

Implementing a high-throughput parallel CRISPRi screening platform to identify functional lncRNAs

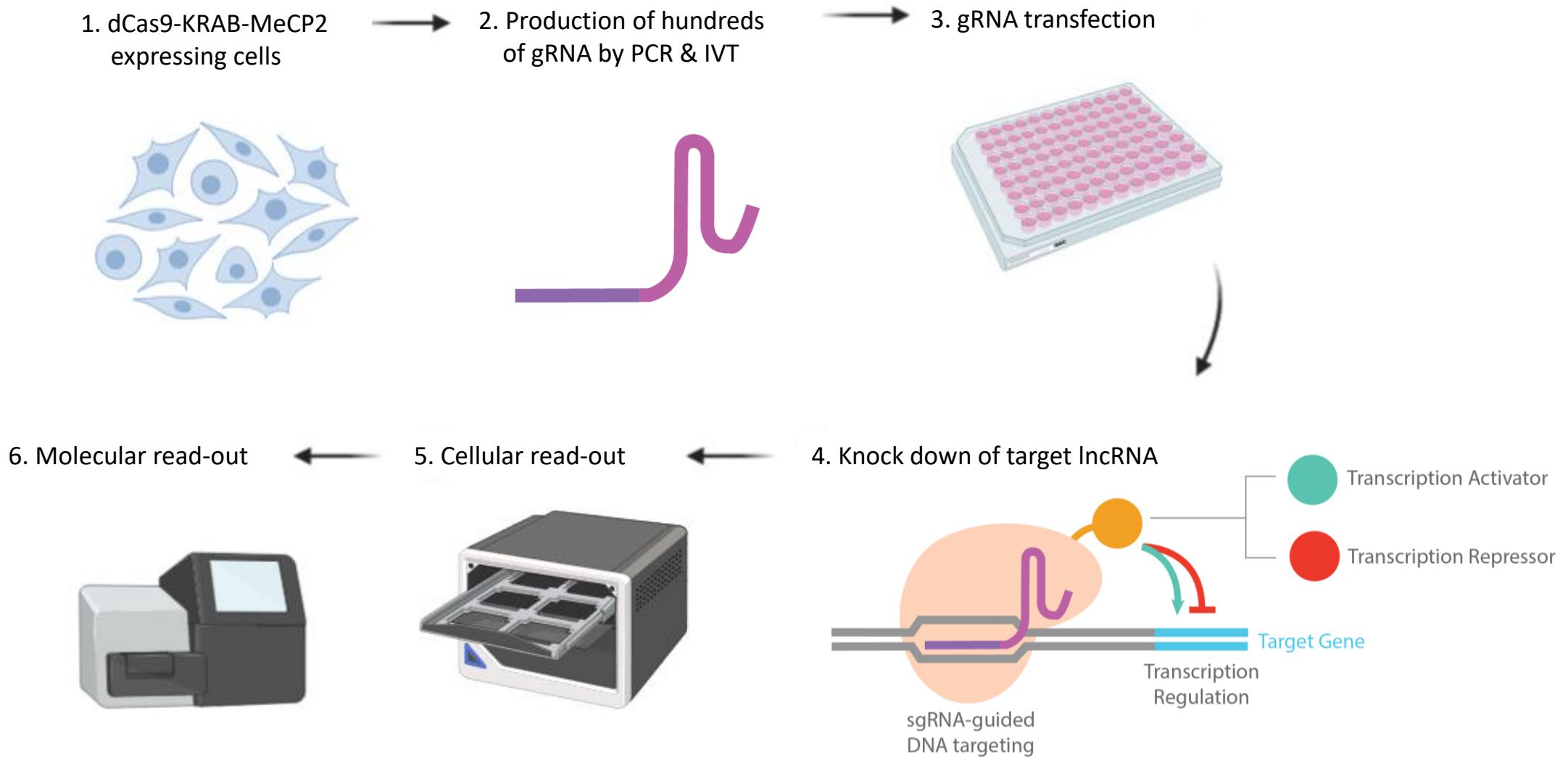
Fien Gysens

25/03/21

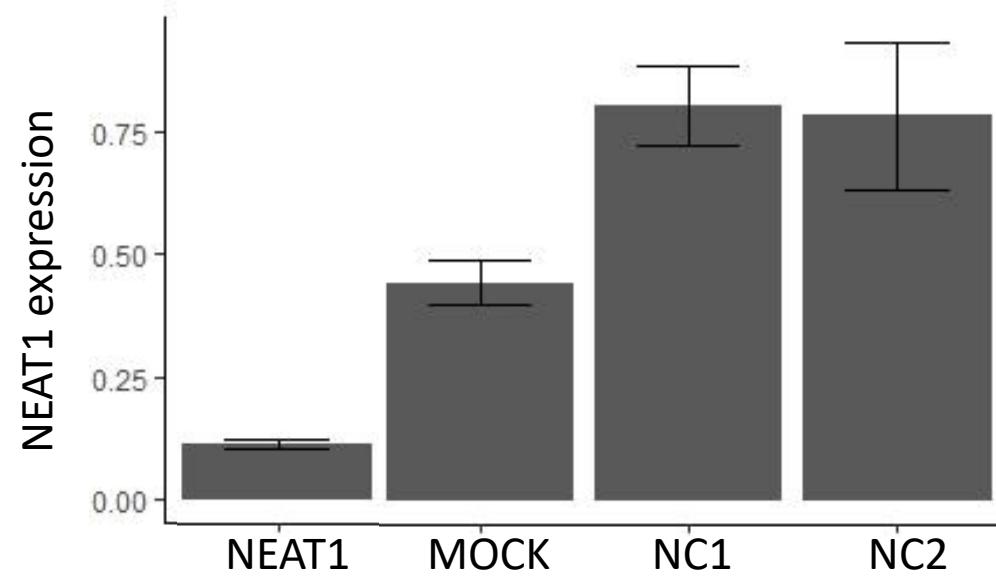
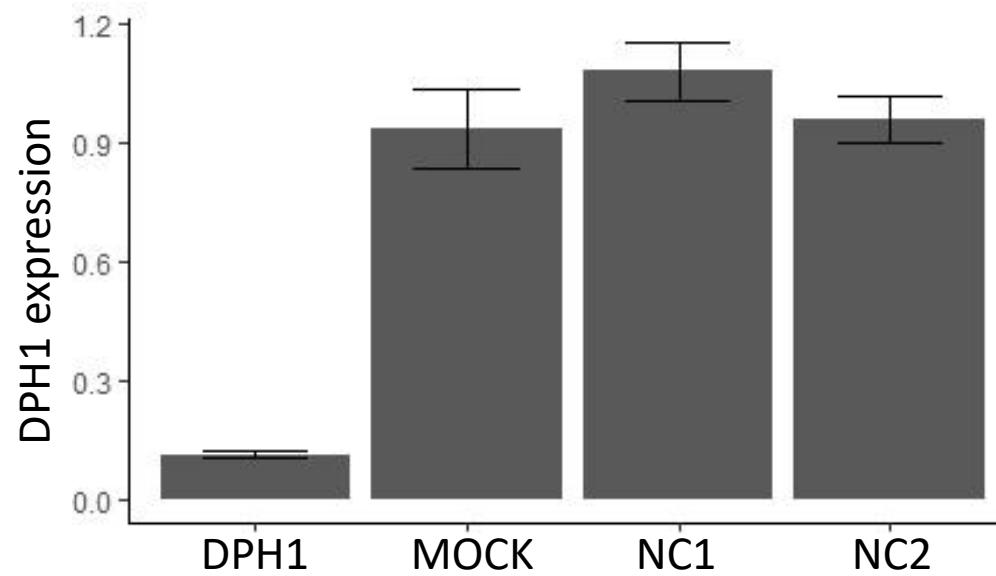
Important to investigate the function of lncRNAs

- 15 000-100 000 lncRNAs vs 21 000 protein coding genes
- Minor part studied on a functional level
- Are they all functional?
 - > Need for a high-throughput platform to probe functionality

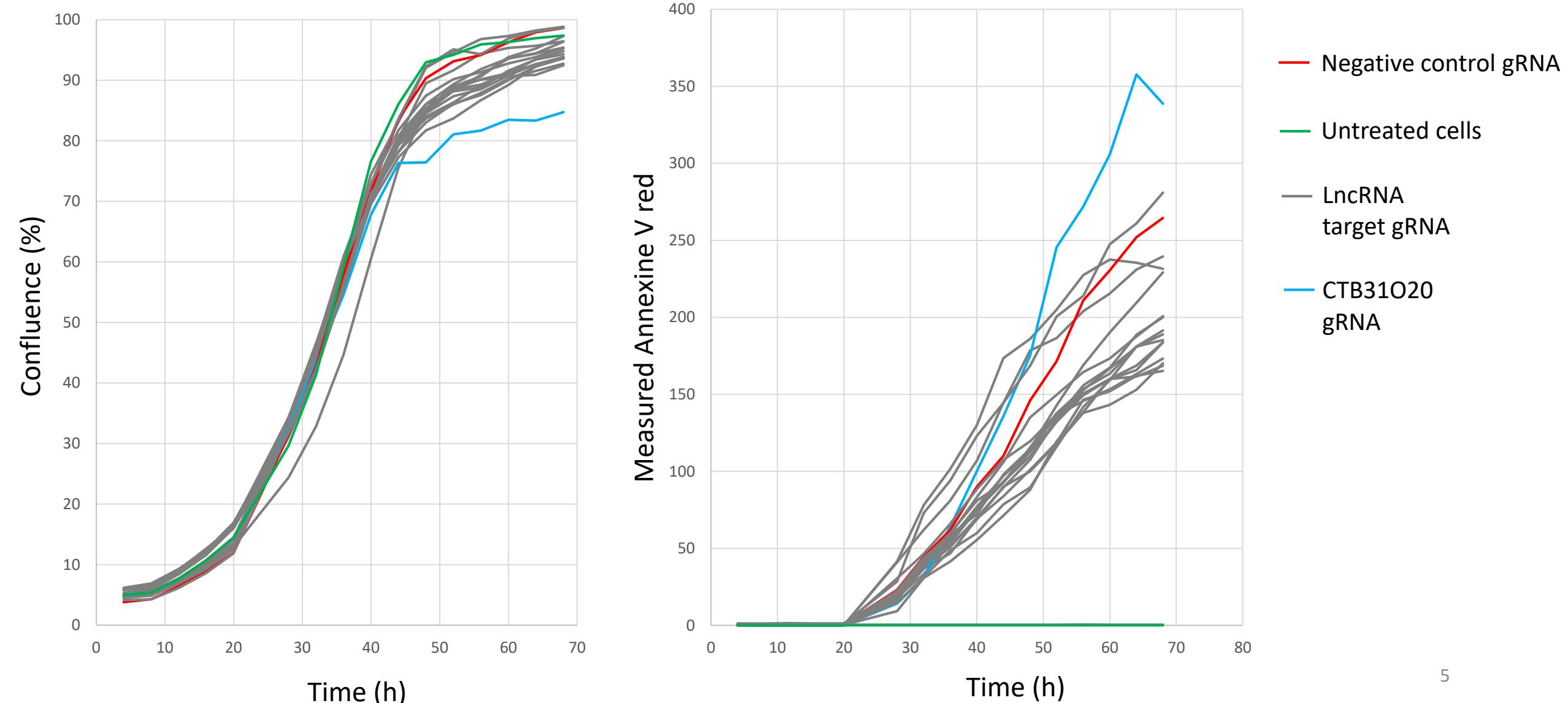
Workflow of a high-throughput parallel CRISPRi screen



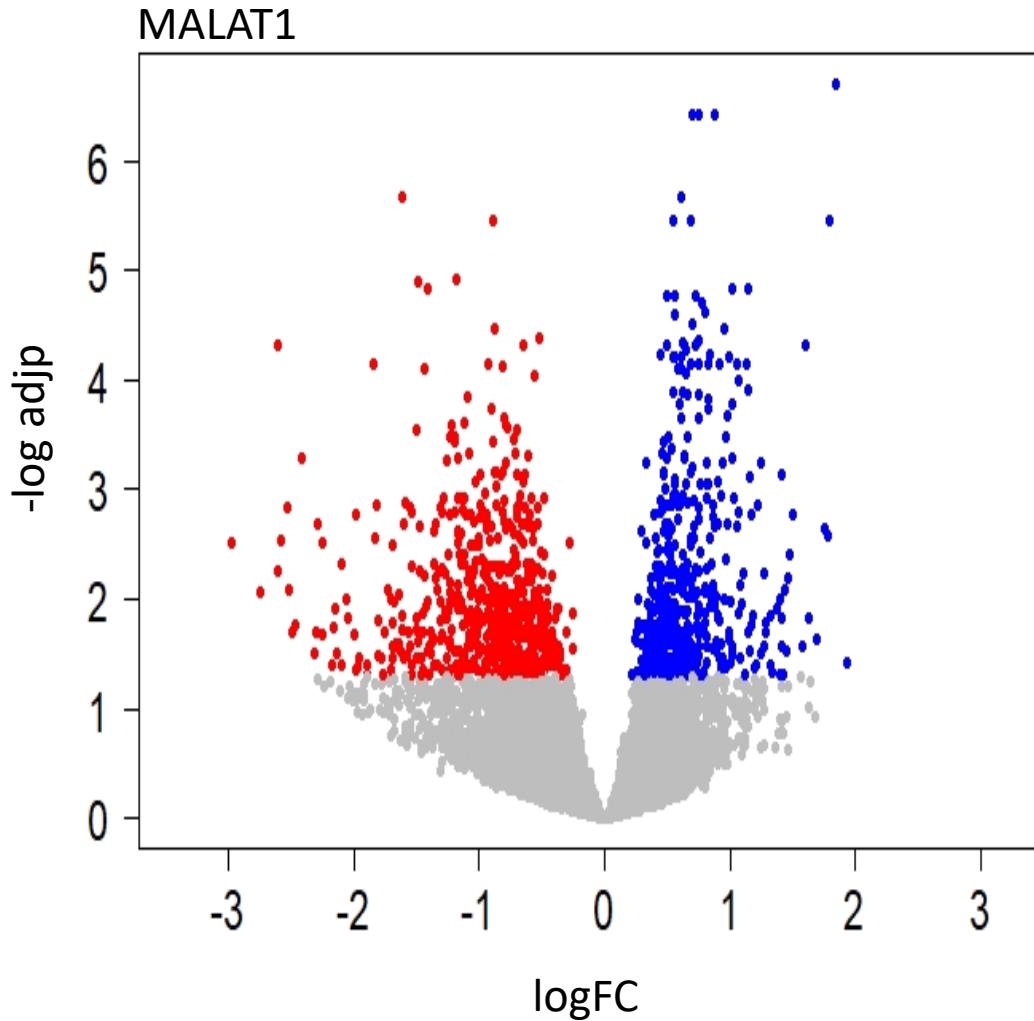
Functional CRISPRi machinery



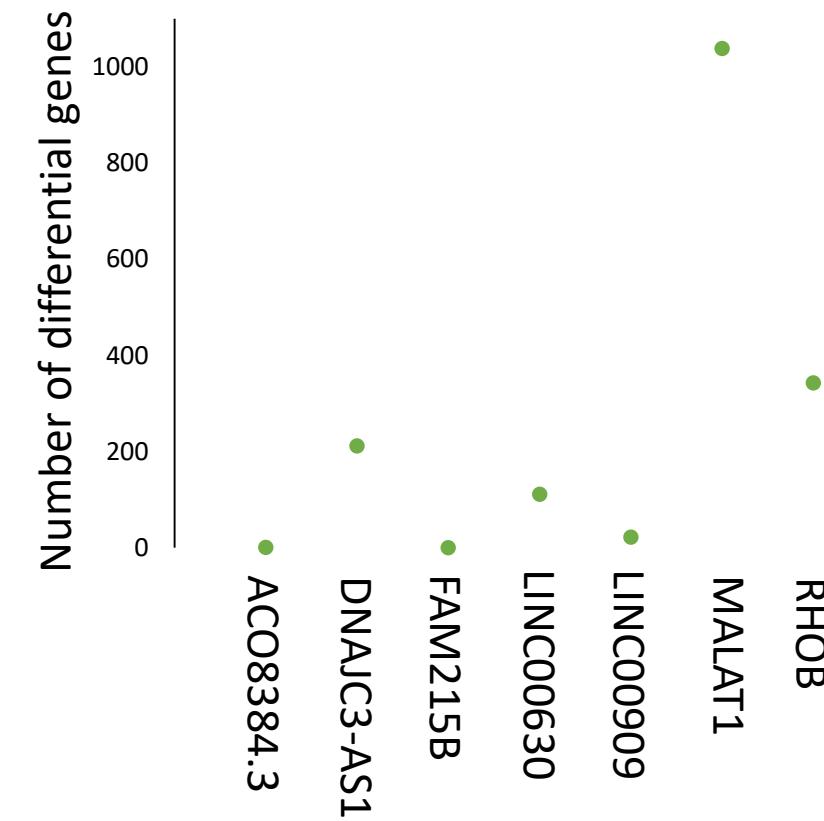
Cellular read-out: Real-time monitoring reveals that lncRNAs affect proliferation and/or apoptosis



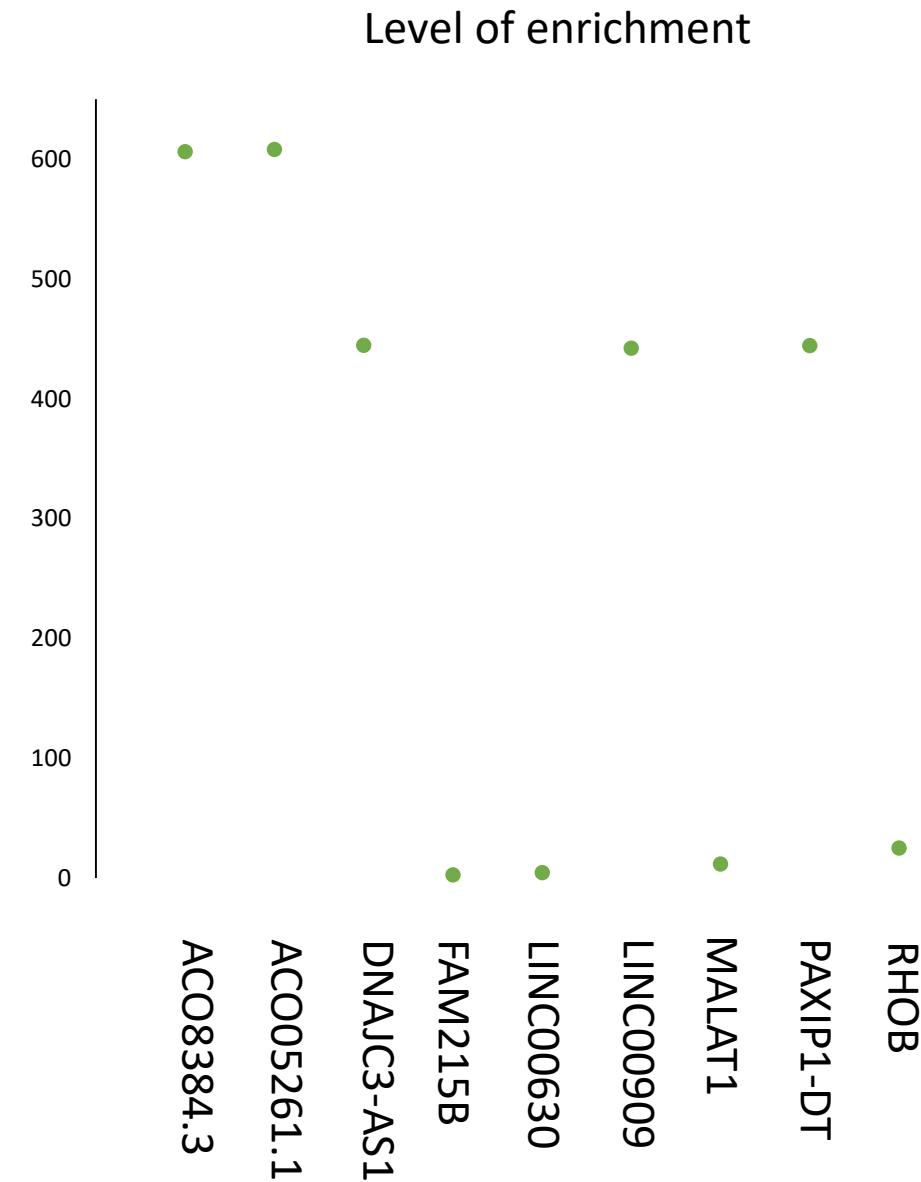
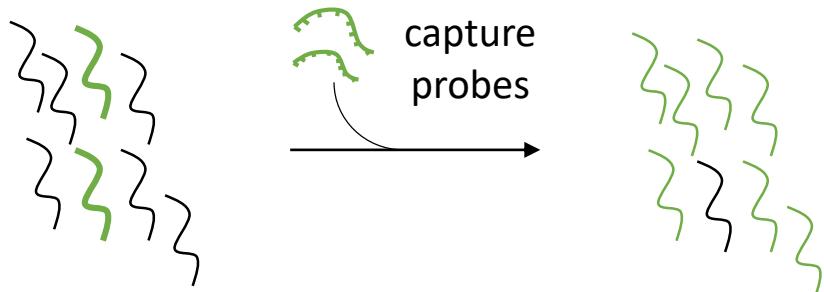
Molecular read-out: Differentially expressed genes were detected after KD



- 1038 differentially expressed genes with MALAT1 KD
- 12 significantly affected gene sets with MALAT1 KD (GSEA analysis)



Using capture sequencing to detect KD of lncRNAs



What is next?

- Plan to screen for an extra 144 targets in HEK293T
- What with difficult to transfect cell lines
 - High-throughput lentiviral gRNA production
- Not limited to study lncRNAs
- Set-up an arrayed screen in different cell lines
 - Respiratory diseases
 - Immuno-ocology
 -

Acknowledgement

CMGG

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