

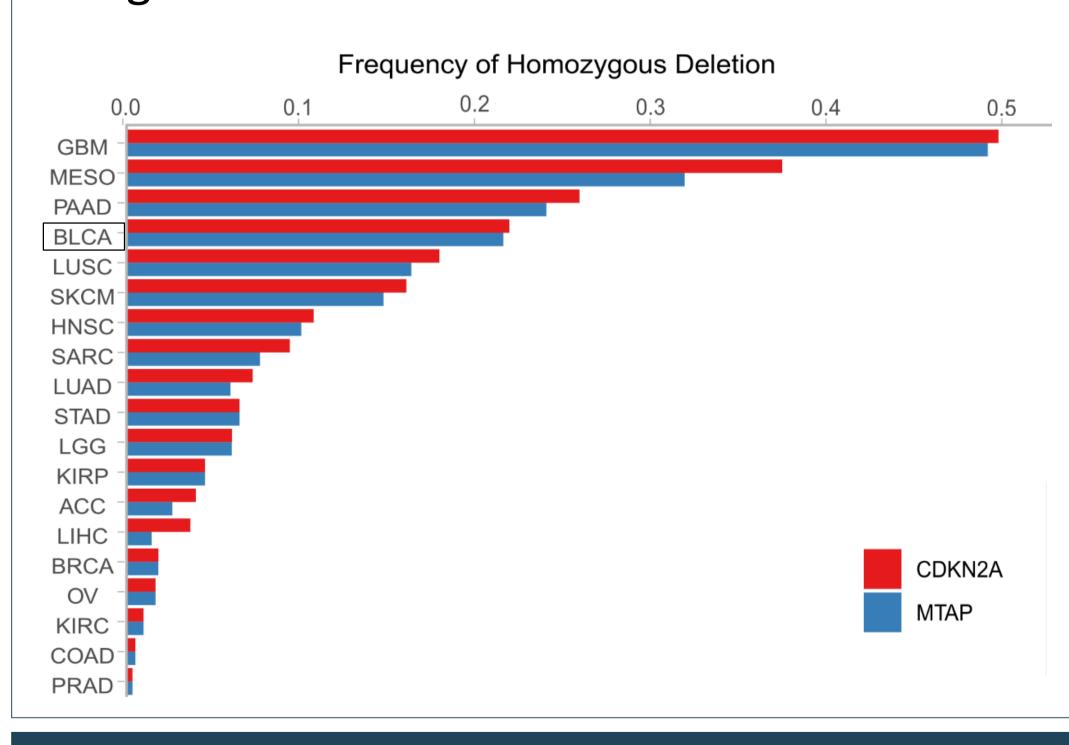
# Exploration of therapeutic vulnerabilities exposed by 9p21 loss in bladder cancer cell lines

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## Background

The deletion of chromosome 9p21 locus is the most frequent homozygous deletion in bladder cancer<sup>1</sup>. It causes loss of the tumor CDKN2A/2B and metabolic gene MTAP, involved in the methionine and adenine salvage pathway. Large-scale shRNA screens have shown enhanced dependency of MTAP-deleted cells on PRMT5 and MAT2A, which led to the development of highly specific inhibitors (i.e. MRTX1719 for PRMT5 and AG270 for MAT2A)<sup>2-6</sup>. Here, we performed a multiparametric drug screening to uncover new pharmacological vulnerabilities of 9p21deleted bladder cancer cells, a disease with limited treatment options in the advanced stages<sup>7</sup>.



#### References

- 1. Robertson, A. G. *et al.*Comprehensive Molecular Characterization of Muscle-Invasive Bladder Cancer. *Cell* **171**, 540-556.e25 (2017).
- 2. Mavrakis, K. J. *et al.* Disordered methionine metabolism in MTAP/CDKN2A-deleted cancers leads to dependence on PRMT5. *Science* **351**, 1208–1213 (2016).
- Kryukov, G. V. *et al.*MTAP deletion confers enhanced dependency on the PRMT5 arginine methyltransferasein cancer cells. *Science* 351, 1214–1218 (2016).
   Marion, K. *et al.* MTAP Deletions in Cancer Create Vulnerability to Targeting of the
- MAT2A/PRMT5/RIOK1 Axis. *Cell Rep.* **15**, 574–587 (2016).

  5. Kalev, P. *et al.* MAT2A Inhibition Blocks the Growth of MTAP-Deleted Cancer Cells by Reducing PRMT5-Dependent mRNA Splicing and Inducing DNA Damage. *Cancer Cell* **39**, 209-224, e11 (202).
- PRMT5-Dependent mRNA Splicing and Inducing DNA Damage. *Cancer Cell* **39**, 209-224.e11 (2021).

  6. Smith, C. R. *et al.* Fragment-Based Discovery of MRTX1719, a Synthetic Lethal Inhibitor of the PRMT5•MTA Complex for the Treatment of *MTAP* -Deleted Cancers. *J. Med. Chem.* **65**, 1749–1766 (2022)
- 7. Faltas, B., Prandi, D., Tagawa, S. *et al.* Clonal evolution of chemotherapy-resistant urothelial carcinoma. *Nat Genet* **48**, 1490–1499 (2016).
- Alhalabi, O. et al. MTAP deficiency creates an exploitable target for antifolate therapy in 9p21-loss cancers. Nat. Commun. 13, 1797 (2022).

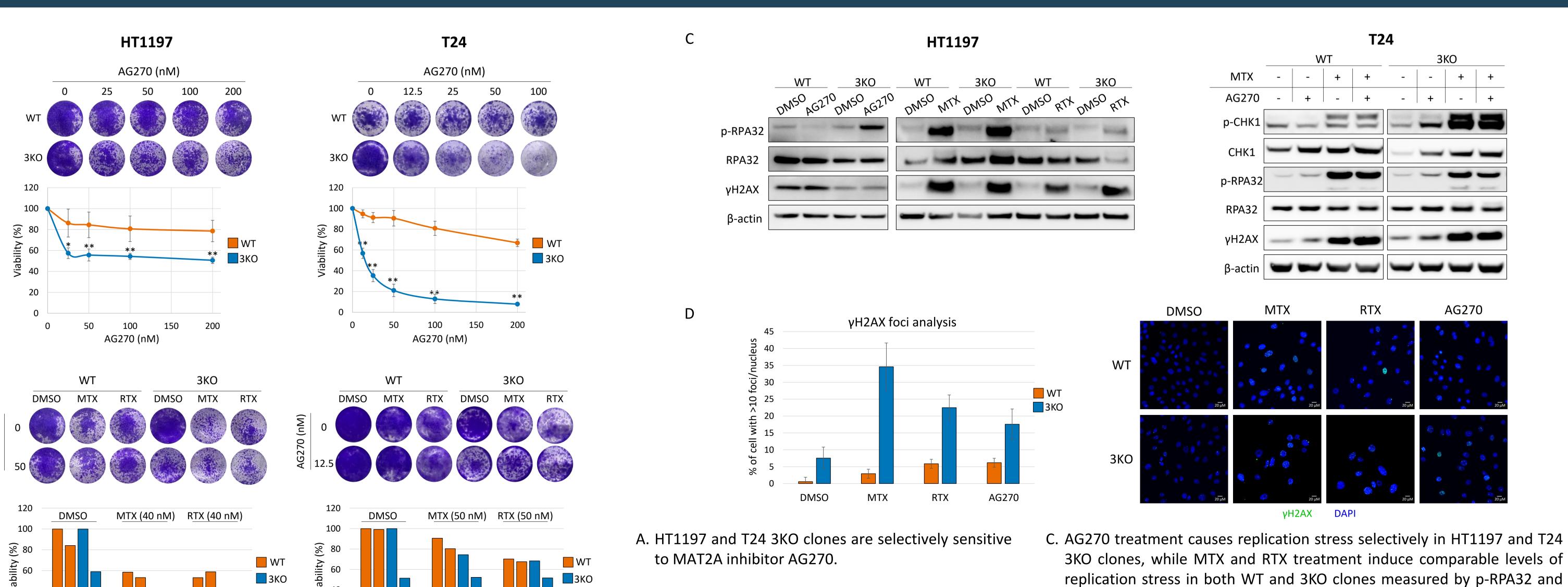
#### Drug screening workflow and validation Generation and characterization of 9p21 locus isogenic pairs Chr9: 20,900,00-23,900,000 (hg19 • The HT1197 isogenic pair was used to perform a multi-parametric drug screening testing 2351 compounds. DMRTA1 • 18 drugs that selectively affected 3KO clone cell number were found. • Methotrexate (MTX) and raltitrexed (RTX) were confirmed in doseresponse viability assays performed with several HT1197 WT and 3KO However, T24 3KO clones do not show enhanced sensitivity to MTX and RTX treatment compared to WT cells. ЗКО 🍎 👉 🧓 – MTAP-/-, CDKN2A/2B-/- (3KO) Competitive cell growth assay Symmetric Di-Methyl Arginine (SDMA) levels HT1197 3KO clones have higher proliferation rates SDMA levels are lower in 3KO clones respect to WT clones; compared to WT clones, while T24 clones proliferation is PRMT5 activity is partially impaired in HT1197 and T24 3KO clones not altered by 9p21 loss

p-CHK1 levels by Western Blot.

D. MTX, RTX and AG270 induce higher DNA damage levels in HT1197 3KO

cells compared to WT when considering vH2AX foci formation by IF.

## Dependency of 9p21 KO cells on MAT2A



T24 clones.

B. Combination treatment of MAT2Ai with MTX or RTX

in proliferation assays does not increase the effect

of either drug used alone neither in HT1197 nor in

# 5 Conclusions and future plans

We successfully generated isogenic bladder cancer cell lines (9p21 locus WT and 3KO) that recapitulate literature findings in terms of proliferation rate, reduced SDMA levels and sensitivity to MAT2A inhibition. Our drug screening nominated the antifolates agents MTX and RTX as therapeutic vulnerabilities of *MTAP*-deleted cells. Our findings are in line with a recent study showing that the antifolate agent pemetrexed is selectively effective in *MTAP*-deficient bladder cancer patients and preclinical models<sup>8</sup>. We are currently testing new combinations of drugs in order to increase treatment effectiveness in *MTAP*-deleted cells and patient-derived organoids.

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