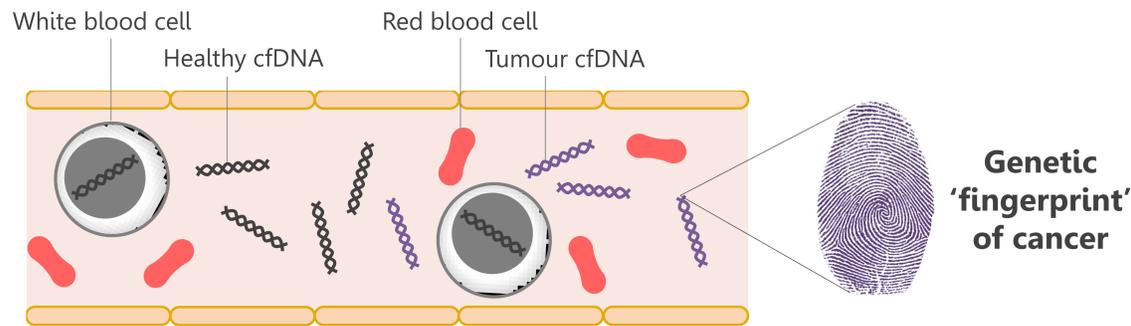


# Circulating free DNA: a 'liquid biopsy' for the early detection of cancer?

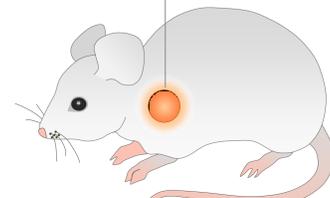
## Circulating free DNA



- Blood contains tiny amounts of free-floating DNA, called circulating free DNA (cfDNA)
- It is thought to leak into the blood from normal cells when they die
- In patients with cancer, DNA leaks into the blood from dying tumour cells
- Cancer is a genetic disease – thus, tumours release their genetic 'fingerprints' into the blood

## Mouse models

Tumour growth in organ of choice



- Early-stage cancers often go undetected, so it is difficult to study cfDNA in these patients
- Mouse cancer models can mimic the human disease and could help determine **how early** cfDNA reveals signs of disease

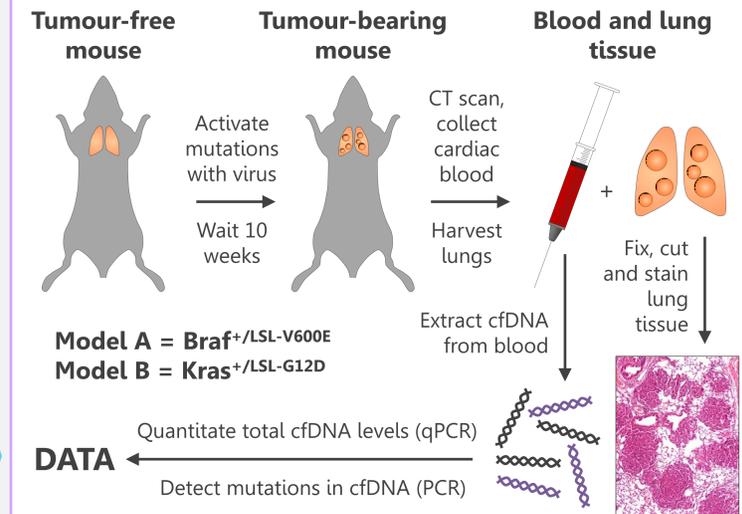
## Research question

Can we use cfDNA as a non-invasive 'liquid biopsy' to detect cancers before other tests?

## Study aim

To determine whether cfDNA can be isolated and quantitated from small volumes of blood in two well-characterised mouse models of lung cancer

## Methods



## Conclusions

- This study has detected higher levels of cfDNA in the blood of mice with pre-cancerous lung lesions, and has identified the lesion-causing mutations in the cfDNA
- An appropriate next step would be to allow the lung lesions to develop over a longer period of time, to allow them to become cancerous
- Collection of blood over a series of time points could determine if the cancerous lesions leak more mutated DNA as they grow larger

## Acknowledgements

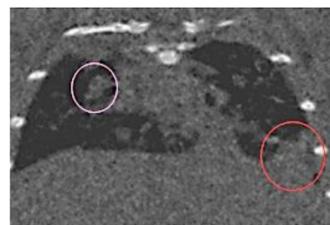
Many thanks to:

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- my PhD colleague, Callum Rakhit, for his input into the project
- Susan Giblett for her kind technical support
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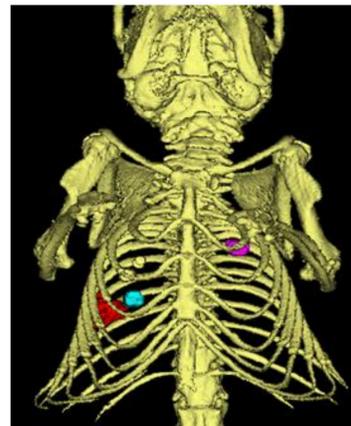
## Results

### 1) Pre-cancerous lung lesions were identified at 10 weeks by CT scanning

**Right:** circles show pre-cancerous lung lesions after 10 weeks by CT scanning.

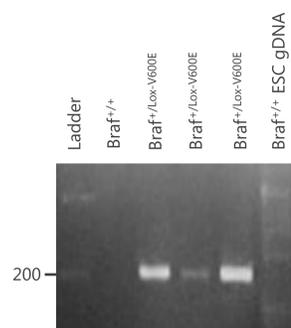


**Right:** many images produced by the CT scanner can be combined into a 3D image. The coloured spots indicate pre-cancerous lung lesions.

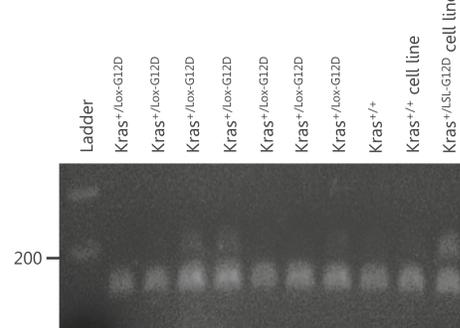


### 2) The tumour-causing mutations were identified in cfDNA of both models at 10 weeks, but not in cfDNA of healthy mice

Mouse model A

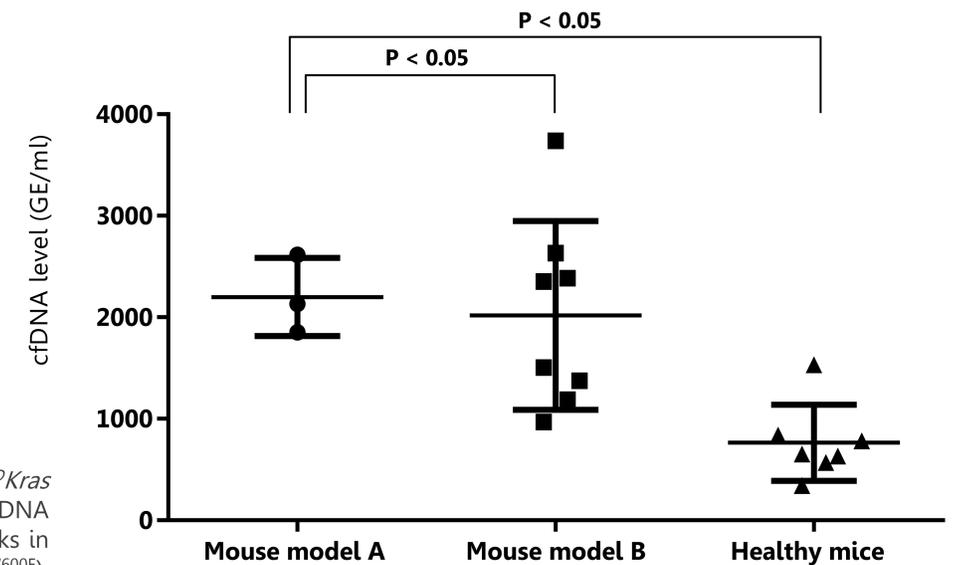


Mouse model B



**Left:** the mutant *V600EBraf* and *G12DKras* alleles can be detected in cfDNA isolated from plasma after 10 weeks in some mice. In model A (*Braf*<sup>+/LSL-V600E</sup>), PCR primers amplify a ~ 200 bp region specific to the mutant allele, but not the WT. WT mouse embryonic stem cell (ESC) DNA did not amplify. NTC, no template control. In model B (*Kras*<sup>+/LSL-G12D</sup>), PCR primers amplify ~ 140 bp and ~ 180 bp regions corresponding to the WT and mutant alleles, respectively. The mouse genotype is shown above each lane.

### 3) cfDNA levels were elevated in both models at 10 weeks, compared to healthy mice



**Above:** After 10 weeks, model A mice (*Braf*<sup>+/Lox-V600E</sup>; n = 3) had an average cfDNA concentration of 2200 genome equivalents (GE)/ml blood, and model B mice (*Kras*<sup>+/Lox-G12D</sup>; n = 8) had an average cfDNA concentration of 1970 GE/ml blood. Healthy controls (WT; n = 7) had an average cfDNA concentration of 760 GE/ml blood. Lines represent the mean +/- standard deviation. GE = genome equivalent = 6.6 pg DNA/cell.