

TEAM

Maxim Rossmann (primary contact), University of Cambridge CIMR, Department of Haematology,

mr579@cam.ac.uk. Maxim will contribute to the project with his knowledge of biophysics and will work on the development of the assay and device design. Maxim has experience in project management and also has worked on the development of commercial diagnostic tests.



Shengjiang Tan, University of Cambridge CIMR, st299@cam.ac.uk

Shengjiang will contribute to the project with his knowledge of cell biology and expertise in the ribosomopathies. He will work on the assay design and refinement.



Norberto Escudero, University

of Cambridge CIMR, ne272@cam.ac.uk
Norberto has expertise in molecular biology and genetic and will be working on the assay development and optimization.



Christine Hilcenko, University of Cambridge CIMR, Department Haematology, of ch@mrc-Imb.cam.ac.uk. Christine is an experienced protein chemist and work on the assav development. She will contribute by providing high quality proteins and ribosomes.



Christie **Nel**, Scaturio Ltd., Cambridge, christie.nel@gmail.com. Software engineer experienced in development of LabView applications and instruments based on platform. Christie will support the development electronics software. and



SUMMARY

Our lab works on understanding how ribosomes are put together from their component parts and how this process is regulated by the cell. In this project we aim to design a tool for monitoring protein synthesis (translation) *in vitro*. We aim to design a simple device that will allow for the tracking of translation dynamics, which can be used for the discovery of novel translation-inhibiting or activation molecules and ribosome profiling for biophysical or structural studies. The device could also be used for teaching purposes and will serve as a prototype for a diagnostic device for the preliminary/quick detection of ribosomopathies,

i.e. inherited disorders characterized by the disruption in the ribosome assembly pathway and low protein synthesis capacity - an emerging class of human developmental and cancer predisposition disorders.

PROPOSAL

i) Problem

Ribosomes and protein synthesis in general are subjects of intensive research. But measuring ribosomal activity is cumbersome and requires expensive reagents and equipment. A robust and low-cost system is needed [**Ref. 1-2**] to perform basic experiments for characterisation of ribosomes and investigation of translation cycle effectors.

ii) Biological system

To develop a prototype we will use commercial in vitro translation system PURExpress kit (NEB) as a characterized ribosome source. A liquid sample containing DNA or mRNA encoding for the eGFP protein will be mixed with the kits components, which will result in the formation of fluorescent protein. Fluorescence intensity will be monitored in real-time and used to quantify the fraction of active ribosomes. Normalised translational activity will be used to draw conclusions about the ribosome number and performance.

iii) Design goals for the hardware

We aim to perform reaction and analysis using commercially available disposable 10 μ L slides Countess (Invitrogen) (**Fig. 1-2**), which ensure precise reaction volume control. Upon mixing the reagents fluorescent eGFP will be produced and the intensity of the fluorescent light will be measured in real-time by a RGB colour sensor with I2C interface. A UV LED will be used as the source of incident light. We will decide whether to use any additional optical filters or multiple channels during the trials involving the model and reference samples. The data logging and primary processing will be done on the PC using the colour sensor manufacturers software provided with the demo boards. The device will be assembled in a 3D printed enclose (**Fig. 3**), controlled by Arduino and operated via a touch screen provided in the starter kit.

iv) how you plan to implement the project

We will start the project with the development of the detector unit. Using standard fluorescent solutions we will determine the optimal distance between the RGB detector and sample and sample and the UV lamp and learn how to read the output. In the next step we will work on the optimal reaction condition using standard plate reader and then translate knowledge onto the prototype. Once we have a robust and consistent output we will engineer the enclosure and let it print using 3D printing service at the Addenbrookes Hospital.

v) proposed outcomes and benefits

In the first place we plan to use the developed device for the analysis of the ribosome preparations. The device will also be used to measure translation inhibition and activation in microbiology, cell- and plant biology, drug discovery projects targeting ribosomes and could serve as a prototype for a simple diagnostic device to detect ribosomopathies, such as for example SBDS (Shwachman-Diamond Syndrome) [Ref.3].

Budget:

Total	715		
Disposable <i>Countess</i> slides	30		
Expression plasmid	90		
RGB Sensor TCS3472 I2C 3D printing costs PURExpress Kit E6800S Reagents	55 200 190 100		
		LED UV5TZ-390-30:	50

Fig 1. RiboReader - general design.

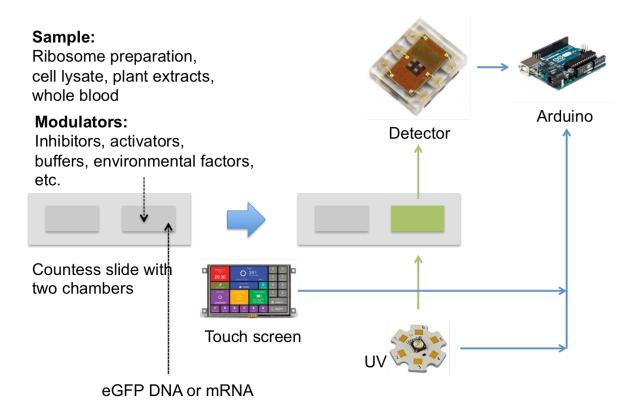
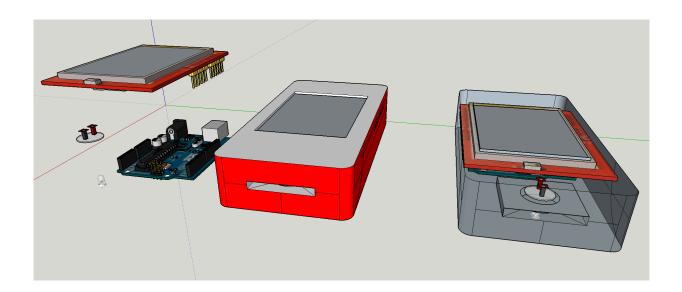


Fig 2. Countess disposable 10uL slide chamber



Fig 3. RiboReader.



References:

- (1) Rosenblum G., 2012
 (2) Capece CM, 2015
 (3) Wong CC., 2011