

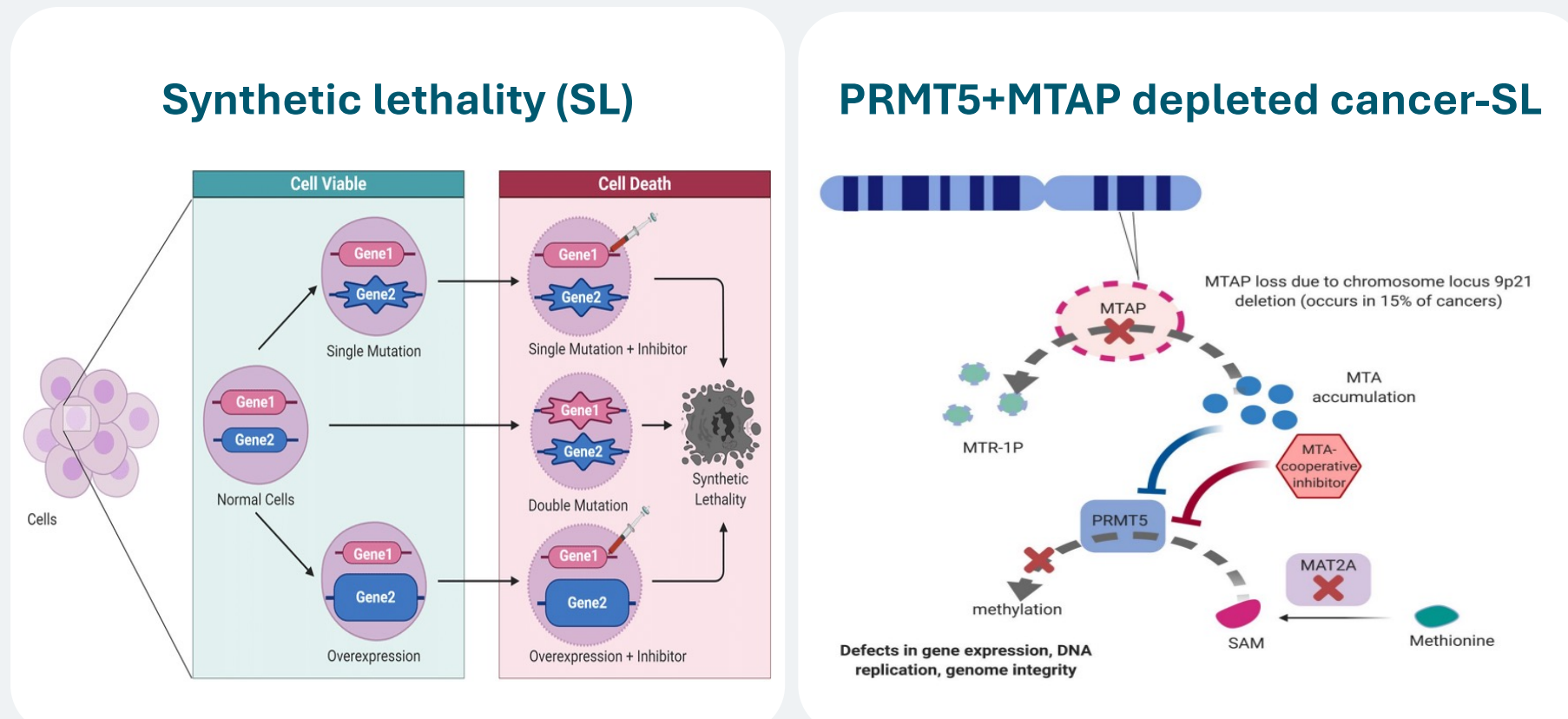
Exploring the Clinical Translation of Synthetic Lethality, PRMT5 Inhibitors in MTAP Depleted Cancers: A Scoping Review

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INTRODUCTION

- Targeting the PRMT5/MTAP pathway exploits synthetic lethality, enhancing the susceptibility of MTAP-deficient tumors to PRMT5 inhibition, thus providing a strategic approach for cancer treatment.
- MTAP deficiency leads to elevated levels of methylthioadenosine (MTA), which inhibit PRMT5 through the formation of a PRMT5-MTA complex, thereby increasing the vulnerability of MTAP-depleted tumors to PRMT5 inhibitors.
- This review examines the current translational research landscape and the future potential of PRMT5 inhibition as a therapeutic strategy for cancers characterized by MTAP deficiency.



Abbreviations – MAT2A: Methionine adenosyltransferase 2A, MTA: Methylthioadenosine; MTAP: Methylthioadenosine phosphorylase; PRMT-5: Protein arginine methyltransferase 5; SAM: S-adenosylmethionine; SL: Synthetic lethality

OBJECTIVE

- To synthesize current evidence (preclinical and clinical) on PRMT5 and MTAP Synthetic Lethality combination.
- To provide detailed Challenges For Clinical Translation of Synthetic Lethality, PRMT5 Inhibitors in MTAP Depleted Cancers.

METHOD

Literature Search

- Databases: PubMed, Google Scholar, clinicaltrials.gov
- Keywords & MeSH terms based on PCC format
- Machine assisted expedited search with support of MAIA Evidence module

Inclusion Criteria

- Studies: Clinical trials, observational studies, reviews, meta-analyses & Timeframe: Last 10 years (up to April 30, 2024)
- Focus: PRMT5 inhibitors in MTAP-deleted cancer cells

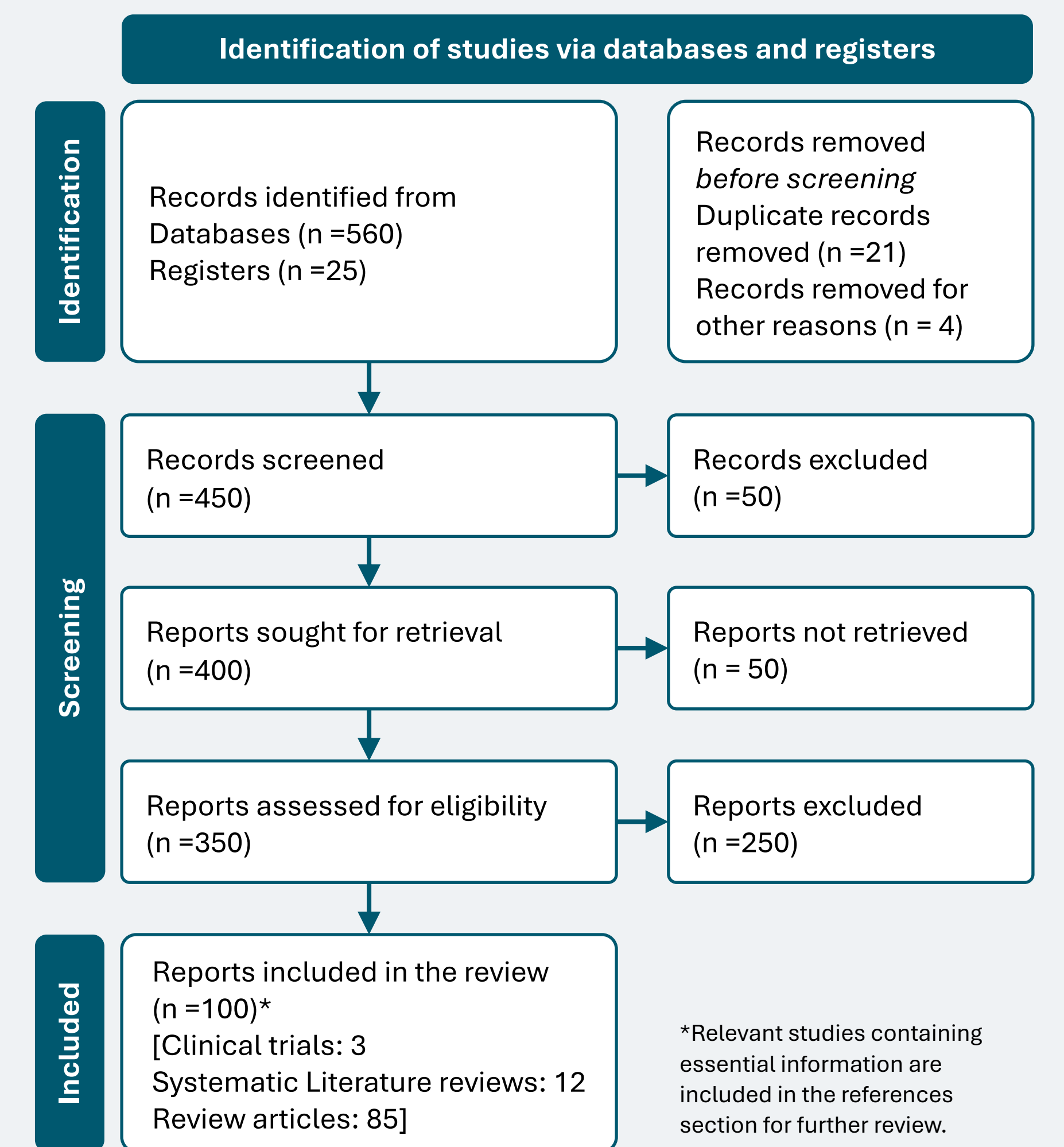
Method

- Adherence to JBI guidelines
- PRISMA-ScR checklist compliance

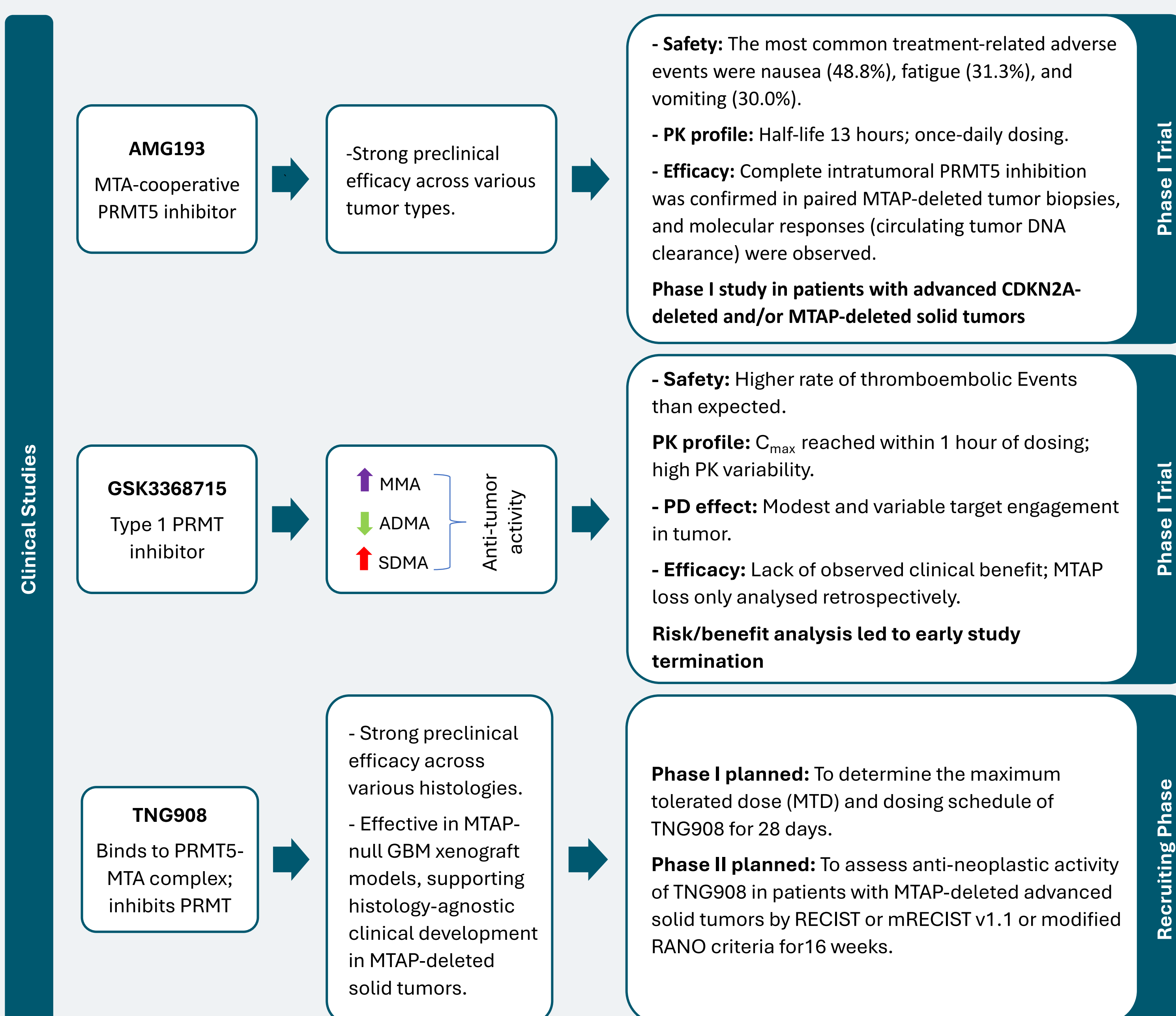
Data Charting

- Tool: Microsoft Excel (JBI ScR template)
- Elements: In vitro/preclinical data (IC50, cytotoxicity, etc.), clinical trial info (molecule, tumor type, phase, NCT number), and conclusions/remarks

Figure 1. PRISMA Diagram



RESULTS



Challenges For Clinical Translation of Synthetic Lethality, PRMT5 Inhibitors in MTAP Depleted Cancers



Table 1: Oncology drug molecules under pre-clinical development

Inhibitor	Mechanism of action	Tumor model/ In-vitro cell line	Key findings	Source
MRTX-1719	Selectively inhibited PRMT5 in the presence of MTA Inhibition of PRMT5-dependent SDMA modification in MTAP del tumors	Human colorectal cancer HCT116 cell line Lung tumor xenografts	Differentiated binding mode leverages the elevated MTA in MTAP del cancers Dose-dependent antitumor activity and inhibition of PRMT5-dependent SDMA modification in MTAP del tumors	Engstrom, Lars D., et al 2023
AG-270	Reduced protein arginine methyltransferase 5 (PRMT5) activity and splicing perturbations	Patient-derived xenograft models	Potent reduction in intracellular SAM MTAP selective antiproliferative activity Treats tumors with homozygous MTAP deletion	Kalev, Peter, et al 2021 Konteatis, Zenon, et al 2021
AMI	Down-regulation of eIF4E and targeting PRMT5 Reduce the symmetric demethylation expressions of PRMT5, eIF4E, histone 3, and histone 4	Lung cancer cells	Apoptosis of lung cancer cells. Down-regulation of eIF4E and targeting PRMT5	Chen, Yingqing, et al 2021
LLY-238	Binds in the SAM pocket of PRMT5 in breast, gastric, and hematological tissues	Mouse xenografts	Highly potent and cell-permeable with well-defined PK characteristics	Bonday, Zahid Q., et al 2018
HLCL-61	Increased expression of miR-29b and consequent suppression of Sp1 and FLT3	In-vitro activities, yet to start	Potent and selective PRMT5 inhibitor for treatment of AML	Tarighat, Somayeh S., et al 2016
EPZ015666	Binds selectively to the SAM-PRMT5 complex via a cation-pi molecular interaction	Panel 64 cell lines	Growth inhibition upon pharmacologic inhibition of PRMT5 with EPZ015666 was not selective for the MTAP-/- genetic background	Marjon, Katya, et al. 2016

Abbreviations- AML: Acute myeloid leukemia; HCT116: Human colorectal carcinoma cell line; MTA: Methylthioadenosine; MTAP: Methylthioadenosine phosphorylase; PK: Pharmacokinetics;

PRMT-5: Protein arginine methyltransferase 5; SAM: S-adenosylmethionine; SDMA: Symmetric dimethylarginine.

CONCLUSIONS



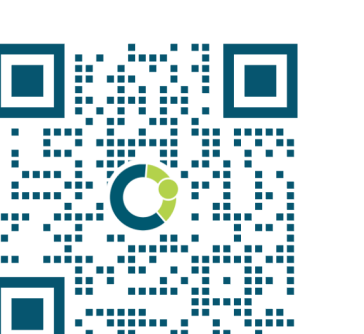
- Strong evidence supports early drug development targeting PRMT5 in MTAP-deleted cancers
- A deeper understanding of molecular changes post-inhibition is crucial for identifying patients who will benefit from MAT2A/PRMT5 inhibitors
- Combining PRMT5 and MTAP synthetic lethality strategies with chemotherapy can enhance DNA damage, increasing cancer cell sensitivity and overcoming drug resistance
- Further research is needed to clarify molecular mechanisms and enhance the clinical viability of this synthetic lethality combination

REFERENCES

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