

Coronavirus 19

Understanding Breath Aerosol Transmission Risk

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Northumbria
University
NEWCASTLE

Overview

- **Breath for Coronavirus screening.**
 - Scientific evidence.
 - Key unknowns.
- **Exhaled Breath Diagnostics.**
 - Opportunities.
 - Challenge.
 - Competition.
- **The PBM-HALE™ approach.**
 - Platform IP.
 - Supporting key data.
- **The proposition.**



Coronavirus 19 (COVID-19) in breath: Confidence in rationale



Biology

- SARS-CoV-2 binds ACE2 receptor.¹
- ACE2 protein levels highest in lower lung.²
- Aerosols (<5 μm) best to reach lower lung (drug delivery science).³

Pathology

- Disease of the lower lung: respirator need.
- Proposed transmission routes: fomites, droplets (cough, >5 μm), but:
 - Models & data⁴ show transmission without symptoms (no cough!).
 - Aerosol science used in epidemiology out of date.⁵

Coronavirus 19 (COVID-19) in breath: Confidence in rationale



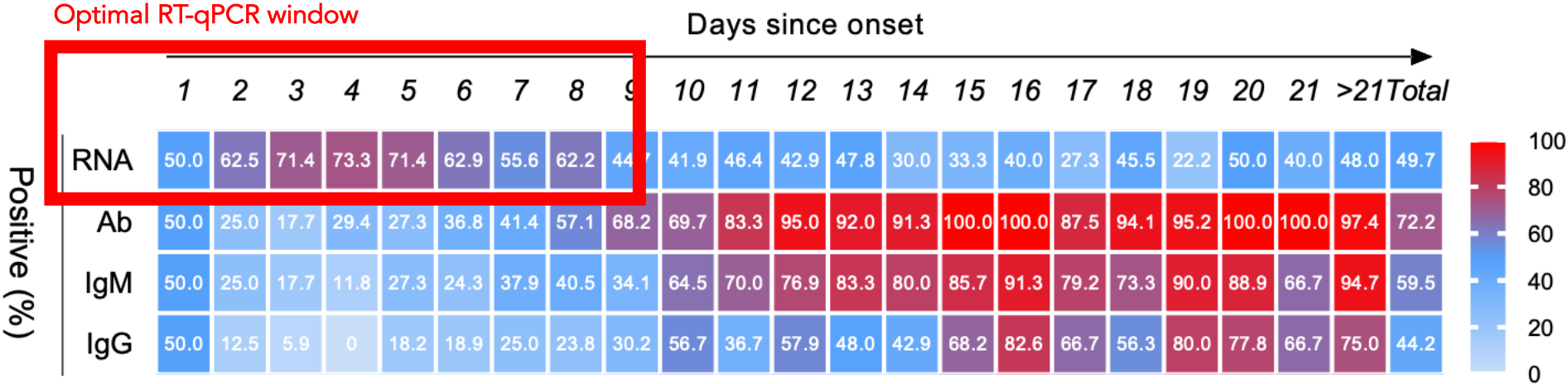
Clinical evidence

- Transmission occurs up to 1 week before symptoms (peak @ -2.9 days).¹
- Virus genome levels max in lower lung samples > nose > throat.²⁻⁵
 - <42% false negative oral swabs.
 - 10-25% false negative nasal swabs.
 - Viable virus levels low in nasal swabs.
 - Nasal detection ~70% days 0-5 from symptoms⁶

Experimental evidence:

- COVID-19 ward aerosol gel traps –ve, but ceiling air vents +ve:⁷ Droplets pulled by gravity, aerosols pulled by air flow.⁸
- Aerosolised virus infectious for 16hrs after mechanical generation.^{9, 10}
- Other coronaviruses can naturally aerosolize (n=3000).¹¹

Coronavirus 19 (COVID-19) in breath: Confidence in rationale



RNA	2	8	14	15	21	35	27	37	38	31	28	35	23	20	18	20	11	11	9	6	5	25	439
Ab	2	8	17	17	22	38	29	42	44	33	30	40	25	23	24	25	24	17	21	9	6	39	535
IgM	2	8	17	17	22	37	29	42	44	31	30	39	24	20	21	23	24	15	20	9	6	38	518
IgG	2	8	17	17	22	37	28	42	43	30	30	38	25	21	22	23	24	16	20	9	6	36	516

Sample no.

Detection of SARS-CoV-2 in nasal swabs (RNA) or blood (IgX) from symptom onset

Coronavirus 19 (COVID-19) in breath: Confidence in rationale



Our hypothesis

- Disease is a function of amount of virus reaching the lower lung.
- Achieved mainly by breath aerosols (or poor immune system).
- Explains close contact transmission chains.

We need to test breath aerosols for:

- *The amount of virus present (genomes).*
- *Infectivity (viruses).*

Diagnosing from Exhaled Breath Condensates (EBC)



Breath is 95% hydrated:

- Volatile compounds (smells, eg garlic, alcohol).
- Vapour & aerosols.
- Biological molecules.

Health and Disease indicators:

- Lung infections.
- Liver diseases.
- Multiple cancers:
 - Blood.
 - Breast.
 - Brain.

Challenges to clinical use

- Reproducibility.
- Contamination:
 - Saliva.
 - Ambient.
- Sample loss.
- Safety.
- Upper vs deep lung separation.

RTube™



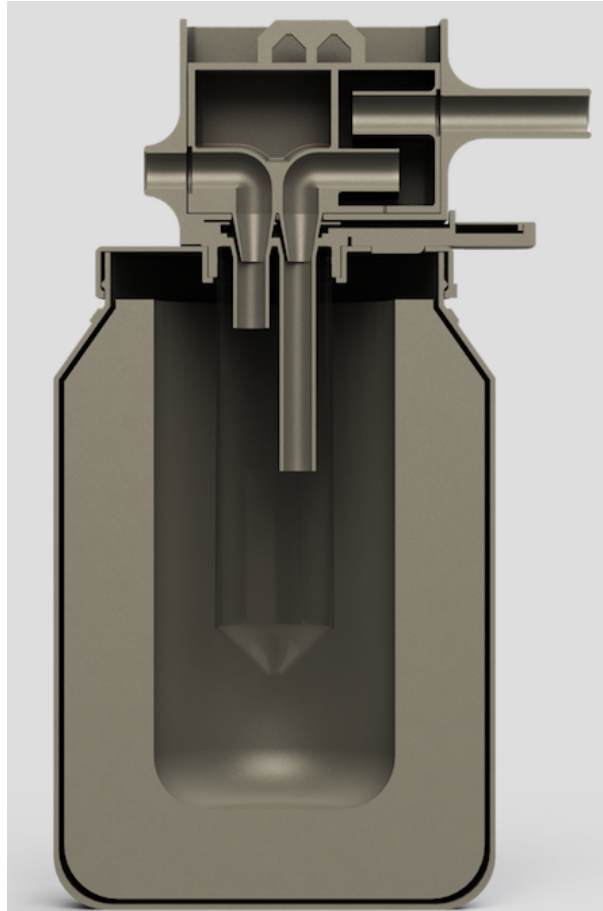
Poor process control

EcoScreen™



Sample lost in black tube
17Kg + weight

PBM-HALE™: the platform



EBC collector:

- Volatiles and
- Proteins.
- DNA.
- RNA.
- Lipids.
- Medications

Solves key problems:

- Reproducibility.
- Contamination.
- Sample loss.
- Safety.

Cold Chain Dependent:

- Uses dry ice powder (CO₂) to collect sample reliably.
- Dry ice replenished every 1 hr from compressed gas cylinder.
- Sample needs on the spot test or frozen transfer to lab.

PBM-HALE™: the platform



PBM-HALE™

EBC collector:

- Volatiles and
- Proteins.
- DNA.
- RNA.
- Lipids.
- Medications.

Solves key problems:

- Reproducibility.
- Contamination.
- Sample loss.
- Safety.

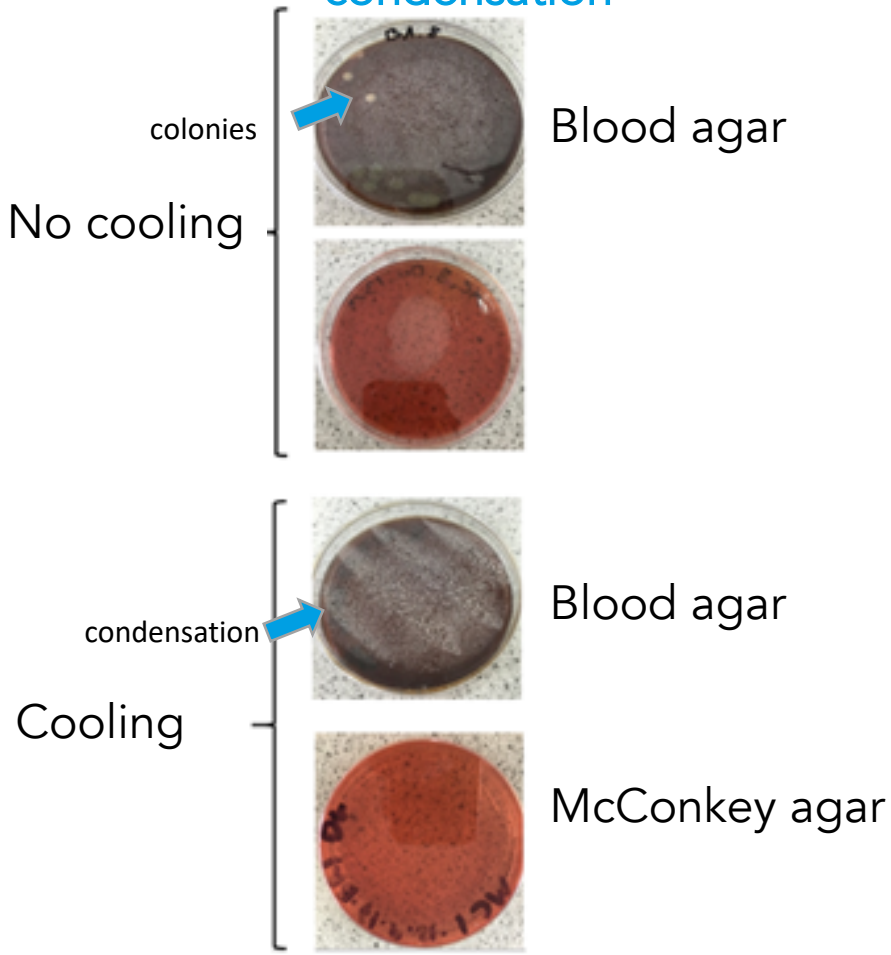
Path to removing the cold chain:

- Proprietary coating to remove need for dry ice.
- Stabilisation material to remove freezer storage.

Experiments under way

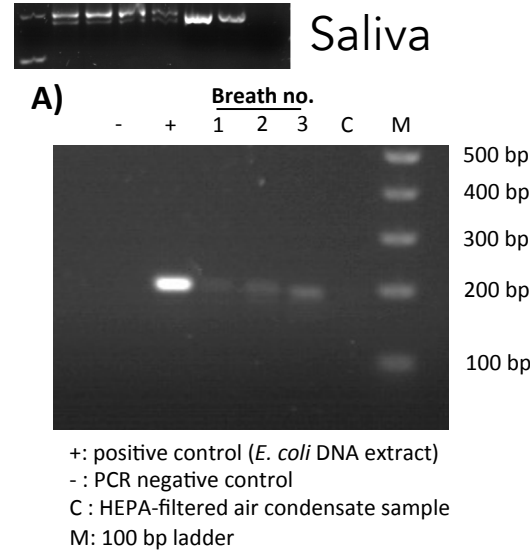
Preliminary data: pathogen DNA

Bacteria & fungi die by dry ice condensation

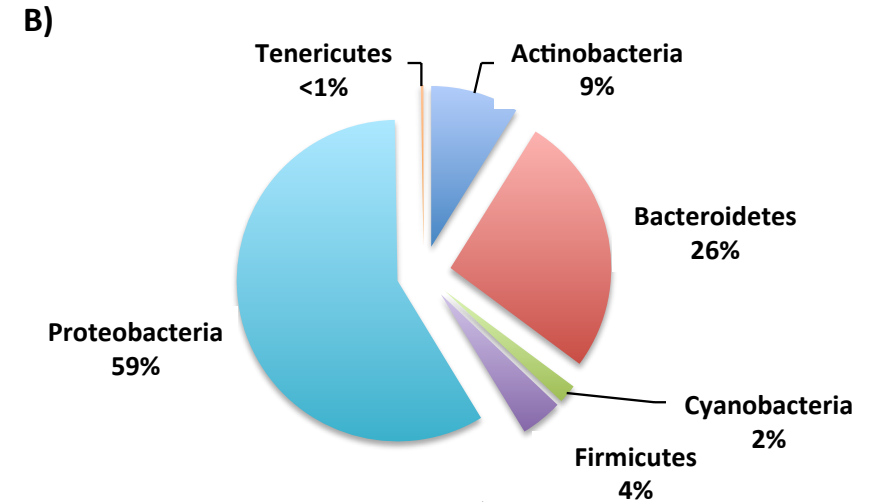
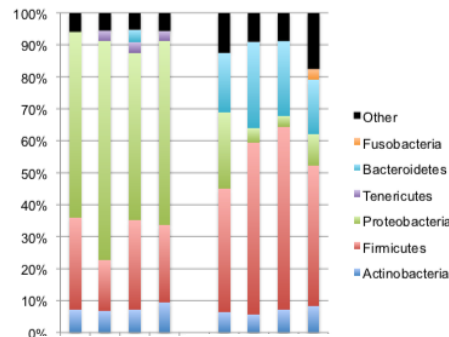


NB: no colonies on plates, white marks are condensation

COVID-19-like test works with 1-3 breaths for bacteria

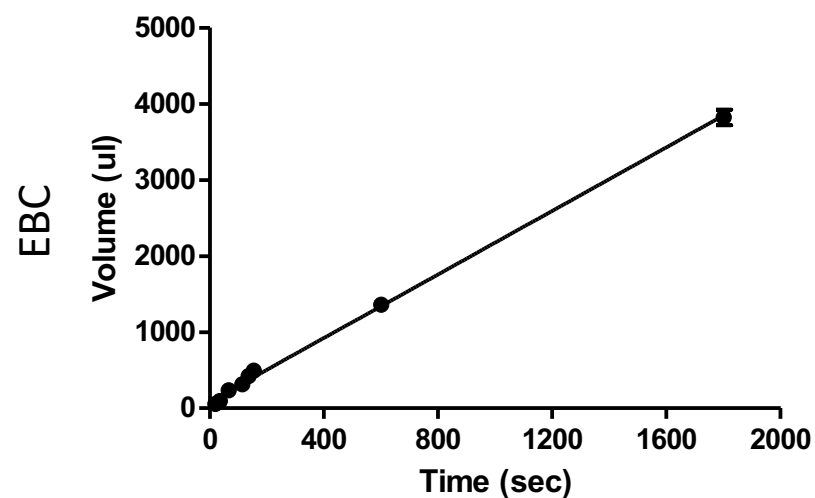
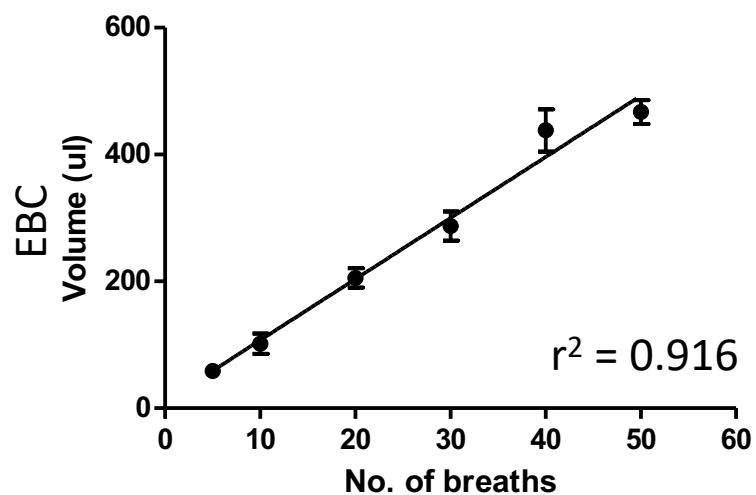


EBC is distinct to saliva



Closest sample like this comes from lung surgery samples only (60% proteobacteria)
 (Sze MA et al. Am. J. Resp. Crit. Care Med. 2012)

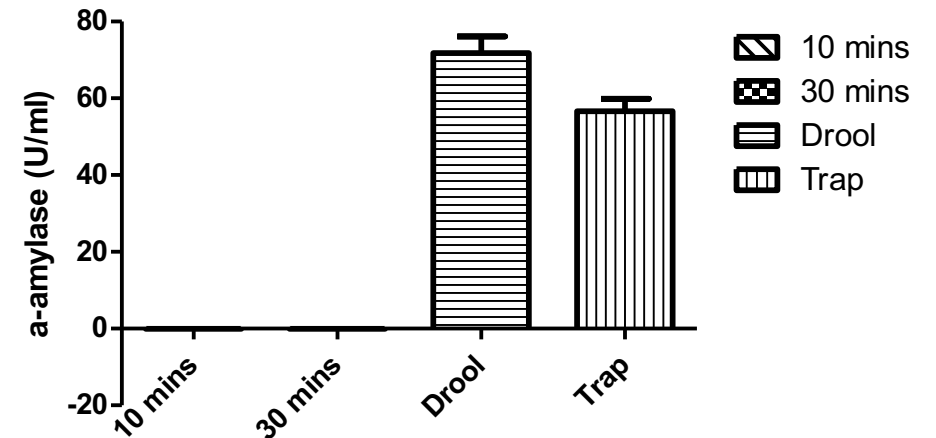
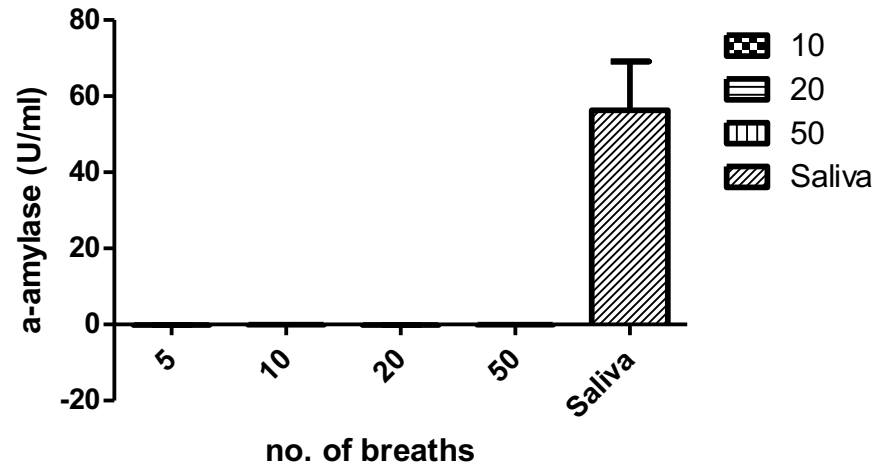
Prototype: highly consistent sampling



Whether 5 breaths (25 sec; e.g. screening) or 30 min of sampling (e.g. discovery)

R^2 range: 0.88 to 0.95, $n = 5$.

Prototype: no salivary contamination

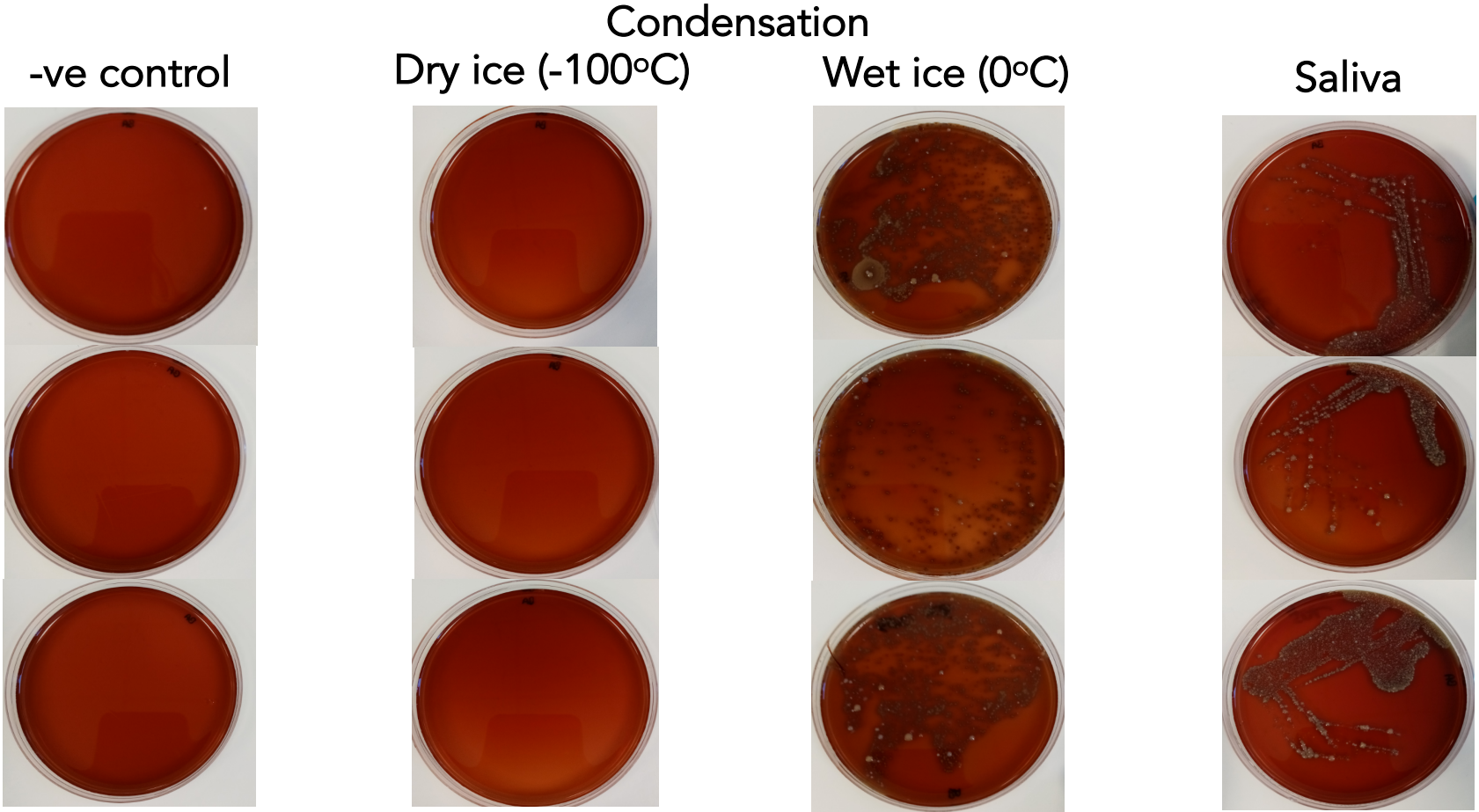


Saliva enzyme levels below limit of detection in EBC:

At least 5000x less in EBC than in saliva (drool) or device saliva trap levels even after 30 min sampling.

n = 5.

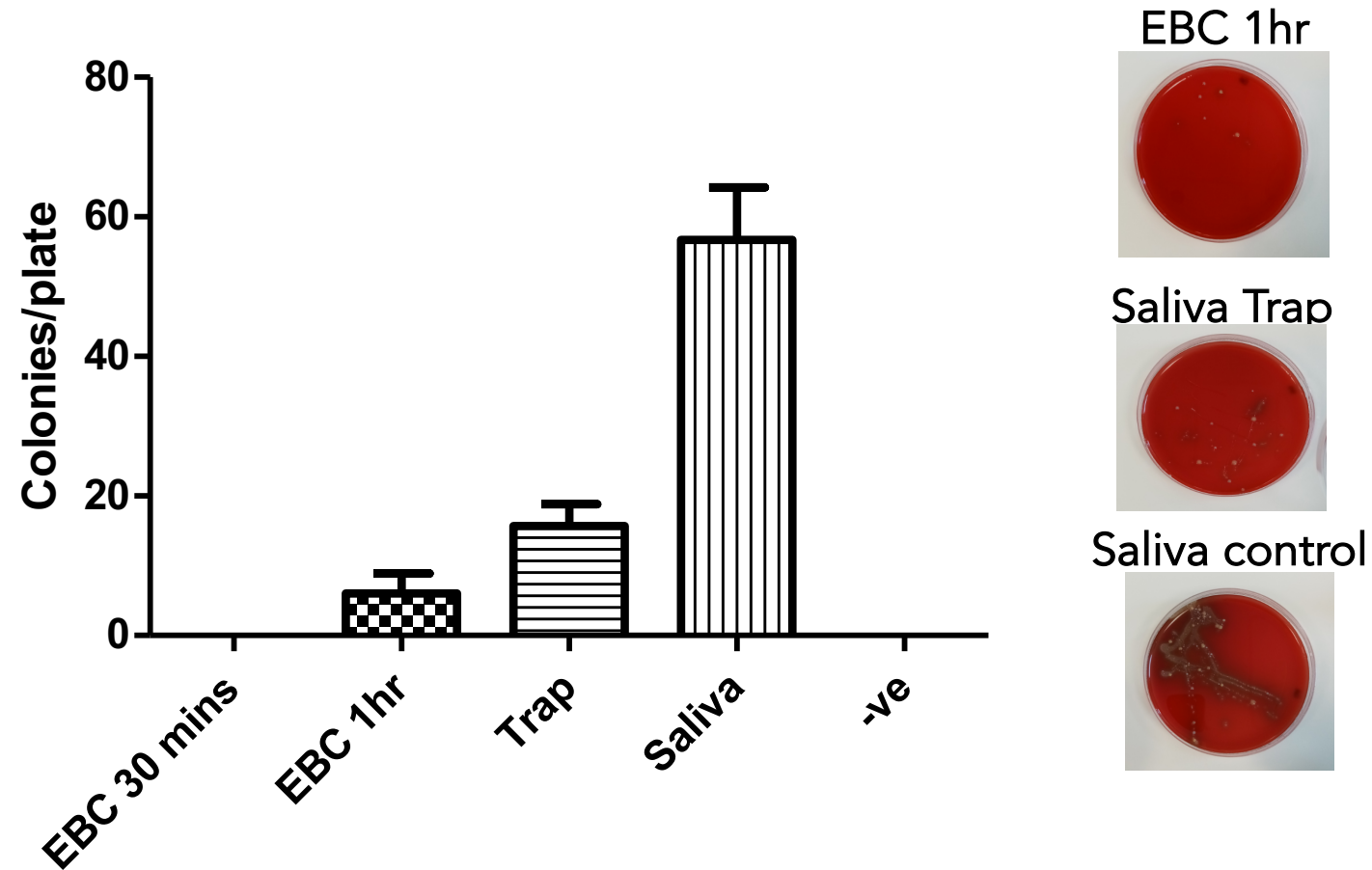
Prototype: No microbial growth due to dry ice condensation (blood agar).



2 min sampling period
(2x target sampling period for COVID-19 screening use).

n = 5

Prototype: microbial growth only after loss of dry ice cooling efficacy

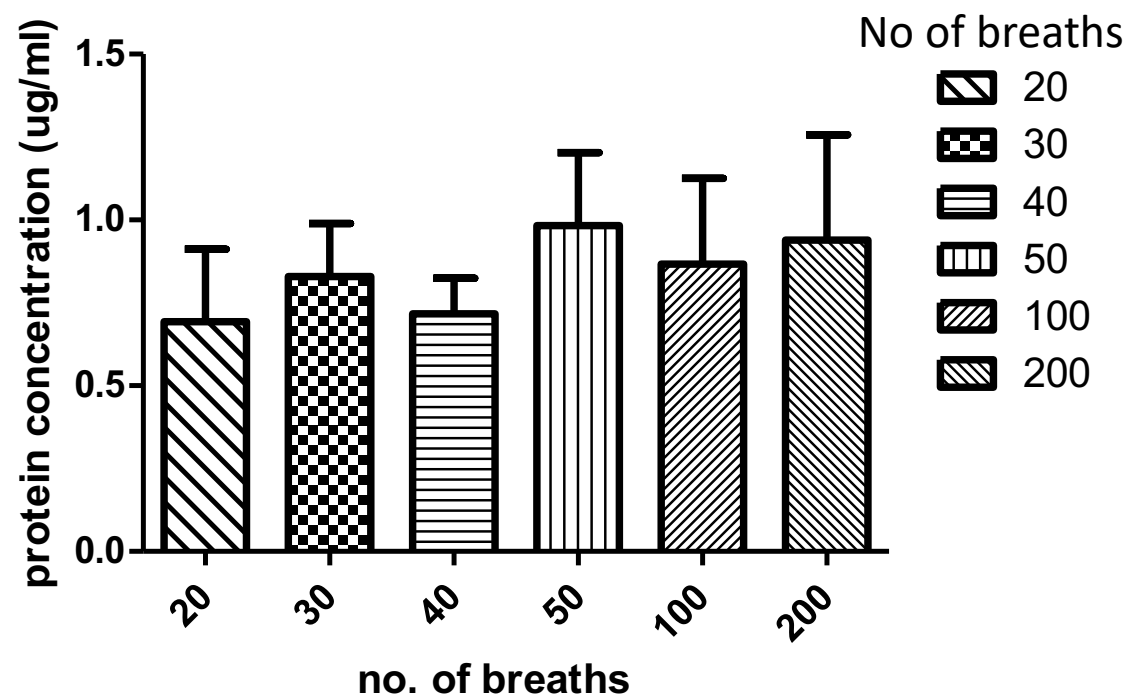


Colonies from dry ice-captured EBC cultured on blood agar.

Lateral contact of sampling tube to dry ice lost ~40 min after continuous sampling.

n = 3.

Prototype: consistent [protein] in EBC

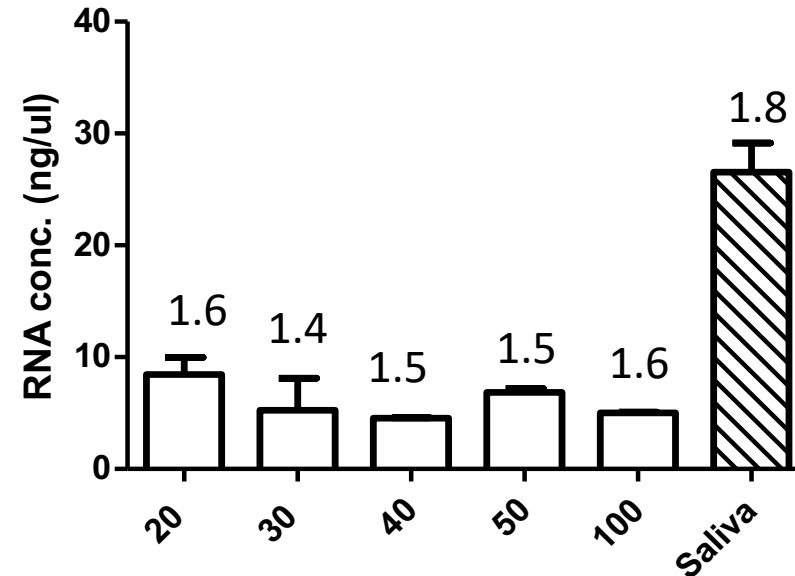


Samples lyophilized and re-constituted in 1/5th of original volume:
No statistically significant difference in concentration over time by micro BCA (data close to LLOD).

No concentration increase anticipated.

n=5

Prototype: consistent [RNA] in EBC



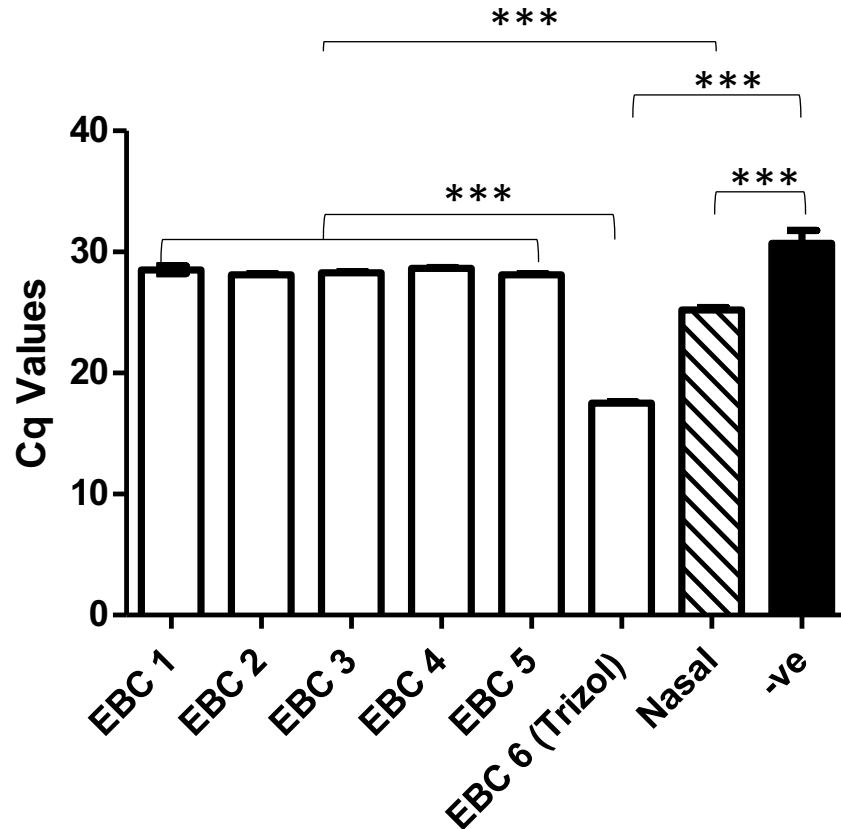
EBC breaths, n=6

EBC 30-100 normalized to 20 breath
sample volume, Trizol extraction

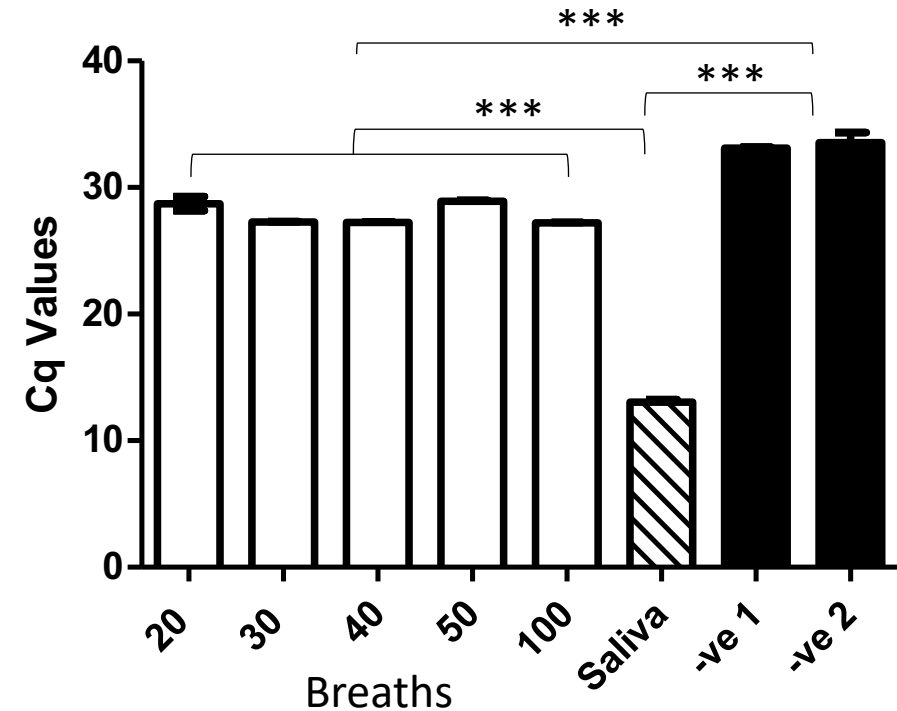
No statistically significant difference in concentration.

260/280 ratios reported per column

Prototype: 18S by PCR in EBC RNA

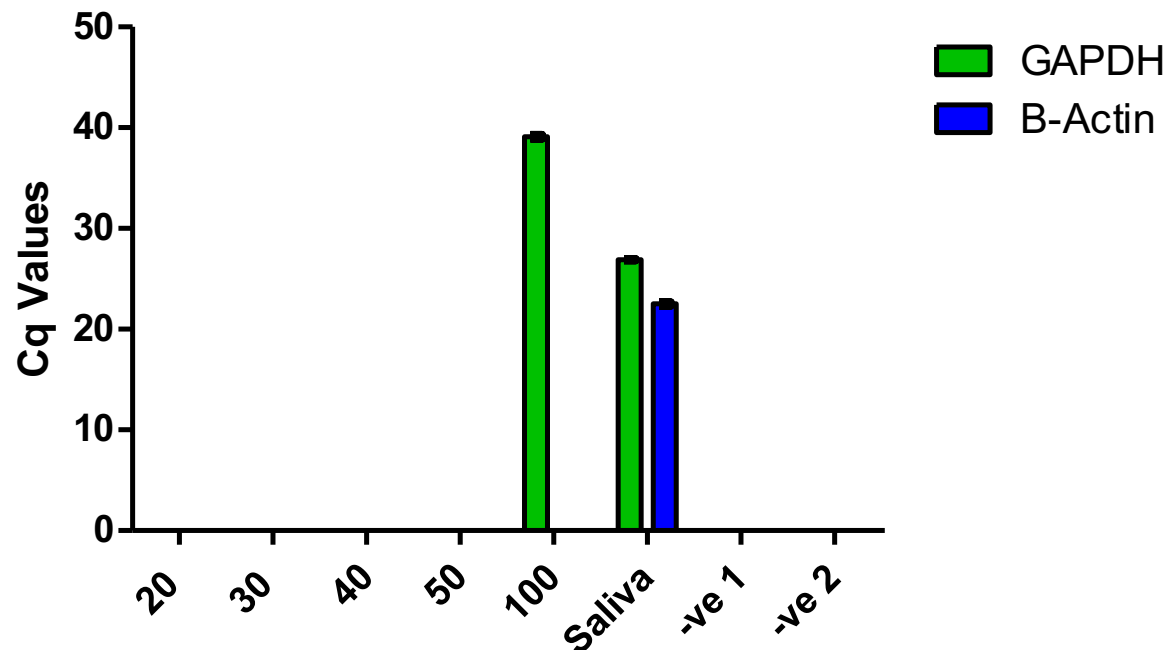


2 step SYBR Gold RT-qPCR (triplicate)
EBC1-5: RNeasy kit 20 breaths
EBC6: Trizol 30 min sample
Nasal swab.



2 step SYBR Gold RT-qPCR (triplicate)
-ve 1: No RT control
-ve 2: no cDNA
EBC 30-100 normalized to 20 breath sample volume
P<0.01 Saliva vs EBC, EBC vs -ve, Saliva vs -ve.

Prototype: GAPDH & β -actin in EBC RNA



2 step SYBR Gold RT-qPCR (triplicate)

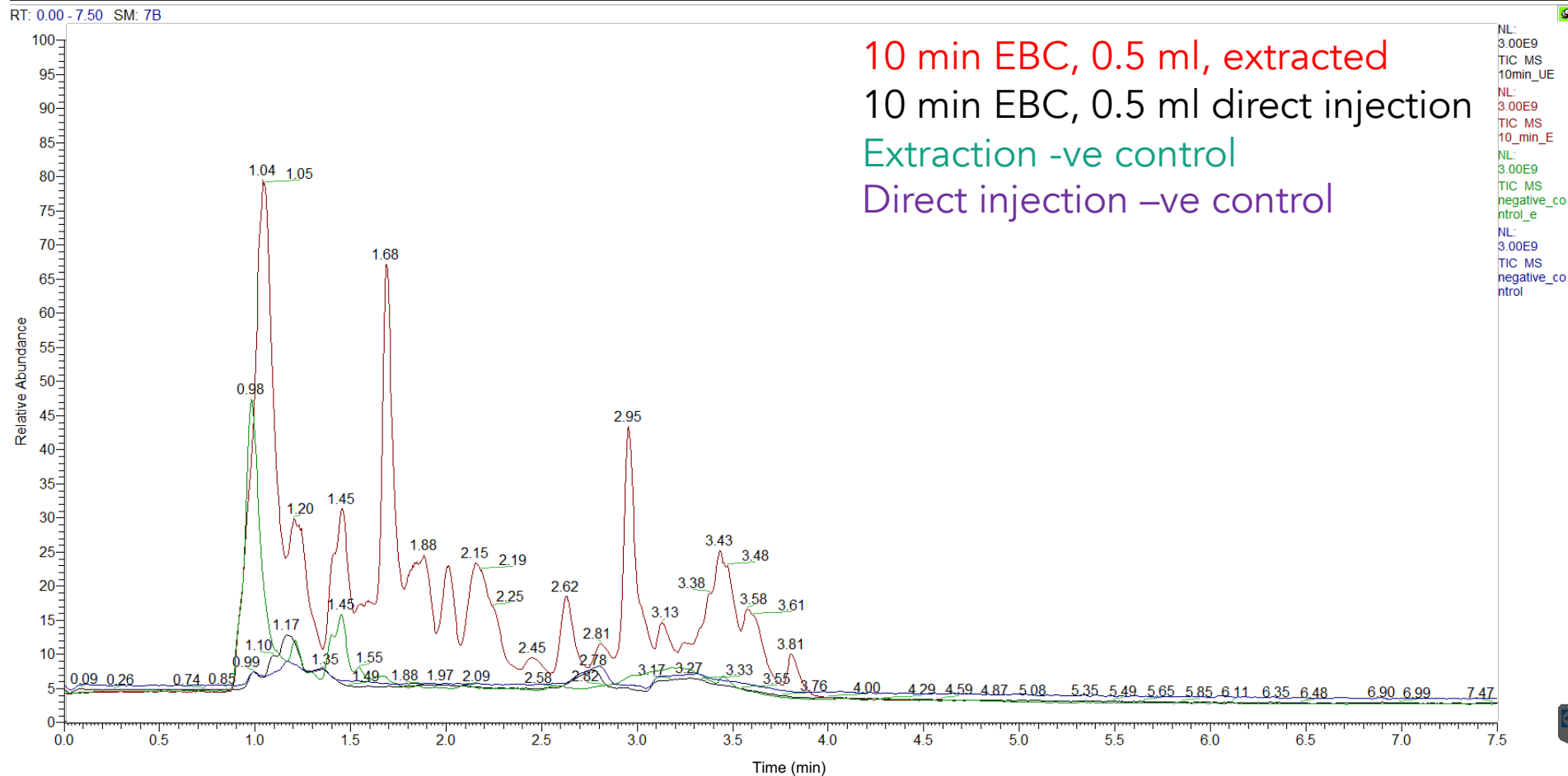
-ve 1: No RT control

-ve 2: no cDNA

EBC 30-100 normalized for EBC 20 sample volume

- EBC not classified as human tissue.
 - Human Tissue Act 2004.
 - Cells, DNA, or RNA.
 - EBC explicitly excluded.
- Early data: EBC is 18S+
 - Validation round under way with Taqman® assay.
 - ITS SEQ run planned (human vs fungal).

Prototype: Metabolomics in EBC



Metabolite profile in EBC after 5x concentration by lyophilization

n=5

Data generated at the Northumbria University Metabolomics Core Service

Prototype: Metabolomics in EBC



Compound	RMM (g/mol)	RT [min]	Relative ion abundance
1-hexadecyl-glycero-3-phosphate	396.3	1.002	810,094
monoacylglyceride	352.3	1.02	281,866
LysoPA	410.2	1.032	968,316
Palmitoleylethanolamide	297.3	1.047	187,282
eicosatetraenoate	335.2	1.054	348,544
Linoleamide	279.3	1.061	216,809
Cuscohygrine	224.2	1.067	723,759
N-Decanoylglycine	229.2	1.156	2,612,124
N-Nonanoylglycine	215.2	1.198	1,942,872
cis-3-Hexenyl b-primeveroside	394.2	1.221	160,089
N-Lauroylglycine	257.2	1.923	286,977
N-Undecanoylglycine	243.2	2.072	227,826
phosphatidylethanolamine	837.5	2.388	381,518
Gambogic acid	628.3	2.536	416,778
2-Hexenoylcarnitine	257.2	3.062	994,821
L-argininium	175.1	3.367	502,141
N-Acetylputrescine	130.1	3.519	192,382

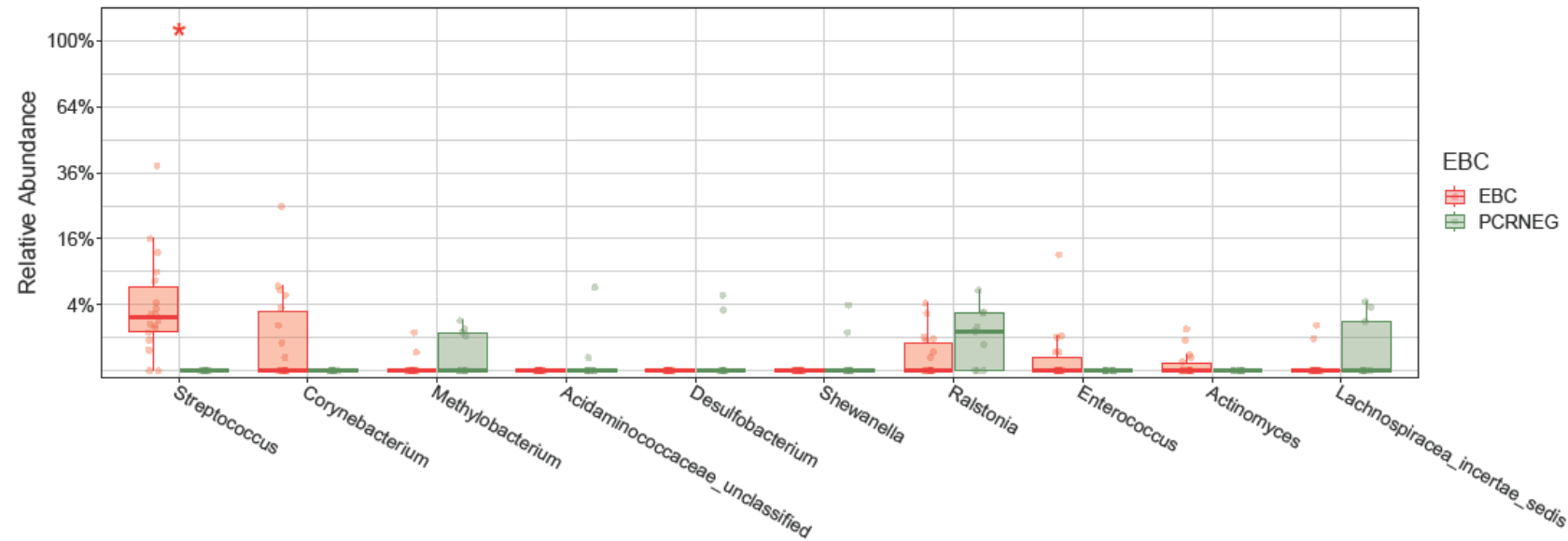
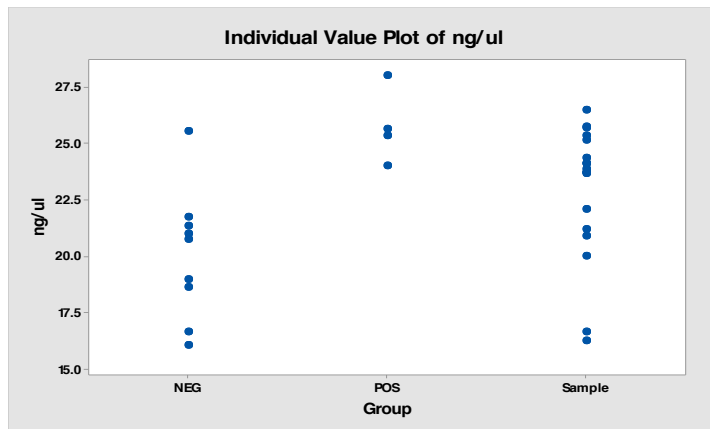
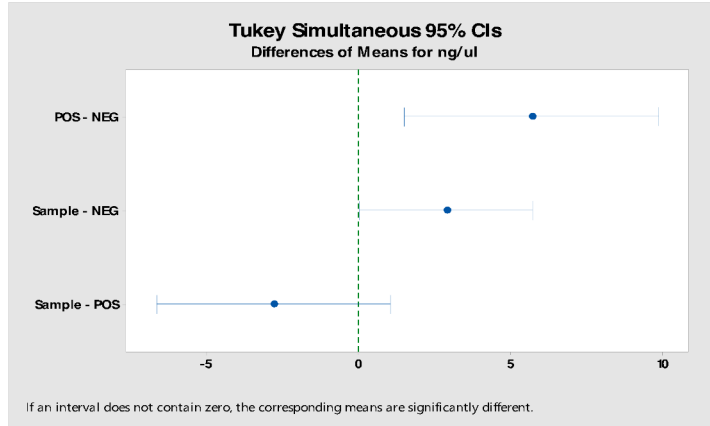
Compounds detected by MS1:

- C6-C24 fatty acids.
- Phospholipids & precursors.
- Glycans.
- Medications.
- Drugs of abuse.
- Dietary compounds.

Additionally:

- 20 multiple HDBM hits.
- 104 novel compounds.

Prototype: 16S Microbiomics



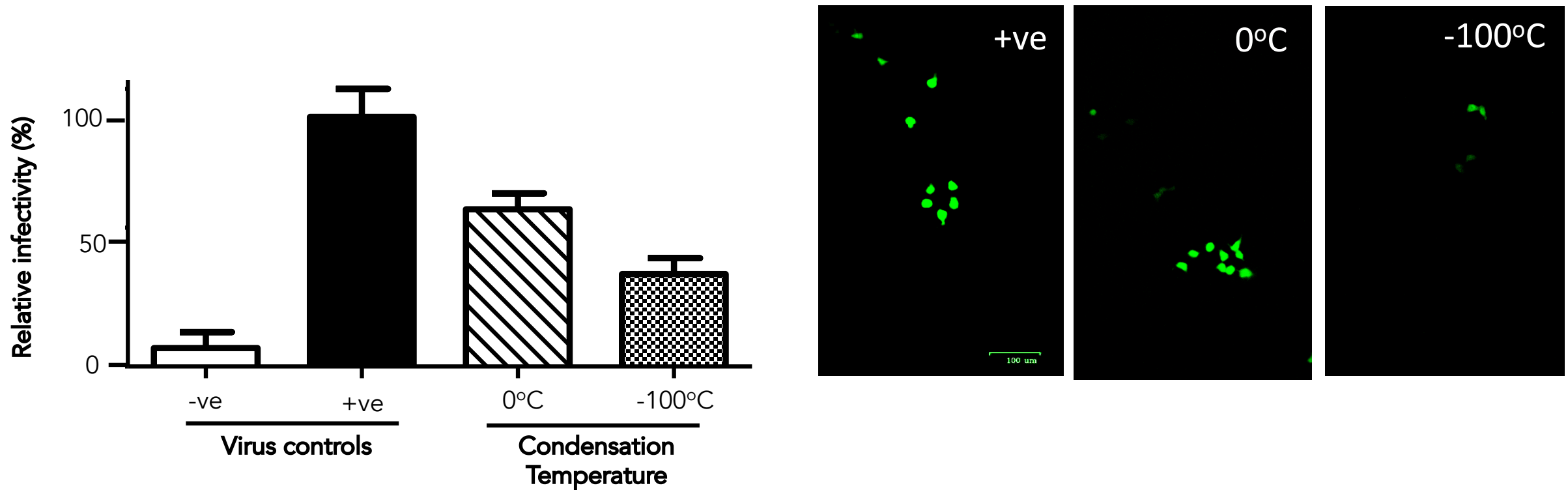
Higher DNA content vs background controls.

Detection of *Streptococcus* (BI FDR $q = 0.019$); amplicon generation w/out extraction.

Detailed work under way to optimize processes, polymerases, pipelines.

Prototype: Detection of aerosolized virus

Efficient capture of aerosolized virus, halves infection risk.



GFP-expressing VSV-pseudotyped lentivirus nebulized using PARI TurboBoy SX and captured using PBM-HALE™ (15 min). Condensates seeded on 10,000 HEK-293T's and GFP expression measured at 72hrs by FACS, visualized by fluorescent microscopy. Bar = 100 µm

Prototype: Clinical Pilot Update.



NO FALSE POSITIVES:

- COVID19 patients (n=12).
 - Nasal swab negative.
 - Week 2-3 of symptoms, known nasal -ve period.
 - Antivirals / hydroxychloroquine.
 - (dyspnoea) 25-30 breaths/min.
- 5-20 min sampling.
- In COVID19 wards.
- Blinded analysis.
- N=60 study actively recruiting.
 - Must be in symptoms week 1.
 - Must be nasal positive.
 - Interim data release: n=30.

We believe we can detect COVID-19 possibly with a 1 min sample



- **Safely:** Kill the virus.
- **More reliably:** Larger sample than nasal swabs.
- **Simply:** With no skills needed: just breathe out.
- **Using mass screening:** by mass production of plastic.
- **Where patients are:** using *any* point of need testing system.
- **With current gold standard tests:**
 - e.g. Abbott ID NOW[®]: 5 min test.
 - e.g. DeepVerge MicroTox BT: 4 sec test.

We believe we can detect COVID-19 possibly with a 1 min sample



- Confirm infectious virus load
 - By source of virus (oral, lung, nose)
 - By particle size (droplet, aerosol)
 - Optimise sampling maneuver.
- Determine the smallest sample amount needed for RT-PCR detection.
- Expand to pre/asymptomatic contacts / time course
- Deliver Emergency Use Authorisation (USA, UK).
- Produce >50,000 units.
- Supply at no profit basis under development funding.

How do I use the device?



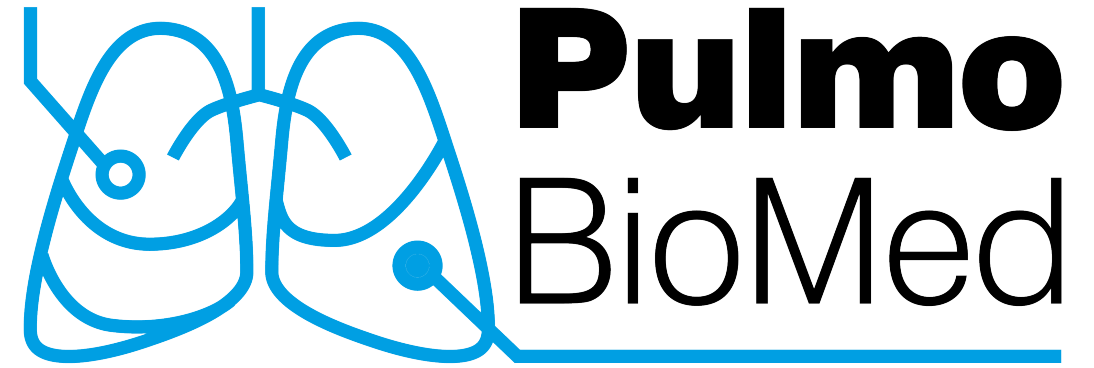
1. Device use SOP: <https://youtu.be/h6tLt9u-rWU>
2. Lay explanation of use: https://youtu.be/TkQEj-KN_os



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Pulmo
BioMed

Your Breath: Your Health™

Scientific and Technical Team

Saqib Ali (Lead Design Engineer): Design, modification, assembly, production oversight.

Dr Theodora Mantso (Biologist): Device testing and wet biology, microbiology.

Dr Andrew Nelson (Senior Biologist): Next Generation Sequencing.

Dr William Cheung (Senior Biologist): Proteomics & Metabolomics.

Adam Cosheril (Fabrication Lead): In-house 3D Printing

Paul Broom(Health & Safety Lead, technical lead).

PulmoBioMed™ Ltd. is a technology spin-out of Northumbria University

Company no. 12552857

This slide deck is live and will be updated on an ASAP basis.