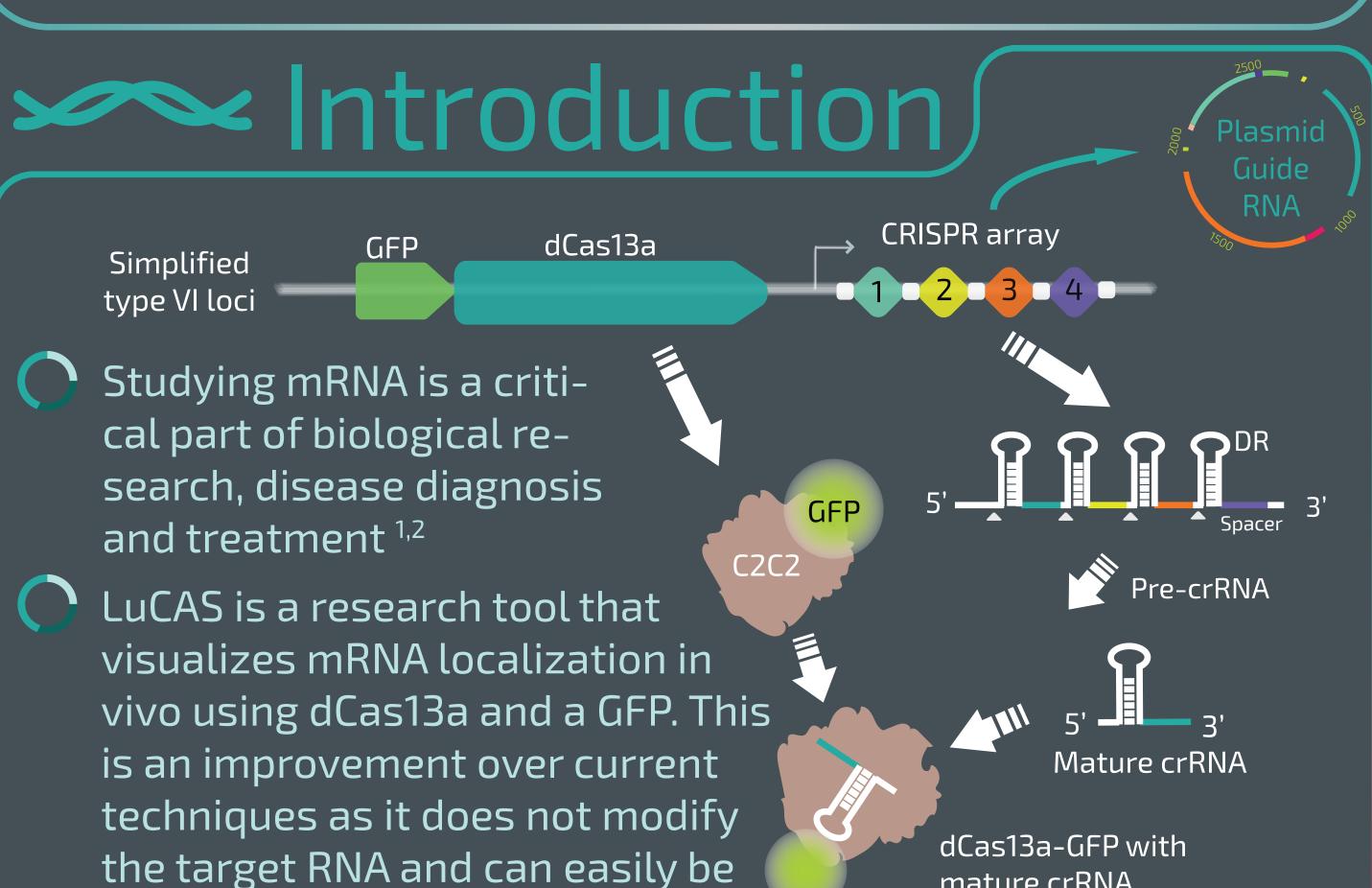
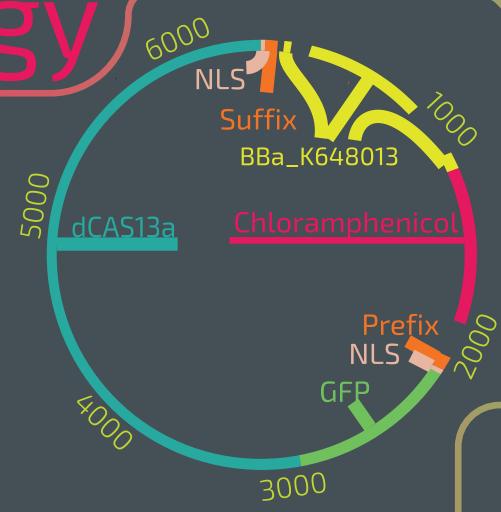


"A novel way to track mRNA localization in vivo"



Gibson Assembly of pSB1C3 with GFP (BBa\_K648013) and 'dCas13a' derived from L. Buccalis containing four point mutations resulting in a HEPN nuclease domain 1 and 2 inactive mutant, yielding our BioBrick (B-Ba\_K2340000)

adjusted for another target



mature crRNA

Design of fusion protein for mammalian expression:

LS = Linker sequence NLS = Nuclear localisation sequence

Ligation of fusion protein via HindIII and amplification

Design of guide RNA sequences (BBa\_K2340001 -BBa\_K2340011) for crRNA synthesis derived from human pkp4, Rab13, inpp1, and  $\beta$ -Actin expressing mRNA:

RCS Suffix SLS = Serine linker sequence | RCS = RNA coding sequence | 6T = Terminator

Ligation of whole fusion protein with a) pSB1C3 bacterial expression vector and b) pcDNA 3.1 mammalian expression vector via EcoRI and Xhol

Transformation of bacterial fusion protein construct into DH5α cells and transfection into HEK293 cells followed by imaging

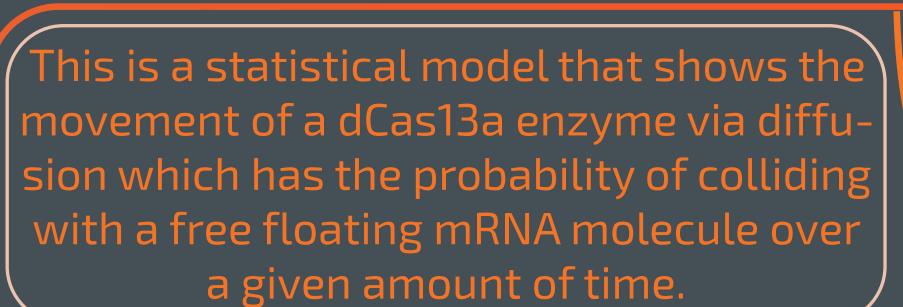
# Modelling











Probability of enzyme-mRNA molecule collision using a normalized integral

T is defined as the average amount of time for a given enzyme to collide  $T = \frac{1}{2.8}$  milliseconds

Maximum no. of enzymes in the given amount of time yielding a 100% collision rate for the fibroblast and neurons was **24**.

Einstein-Stokes theorem to find the diffusion constant

Raleigh criterion was used to distiguish between two GFP molecules

Where  $\lambda$  is 510nm, The minimum distance we can resolve for 2 GFP molecules is: R = 256 nm

Association of dCas13a-GFP with mRNA target Graph. 1 How long does it take dCas13a to find its target



Liaising with the public was incredibly important to us and we wanted to exercise the connections we had to our community

through different outreach opportunities.

Human Practices

Family Festival: Created a science stall for

children where we showcased different experiments and activities whilst gathering survey data from their parents



Open days:

Spoke with aspiring science students about our project

Gathered data to integrate into our project through surveys and communication

Consulted with professionals in their field

Gained feedback through presenting in the Bioscience Symposium

## Collaboration Co.C



our dCas13a

We collaborated with team udd\_UK on the interlab stud

We provided lab space and

equipment required for the standardized study Judd provided DNA from

Distribution Kit plate 6 as we weren't successful in transforming the DNA from our kit



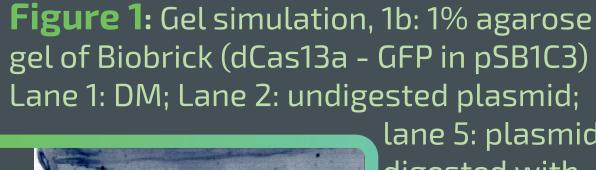


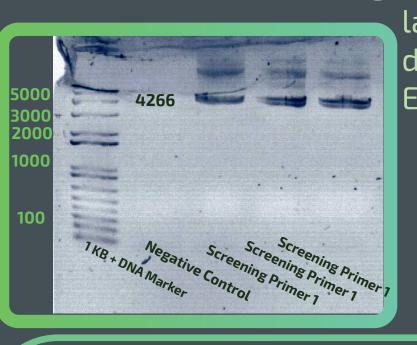
— We built one biobrick, validated by screening primers (Fig 1 and 2), by fusing together a wtGFP gene using NEB Gibson Assembly<sup>3</sup> a 'dead' Cas13a gene, adding a flexible linker and two nuclear localization sequences to improve stability and reduce background noise.

— We transfected our construct (BBa\_K2340000) with a guide RNA plasmid (BBa\_K2340001 - BBa\_K2340011) (1:2 ratio) into HEK293<sup>4</sup> cells using Lipofectamine™ 2000 in a 24 well plate in media and incubated for 24 hrs at 37°C.

The function of our construct was not proven (Fig 3) perhaps due to the poor folding of the wtGFP at 37°C and the short time frame.

— Furthermore, we designed 11 biobricks which could not be sent due to time constraits.





igested with ECO RI and Pst



### ~~ References E

1. Briley WE, Bondy MH, Randeria PS, Dupper TJ, Mirkin C a. Quantification and real-time tracking of RNA in live cells using Sticky-flares. Proc Natl Acad Sci. 2015;112(31):201510581. doi:10.1073/pnas.1510581112.

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**Figure 3:** Brightfield microscopy. A-19:

Cas13 +  $\beta$ -actin (1); no expression after 24

hrs. After 30/48 hours - no expression (not

shown), B-18: GFP Control; GFP expression





