## Optimization of logic gates for one-step detection of microRNAs via Split loop-mediated isothermal amplification (Split-LAMP)

Medha Shridharan

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#### Introduction

#### Development of a diagnostic miRNA biomarkers To be used in point-of-care test (POCT)

- High specificity
- High stability

Most diagnostically valuable in panels

Easy accessibility from patient samples

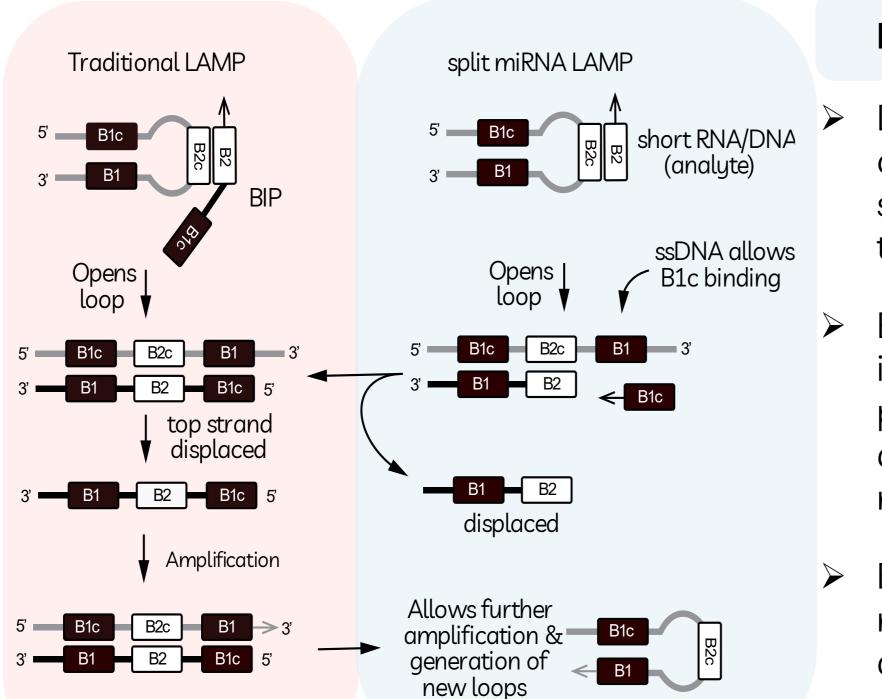
- Easy-to-use
- Rapid
- Customisable to any target disease by detecting miRNA concentration

Integrates signals from miRNA panel on a molecular scale into a single diagnostic readout

**Aim:** Show proof-of-concept for an easy-to-use, rapid, customizable novel miRNA POCT platform (Split-LAMP)

#### **Traditional LAMP**

- Detects concentrations of long DNA template
- ➤ Uses BIP (B1c & B2 concatemer) and FIP (F1c & F2 concatemer) for strand invasion and amplification
- LAMP cannot detect shorter miRNA targets



#### **Novel Split-LAMP**

- Detects concentrations of shorter miRNA target
- BIP and FIP are split into component primers: B1c & B2 and F1c & F2 respectively
- B2 and F2 act a miRNAs to be detected

#### **Key Insight:**

In Split-LAMP, amplification depends on the presence of short miRNA analytes that act as **primers** as opposed to templates.

#### **Key benefits of Split-LAMP:**



Split-LAMP is a more easily customizable platform than Traditional LAMP, which can be designed to detect miRNA biomarkers specific to various target diseases by varying the B2c/F2c sequence in the template.



Split-LAMP displays a positive readout only when two miRNA signals are present concurrently. It integrates multiple signals from a diagnostically valuable miRNA panel for diagnosis.



Adding B1c and B2 as separate oligonucleotides rather than a single BIP primer results in faster and stronger amplification due to more strand invasion and amplification.

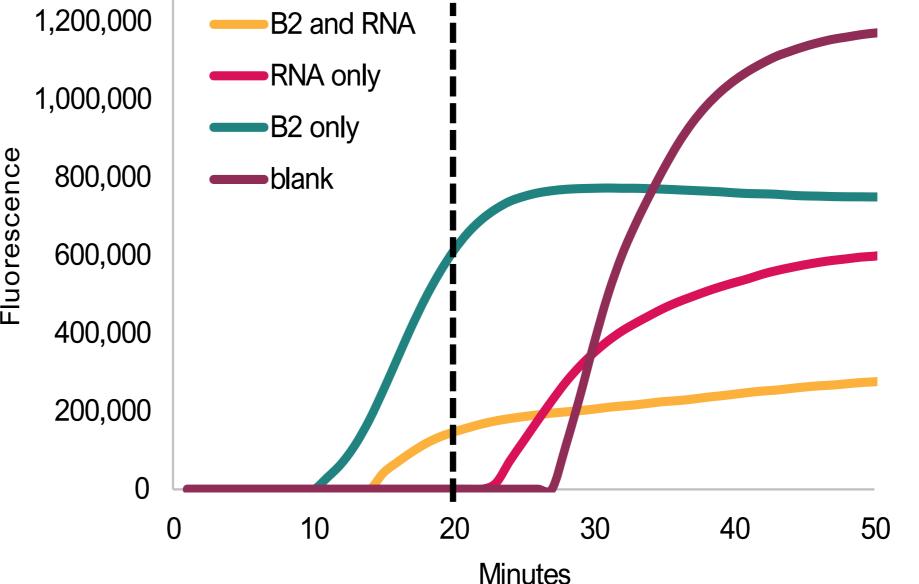
## A) B2 and F2 can be detected concurrently for diagnosis

- Preliminary experiments were conducted with DNA rather than RNA primers.
- 2 miRNAs modelled by B2 and F2.

Optimal primer concentrations to produce a signal most responsive to a 100fold change in concentrations of B2 and F2 were empirically identified.

Nucleic acid component	Template	B1C	F1C	LB	LF
Empirically optimized concentration	53pg/ml	100nM	100nM	10nM 100nM	0nM

# Split-LAMP is highly specific



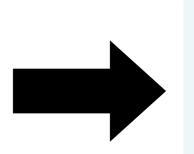
- B2 target is detected even in the presence of background total-cell RNA
- Samples containing background RNA can be clearly distinguished from those containing B2 target

Split-LAMP is **highly specific** with low risk of false positives.

**Fig. 2** Real-time fluorescence curves aroused by B2 (target analyte) and HEK-293 total cell RNA (background RNA) in split-LAMP.

### Bayesian Optimization (BO) identifies new points in the solution space

Need for further finetuning of difference between F2 amplification curves from (A)



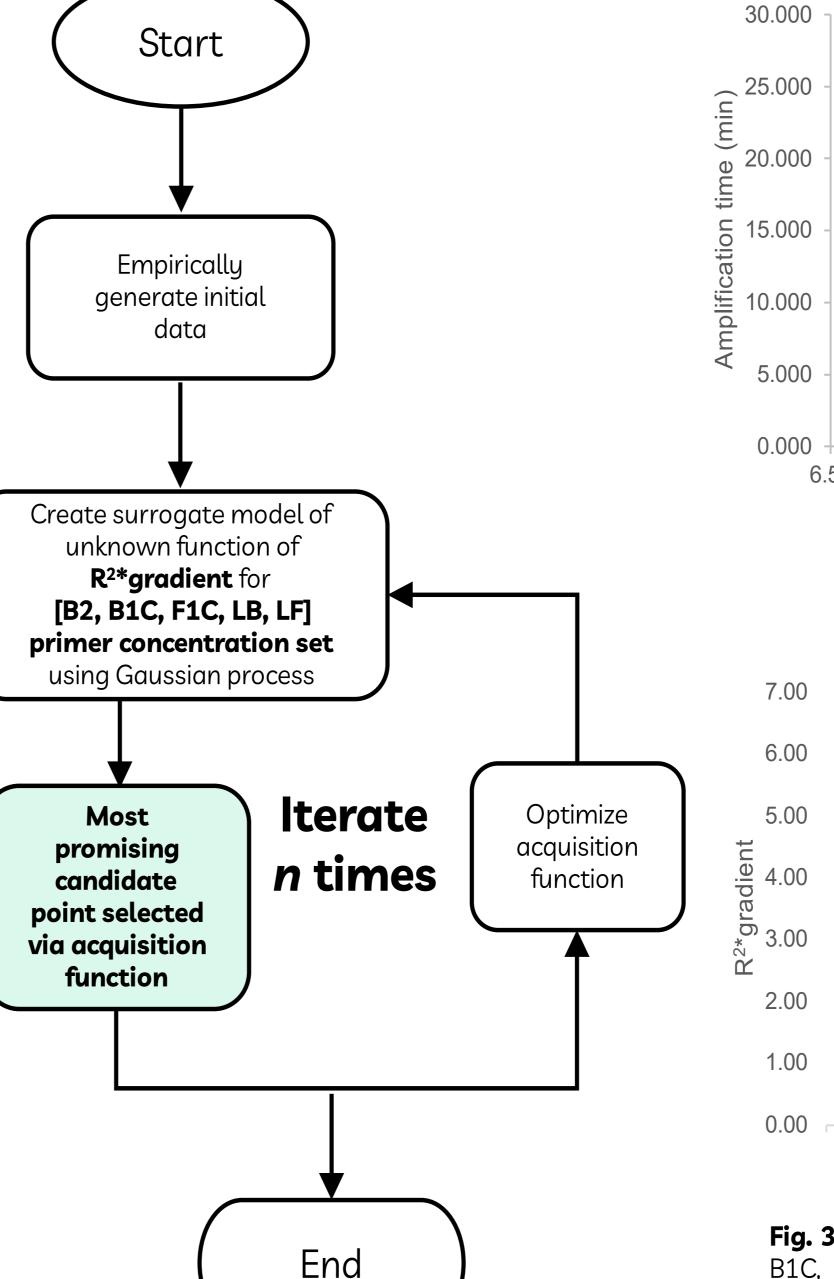
(A)

Implemented BO to introduce novel points in the space of **primer** concentration sets which we did not investigate from empirical data from (A)

Test 1

Test 2

• Test 3



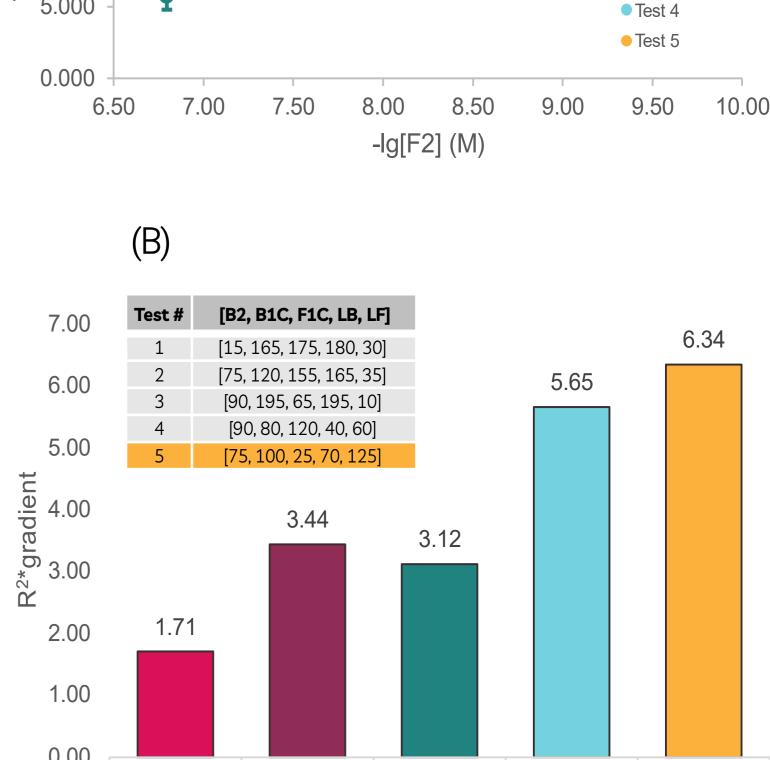


Fig. 3 (A) Standard plots for F2 serial dilutions at 5 different [B2, B1C, F1C, LB, LF] primer concentration sets generated by algorithm. (B) Bar graphs of R<sup>2</sup>\*gradient values derived from standard plots in (A) with [B2, B1C, F1C, LB, LF] primer concentration set values (nM) in each test shown in table.

Test number

Although the points introduced by the preliminary BO model did not outperform the empirical data from (A), the model outputs competitive points in the space of primer concentration sets. The model shows promise in outperforming empirical data if fine tuned with more complexity and iterations.

#### **Conclusions and Future Work**

The effective dynamic range of Split -LAMP for B2 and F2 is  $\sim 1-200$  nM. 900,000 1,000,000 (B) (A)800,000 900,000

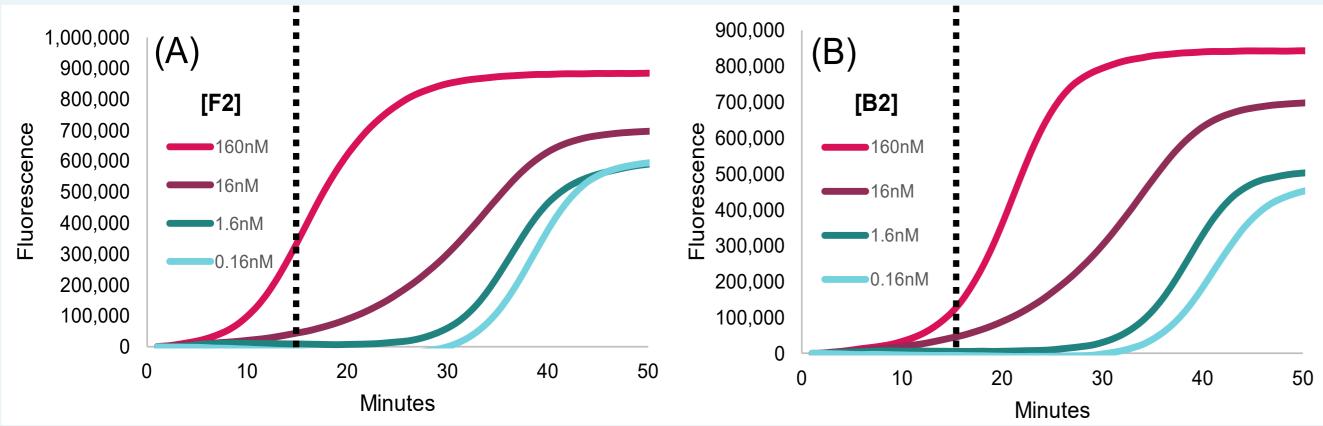


Fig. 1 (A) represents real-time fluorescence curves aroused by varying [F2] in a serial dilution, where [B2] = 16nM. (B) represents similar curves aroused by varying [B2] in a serial dilution, where [F2] = 16nM. Suitable cutoff amplification time (dotted line) can be applied to determine is readout is positive or negative.

A single cut-off amplification time can be applied such that positive readout is observed only if both B2 and F2 are above threshold concentrations. This ensures that B2 and F2 can be detected **concurrently for diagnosis.** 

Showed proof-ofconcept for Split-LAMP

- Integrates multiple miRNA signals into a single readout
- Customizable to detect any target disease
- Highly specific with low rate of false positives

To identify a potential disease diagnostic application

To develop a POCT device for a target disease

Development of OR/NOT logic gates

[6] Rasmussen, C. E. Gaussian processes for machine learning. MIT Press, 2006

- Easy-to-use
- Rapid
- Low cost

All media objects are self-created.

[1] Abdullah AL-maskri, A. A., Ye, J., Talap, J., Hu, H., Sun, L., Yu, L., Cai, S., & Zeng, S. (2020). Reverse transcription-based loop-mediated isothermal amplification strategy for real-time miRNA detection with phosphorothioated probes. Analytica Chimica Acta, 1126, 1-6. https://doi.org/10.1016/j.aca.2020.06.007 [2] Biolabs, N. E. (n.d.). Loop-mediated isothermal amplification. NEB. Retrieved November 17, 2022, from

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