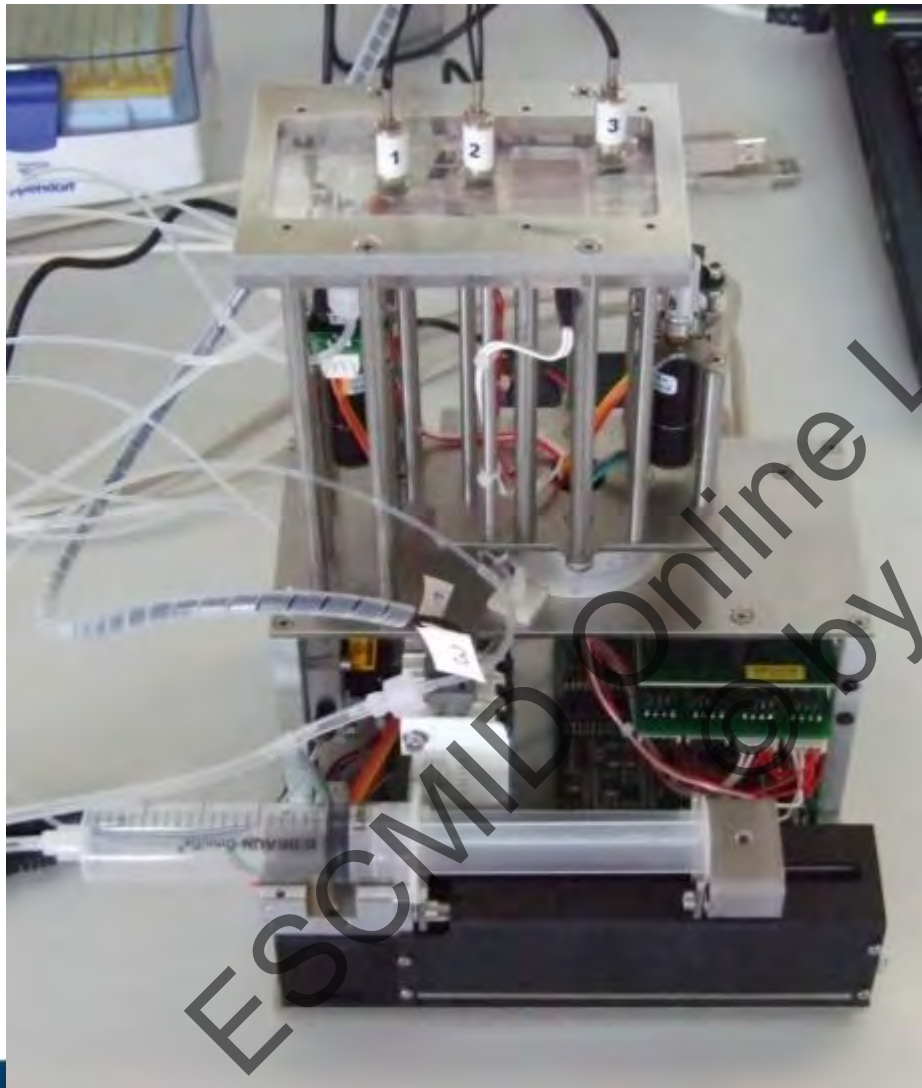
Chip design

Reaction steps





Semi-automated demonstrator setup





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ESCMID

EUROPEAN SOCIETY
OF CLINICAL MICROBIOLOGY
AND INFECTIOUS DISEASES

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RAPP-ID: Public Private Partnership



www.rapp-id.eu

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PROJECT WEBSITE: www.rapp-id.eu

Development of Rapid Point-of-Care Test Platforms for Infectious Diseases



RAPP-ID will develop a Point-of-Care Test (POCT) for rapid (hospital <2h, primary care <30min) detection of **bacteria, mycobacteria, fungi**, as well as **viruses** and **host biomarkers** by combining novel specific probes, novel methods of sample preparation, and demonstrated ultra-high sensitive detection methods. The platforms will also determine **resistance** to antimicrobial drugs

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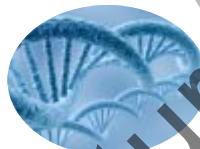
A plea for an integrated approach!!



Biotechnologies

- Sample preparation
- Protein chemistry
- Novel surface chemistry

Biological Sciences



Clinical practice

- Selection of relevant targets and applications
- Probe design
- Validation of analytical, clinical performance



Physical Sciences



Clinical Practice



Innovative Medicines Initiative

RAPP-ID

Microtechnology

- Microfluidics, LOC
- Photonics
- biosensors

FP7 projects

