

## Characterising the sensitivity of super-enhancer hijacked lymphoid malignancies to epigenetic perturbation

Newcastle University

of A-485 treatment.

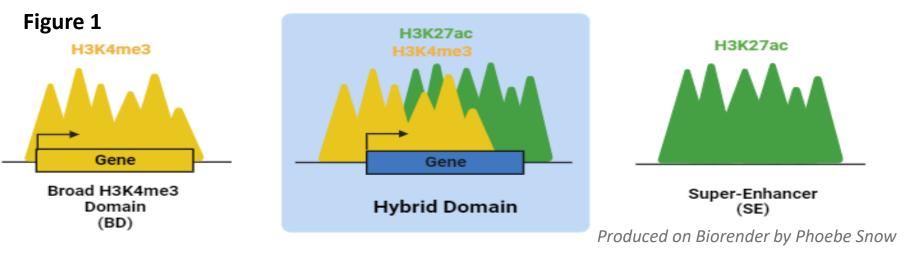
Claire Mowbray, c.mowbray2@newcastle.ac.uk

Marchetti and Phoebe Snow

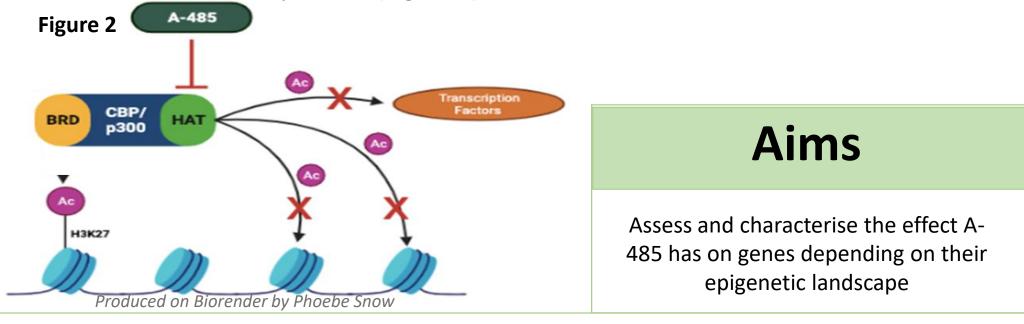
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## Background

- Acute lymphoblastic leukaemia (ALL) is a cancer of lymphoid cells in the blood and bone marrow.
- ALL is treated with chemotherapy and efforts are being made to increase our understanding of the genetics fuelling the disease.
- The epigenetic landscape forms a layer (marks) over the genetic code, altering the level of activity of certain genes.
- 'Marks' on this landscape, are linked to the activity of a gene H3K4me3 denotes promoter activity. When these marks are stretched, they form a 'broad domain', H3K27ac denotes a 'super enhancer'.
- These two marks can overlap to form a hybrid domain (Figure 1).



- Super enhancers are known to be hijacked in cancer cells and can fuel the progression of disease.
- The drug A-485 acts to inhibit the enzyme EP300, which has a region that deposits H3K27ac onto histone proteins (Figure 2).



## Methods

- 4 flasks of GM12878 cells (healthy lymphoblasts) were set up in sterile conditions a negative control and increasing concentrations of A-485. these flasks were left for 48/96 hours.
- Cell viability was measured, and RNA extracted.
- RNA was converted to cDNA, and gene quantities analysed by qPCR.

# Figure 3 A485 48/96 hours PNA Produced on Pierender

- The entire experiment was performed three times to produce a set of biological triplicates and analysed by the 2-way-Anova test.
- 1.NHS Acute ;lymphoblastic leukaemia. https://www.nhs.uk/conditions/acute-lymphoblastic-leukaemia/
- 2.. Transcriptional regulation by Fos and Jun in vitro: interaction among multiple activator and regulatory domains. Mol Cell Biol. 1991 Jul; 11(7): 3624–3632

#### Results 'Normal' genes show no change or increase in response Figure 4 Figure 5 A-485 treatment does not affect cell viability to A-485 Comparison of the Gene without 100% percentage of live hybrids, broad cells and the 95% domains or percentage of dead 90% enhancers. cells after drug Significant results 85% treatment are indicated by 80% bars with stars. Significance was 75% calculated using the **DMSO** $3\mu M$ $5\mu M$ 2-way-Anova test. ■ Live ■ Dead Some genes decrease in response to 48 hours of A-485 treatment. Others show not significant changes Figure 6 Hybrids with Fold change Gene changes in response to 48 hours of A-485 treatment. Figure 7 Some genes decrease in response to 96 hours of A-485 treatment. Others show not significant changes old change Gene changes in response to 96 hours

## Discussion

- The results from qPCR analysis showed that some genes, such as PIK3R5 were significantly downregulated after 48/96 hours A-485 treatment, however there was no trend between the 'normal', broad domain only or hybrid groups. Neither was there significant changes in gene expression between 48 and 96 hours of drug treatment.
- The two genes that showed a significant increase in expression due to A-485 treatment are response genes that increase expression due to a range of cellular stimuli<sup>2</sup>. The increase of these genes JUN and FOS show in this context, that there is indeed something occurring in the cells due to A-485 exposure.
- Promoter capture high seq data was used to profile the interactions each gene had with other areas of the genome. Preliminary data suggests that the genes that showed significant downregulation did not interaction with other broad domains and/or hybrid domains around the genome.
- Unfortunately, there was wide variation between the fold change of genes between the three biological replicate experiments, limiting the extent analysis can be performed.

## **Next Steps**

- RNA from some of the samples have been sent off for RNA sequencing.
- Other members of the group will complete the analysis of A-485 treated Z-138 cells

## Summary

Genes with different epigenetic profiles respond differently to EP300 inhibition. There is strong potential to develop this study further in the hope to improving treatment of ALL.