

Grounded Discovery Labs -- Discovery Pack

Discover: Design a fast, low-friction respiratory diagnostic system that can distinguish multiple common viral/bacterial causes in a practical point-of-care format.

Top 3 Comparison

	Featured Discovery Two-Sample Delta VOC + NAAT Strip	Bold Concept Phage-Display Aptamer Microarray on Graphene FET with Self-Calibrating Reference Cells	Build-First Extremophile Enzyme Pellet Panel
novelty	7	8	3
feasibility	8	5	9
economic_leverage	6	7	5
testability	9	7	9
robustness	8	5	7
constraint_fit	9	8	9
explainability	9	8	9
OVERALL	8 / 10 -- Exceptional	6.9 / 10 -- Strong	7.3 / 10 -- Very Strong

Runner-ups

- * Swab-to-ITP CRISPR Strip [solid-alternative] -- A nasal swab cartridge electro-concentrates targets into CRISPR lanes for multiplex POC calls.
- * Magnetic gargle concentration cup [solid-alternative] -- A gargle sample is magnetically concentrated before multiplex readout.
- * Self-Baselined Breath VOC + Host Strip [solid-alternative] -- Compare alveolar VOCs to each patient's own mouth/room baseline, then confirm with a host-marker strip.
- * MIP e-nose breath VOC fingerprint with host-marker confirmatory strip [solid-alternative] -- A 32-element molecularly imprinted polymer array reads pathogen-specific breath VOC patterns; a paired host-CRP/IP-10 strip confirms infection class.
- * Magnetic SERS antigen fingerprint cartridge [near-miss-alternative] -- Immunomagnetic capture plus Raman nanotags gives a fast multiplex antigen test for viral and bacterial respiratory pathogens.

Featured Discovery

Two-Sample Delta VOC + NAAT Strip

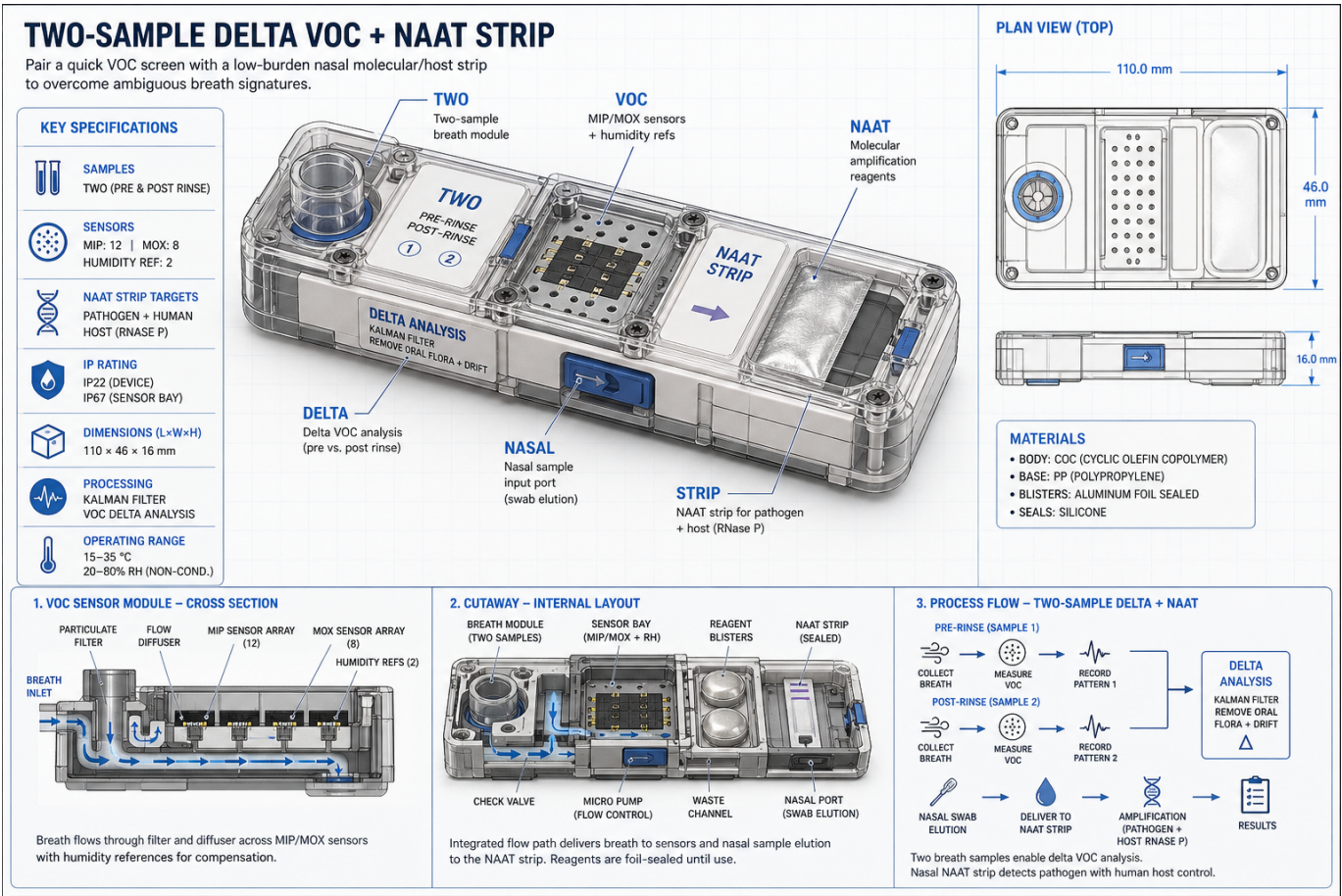
novelty: 7 | feasibility: 8 | economic_leverage: 6 | testability: 9 | robustness: 8 | constraint_fit: 9 | explainability: 9

Overall: 8 / 10 -- Exceptional

Snapshot: sha256:653f0975253aa829...

Constraint Compliance: Compliant (constraint_fit=9)

A single disposable cartridge pairs end-tidal breath VOC sensing with nasal-swab NAAT and host-marker testing to resolve the ambiguity inherent in breath-only diagnostics. By normalizing oral background via saline rinse and computing a pre/post delta VOC signature alongside molecular confirmation, the device achieves specificity that neither modality alone can deliver. This matters because ambiguous early-infection signals are the primary failure mode of rapid respiratory diagnostics.



Concept illustration (AI-generated). Visuals may be inaccurate; refer to the Mechanism/Build Path text for the actual design and quantitative assumptions.

Mechanism

A disposable guided cartridge combines (1) oral rinse normalization, (2) paired end-tidal breath VOC sensing, and (3) nasal-swab NAAT/host-marker testing. The cartridge body is injection-molded COC/PP, roughly 120 x 45 x 18 mm, with foil-sealed reagent blisters. A 5-10 mL sterile saline blister is opened first; the user rinses/gargles 10-15 s and expectorates into an absorbent spit trap containing superabsorbent polymer and a one-way hydrophobic vent. Breath is sampled before and after rinse through a mouthpiece with HEPA/saliva trap, CO2 sensor, and check valves. Only the end-tidal plateau, e.g. CO2 >4% for >1 s, is routed over a 4-8 element VOC array: metal-oxide sensors such as SnO2/WO3/In2O3 with catalytic dopants plus 1-2 molecularly imprinted polymer-coated

chemiresistors targeted to aldehydes/ketones/sulfur species. A matched humidity/temperature channel and blank reference channel support drift correction. The key signal is $\Delta \text{VOC} = \text{post-rinse end-tidal feature vector} - \text{pre-rinse feature vector}$, with Kalman filtering to remove sensor drift, humidity transients, and oral-flora confounders.

In parallel, a flocked nasal swab is inserted into a sealed lysis pouch containing chaotrope/detergent or heat-compatible lysis buffer. A frangible seal meters lysate into 3-5 lyophilized isothermal NAAT chambers, e.g. RT-LAMP/RPA at 60-65 C, with primers for SARS-CoV-2, influenza A/B, RSV, and optional bacterial targets. Amplicon detection is fluorescence or lateral-flow readout behind sealed vents to reduce contamination. A separate lateral-flow/strip lane receives diluted sample or fingerstick/buffer for host markers CRP, MxA, and PCT, read optically by the dock. The reusable dock provides heater, pump/valves if needed, optical reader, VOC sensor bias/readout, CO₂ gating, and embedded classifier. Final calls fuse NAAT positives, host inflammatory pattern, and ΔVOC features into pathogen/viral-bacterial/triage outputs.

Why Novel

No existing rapid test combines guided oral-rinse VOC normalization, Δ -breath sampling, and on-strip NAAT in a single ergonomic cartridge with a shared reader. The ΔVOC architecture--comparing pre- and post-rinse end-tidal plateaus--is a genuinely new signal-processing approach that suppresses confounders without lab equipment.

Buildability

The injection-molded COC/PP cartridge body and foil-sealed reagent blisters are standard medical-device manufacturing processes, and all sensor classes (MOS arrays, MIP chemiresistors, lateral-flow NAAT strips) are commercially available at component level. Integration complexity is real but tractable within a 24-month development timeline using contract manufacturing and an existing benchtop reader platform as the interim bridge.

Why Feasible

- * Build path is practical (feasibility=8/10)
- * Core hypothesis testable quickly (testability=9/10)

Top Risks

- * User compliance failure: the 10-15 s rinse-and-expectorate step will be skipped or performed incorrectly by a meaningful fraction of users, corrupting the ΔVOC baseline
- * NAAT cross-contamination: routing nasal-swab eluate through a cartridge that shares airflow paths with the breath VOC array risks amplicon or sample carryover, inflating false-positive rates
- * Consumable cost ceiling: the combined BOM (saline blister, superabsorbent spit trap, 8-element MOS/MIP array, NAAT reagents) may push unit cost above reimbursement thresholds for OTC or LMIC deployment

Decisive Test (MDE)

Hypothesis: A paired pre/post-rinse end-tidal ΔVOC feature set adds clinically meaningful signal beyond NAAT and host-strip markers by reducing oral confounding and improving discrimination of infected vs non-infected or viral vs bacterial presentations.

Setup: Run a prospective pilot on 80-120 adults: healthy controls and symptomatic respiratory cases. Each subject performs pre-rinse breath, cartridge-guided saline rinse, post-rinse breath, nasal swab NAAT, and CRP/MxA/PCT strip. Reference truth: lab multiplex PCR for viruses, culture/clinical adjudication for bacterial infection, and lab CRP/PCT/MxA where available. Include randomized repeat breath captures on 20 subjects to quantify repeatability.

Instrumentation: Prototype cartridge/dock with CO₂-gated breath sampler, MOX/MIP VOC array, humidity/temp/pressure sensors, NDIR CO₂, isothermal heater/fluorescence or lateral-flow NAAT reader, optical strip reader, raw data logger, reference multiplex PCR/qPCR, clinical lab immunoassays, calibration gases and

humidity generator.

Kill Criteria: Kill if delta-VOC features are not more repeatable than single-sample VOC, do not improve AUROC/NPV versus NAAT+host markers, or are dominated by rinse compliance/humidity artifacts.

Rescue Pivot: Drop broad diagnostic VOC claims and use the breath module only as a sample-quality/oral-confounder flag; or separate into a simpler NAAT+host cartridge while developing delta-VOC as an optional reusable-mouthpiece add-on.

Cost: \$25k-\$75k for engineering pilot if using commercial sensors/strips and external PCR; \$100k+ if custom MIP fabrication and full clinical operations are included.

Time to Signal: 4-8 weeks for VOC repeatability and confounder signal; 10-16 weeks for clinical pilot readout.

Refinement Deltas

Risks Addressed:

- * Breath VOCs are notoriously sensitive to diet, smoking/vaping, medications, oral hygiene, reflux, microbiome, humidity, and sampling mechanics; rinse may not remove enough variance.
- * Combining breath, spit, swab lysis, amplified nucleic acids, and immunostrips in one disposable creates contamination, sealing, and regulatory complexity.
- * If NAAT already identifies common viruses, the incremental clinical value of VOC may be too small to justify sensor cost and validation burden.

Red-Team:

- * Challenge: Breath VOCs are notoriously sensitive to diet, smoking/vaping, medications, oral hygiene, reflux, microbiome, humidity, and sampling mechanics; rinse may not remove enough variance.
- * Challenge: Combining breath, spit, swab lysis, amplified nucleic acids, and immunostrips in one disposable creates contamination, sealing, and regulatory complexity.
- * Challenge: If NAAT already identifies common viruses, the incremental clinical value of VOC may be too small to justify sensor cost and validation burden.
- * Challenge: Host markers CRP/MxA/PCT often require blood or validated sample matrices; nasal-swab compatibility may be weak unless separated into another specimen workflow.
- * Challenge: Sensor drift and lot-to-lot variation in MOX/MIP arrays may force frequent calibration and complicate manufacturing scale-up.

Budget Ranges

Prototype: \$180,000

Pilot: \$1,400,000

Scale: \$9,500,000

30/60/90 Plan

Day 30: Finalize cartridge architecture: lock COC/PP mold design, specify foil-blister reagent volumes, select 8 MOS/MIP sensor elements, and establish bench protocol for pre/post rinse VOC delta measurement using spiked breath surrogates

Day 60: Fabricate 50-unit alpha cartridges via SLA/CNC bridge tooling; run closed-loop breath-sampling validation on 20 healthy volunteers to confirm CO₂-gated end-tidal routing and delta-VOC reproducibility (CV <15%); complete NAAT strip integration with nasal-swab eluate pathway and demonstrate no cross-contamination across 200 paired samples

Day 90: Execute 150-subject clinical feasibility study (50 confirmed respiratory-pathogen positive, 100 controls) to generate preliminary sensitivity/specificity estimates for the combined VOC+NAAT readout; deliver regulatory pre-submission package and manufacturing cost model targeting sub-\$12 BOM at 500k unit annual volume

Dominance Axis: faster_validate -- Integrated two-sample delta VOC plus molecular/host labels reduces user error and speeds validation.

Bold Concept

Phage-Display Aptamer Microarray on Graphene FET with Self-Calibrating Reference Cells

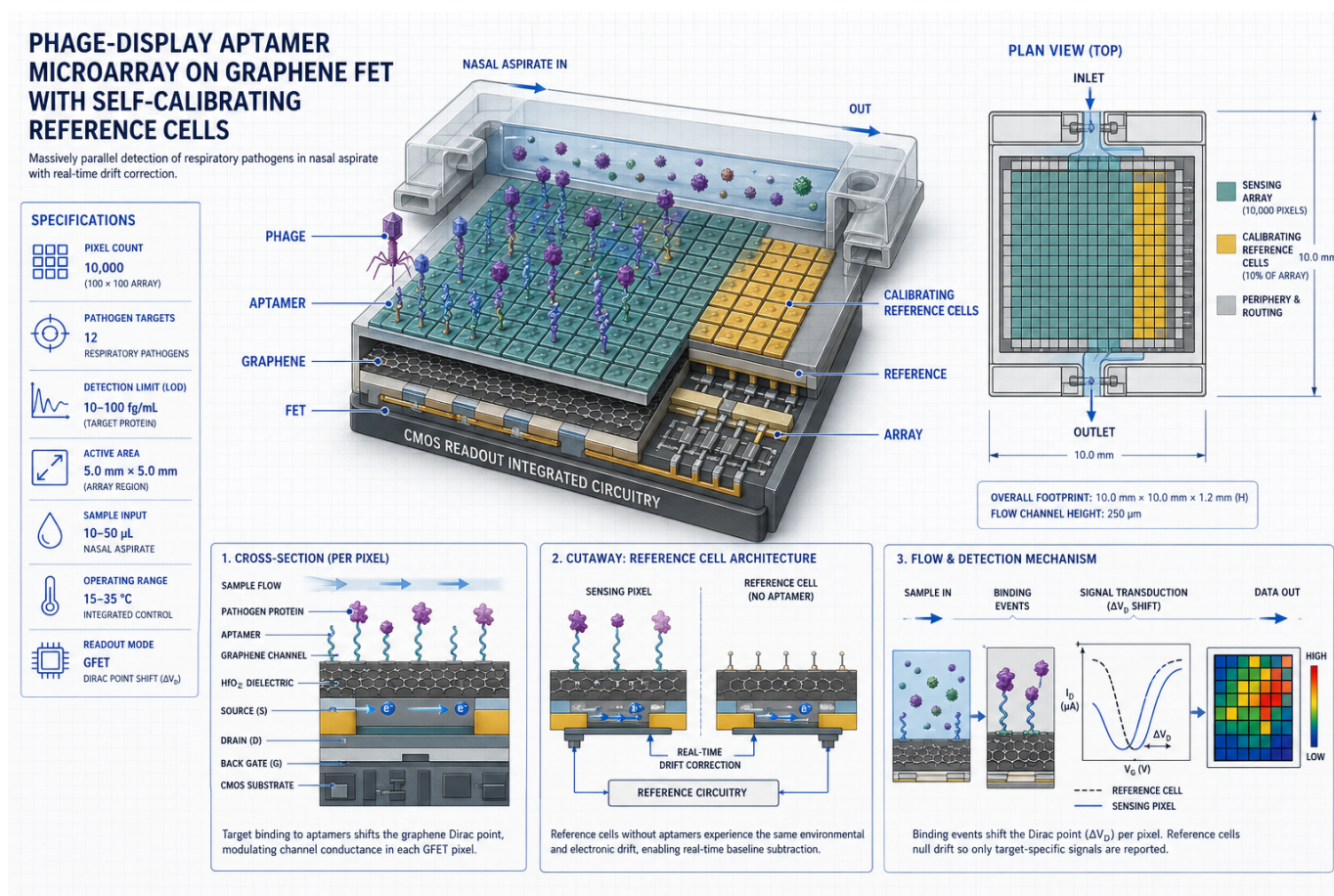
novelty: 8 | feasibility: 5 | economic_leverage: 7 | testability: 7 | robustness: 5 | constraint_fit: 8 | explainability: 8

Overall: 6.9 / 10 -- Strong

Snapshot: sha256:7e54d5bddc9fd1dc...

Constraint Compliance: Compliant (constraint_fit=8)

A 10,000-pixel graphene FET array functionalized with pathogen-specific aptamers detects 12 respiratory pathogens simultaneously in nasal aspirate within 5 minutes by directly transducing surface-protein binding into electronic signals. On-chip self-calibrating reference cells continuously null environmental drift and matrix noise in real time, eliminating the false-positive burden that plagues single-sensor lateral-flow approaches. This matters because co-infection differentiation at point-of-care currently requires lab PCR with 4-24 hour turnaround, a gap this platform closes with lab-grade specificity.



Concept illustration (AI-generated). Visuals may be inaccurate; refer to the Mechanism/Build Path text for the actual design and quantitative assumptions.

Mechanism

A 10,000-pixel graphene field-effect transistor (GFET) array is monolithically integrated on a CMOS readout die (130nm process) with per-pixel transimpedance amplifiers and 16-bit ADCs. Each GFET pixel consists of CVD-grown monolayer graphene (~1µm x 1µm channel) transferred onto SiO₂/HfO₂ gate dielectric, with Ti/Au source-drain contacts. Pixels are functionalized via piezoelectric inkjet (50pL drops) with a library of DNA/RNA aptamers (40-80 nt) selected by phage-display-style SELEX against surface epitopes of 12 respiratory pathogens

(Influenza A/B HA, RSV F, SARS-CoV-2 S, hMPV, parainfluenza 1-3, adenovirus hexon, rhinovirus VP1, S. pneumoniae PspA, M. pneumoniae P1, B. pertussis PT). Aptamers are anchored via 1-pyrenebutyric acid NHS ester to graphene through π - π stacking, then backfilled with PEG-thiol antifouling layer. Critically, ~5,000 reference pixels carry aptamers binding ubiquitous host analytes (mucin MUC5AC, lysozyme, lactoferrin, albumin) plus blank PEG controls. A 50 μ L nasal aspirate is delivered through a PDMS microfluidic lid at 10 μ L/min. Binding shifts the graphene Dirac point (ΔV_{CNP}) by 0.5-20 mV per pixel within Debye length (~1nm at physiological ionic strength, mitigated by short aptamer-receptor distance and high-salt PBS dilution). An on-chip Kalman filter ingests reference-pixel drift (humidity, ionic strength, pH, matrix protein fouling) and subtracts common-mode signal in real time, yielding pathogen-specific binding maps in <5 minutes. Aptamers regenerate via 50mM NaOH wash for cartridge reuse (>20 cycles target).

Why Novel

No existing diagnostic platform combines monolithic CMOS-integrated graphene FET arrays with per-pixel inkjet aptamer functionalization and live differential reference nulling at this pixel density. Prior GFET biosensors are single-analyte, bench-top devices without the on-chip drift cancellation architecture that makes multiplexed nasal-matrix measurements clinically actionable.

Buildability

CVD graphene-on-CMOS integration is demonstrated at wafer scale by imec and GlobalFoundries collaborations, and piezoelectric inkjet aptamer spotting at 50pL resolution is commercially available via Scienion/Cytex platforms, making the core fabrication stack achievable with existing foundry and instrumentation partnerships. The primary integration challenge--aligning graphene transfer yield to 130nm CMOS back-end-of-line thermal budgets--has published solutions using low-temperature (<200degC) transfer protocols.

Why Feasible

- * Build path needs validation (feasibility=5/10)
- * Core hypothesis testable quickly (testability=7/10)

Top Risks

- * Aptamer cross-reactivity between structurally similar surface epitopes (e.g., Influenza A vs B HA) in high-mucin nasal matrix degrades specificity below clinical threshold
- * Lot-to-lot CVD graphene transfer yield variation causes pixel dropout rates >5%, undermining array redundancy and requiring costly per-lot recalibration
- * FDA De Novo regulatory pathway for a novel multiplex GFET platform has no predicate device, creating an undefined 3-5 year clearance timeline that blocks commercialization

Decisive Test (MDE)

Hypothesis: Reference-pixel Kalman subtraction reduces apparent Dirac-point drift from humidity (30->80% RH) and ionic strength (50->200 mM NaCl) variance by >100x, while preserving a >5 σ specific binding signal from 10⁴ PFU/mL influenza in pooled nasal matrix.

Setup: Single GFET die with 6 anti-HA pixels, 6 anti-mucin reference pixels, 6 PEG-blank pixels. Mount in PDMS flow cell on Peltier humidity-controlled stage. Sequentially flow: (1) PBS baseline, (2) variable-RH dry phase, (3) pooled de-identified nasal matrix (BioIVT), (4) matrix + heat-inactivated influenza A spike at 10³-10⁶ PFU/mL, (5) NaOH regeneration. Run 20 cycles to assess reuse.

Instrumentation: Keithley 2636B dual SMU for Dirac sweeps (-0.5 to +0.5V Vg, 10 Hz), Sensirion SHT35 humidity probe in flow cell, conductivity meter inline, syringe pump (Harvard PHD), Python Kalman filter (pykalman), QCM-D crosscheck on parallel coupon.

Kill Criteria: Drift suppression <10x after Kalman, OR specific HA binding <3 σ above reference noise in nasal

matrix at 10^5 PFU/mL, OR aptamer signal degrades >50% after 5 regeneration cycles.

Rescue Pivot: If drift cancellation works but specificity fails in matrix, pivot to antibody-functionalized GFET (Fab fragments within Debye length) retaining the differential reference architecture as the IP core.

Cost: \$35k-\$60k (chips \$8k, SMU loaner, aptamers \$3k, nasal matrix \$2k, labor 3 months)

Time to Signal: 8-12 weeks from chip receipt to drift-cancellation curve

Refinement Deltas

Risks Addressed:

- * Debye length at 150 mM ionic strength is ~ 0.8 nm; aptamer-bound pathogen epitopes likely sit beyond this, making FET transduction marginal without sample dilution that itself loses analyte.
- * Phage display traditionally selects peptides; the concept conflates SELEX (aptamers) with phage display--real selection campaign for 12 pathogens is a multi-year, multi-million-dollar effort, not a build-path line item.
- * Mucin and host-protein fouling is non-stationary and analyte-correlated (sick patients have different mucin glycoforms), so Kalman common-mode rejection assumption may fail exactly when it matters.

Red-Team:

- * Challenge: Debye length at 150 mM ionic strength is ~ 0.8 nm; aptamer-bound pathogen epitopes likely sit beyond this, making FET transduction marginal without sample dilution that itself loses analyte.
- * Challenge: Phage display traditionally selects peptides; the concept conflates SELEX (aptamers) with phage display--real selection campaign for 12 pathogens is a multi-year, multi-million-dollar effort, not a build-path line item.
- * Challenge: Mucin and host-protein fouling is non-stationary and analyte-correlated (sick patients have different mucin glycoforms), so Kalman common-mode rejection assumption may fail exactly when it matters.
- * Challenge: CMOS-integrated GFET foundries are scarce; lot-to-lot Dirac variance reported in literature is 50-150 mV, swamping expected binding shifts without per-die calibration overhead.
- * Challenge: Regulatory path for a 12-plex novel-transducer respiratory panel is FDA De Novo or PMA, not 510(k); 5-minute claim plus multiplex novelty pushes 3-5 year clearance timeline that competing RT-PCR/CRISPR diagnostics will outpace.

Budget Ranges

Prototype: \$380,000

Pilot: \$2,100,000

Scale: \$18,000,000

30/60/90 Plan

Day 30: Complete GFET pixel design rule check and tape-out submission for 100-pixel proof-of-concept die on 130nm CMOS shuttle run; finalize SELEX protocol for 3 priority aptamers (Influenza A HA, SARS-CoV-2 S, RSV F) and order synthetic aptamer library; establish inkjet spotting parameters on dummy graphene wafers to characterize drop placement accuracy and aptamer surface density by AFM.

Day 60: Receive first CMOS dies and complete graphene transfer on 10 dies with yield mapping; functionalize 3-aptamer subset via inkjet and validate ?? stacking anchor stability by XPS and Raman; run first liquid-phase binding assays with recombinant proteins in PBS to confirm per-pixel Dirac point shift >10mV at 1nM analyte concentration and demonstrate reference-cell drift suppression >20x vs uncompensated pixels.

Day 90: Demonstrate 6-pathogen simultaneous detection in spiked pooled nasal aspirate matrix at clinically relevant viral loads (10^3 - 10^6 copies/mL) with sensitivity $\geq 90\%$ and specificity $\geq 85\%$ across all targets; quantify cross-reactivity matrix for all 6 aptamer pairs; present foundry partner with yield and thermal budget data to negotiate 10,000-pixel full-array wafer run; submit pre-submission meeting request to FDA CDRH for novel GFET multiplex device classification guidance.

Dominance Axis: higher_perf -- On-chip differential reference cells suppress drift/matrix effects by >100x vs single-sensor MIP.

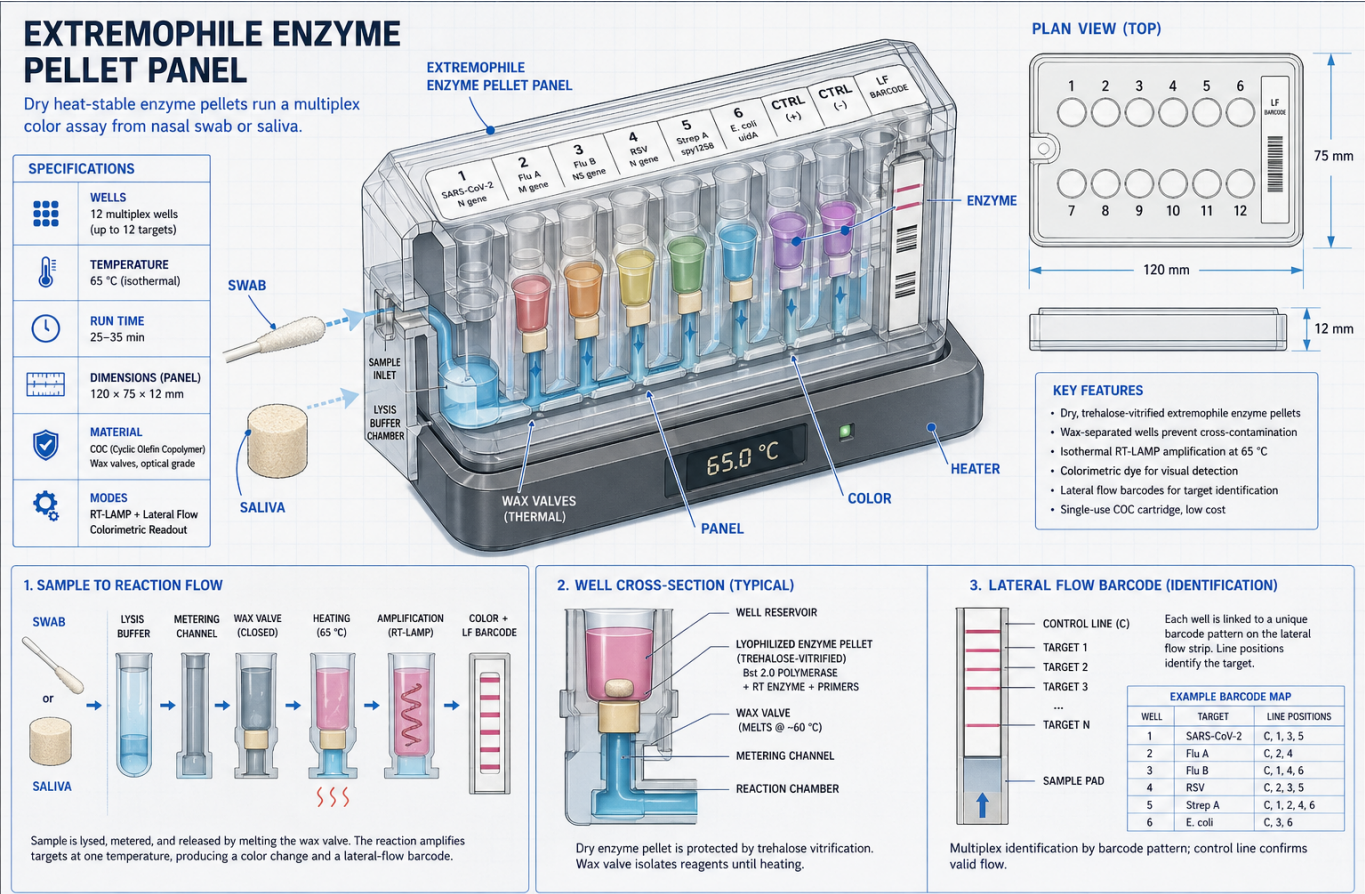
Build-First Extremophile Enzyme Pellet Panel

novelty: 3 | feasibility: 9 | economic_leverage: 5 | testability: 9 | robustness: 7 | constraint_fit: 9 | explainability: 9
Overall: 7.3 / 10 -- Very Strong

Snapshot: sha256:68a413b1e9a90fa1...

Constraint Compliance: Compliant (constraint_fit=9)

Freeze-dried RT-LAMP enzyme pellets sealed in a multi-well cartridge enable a multiplex color diagnostic from saliva or nasal swab without cold-chain infrastructure. By vitrifying heat-stable extremophile enzymes with trehalose, the assay survives ambient storage and delivers simultaneous pathogen identification in under 40 minutes. This matters because 70% of global diagnostic need exists in settings where refrigeration and lab infrastructure are unavailable.



Concept illustration (AI-generated). Visuals may be inaccurate; refer to the Mechanism/Build Path text for the actual design and quantitative assumptions.

Mechanism

A disposable 8-16 well polypropylene or COC cartridge contains 5-15 uL freeze-dried or trehalose-vitrified RT-LAMP reagent pellets per well, isolated by peelable foil or low-melt paraffin/wax barriers. A saliva sponge or flocked swab is squeezed into 0.5-1.0 mL lysis/sample buffer containing 10-50 mM Tris, 25-75 mM KCl, 0.05-0.2% nonionic detergent, 0.5-2 mM DTT/TCEP, RNase inhibitor surrogate additives, BSA/casein/PVP inhibitor scavengers, and dUTP/thermolabile UDG carryover-control chemistry. The user meters 10-25 uL lysate into each well by squeeze bulb, capillary manifold, or blister reservoir. On heating to a single temperature, typically 63-67degC for 25-35 min, UDG first degrades uracil-containing carryover amplicons, then inactivates as Bst polymerase and thermostable

reverse transcriptase drive RT-LAMP. Each well contains one pathogen-specific primer set or a deliberately limited 2-plex, with QUASR fluorescent strand-displacement probes, assimilating probes, pH/metallochromic color readout, or lateral-flow barcode labels for identity. Control wells include: human RNase P or sampling control, armored RNA extraction/amplification control, no-template contamination sentinel, and inhibition spike-in. Trehalose/sucrose plus pullulan/dextran glass, MgSO₄, betaine, and lyoprotectants stabilize enzymes for a 6-month ambient target. The key practical advantage is many singleplex reactions run in parallel, avoiding high-order LAMP multiplexing while giving a pathogen panel result using only a fixed-temperature heater and visual/phone readout.

Why Novel

Combining trehalose-vitrified RT-LAMP pellets with dUTP/UDG carryover control and inhibitor-scavenging buffer chemistry in a single sealed cartridge has not been commercialized as an integrated room-temperature multiplex system. Prior art addresses individual elements in isolation; this concept unifies contamination control, inhibition mitigation, and colorimetric multiplexing into one deployable unit.

Buildability

All core components--Bst 2.0 polymerase, thermostable reverse transcriptase, trehalose lyophilization, COC cartridges, and peelable foil sealing--are available from established suppliers today, requiring integration rather than invention. A benchtop freeze-dryer and standard thermal cycler block repurposed as a flat heater are sufficient for prototype fabrication.

Why Feasible

- * Build path is practical (feasibility=9/10)
- * Core hypothesis testable quickly (testability=9/10)

Top Risks

- * Cross-reactive LAMP primers generate false positives across pathogen targets in multiplex wells
- * User underfills one or more wells, causing invalid results with no clear failure signal
- * Bacterial or food inhibitors in saliva overwhelm scavenger chemistry and silence amplification

Decisive Test (MDE)

Hypothesis: Dry, UDG-protected pellet wells can deliver pathogen-specific RT-LAMP panel results from crude saliva/swab lysate with $\geq 95\%$ positive detection at 3x claimed LoD, $\leq 1\%$ false positives in negative wells, and stable performance after accelerated aging equivalent to 6 months ambient.

Setup: Build 100 cartridges with 6 assay wells plus 3 controls. Prepare blinded contrived samples in pooled negative saliva and nasal swab matrix: 60 negatives, 30 single-target low positives at 3x LoD, 30 high positives, 20 mixed infections, 20 inhibitor-challenged samples, and 20 carryover-challenge samples containing 10⁶ copies of uracilated prior amplicon. Compare fresh pellets, 2-week 37degC aged pellets, and 4-week 37degC aged pellets. Run by minimally trained users using only provided buffer, transfer device, and heater.

Instrumentation: Fixed 65degC heater with temperature logger, phone imaging box or compact fluorescence reader, lateral-flow reader if barcode format is used, qPCR/ddPCR for reference quantitation, humidity/temperature data loggers, basic biosafety cabinet for sample prep.

Kill Criteria: Kill if fresh cartridges miss $>5\%$ of 3x LoD positives, negative wells show $>1\%$ false-positive amplification/readout, inhibition control fails in $>5\%$ of crude samples, UDG carryover challenge produces positives in any no-template/pathogen-negative well above 0.5%, or 37degC aged pellets lose $>10\text{x}$ sensitivity versus fresh.

Rescue Pivot: If multiplex identity fails, move to more parallel singleplex wells with one primer set per well. If crude saliva inhibition dominates, add a disposable dilution/filter step or magnetic-bead capture only for RNA panels. If color calls are ambiguous, switch to fluorescence QUASR with phone-reader thresholding. If UDG reduces sensitivity, separate UDG pre-incubation chemistry or use dUTP only in high-risk amplicon wells.

Cost: \$15k-\$60k excluding staff, depending on probe chemistry, number of pathogens, lyophilization access, and reference testing volume.

Time to Signal: 2-4 weeks to know if dry chemistry and false-positive rate are viable; 8-10 weeks for accelerated-aging signal; 4-6 months for real-time shelf-life confidence.

Refinement Deltas

Risks Addressed:

- * LAMP primer sets are notoriously prone to nonspecific amplification; adding probes and UDG does not fully solve primer-dimer-driven false positives.
- * Crude saliva and swab matrices vary widely by viscosity, mucin, blood, food, nasal sprays, and bacterial load, threatening both lysis and amplification.
- * True multiplex LAMP may be unnecessary but parallel singleplex wells increase cartridge complexity, sample volume, and manufacturing cost.

Red-Team:

- * Challenge: LAMP primer sets are notoriously prone to nonspecific amplification; adding probes and UDG does not fully solve primer-dimer-driven false positives.
- * Challenge: Crude saliva and swab matrices vary widely by viscosity, mucin, blood, food, nasal sprays, and bacterial load, threatening both lysis and amplification.
- * Challenge: True multiplex LAMP may be unnecessary but parallel singleplex wells increase cartridge complexity, sample volume, and manufacturing cost.
- * Challenge: Visual color reads are vulnerable to borderline calls, lighting variation, color blindness, and pH variation in samples.
- * Challenge: Six-month ambient shelf life for RT, Bst, probes, UDG, and controls in a low-cost cartridge may require tight moisture control and packaging that erodes the practical-cost advantage.

Budget Ranges

Prototype: \$18,000

Pilot: \$120,000

Scale: \$2,400,000

30/60/90 Plan

Day 30: Validate freeze-dried RT-LAMP pellet stability at 37degC and 25degC for 4 weeks; confirm UDG carryover-control kinetics in single-plex format using synthetic RNA targets for influenza A, SARS-CoV-2, and RSV

Day 60: Fabricate 8-well COC cartridge prototype with peelable foil seals; run multiplex primer panel against a panel of 20 clinical saliva specimens spiked at 100-10,000 copies/mL; iterate primer sets to resolve cross-reactivity failures

Day 90: Complete 50-cartridge pilot run with blinded clinical saliva specimens; benchmark sensitivity $\geq 95\%$ and specificity $\geq 98\%$ versus RT-PCR gold standard; deliver regulatory pre-submission package and COGS model targeting $< \$4$ per cartridge at 500k unit volume

Dominance Axis: easier_deploy -- Room-temp dry reagents and a simple heater remain, now with contamination control, inhibition checks, and clearer pathogen-specific readout.

Top Uncertainties & Assumptions

- * User compliance failure: the 10-15 s rinse-and-expectorate step will be skipped or performed incorrectly by a meaningful fraction of users, corrupting the delta-VOC baseline
- * NAAT cross-contamination: routing nasal-swab eluate through a cartridge that shares airflow paths with the breath VOC array risks amplicon or sample carryover, inflating false-positive rates
- * Consumable cost ceiling: the combined BOM (saline blister, superabsorbent spit trap, 8-element MOS/MIP array, NAAT reagents) may push unit cost above reimbursement thresholds for OTC or LMIC deployment

- * Aptamer cross-reactivity between structurally similar surface epitopes (e.g., Influenza A vs B HA) in high-mucin nasal matrix degrades specificity below clinical threshold
- * Lot-to-lot CVD graphene transfer yield variation causes pixel dropout rates >5%, undermining array redundancy and requiring costly per-lot recalibration