

Modulating gene expression in human T cells with PETRA



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Motivation

Cellular therapies use patient's immune cells to treat cancer We need to Killing of solid tumour No killing of solid tumours Efficient killing of by T cells with **enhanced function** find a way to due to insufficient T cell function¹ liquid tumours enhance T cell function Patient's T cells targeted to tumour

Problems

of different genes and can be enhanced by modulating gene expression².



Our solution: PETRA

<u>Prime Editing of Transcribed Regulatory elements to Alter expression</u>

PETRA allows us to test the effects of thousands of edits on gene expression, in a single experiment



Edits can be combined for enhanced

effects

X10

Using PETRA we tested the effect of 11,000 edits on the expression of 4 genes with key roles in T cell function

We discovered edits that both increase and decrease gene expression





MYBL2 motif insertion in different positions in 5' UTR of IL2RA. Edits scored with PETRA in Jurkat cells. Datapoints show replicate score, bars represent mean score between replicates. p<0.05 for all pairwise comparison between groups, Student t-test. Combining PETRA scores with AI models reveals biological mechanisms underlying edit effects



Implications and future applications

PETRA generates large datasets linking edits to their effects, which provide a 🖉 valuable source of information about gene expression...

...and can be used to train new Machine Learning models to better predict edit effects on gene expression in the future.



Edits discovered using PETRA could provide a solution to enhance T cell function and improve cellular therapies.



