M1774, a novel potent and selective ATR inhibitor, shows antitumor effects as monotherapy and in combination

Astrid Zimmermann, Heike Dahmen, Thomas Grombacher, Ulrich Pehl, Andree Blaukat, Frank T Zenke the healthcare business of Merck KGaA, Darmstadt, Germany

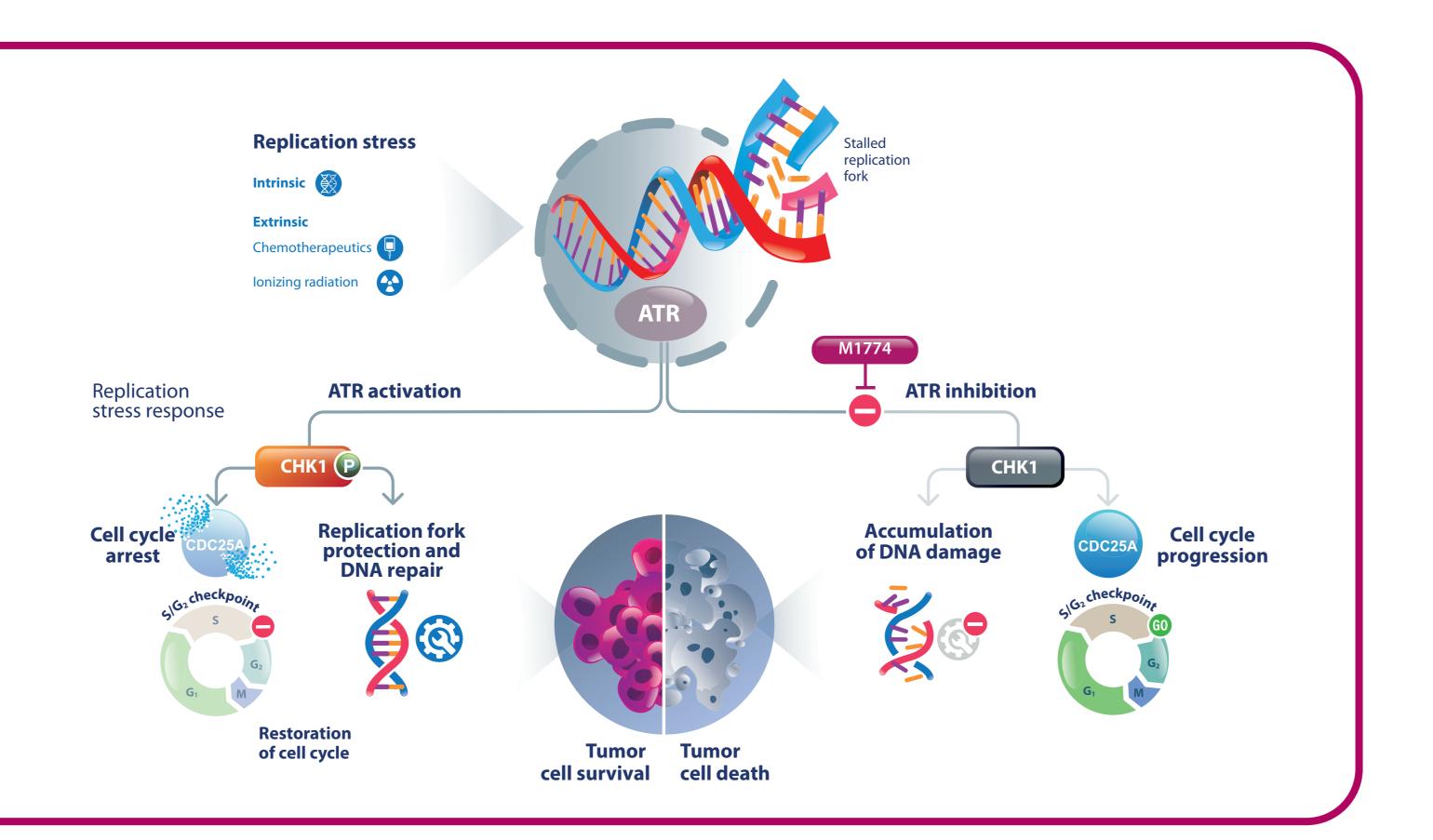
CONCLUSIONS

- M1774 is a potent, orally administered, selective ataxia telangiectasia-mutated and Rad3-related (ATR) kinase inhibitor with differential anti-proliferative activity in tumor cells
- M1774 is active in xenograft models harboring ataxia telangiectasia-mutated (ATM) or AT-rich interaction domain 1A (ARID1A) mutations
- Synergistic activity with specific DNA-damaging agents, poly adenosine diphosphate-ribose polymerase (PARP) inhibitors, and ATM inhibitors was demonstrated



INTRODUCTION

- ATR is activated by single-stranded DNA and recruited to stalled replication forks in response to replication stress¹
- Activated ATR phosphorylates and activates checkpoint kinase 1 (CHK1), which in turn induces S/G2 cell cycle arrest through degradation of cell division cycle 25A (CDC25A) and promotes DNA repair, ultimately restoring replication fork progression¹
- ATR inhibition by small molecule inhibitors such as M1774 abrogates the S/G2 cell cycle checkpoint as well as the restoration of replication forks, resulting in tumor cell death^{1,2}



ATR, ataxia telangiectasia-mutated and Rad3-related

RESULTS

Table 1. Structure, activity and selectivity of ATR inhibitor M1774

Kinase assays	Ki [nM]
ATR	<0.2
ATM	4040
DNA-PK	290
Cell-based assays	IC ₅₀ (nM)
ATR: 4NQO-induced pSer345 CHK1 in HT29	5
ATM: NCS-induced pSer139 H2AX in MO59J (DNA-PK null)	19.800
DNA-PK: NCS-induced pSer139 H2AX in HT144 (ATM null)	2300
HCT116, proliferation inhibition; cytotoxicity	24, 26
HFL1, proliferation inhibition; cytotoxicity	1.870; 11.000
Cancer cell line panel (#93)	23 – >1.000 (median 120
Selectivity	Kinases <175-fold split
Kinase selectivity*	4/321 tested
ATP stayis tolangiostasis-mutated and Pad3-related; ATM stayis tolangiostasis mutated; D	NA-BK DNA-dependent protein kinase

ATR, ataxia telangiectasia-mutated and Rad3-related; ATM, ataxia telangiectasia mutated; DNA-PK, DNA-dependent protein kinase

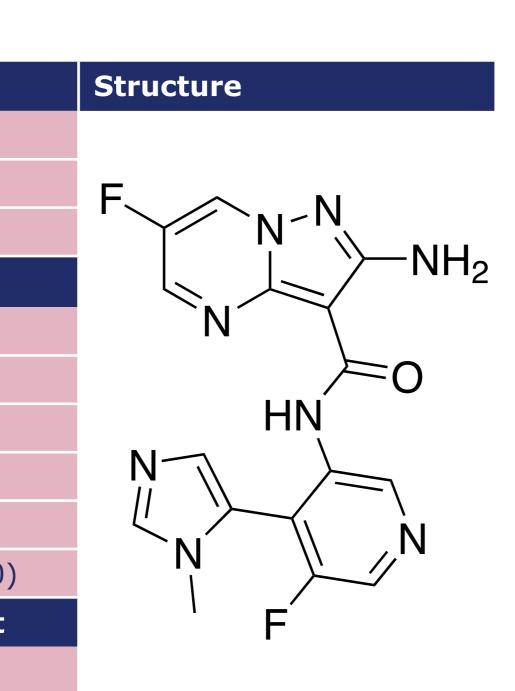
1. Blackford N, Jackson P. *Mol Cell.* 2017;66:801–817; 2. Yap TA, et al. *J Clin Oncol.* 2021;39:15(suppl TPS3153).

The corresponding author for this presentation is Frank T Zenke (frank.zenke@emdgroup.com)

All authors are employees of the healthcare business of Merck KGaA, Darmstadt, Germany, which funded this research (CrossRef Funder ID: 10.13039/100009945) The authors appreciate the contributions from Vertex Pharmaceuticals in the characterization of the ATR inhibitor M1774. Editorial assistance was provided by Mario Pahl of Bioscript Stirling, Macclesfield, UK, and funded by the healthcare business of Merck KGaA, Darmstadt, Germany. Presented at the AACR Annual Meeting 2022 | April 8–13 | New Orleans, Louisiana, USA

Copies of this e-poster btained through QR codes are for personal use only and may not be reproduced without written permission of the authors.

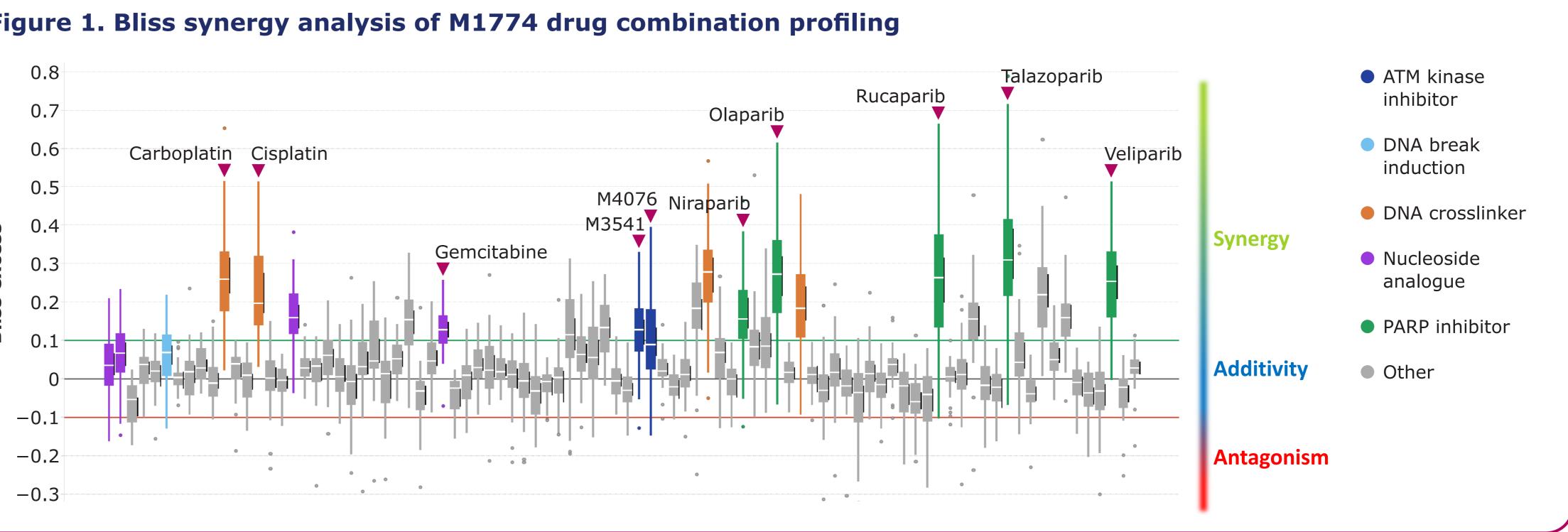
- In combination with DNA-damaging agents (cisplatin, gemcitabine) and DNA damage response (DDR) inhibitors (niraparib, M4076) strong anti-tumor activity was observed in xenograft models
- M1774 is currently being investigated in a Phase I clinical trial in patients with advanced solid tumors (DDRiver Solid Tumors 301, NCT04170153)



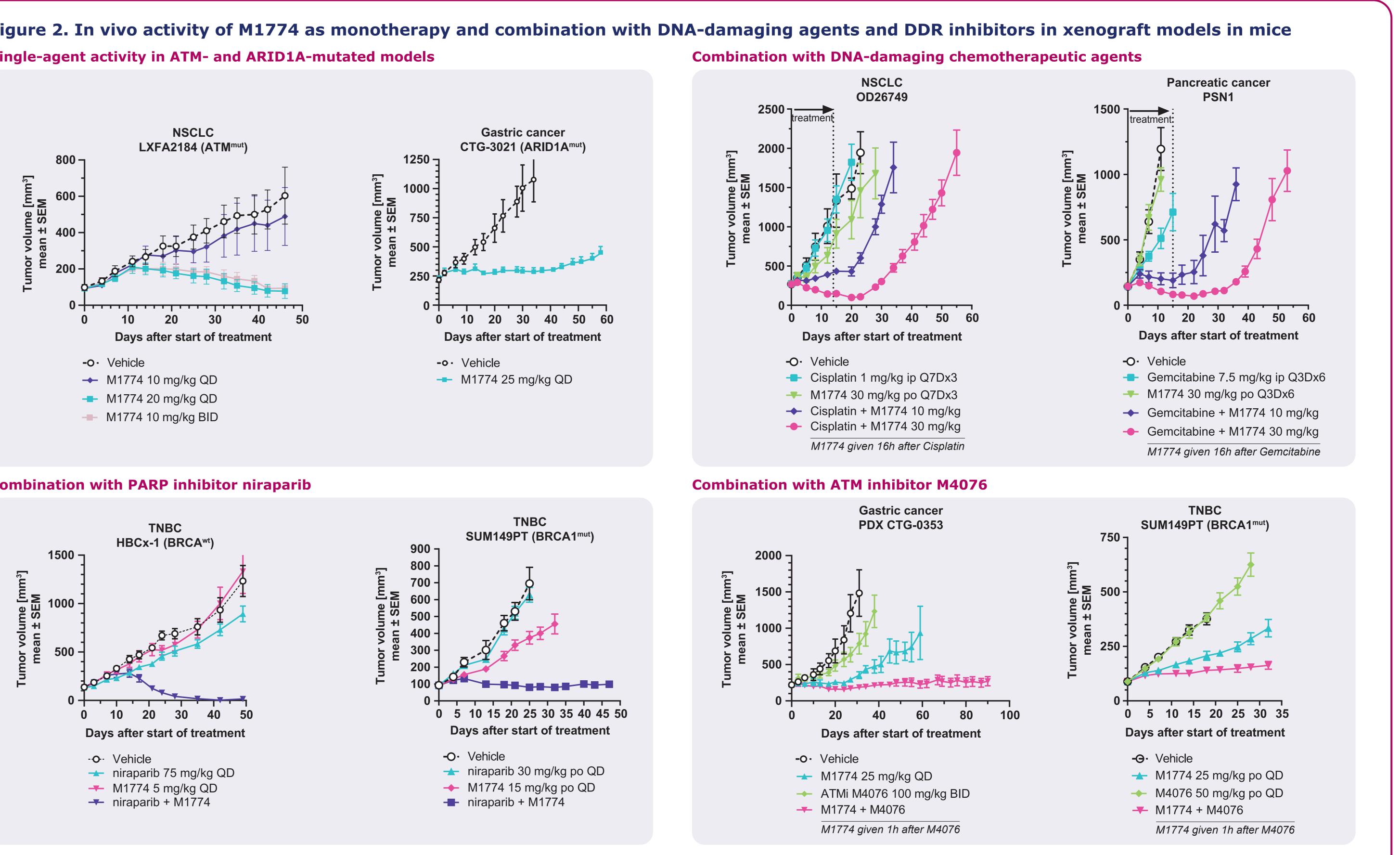
- M1774 is a potent inhibitor of ATR
- In cell-based assays, M1774 has >100-fold selectivity for ATR over ATM and DNA-dependent protein kinase (DNA-PK)
- In large panel kinase profiling, minimal inhibitory activity was observed against unrelated kinases, with at least 175-fold selectivity for ATR over 317 out of 321 kinases tested
- A broad range of antiproliferative activities (IC₅₀ 23 nM – 1 μ M) was observed in a panel of 93 cancer cell lines

RESULTS

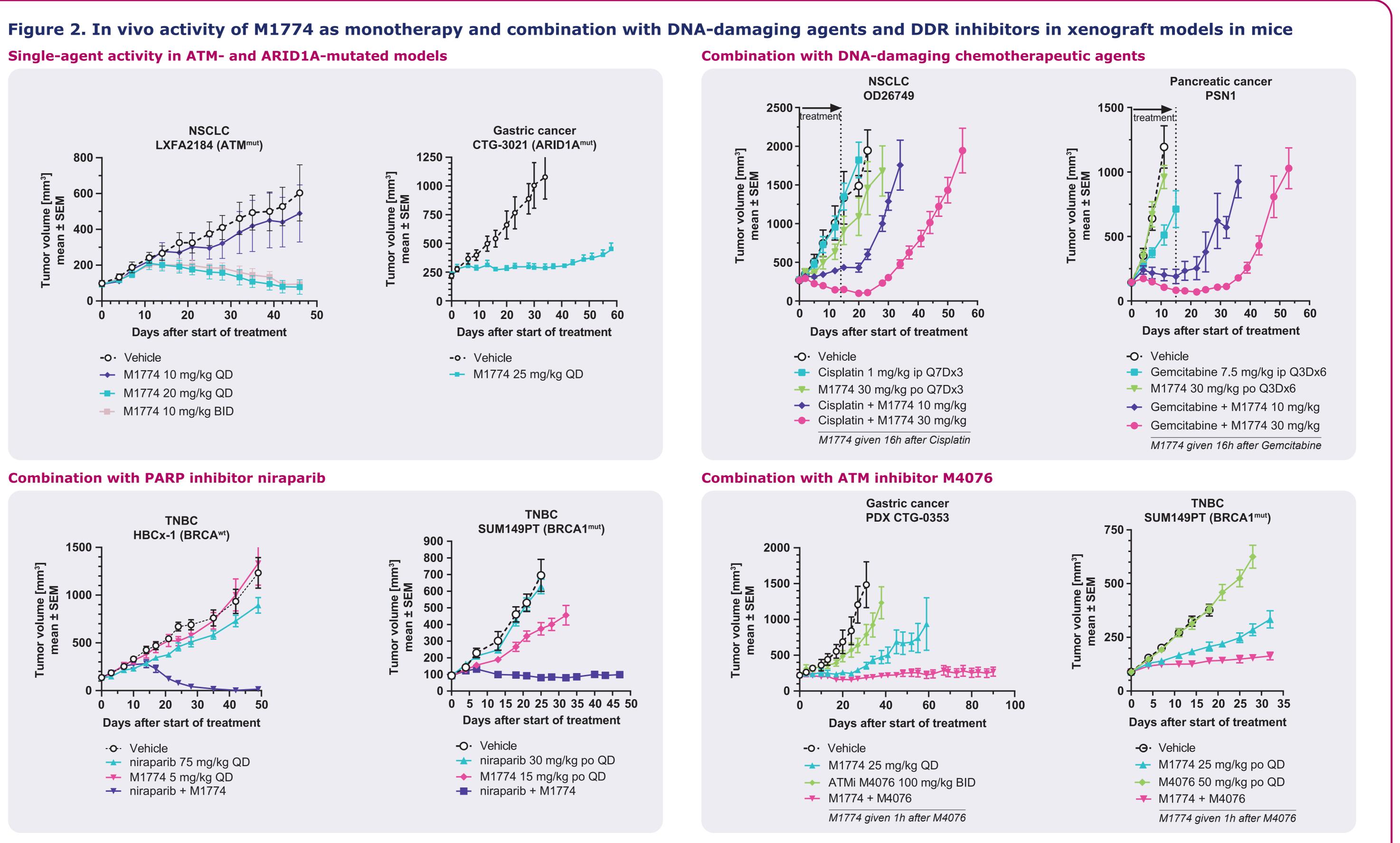
Figure 1. Bliss synergy analysis of M1774 drug combination profiling



Single-agent activity in ATM- and ARID1A-mutated models



Combination with PARP inhibitor niraparib



ARID1A, AT-rich interaction domain 1A; **ATM**, ataxia telangiectasia mutated; **BID**, twice a day; **BRCA**, breast cancer gene; **DDR**, DNA damage response; **ip**, intraperitoneal administration; **NSCLC**, non small cell lung cancer; **PARP**, poly adenosine diphosphate-ribose polymerase; po, oral administration; QD, every day; SEM, standard error of the mean; TNBC, triple negative breast cancer

• M1774 was usually well tolerated with no additional body weight loss compared to chemotherapeutic agent or DDR inhibitor alone

• Combination of M1774 with niraparib at 75 mg/kg resulted in body weight loss in a fraction of mice (up to 20% in 3/14 mice treated with combination regimen), while combination with niraparib at a dose of 30 mg/kg was well tolerated

- Pairwise combination of M1774 with 90 standard of care and investigational agents for oncology
- PARP inhibitors, platinumcontaining anti-cancer drugs, antimetabolites, and ATM inhibitors identified as synergistic combination partners
- Combination data were generated in 35 cancer cell lines (5-day compound incubation; readout: sulforhodamine B staining)
- Box plots show the distribution of Bliss scores for each combination