

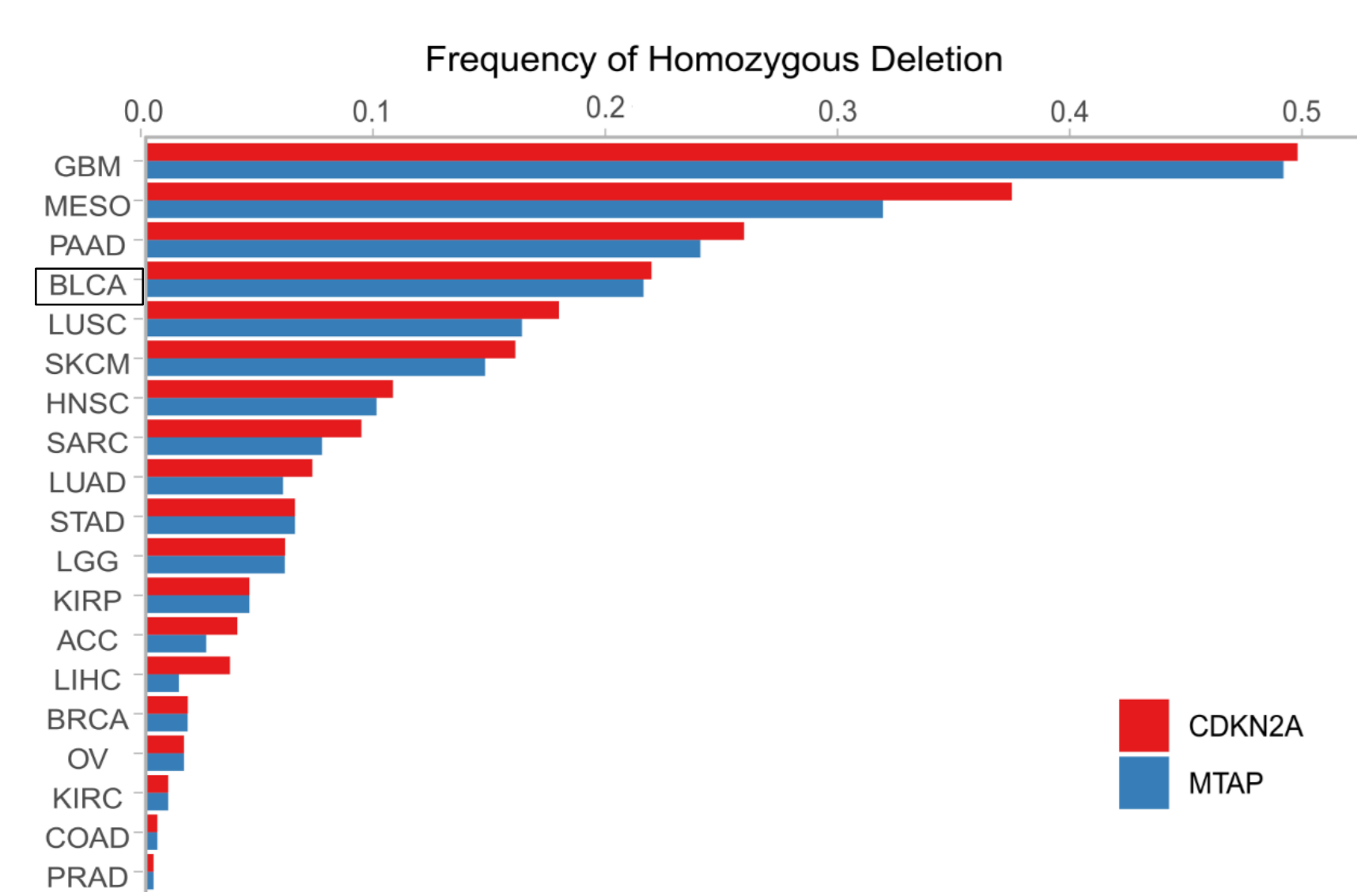
# Investigation of novel pharmacological vulnerabilities of 9p21-deleted bladder cancer cells

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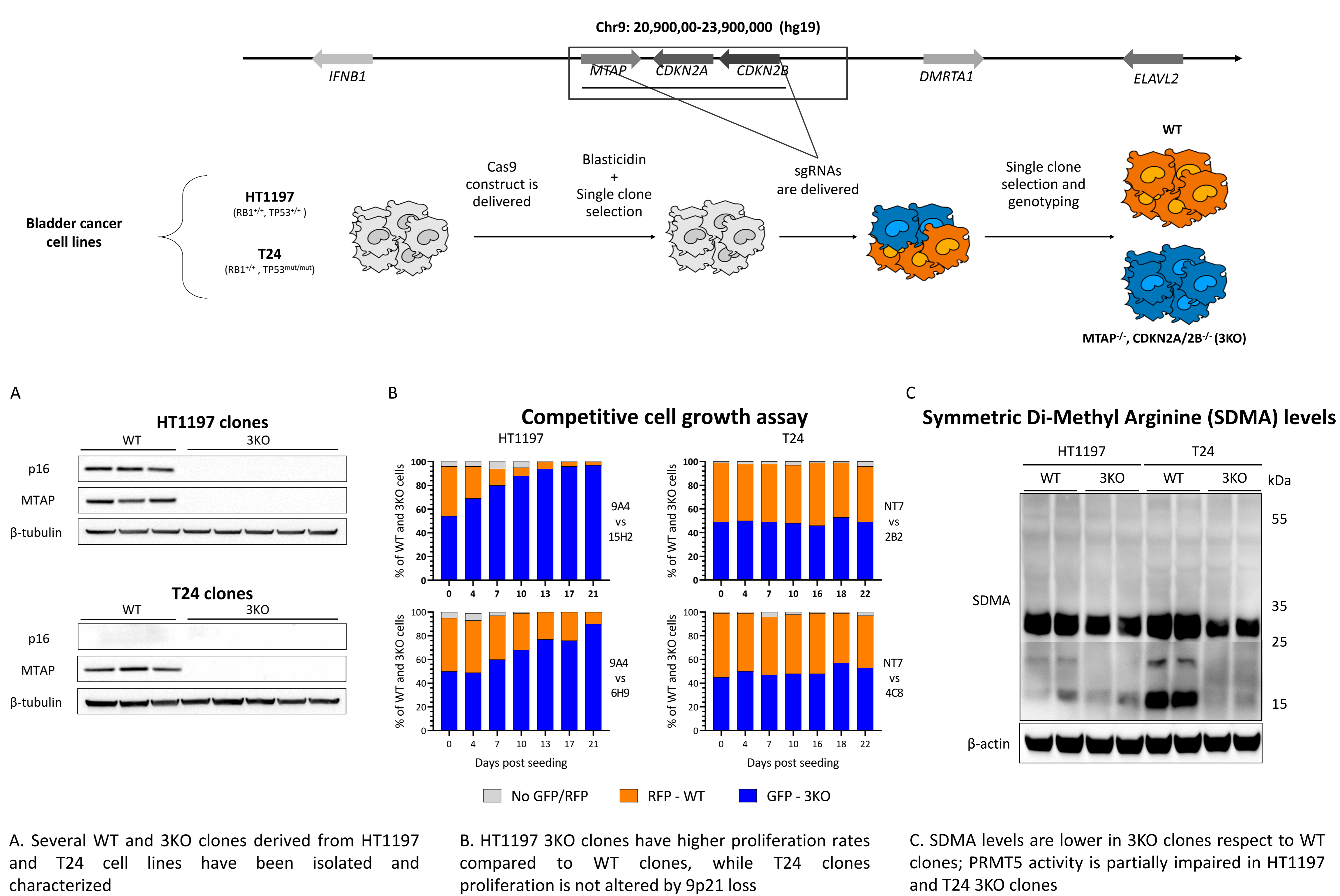
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## Background

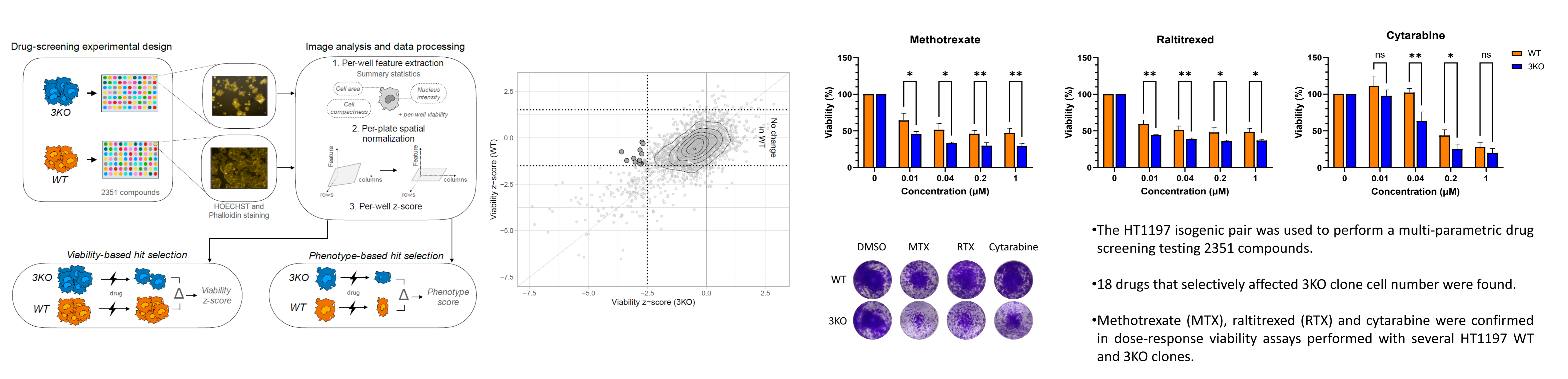
Deletion of the chromosome 9p21 locus is the most frequent copy number alteration in bladder cancer, identified in 23% of the TCGA muscle-invasive bladder cancer cohort<sup>1</sup>. It causes loss of the tumor suppressors *CDKN2A/2B* and of the metabolic gene *MTAP*, involved in the methionine and adenine 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 multi-parametric drug screening to uncover new pharmacological vulnerabilities of 9p21-deleted bladder cancer cells, a disease with limited treatment options in the advanced stages<sup>7</sup>.



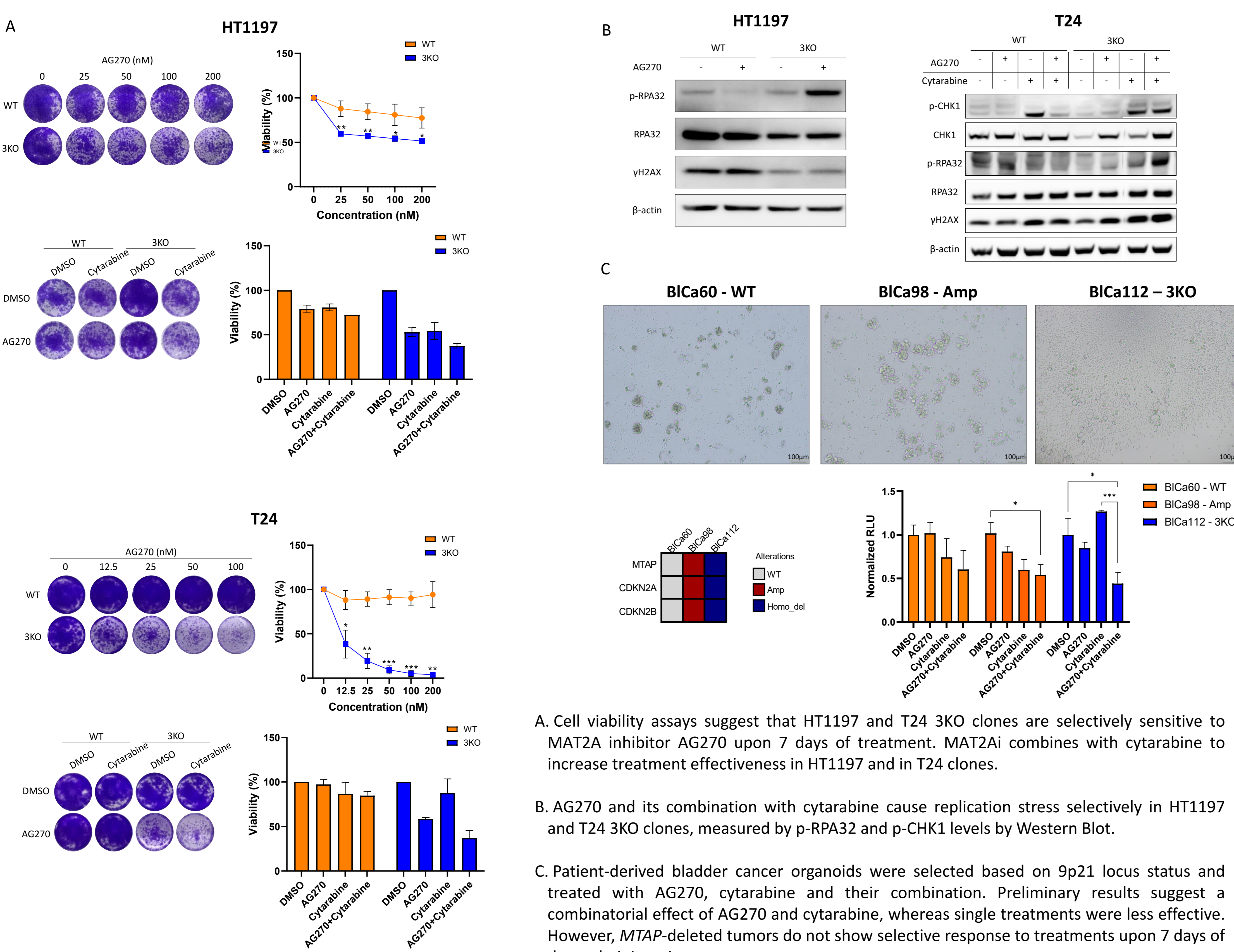
## Generation and characterization of 9p21 locus isogenic pairs



## Drug screening workflow and validation



## Dependency of 9p21 KO cells on MAT2A



## 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 two antifolates agents (MTX and RTX) and cytarabine 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>. AG270 effectively combines with cytarabine in isogenic bladder cancer cells. Preliminary results on patient-derived bladder cancer organoids suggest a combinatorial effect of AG270 and cytarabine but do not show clear genotype selectivity.

## References

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