

# Unravelling genetic factors underlying cancer cell adaptation - a foundation for developing cancer therapies





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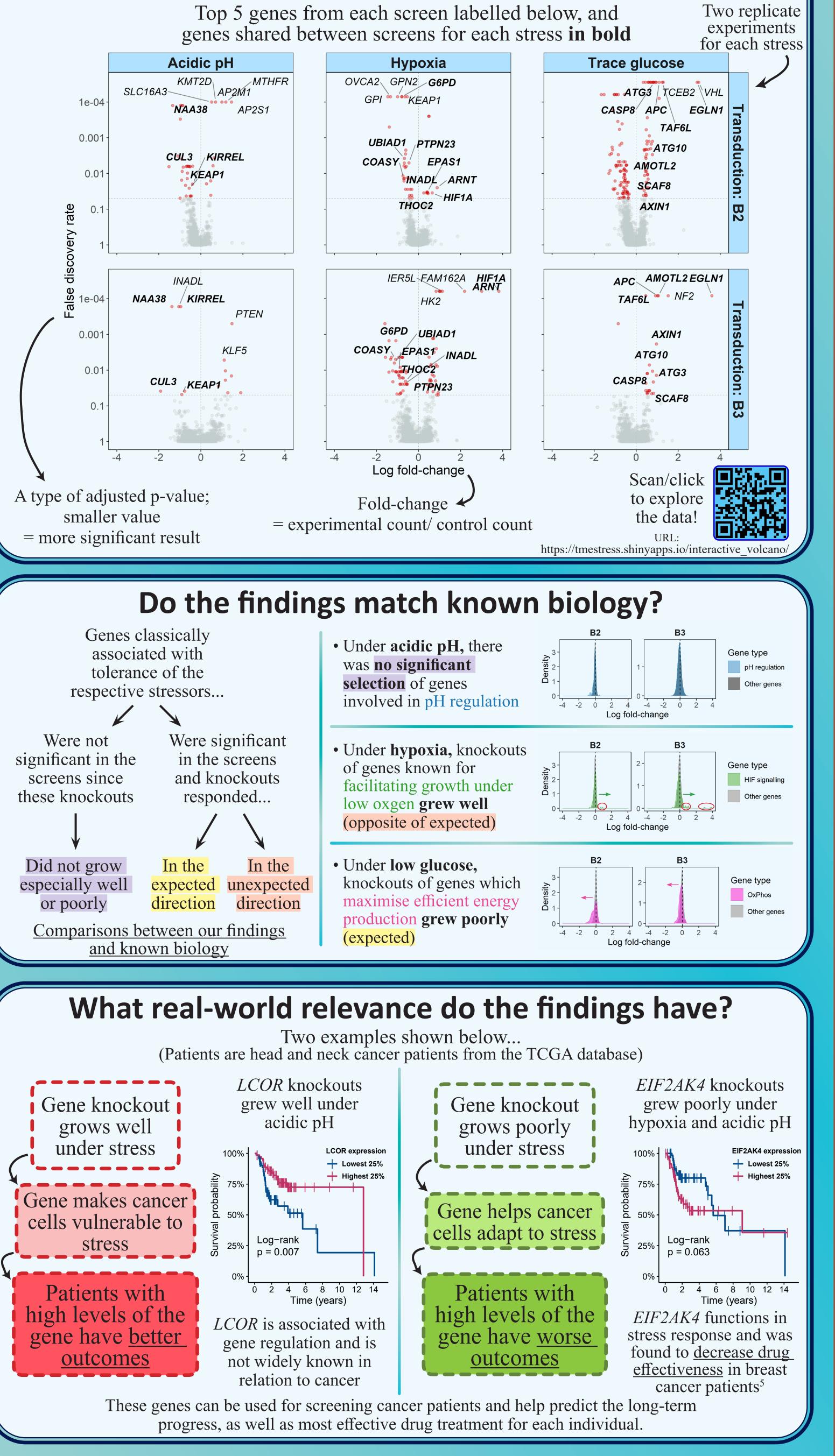
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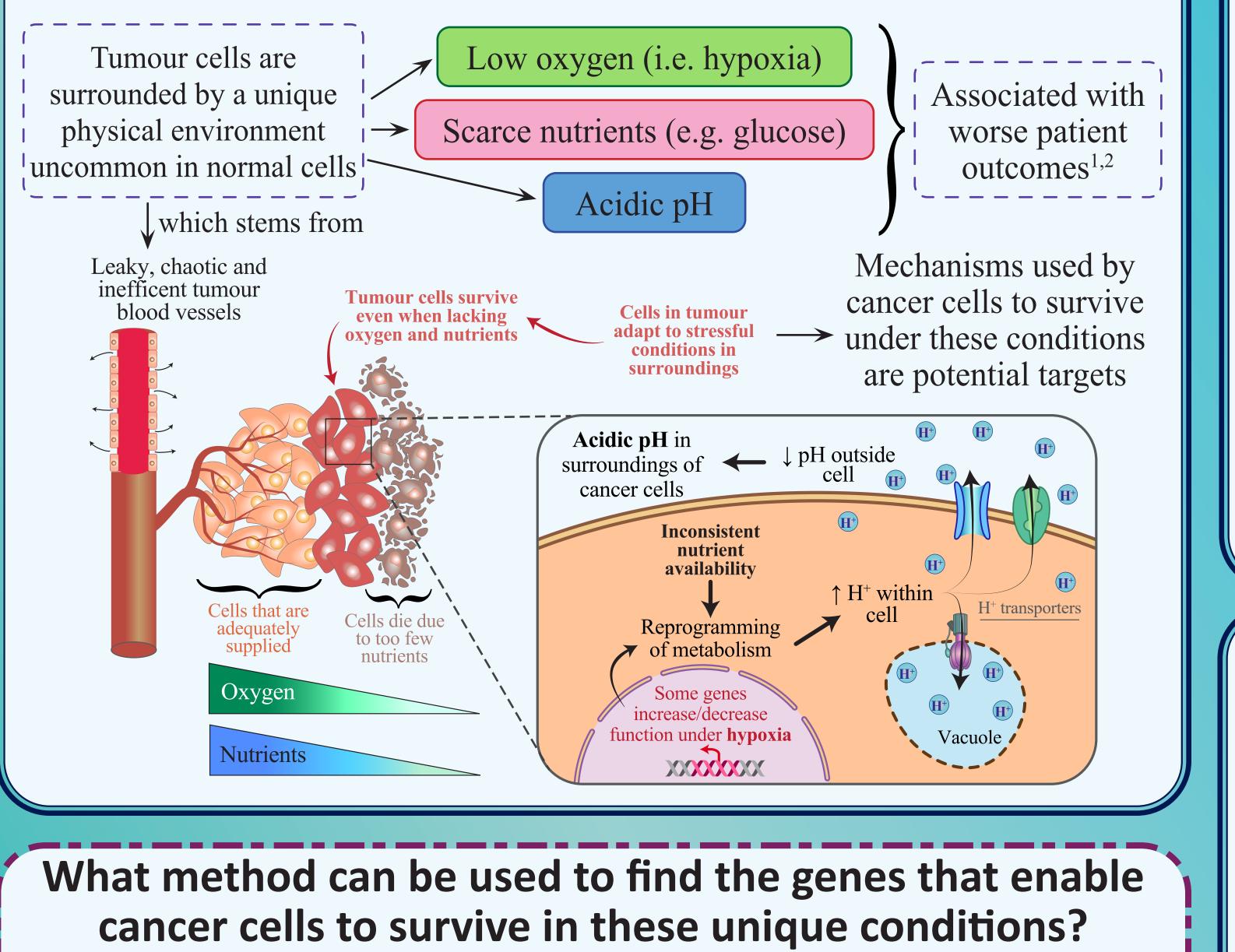
## Main questions

Which genes are important for cancer cell adaptations?
How can these results be used to benefit cancer patients?

How can we target cancer cells specifically?

#### Which genes were most important in the screens?





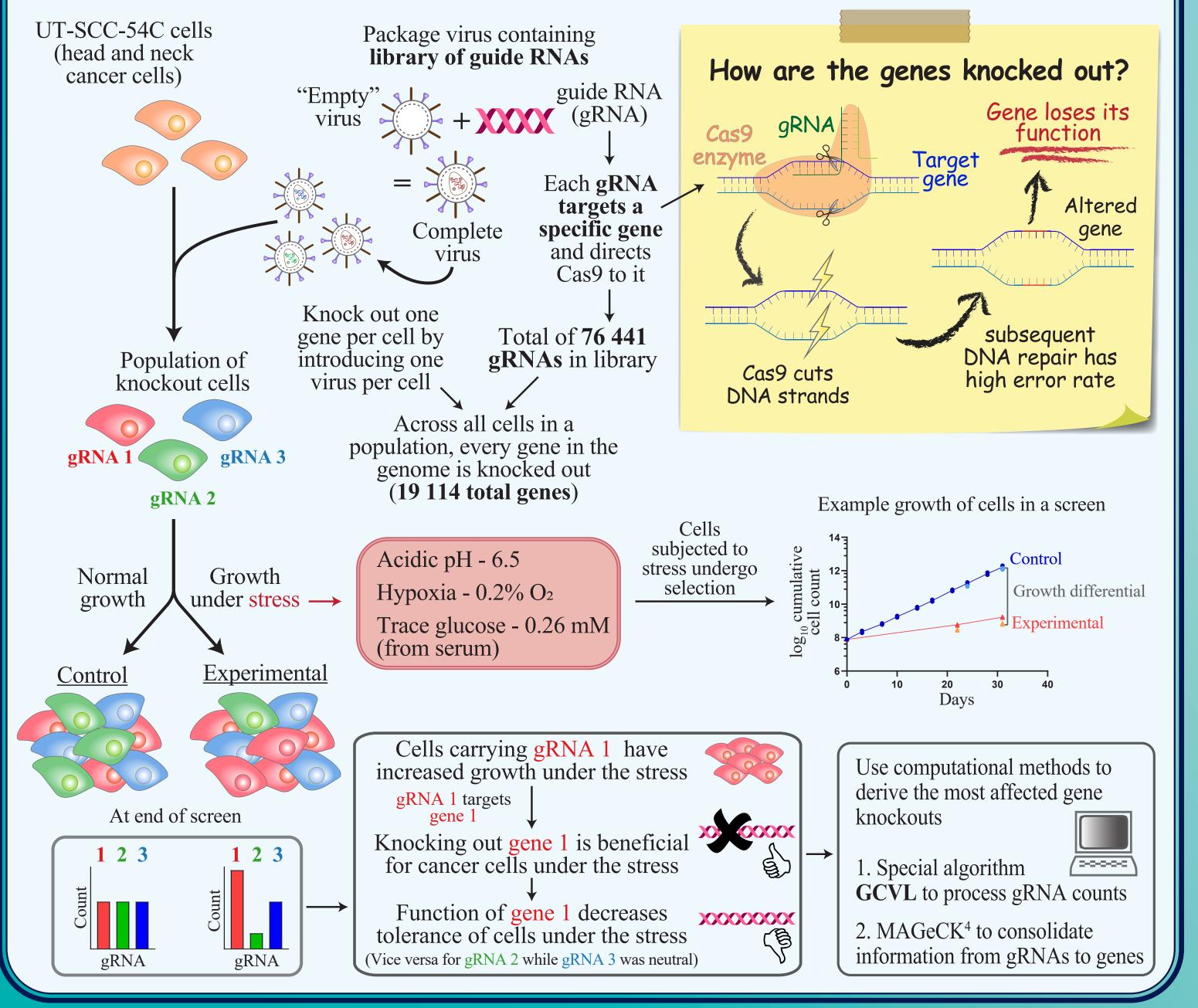
Test how disrupting (i.e. knocking out) each gene affects cancer cell survival Thousands of genes exist, so testing one gene at a time will take too long Instead we can do so for all genes across the genome in parallel using genomic knockout screens<sup>3</sup>

#### Why is this knowledge useful?

By identifying the genes that are essential for cancer development, we can potentially

- ✓ Improve our knowledge of the biology underlying this stress
- ✓ Develop new anti-cancer drugs
- $\checkmark$  Use them for genetic screens to match cancer drugs to cancer patients

### What steps are involved in a genomic knockout screen?



Figures created with Adobe Illustrator 24.3, graphs plotted with R (v4.0.3; RStudio) and statistical tests carried out in R.

#### **Overall takeaways**

- Our genomic knockout screens successfully identified genes required by cancer cells to adapt to their stressful surroundings
- ✓ Many gene knockouts significantly affected cancer cell survival under stress further study into these genes may lead to future cancer therapies
- ✓ Some findings are not consistent with the biology that is currently known, constant ongoing research is thus important
- ✓ Several genes were associated with difference in survival and treatment response of cancer patients, and may be useful for applications in the clinic

#### References

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- 4. Li, W. et al. (2014) MAGeCK enables robust identification of essential genes from genome-scale CRISPR/Cas9 knockout screens. Genome Biology, 15(12).
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