

Background

- TEAD transcription factors are the major effectors of the Hippo-YAP/TAZ-TEAD pathway involved in the regulation of cell proliferation, survival, and cell BRAF inhibitors (encorafenib, dabrafenib), MEK inhibitors (trametinib, cobimetinib), and mTOR inhibitor (everolimus). migration.
- TEAD auto-palmitoylation has been shown to be required for TEAD interaction with coactivator YAP/TAZ and hence activation of transcriptional activity.
- There are multiple reports that YAP/TEAD activation provides the essential survival signal in drug-tolerant persister/dormant cells in cancer in response to treatment with targeted therapies, leading to drug resistance and cancer relapse in BRAF-mutant, KRAS-mutant, EGFR-mutant, and ALK-rearranged non-small efficacy and durability of treatment. cell lung cancers (NSCLC).
- VT3989, with similar biological activity in vitro and in vivo as the potent small molecule TEAD auto-palmitoylation inhibitors reported (Tang et al, 2021, Mol Ther. 20(6):986-998), disrupts YAP/TAZ-TEAD protein interaction, Cancer suppresses TEAD transcriptional activity, and selectively blocks proliferation of NF2-deficient mesothelioma in vitro and inhibit NF2 mutant xenograft tumor growth in vivo.
- VT3989, currently in clinical testing, was evaluated for effect on emerging drug tolerant persister/dominant cancer cells and drug resistance.



Tang et al., Mol Cancer Ther, 2021 Chan et al., Nat Chem Biol, 2016 Noland et al., Structure, 2016 Pobatti et al., Structure, 2015

Palmitoylation of a conserved cysteine is

YAP-binding domain of TEAD.

palmitate is buried in a central pocket in the

FAD

2 3 4 1 2 3 4 M 80 kD required for TEAD interaction with YAP. The TEAD1-PEG 65 kD TEAD1 —

> VT3989 and other Vivace TEAD inhibitors occupy the palmitate pocket in TEAD, inhibiting auto-palmitoylation and thereby inhibiting transcription function of YAP-TEAD.

VT3989, a clinical TEAD palmitoylation inhibitor, enhances the efficacy and durability of multiple targeted therapies of the MAPK and PI3K/AKT/mTOR pathways

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Summary and Conclusions

- Study started with n=12 mice per arour Day 38 and Day 54 were the days when average tumor volume went back to be a set of the days and back to be a set of the days when average tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to be a set of tumor volume went back to the starting point for osimertinib alone and combination with 10mg/kg VT3989 \: mouse sacrificed or found dead Figure 1. VT3989 and EGFR small molecule inhibitor combination studies in vivo continuous throughout the entire study in the study shown in (A). Treatments in the studies shown in (B, C, and D) were stopped at the indicated time points. Tumor volumes were measured throughout the studies to monitor tumor regrowth.VT3989 shows strong synergy in combination with EGFR inhibitors and significantly delay the tumor regrowth in combination. **Combination with KRAS-G12C inhibitor** SW837 CRC CDX Mode CR6243 CRC PDX Mode -VT104 10 mg/kg TREATMENT TREATMENT MRTX849 100 mg/ **+**MRTX849 100mg/kg
- ••VT104 10mg/kg + MRTX849 100mg/kg All Treatments (PO, QD) stopped on Day 31 All Treatments (PO, QD) stopped on Day 30 59 days VT104 + 4000-3500- WRTX84 0 5 10 15 20 25 30 35 40 45 50 55 60 65 70 75 80 85 0 5 10 15 20 25 30 35 40 45 50 55 60 Days post Treatment Start Days ofter the start of treatment = mouse sacrificed or found dead

Figure 3. TEAD and KRAS-G12C small molecule inhibitor combination studies KRAS-G12C inhibitor, MRTX869 (adagrasib), was combined with TEAD inhibitors, VT104 (A) or VT3989 (B) in KRAS-G12C CRC models. Treatments in the studies were stopped at the indicated time points. Tumor volumes were measured throughout the studies to monitor tumor regrowth.

2.VT3989 3 µM 3.VT3989 0.3 µM 4.VT3989 0.03 µM

- REFERENCE
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- 2. Kurppa et al., 2020, Cancer Cell 37: 104–122.
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- 5. Tsuji et al., 2020, Nat Commun. 11(1):74.
- 3. Tang et al, 2021, Mol Cancer Ther. 20(6):986-998 6. Nilsson et al., 2020 Sci. Transl. Med. 12, eaaz4589

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Combination with EGFR small molecule inhibitors in EGFR mutant NSCLC models



VT3989 enhances the efficacy and durability of response to clinically approved EGFR inhibitors (osimertinib, lazertinib), MET inhibitors (savolitinib, capmatinib), EGFR-MET bispecific antibody (amivantamab), KRAS-G12C inhibitors (sotorasib, adagrasib),

With the addition of VT3989, combination anti-tumor effect was observed and/or tumor regrowth was delayed compared to targeted therapies alone. The persister cell mechanism seems to work in a very broad range of combinations, and the broad range of combination agents suggests a fundamental role for YAP-TEAD in the development of resistance to targeted therapy.

In addition to Hippo-YAP/TAZ-TEAD pathway driven cancers, such as mesothelioma, where VT3989 can show monotherapy efficacy, future VT3989 clinical studies will include combination therapy with approved targeted therapies to enhance the

EGFR inhibitor, osimertinib (A, B, and C) or lazertinib (D) was combined with VT3989 in EGFR mutant NSCLC CDX models (A and D) and PDX models (B and C). Treatments (>75 days) were



and NSCLC PDX models (C). All treatments were PO, QD throughout the studies.

Poster #B088

Combination with EGFR-MET bispecific antibody +/- EGFR small molecule inhibitor



Combination with MET small molecule inhibitor

Amivantamab 10mg/kg

VT3989 30mg/kg

Days post Treatment Start

Figure 6. VT3989 and mTOR inhibitor combined with VT3989 in mesothelioma CDX model (A) and liver cancer PDX synergy with everolimus. Treatments

and amivantamab in LU-01-1292

combination significantly inhibited

NSCLC PDX model, and the

tumor regrowth.

0 5 10 15 20 25 30 35 40 45 50 55 60 65 70 75 80 85 90 **Days post Treatment Start**

0 15 33 50 68 85 103

Days post Treatment St