



HERA HEALTH EMERGENCY
PREPAREDNESS AND
RESPONSE AUTHORITY

#HealthUnion

HERA Industry days

Health Emergency Preparedness and Response Authority

2 & 3 June 2025, Brussels

Side session

**Advancing diagnostics: scaling
production, overcoming industry
challenges, and addressing future needs
in crises**

Advancing diagnostics: scaling production, overcoming industry challenges, and addressing future needs in crises



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Christian DROSTEN
DURABLE Project



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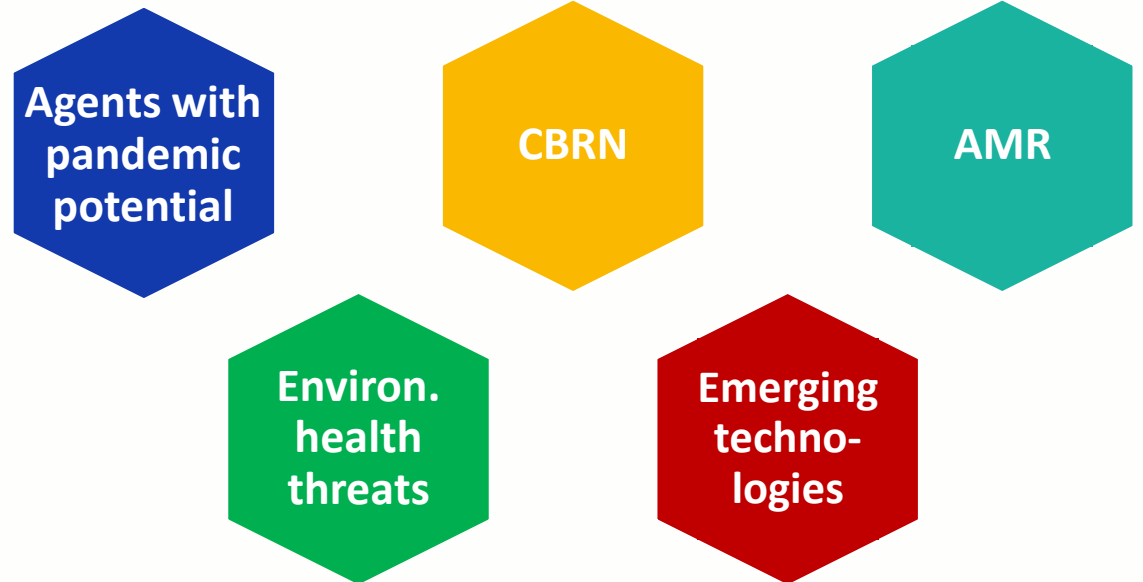
Advancing diagnostics: scaling production, overcoming industry challenges, and addressing future needs in crises

3 June 2025
Cornelius Schmaltz, Head of Unit
HERA.02



- In **2022**, HERA identified its 3 main serious cross-border health threat categories
- In **2023**, HERA identified 2 additional 'threat subtopics', to support the previously identified threat categories

→ **Diagnostics matter for all threat categories**



- Diagnostics are core MCMs within HERA's mandate
- Enable critical decision-making across clinical care, surveillance, and outbreak response
- Bridge preparedness and response—linking threat identification to MCM deployment
- Investing in innovative diagnostic technologies for future threats



Previous funding actions

2022 (EU4Health):

- EUR 25 million: DURABLE lab network
 - Includes work on diagnostics

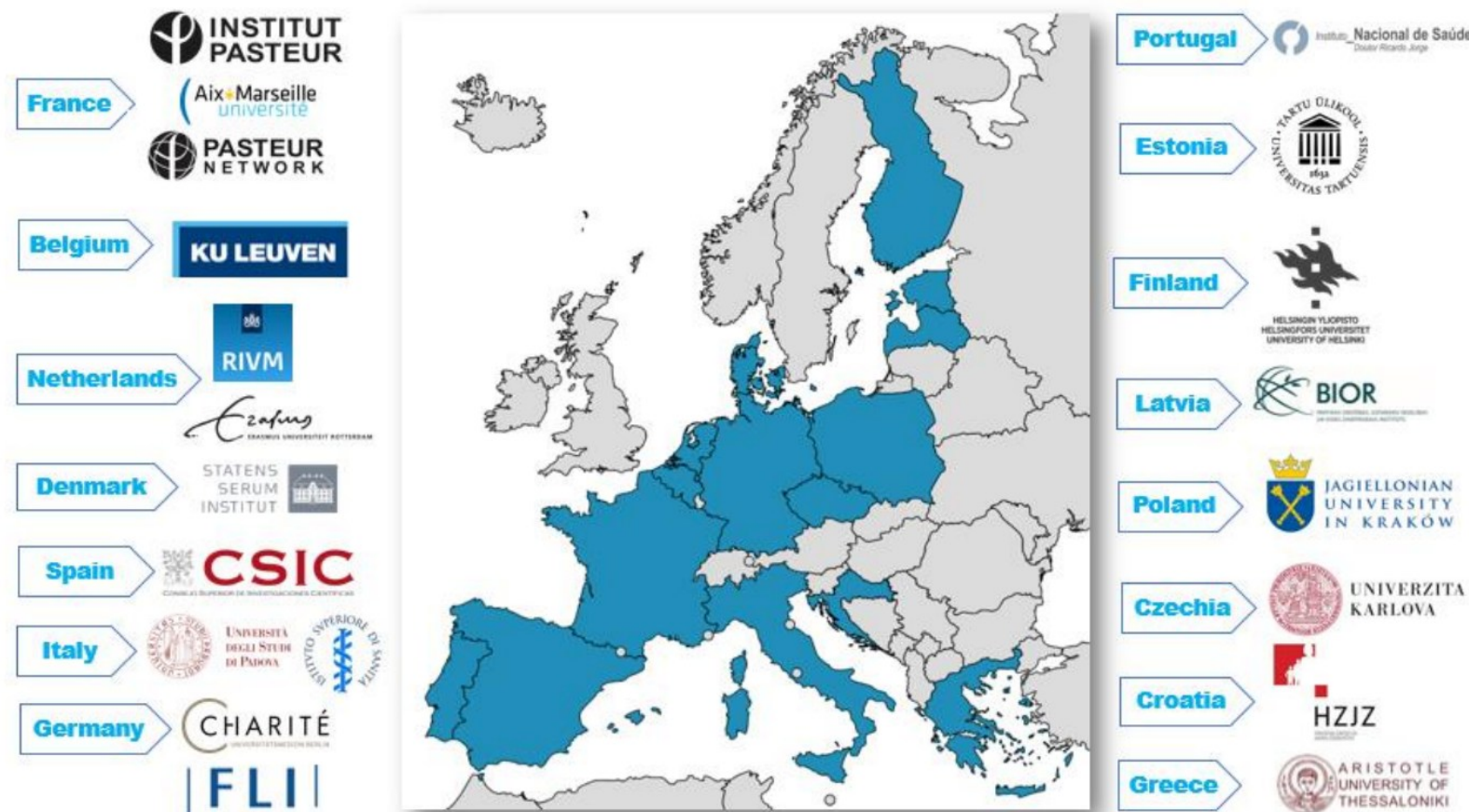


Figure 1: DURABLE consortium

Previous funding actions

2023 (Horizon Europe):

- EUR 40 million: HORIZON-HLTH-2023-TOOL-05-08: Pandemic preparedness and response: In vitro diagnostic devices to tackle cross-border health threats
 - 6 projects funded (UniHealth, B-Path, DECIPHER, PREPARE-TID, PAIR, DRAIGON)

2024 (EU4Health):

- EUR 24 million: Point of care metagenomic sequencing for universal pathogen detection

2025 (EU4Health):

- EUR 12.86 million: Development of a rapid point-of-care antimicrobial susceptibility testing diagnostic medical device
 - Under evaluation



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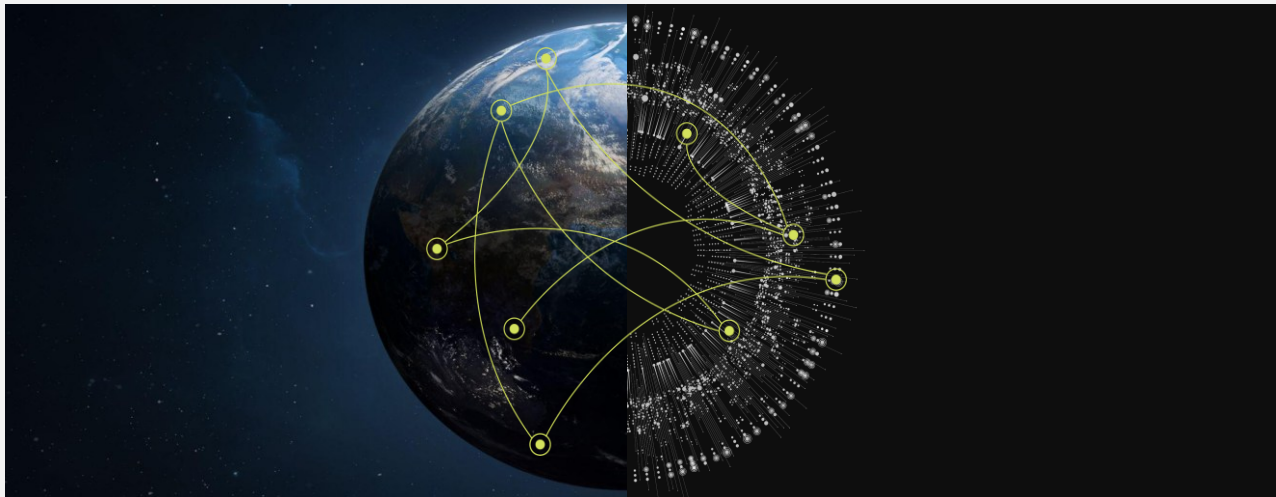
Jeroen NIEUWENHUIS
CEO, Nostics



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DURABLE Project

Ginkgo Biosecurity is powering global biosurveillance and rapid pathogen detection

Ginkgo Biosecurity is building **next-generation biosecurity infrastructure** and deploying the technologies global leaders need to **predict, detect, characterize, and respond** to a wide variety of biological threats.



Global Network

15,300,000+
Samples collected

96,000+
Samples sequenced

24,000+
Pathogen genomes sequenced

11
Key international airports (incl. 4 outside US)

44
Collection nodes (cumulative 2025)

Project RANGER: RApid Next Generation sequencing for Effective medical Response

What is Project RANGER?

A consortium of leading bioscience companies has been assembled to develop an innovative pathogen-agnostic diagnostic platform, led by the biosecurity division of Ginkgo Bioworks.

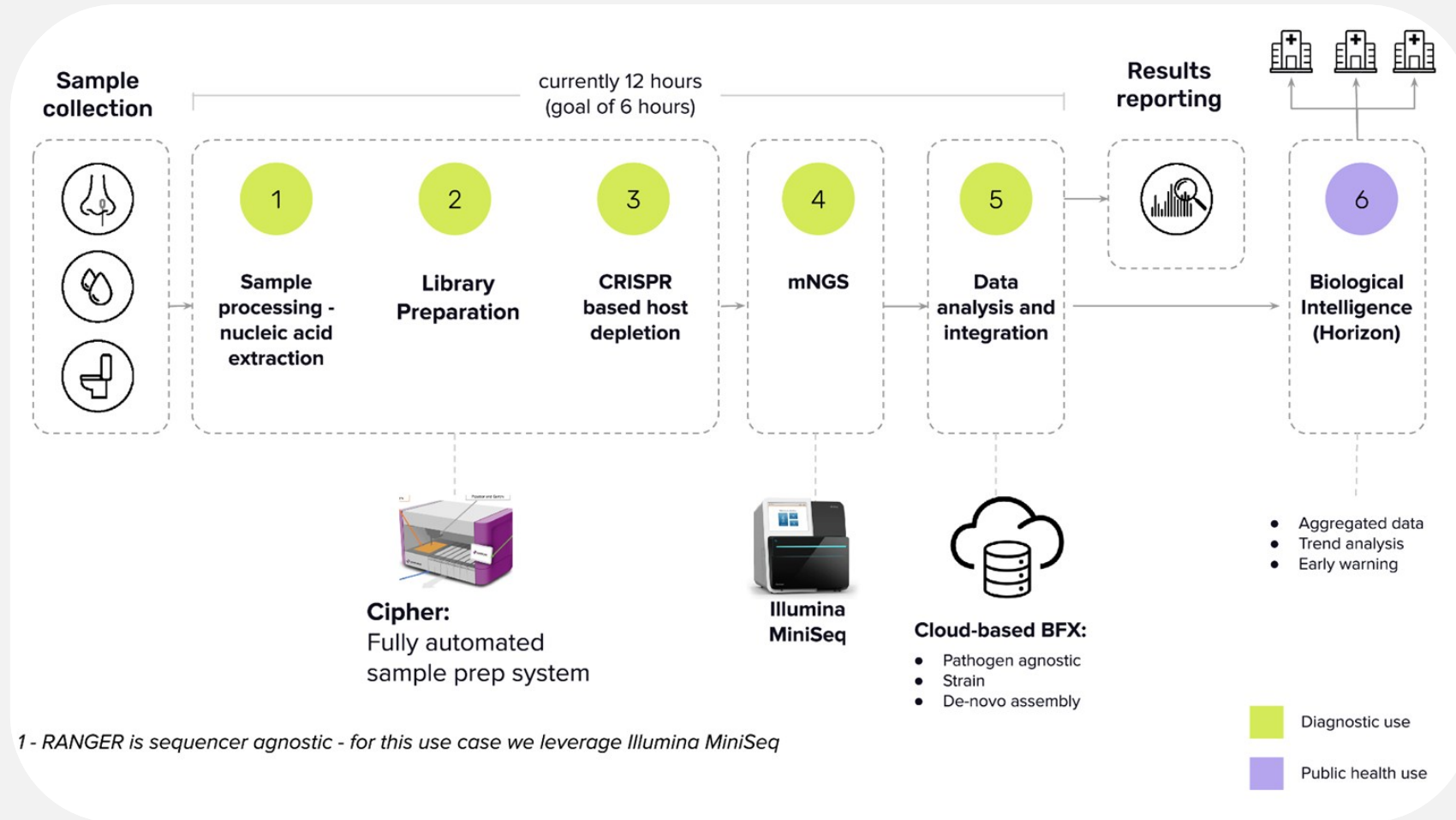
Project RANGER's Purpose:

This collaborative effort brings together industry leaders and cutting-edge technologies to create a comprehensive biosecurity network with global reach.

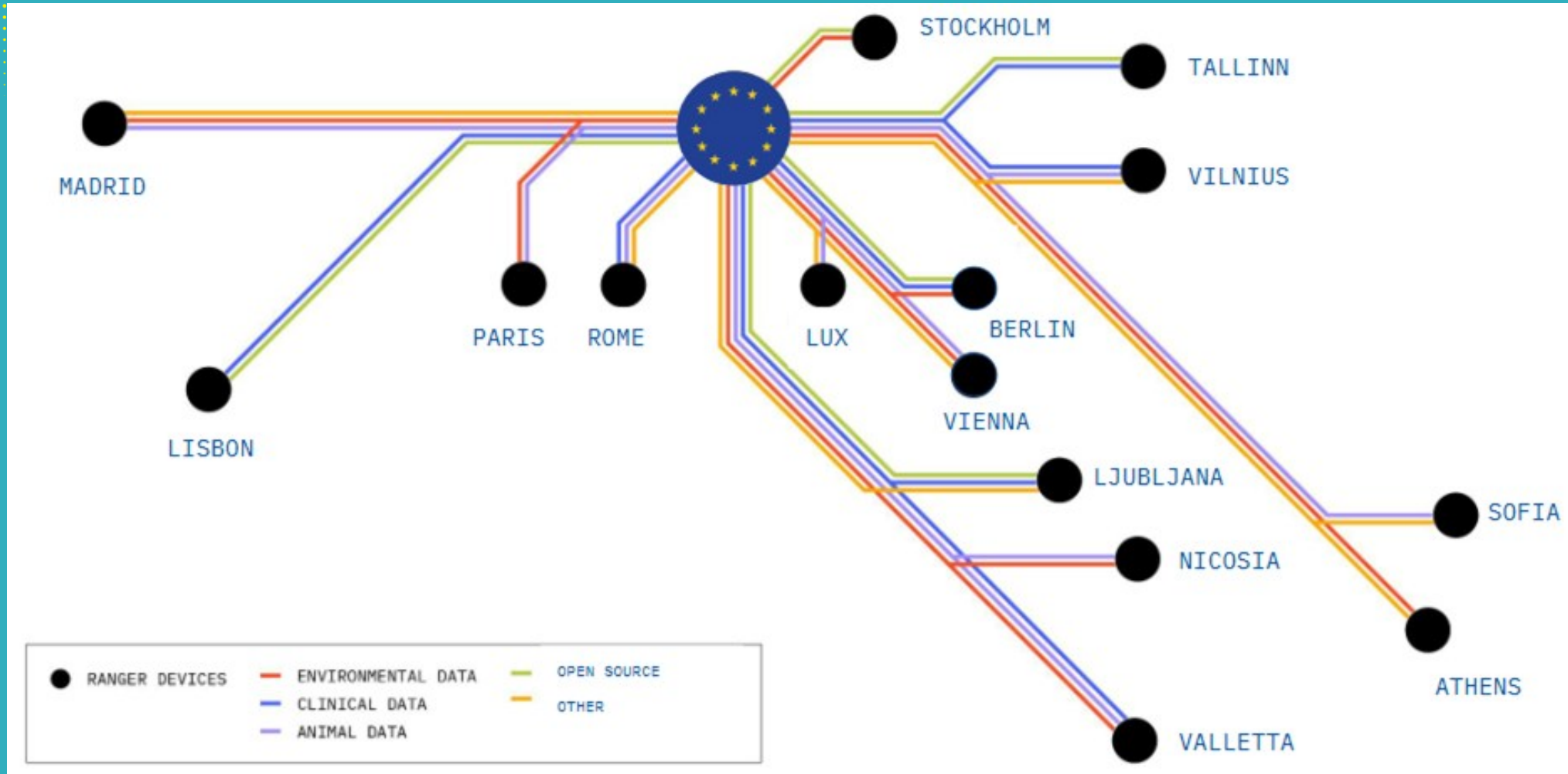


We are developing a system that combines rapid mNGS clinical system with public health data platform

RANGER's objective is to develop and validate a **fully automated, random-access sample preparation instrument** combined with an improved **metagenomic Next-Generation Sequencing (mNGS)** test as a pathogen-agnostic diagnostic for clinical and hospital laboratories.



RANGER's implementation will allow for a widespread mNGS biosurveillance system for the EU



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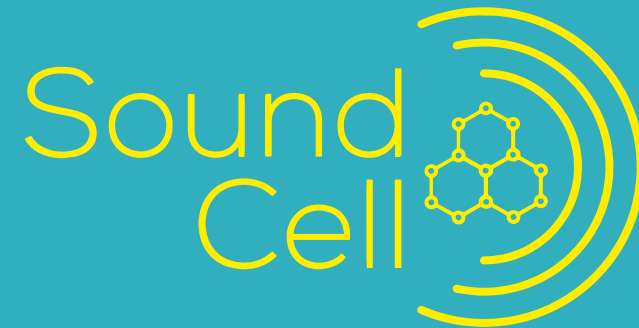
Jeroen NIEUWENHUIS
CEO, Nostics



Christian DROSTEN
DURABLE Project

Graphene single-cell sensors for antibiotic testing with 1hr results

dr. ir. I.E.Rosłoń



Rapid diagnostics. Today.

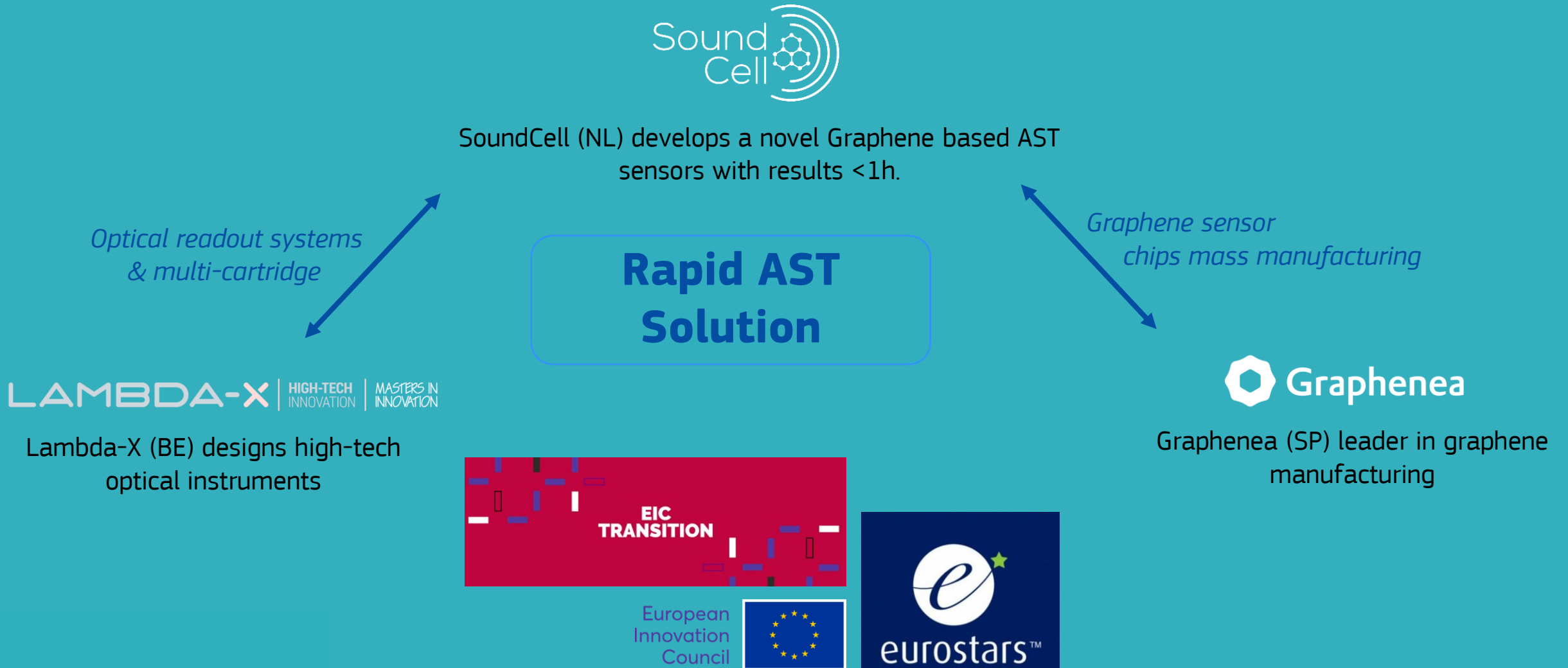


At SoundCell we developed the **MelodyOne**: a rapid AST platform that offers results in one hour.

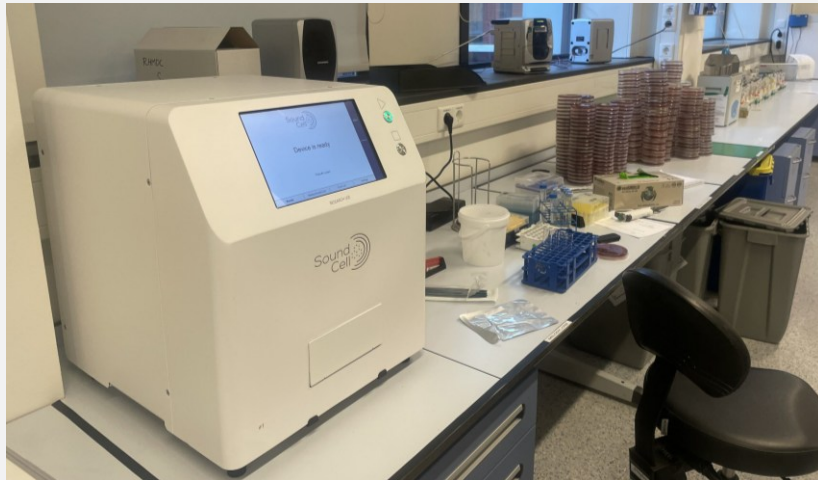
The **MelodyOne** is our cutting-edge nanotechnology platform that detects bacterial resistance within just 1 hour—far faster than traditional methods. Using graphene-based sensors, it measures microscopic bacterial vibrations to determine if antibiotics are working in real time.

- ✓ 1-hour antibiogram from positive blood culture
- ✓ Compatible with bacteria and yeasts
- ✓ Process up to 10 samples a day

EU collaboration towards a solution



The missing piece of the AST puzzle



“Microbiologists urgently need to solve 2 pieces of the puzzle:

- What pathogen am I dealing with?*
- What antibiotic is effective?*

*The first part of the puzzle was solved >20 years ago.
Now, SoundCell is solving the second part of the puzzle.”*

- **dr. Leo Smeets**, microbiologist at Reinier Haga Medical Diagnostic Centre (Delft)

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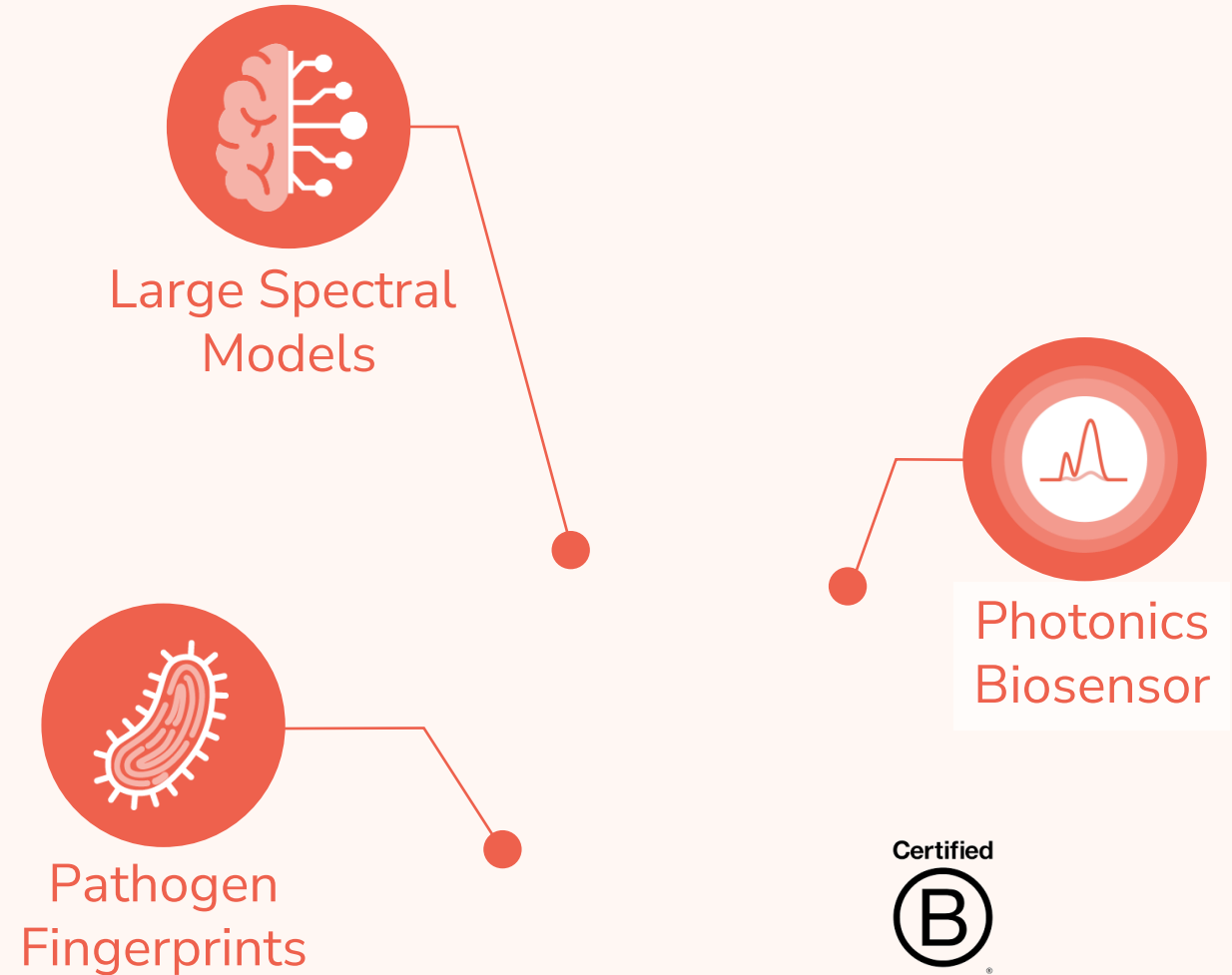
Christian DROSTEN
DURABLE Project



Real-time fingerprinting of living microorganisms for targeted treatment

🚩 Nostics B.V.
Amsterdam, NL

🚩 Nostics Inc.
Cambridge, USA



Solution

Where others need days, we **confirm** infection and **identify** bacteria and fungi in **<10 minutes**

1. Apply sample to cartridge



Easy to use with minimal handling time



2. Automated processing and AI analysis



15 x 15 cm

Rapidly expandable menu by software update, no targets/labels in cartridge



3. Actionable results

Escherichia coli
Klebsiella pneumoniae
Enterococcus spp.
Staphylococcus saprophyticus
Proteus mirabilis
Streptococcus agalactiae
Staphylococcus aureus
Pseudomonas aeruginosa
Enterobacter cloacae

Identification of active infections > living microorganisms

Nostics
guided treatment

Nostics' breakthrough technology

Reading biology as it could not be read before



<10 min directly from samples



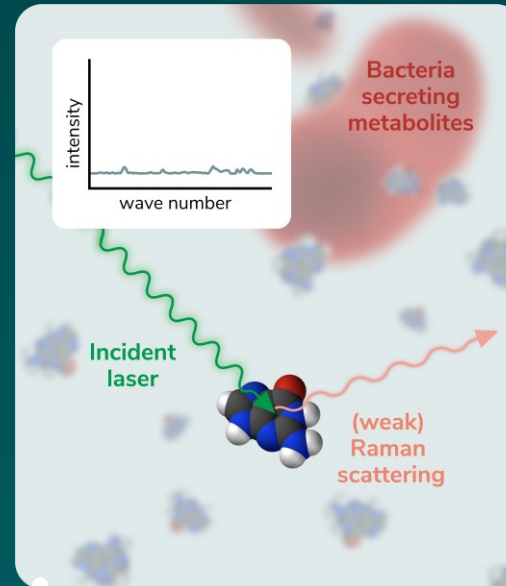
Identification of living microorganisms



Menu expansion by software update

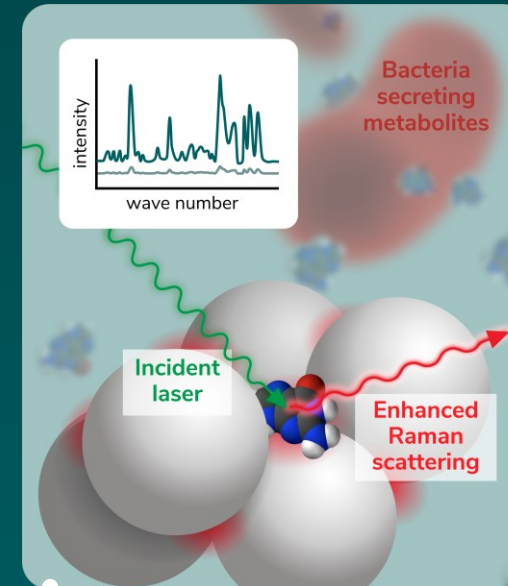


Small footprint and battery-operated



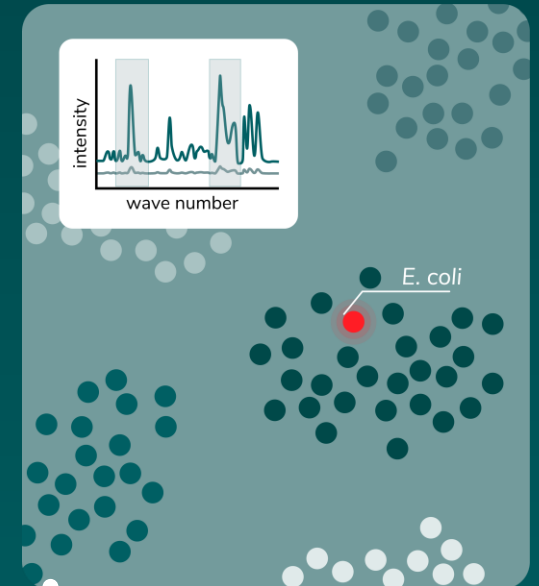
Raman Spectroscopy

Viable pathogens secrete metabolites in a species-specific composition which serve as fingerprints for identification.



Nanomaterials

Nanosubstrate amplifies Raman signals, increasing sensitivity and enabling measurements without culturing.



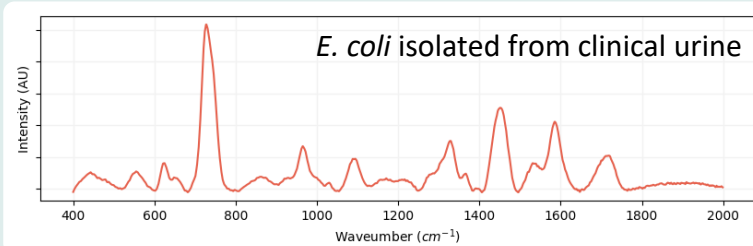
Artificial Intelligence

classification of pathogen fingerprints with proprietary algorithms trained using few-shot learning.

Launch application: Urinary tract infection

Demonstrated capability to expand to further sample types and pathogens

Launch application: Urinary tract infections



UTI panel

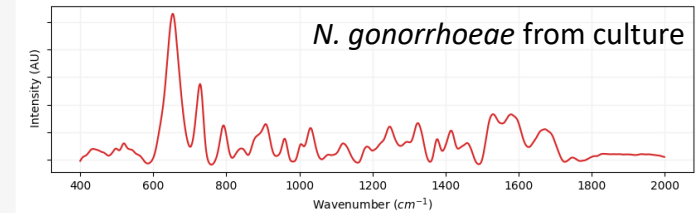
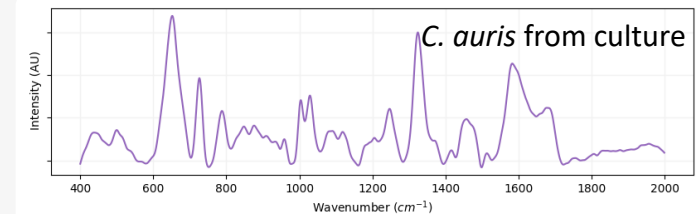
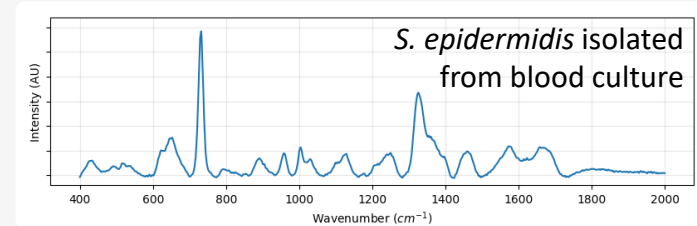
Escherichia coli
Klebsiella pneumoniae
Enterococcus spp.
Staphylococcus saprophyticus
Proteus mirabilis
Streptococcus agalactiae
Staphylococcus aureus
Pseudomonas aeruginosa
Enterobacter cloacae



**Covering > 99%
of all UTI cases
in primary care**

Existing CPT code expedites US launch

Examples of other platform applications:

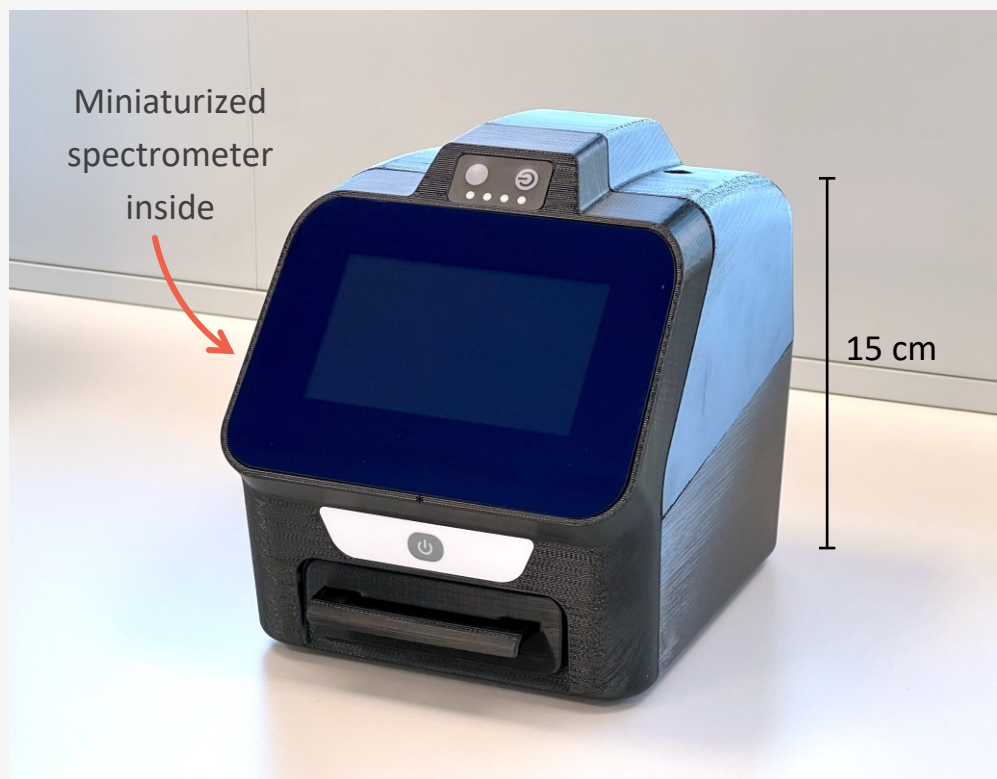


Coming soon

Functional model and high classification performance



Functional model with miniaturized low-cost spectrometer module

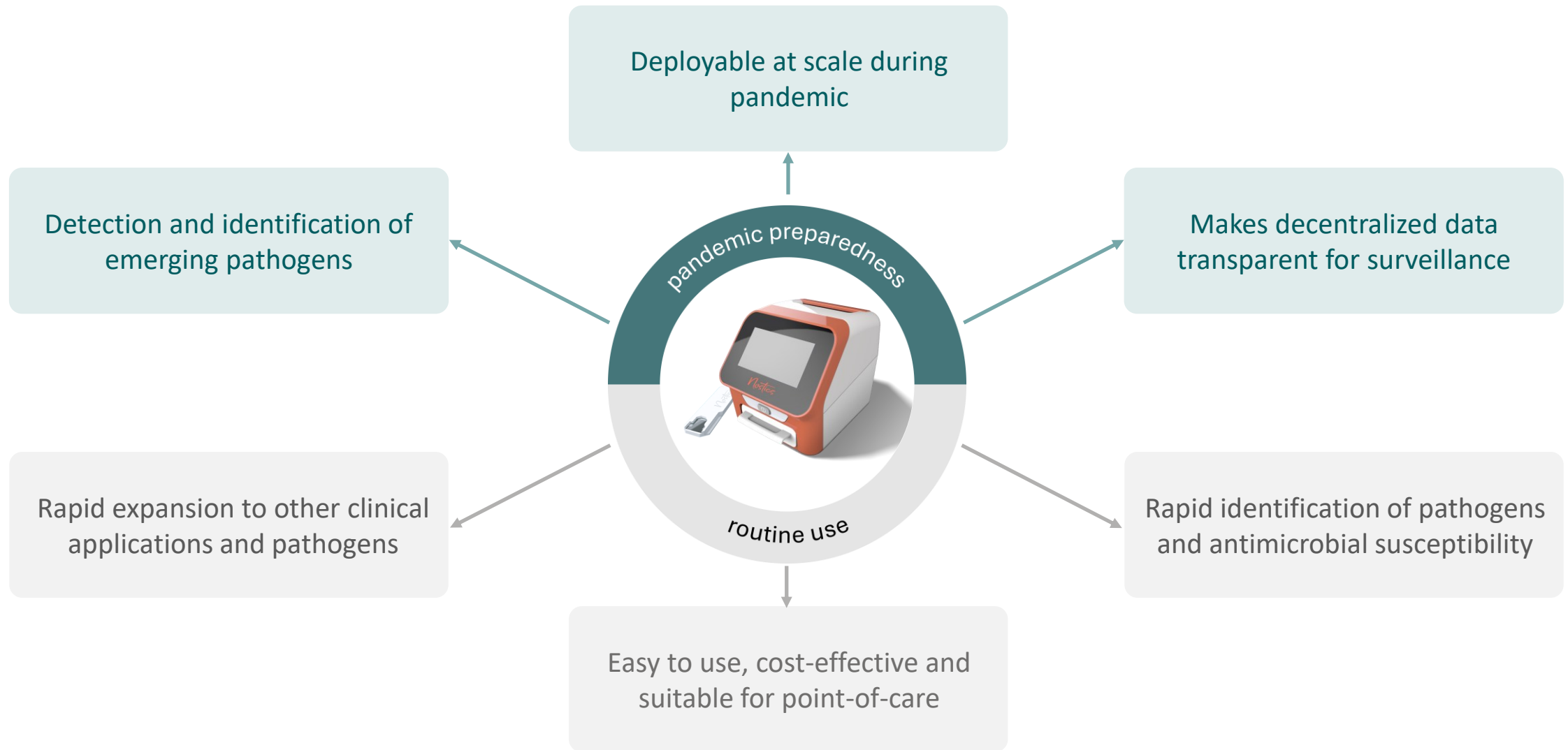


Excellent classification performance on UTI panel with independent datasets

Overall accuracy: 91%
Prevalence weighted: 97%

True	<i>E. coli</i>	<i>K. pneumoniae</i>	<i>P. mirabilis</i>	<i>S. saprophyticus</i>	<i>E. faecalis</i>	<i>E. faecium</i>	<i>S. agalactiae</i>	<i>E. cloacae</i>	<i>P. aeruginosa</i>	<i>S. aureus</i>
	100.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
	11.0	85.3	0.0	0.0	0.0	0.0	0.0	3.7	0.0	0.0
	0.0	0.0	100.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0
	0.0	0.0	0.0	100.0	0.0	0.0	0.0	0.0	0.0	0.0
	0.0	0.0	0.0	0.0	96.3	0.0	0.0	3.7	0.0	0.0
	0.0	0.0	0.0	0.0	5.7	83.3	0.0	11.0	0.0	0.0
	0.0	0.0	0.0	0.0	0.0	0.0	100.0	0.0	0.0	0.0
	6.3	0.0	3.7	0.0	0.0	0.0	17.3	72.3	0.0	0.3
	0.0	0.0	0.0	0.0	0.0	0.0	0.0	0.0	100.0	0.0
	0.0	0.0	3.0	0.0	0.0	0.0	23.7	0.0	0.0	73.3
Predicted										

A decentralized solution to be used as a Medical Counter Measure when needed



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DURABLE Project

HERA/Durable:

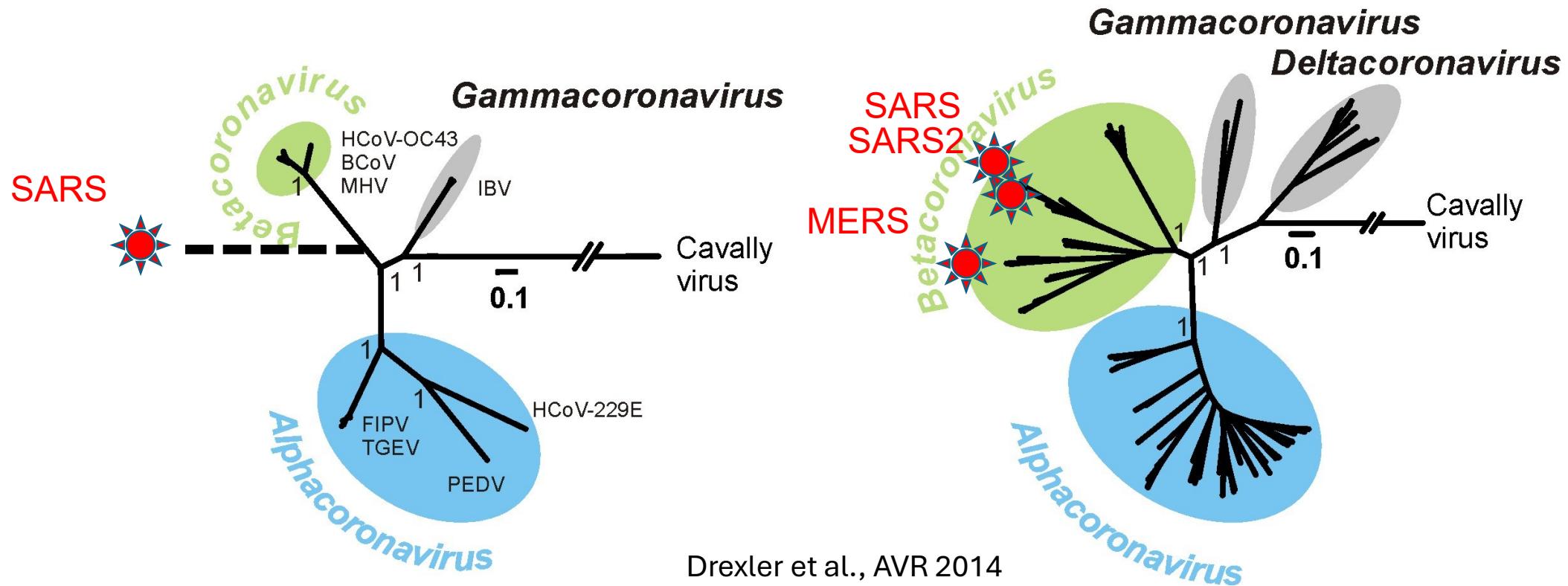
Diagnostics as an immediate
medical countermeasure

Christian Drosten, Charité, Berlin, Germany

Genomic data enables virus recognition

Before SARS

10 years after SARS



Drexler et al., AVR 2014

WH-Human_1|China|2019-Dec
BetaCoV/Wuhan|IPBCAMS-WH-01/2019|EPI_ISL_402123
BetaCoV/Wuhan|IVDC-HB-01/2019|EPI_ISL_402119
BetaCoV/Wuhan|IVDC-HB-04/2020|EPI_ISL_402120
BetaCoV/Wuhan|IVDC-HB-05/2019|EPI_ISL_402121
BetaCoV/Wuhan|WIV04/2019|EPI_ISL_402124
NC_004718 (SARS coronavirus, complete genome)
EU371564 (SARS coronavirus BJ182-12, complete genome)
AY559095 (SARS coronavirus Sin847, complete genome)
FJ882956 (SARS coronavirus ExoN1 isolate P3pp53, complete genome)
FJ882960 (SARS coronavirus ExoN1 isolate P3pp34, complete genome)
FJ882961 (SARS coronavirus MA15 isolate P3pp5, complete genome)
KQ153543 (Bat SARS coronavirus HKU3-8, complete genome)
KY352407 (Severe acute respiratory syndrome-related coronavirus strain BtkY72, complete genome)
NC_014470 (Bat coronavirus BM48-31/BGR/2008, complete genome)

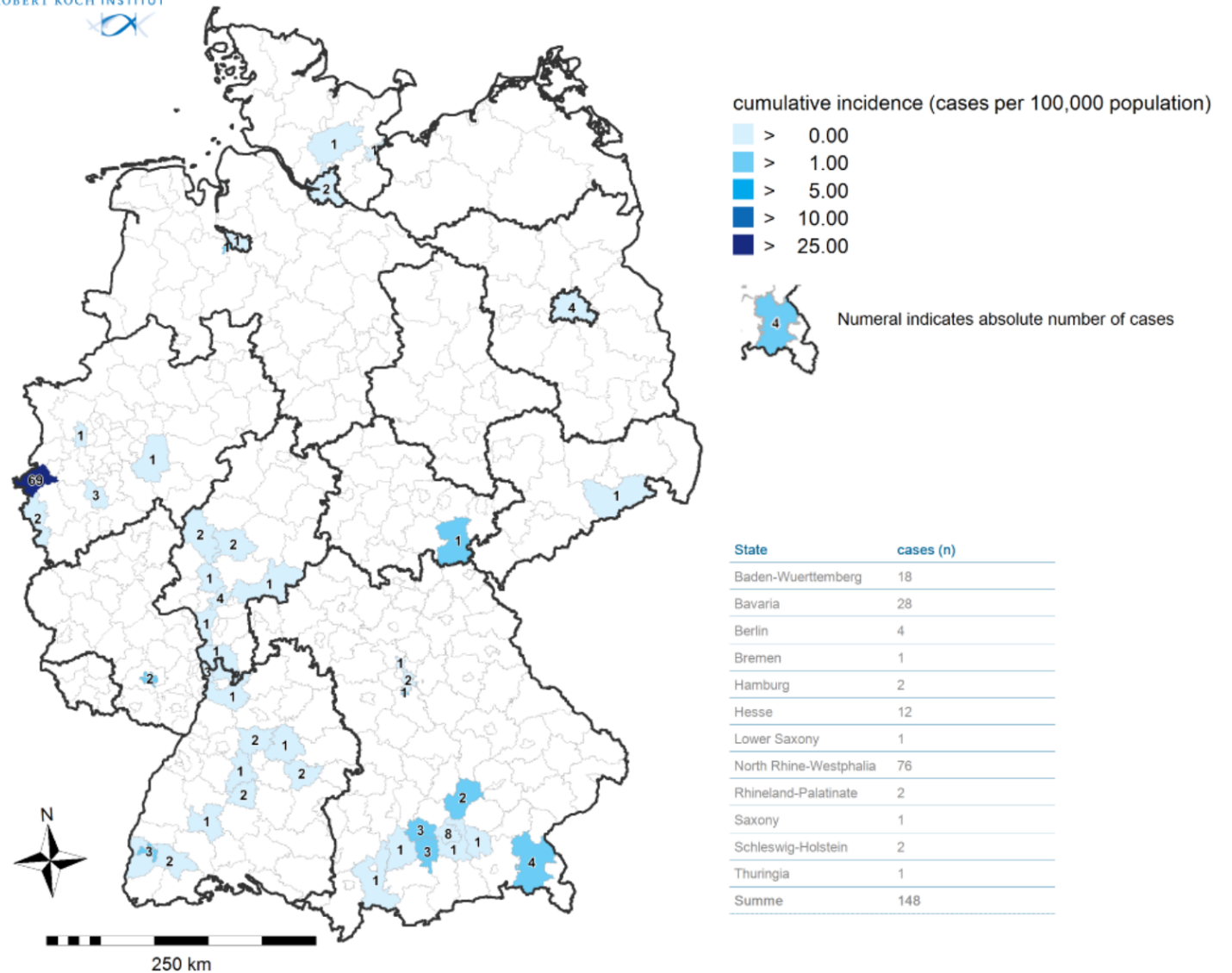
BetaCoV/Wuhan/1PBCAMS-WH-31/2019 [EPI_ISL_452123]
BetaCoV/Wuhan/IVDC HB_01/2019 [EPI_ISL_402115]
BetaCoV/Wuhan/IVDC-IIB-07/2020 [EPI_ISL_402123]
BetaCoV/Wuhan/IVDC-HB-05/2019 [EPI_ISL_402120]
BetaCoV/Wuhan/WIV04/2019 [EPI_ISL_452124]
NC_054113 (SARS coronavirus, complete genome)
DQ322305 (Bat SARS coronavirus HK/3-1, complete genome)
AB585955 (SARS bat coronavirus 3drp gene for RNA dependent RNA polymerase, p...
AB585956 (SARS bat coronavirus 3drp gene for RNA dependent RNA polymerase, p...
DQ412043 (Bat SARS coronavirus Rm1, complete genome)
JX993987 (Bat coronavirus Rp/Shaanxi2011, complete genome)
KF294442 (SARS-related bat coronavirus isolate _longquan_ RNA-dependent RNA po...
KF294453 (SARS-related bat coronavirus isolate Arlung_11 orf1ab polyprotein and ...
KF294455 (SARS-related bat coronavirus isolate Jiyuan-331 orf1ab polyprotein gene, ...
KJ473811 (Bt-RF-BetaCoV/JL20_2, complete genome)
KJ473813 (Bt-RF-BetaCoV/SX2013, complete genome)
KJ473814 (Bt-RS-BetaCoV/HU2013, complete genome)
KU873690 (SARS-related coronavirus isolate F29 RdRp mRNA, partial cds)
MG772846 (Bat SARS like coronavirus isolate bat-SL-CoVDC80 RNA dependent RNA...
MG772849 (Bat SARS-like coronavirus isolate bat-SL-CoVDC84 RNA-dependent RNA...
MG772852 (Bat SARS-like coronavirus isolate bat-SL-CoV22_113 RNA-dependent RNA...
MG772857 (Bat SARS-like coronavirus isolate bat-SL-CoV22_99 RNA-dependent RNA...
MG772870 (Bat SARS-like coronavirus isolate bat-SL-CoV22_76 RNA-dependent RNA...
MG772879 (Bat SARS-like coronavirus isolate bat-SL-CoV22_46 RNA-dependent RNA...
MG772886 (Bat SARS-like coronavirus isolate bat-SL-CoV22_8 RNA-dependent RNA p...
MG772891 (Bat SARS-like coronavirus isolate bat-SL-CoV22_1 RNA-dependent RNA p...
MG772903 (Bat SARS-like coronavirus isolate bat-SL-CoV2K44 RNA-dependent RNA p...
MG772933 (Bat SARS like coronavirus isolate bat-SL-CoV2C45, complete genome)
MG772934 (Bat SARS like coronavirus isolate bat-SL-CoV2XC21, complete genome)
KY352497 (Severe acute respiratory syndrome-related coronavirus strain BtkV72, co...
NC_014470 (Bat coronavirus RM45-31/BGR/2008, complete genome)
KC833195 (Betacoronavirus BtCoV/Whi_fer/117/ITA/2009 RNA-dependent RNA po ym...
KC833200 (Betacoronavirus BtCoV/Whi_fer/117/ITA/2009 RNA-dependent RNA polyme...
KC833202 (Betacoronavirus BtCoV/Whi_fer/BB93-54/BGR/2009 RNA-dependent RNA...
KC833203 (Betacoronavirus BtCoV/Whi_fer/BB93-58/BGR/2008 RNA-dependent RNA...
KC833206 (Betacoronavirus BtCoV/Whi_fer/117/ITA/2009 RNA-dependent RNA po ym...
KC833211 (Betacoronavirus BtCoV/Whi_hip/R37_09/SPA/2010 RNA dependent RNA p...
KC833213 (Betacoronavirus BtCoV/Whi_hip/R46_03/SPA/2010 RNA dependent RNA p...
KC833220 (Betacoronavirus BtCoV/Whi_hip/Slo52/XS_02/2009 RNA-dependent RNA pol...

[illegible]

Corman et al,
Eurosurv 2020

Germany:

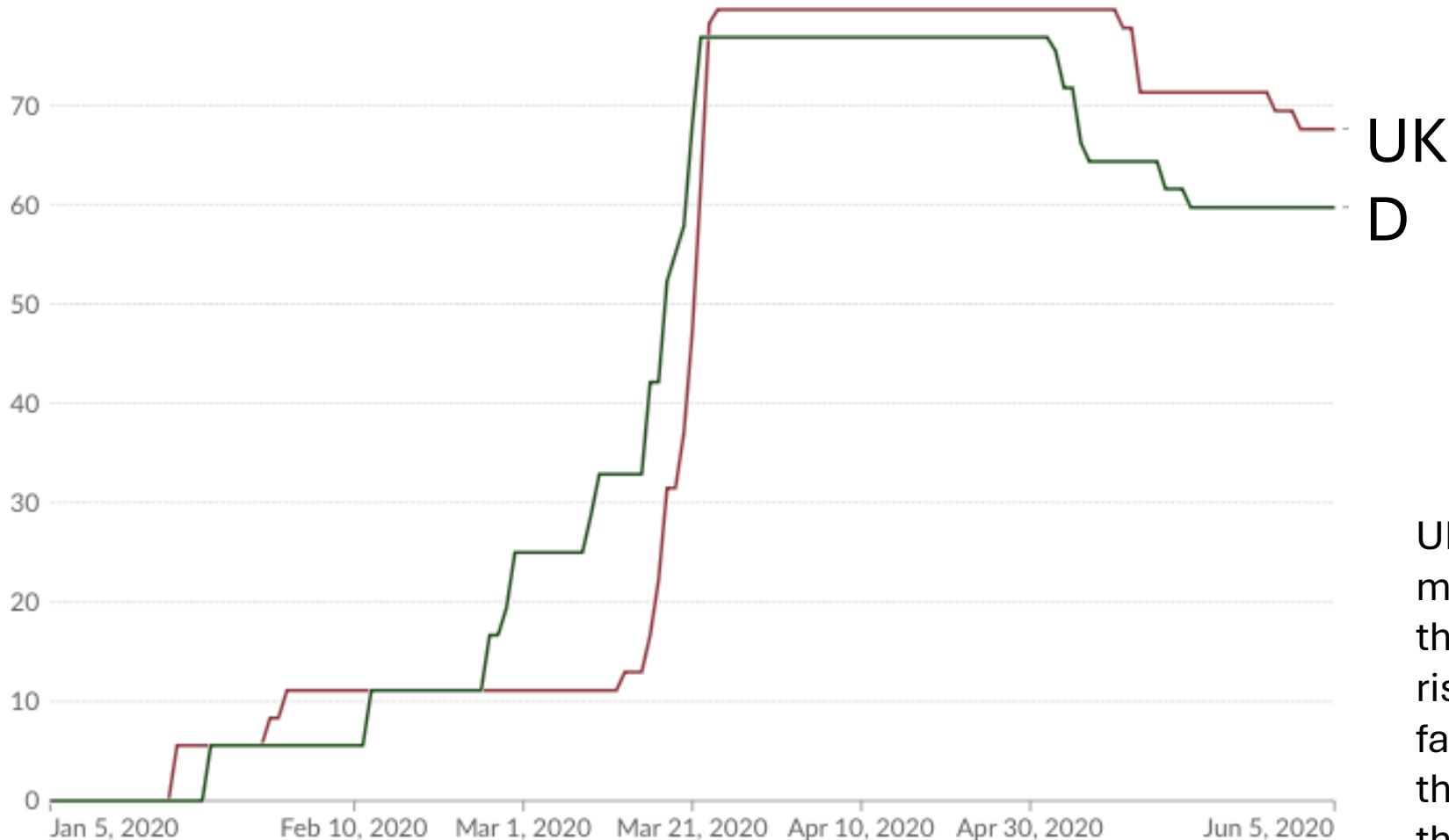
Very early, very broad
availability of
diagnostic testing



COVID-19: Stringency Index

The stringency index is a composite measure based on nine response indicators including school closure, travel bans, rescaled to a value from 0 to 100 (100 = strictest).

Table Chart



UK vs D:
NPI later, stricter, longer

UK initiates contact measures ca. 3 weeks later than Germany (incidence rise from mid February, first fatal case 5th March, more than 100 confirmed cases by that time)

Contact measures during 1st wave, D vs UK

COVID lethality, Germany	111.25 / million pop.
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First wave, Germany (COVID as primary cause of death, through July 2, 2020)

Same for UK	832.47 / million pop.
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Absolute death toll in Germany (Lethality (per million) X 84 million population)	9345
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Death toll assuming UK lethality	69927
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European Centre for Disease Prevention and Control

An agency of the European Union

Evolution of Disease and Laboratory Networks

EVD-Labnet

COVID-19 and Influenza networks

EU Reference Laboratories for Public Health
(EURL) mechanism



**Preparedness
research network
of
20 leading
scientific
institutions**

**located in
15 EU countries**

Challenges

Professional networks connecting research and public health laboratories

IVDR / validation efforts, requirement of formal clinical studies framework

Role for academic consortia in regulation and clearance processes (from samples to cohorts)

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Thank you