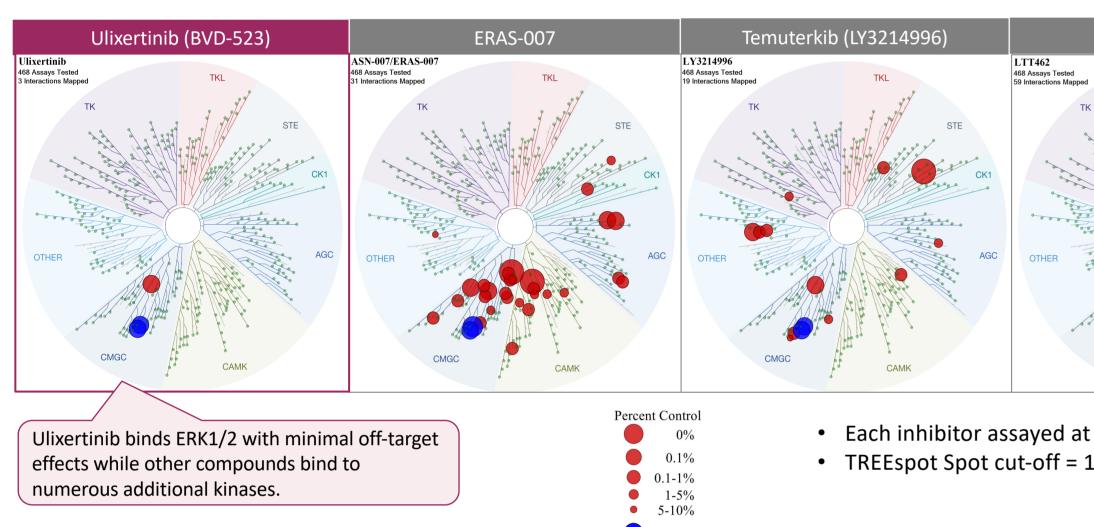
BIOMEDVALLEY D I S C O V E R I E S

Background

- multiple indications^{1,2}.
- observed in patients harboring alterations within the RAS-MAPK pathway.
- Ongoing and planned clinical trials cover both ulixertinib monotherapy and combination settings across a variety of tumor types and mutational drivers.
- pathway re-activation, blockade with an ERK inhibitor at the terminal node of the MAPK pathway should provide durable intervention.

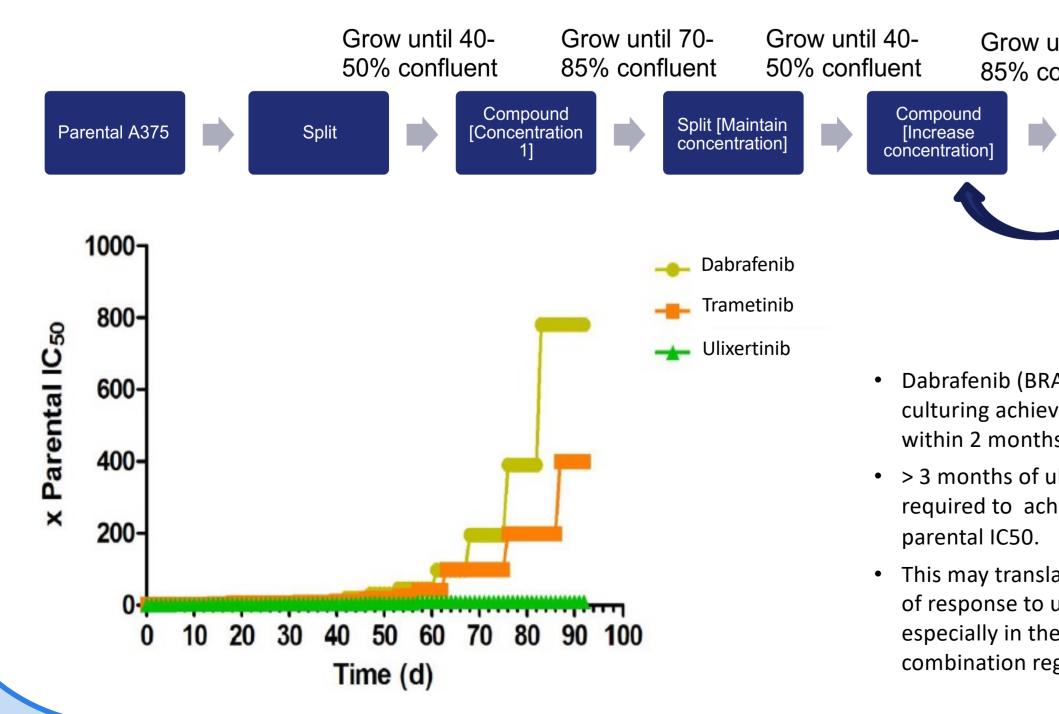
It has been well established that ERK activation ME-010 increases cyclin D1 levels and entry into the cell Melanoma - NRAS^{Q61R} cycle³. The combination of MEKi with CDK4/6i has been 600 RTK shown to be beneficial⁴ and has since been tested in the clinic. We hypothesized that the combination of ERK1/2 and CDK4/6 inhibitors (A) would have synergistic antitumor activity in RAS mutated tumors. • Four xenograft models were utilized to assess the 0 3 6 9 12 15 18 21 combination of ulixertinib and palbociclib, each xenograft harboring a RAS mutation (B). CRC - KRASG12 Ulixertinib monotherapy across all models showed 1200 • modest tumor growth inhibition (TGI) while palbociclib monotherapy showed limited TGI across all models The combination groups demonstrated significant responses ranging from 75% - 90% TGIs. All treatment regimens were well tolerated across all & survival models. Davs Rineterkib (LTT462) ERAS-007 emuterkib (LY3214996 LTT462 468 Assays Tested 59 Interactions Mapped H358 Clinically described mechanisms of acquired resistance to KRAS^{G12C} inhibitors converge on reactivation of the RAS-MAPK NSCLC KRAS^{G12C} pathway (A)⁵⁻⁸. • The efficacy of ulixertinib, in combination with KRAS^{G12C} inhibitor, adagrasib (MRTX849), was assessed in cell line-derived xenograft models harboring KRAS^{G12C} mutations. Models were selected based on response to single agent adagrasib, ranging 1000 from sensitive (H358) to partially responsive $(H2122)^7$. • Combining ulixertinib with MRTX849 (MRTX) resulted in superior tumor growth inhibition compared to dosing of RTKs activation either single agent alone in both models (B-E). In H2122 high-Percent Control MET amplification • Each inhibitor assayed at 2 μ M concentration 0% dose combination treatment (Groups 8 and 9) yielded robust 0.1% • TREEspot Spot cut-off = $10 \% \ge control$ regression compared to all other treatment groups. 0.1-1% • 1-5% • Tumor volume over time is shown for groups dosed to day 21 • 5-10% for H359 (B) and H2122 (C). Tumor volume percent change from baseline at day 21 is shown for H359 (D) and H2122 (E). Visualization of growth is capped at 100% (D, E). • Differential expression analysis was performed on tumors collected on day 21 to identify pathways associated with KRAS G12C amplification tumor growth inhibition (F,G). Consistent with previous studies by others⁹, significantly enriched genesets related to **KRAS G12V** -100 ages generated using TREE*spot*[™] Software Tool and reprinted with BRAF V600 KRAS G12W KRAS signaling, apoptosis, mTOR, MYC, and cell cycle were mission from KINOMEscan[®], a division of DiscoveRx Corporatio ALLMARK KRAS SIGNALING U (RAS G12D revealed. Combination treatment further enriched each of MAP2K1 K57N KRAS Q61R these pathways compared to single agent MRTX849. MAP2K1 Q56P KRAS Q61K MAP2K1 E102 103del • The KRAS signaling up geneset was robustly down-regulated by the MRTX + uli combination groups compared to vehicle in Ulixertinib to other MAPK pathway inhibitors H358 (F), and in H2122 (G). • These data provide a robust rationale for combining MMMMMM ulixertinib with KRAS^{G12C} inhibition in the clinic. Discussions Proliferation are ongoing regarding clinical deployment. uppression of apoptosis Compoun Migration Angiogenesis 5. Ulixertinib clinical trials underway and in development Grow until 40-Grow until 40-Grow until 70-Grow until 70-Pre-clinical / Phase 2 Phase 1 85% confluent 50% confluent 50% confluent 85% confluent rotocol Development Monotherapy Compound [Increase Compound [Concentration Split [Maintain concentration] Split [Maintain concentration] NCT04566393 Expanded Access Protocol (EAP) concentration NCT05804227 Adult low-grade glioma Grow until 40be addressed. 50% confluent Q4 / 2023 Histiocytosis 🚤 Dabrafenib Pediatric low-grade glioma (pLGG) Q1 / 2024 ____ Trametinib activated patient populations. Combinations 🛖 Ulixertinib NCT05221320 (Multiple GI Cancers) + Hydroxychloroquine (HCQ) • Dabrafenib (BRAFi) or Trametinib (MEKi) culturing achieved 100x parental cell IC50 NCT03454035 (NRAS Melanoma) + Palbociclib References within 2 months. + multiple combinations (EAP) NCT04566393 (Solid Tumors) • > 3 months of ulixertinib culturing was + EGFRi, +/- BRAFi required to achieve growth in 15x Q4 / 2023 (CRC) 2. Sullivan et al. Cancer Discov. 2018;8(2):184-195. parental IC50. 3. Ravenhall C, et al. Br J Pharmacol. 2000 Sep;131(1):17-28. Pediatric low-grade glioma (pLGG) Q1 / 2024 • This may translate to improved durability + RAF / Senolytics TBD of response to ulixertinib in the clinic, 6. Zhao Y et al. Nature. 2021 Nov;599(7886):679-683. POC combinations of interest: especially in the setting of an optimal TBD 7. Awad MM et al. N Engl J Med. 2021 Jun 24;384(25):2382 + RAFi, KRASi, JAKi, STATi, MDM2i combination regimen. Time (d) 9. Hallin J et al. Cancer Discov. 2020 Jan;10(1):54-71.



- Ulixertinib, in addition to other ERK1/2 inhibitors, were assayed against the KINOMEscan[®], an active site directed competition binding assay against 468 kinases. Results for ulixertinib, ERAS-007, temuterkib, and rineterkib shown.
- Ulixertinib demonstrated a superior selectivity profile compared to the other ERK1/2 inhibitors.
- In addition to binding to ERK1/2, ulixertinib binds to ERK8 in this assay; 0.15, 0.3 and 0.45 % control, respectively.

Ulixertinib is a highly potent and selective ERK1/2 inhibitor poised for clinical differentiation in • To date, over 500 patients have been treated with ulixertinib. Confirmed responses have been As emerging resistance mechanisms to KRAS^{G12C} inhibitors continue to be dominated by MAPK 1. Ulixertinib is a selective ERK1/2 inhibitor 2. Resistance to ulixertinib is delayed compared

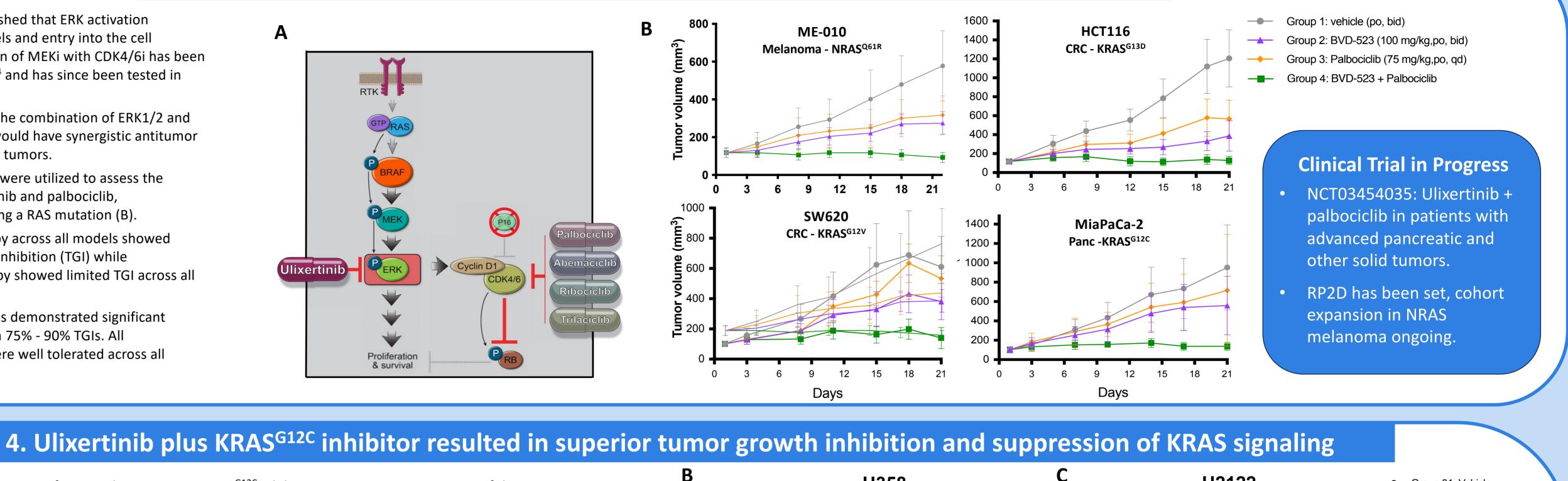
- *In vitro* models of ulixertinib resistance were generated in an endeavor to characterize and predict potential mechanisms of resistance and guide rational combination therapies.
- The BRAF^{V600E} mutant melanoma cell line, A375, was cultured in progressively increasing concentrations of ulixertinib or other MAPK pathway inhibitors (dabrafenib and/or trametinib).

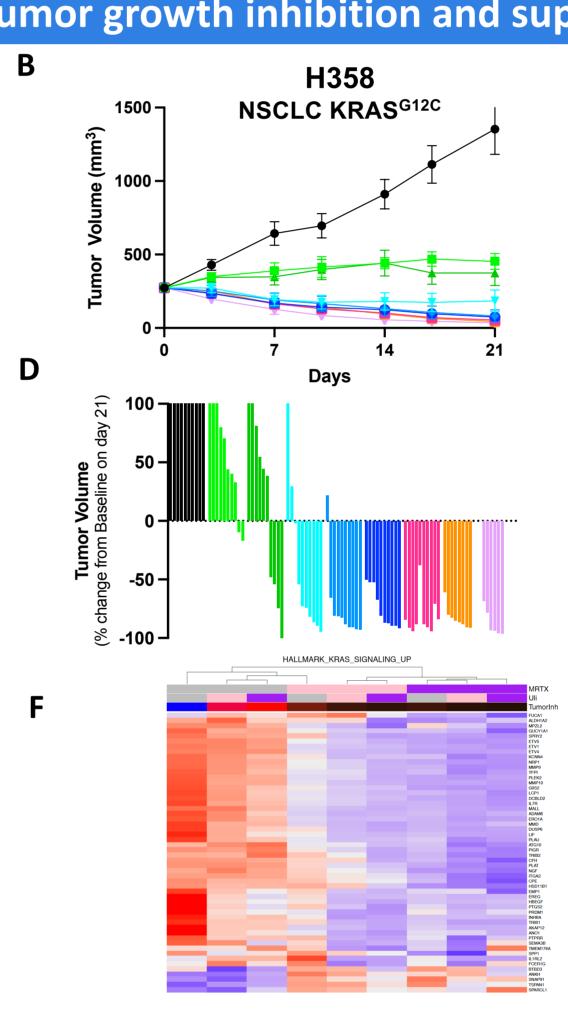


Ulixertinib, a first-in-class ERK1/2 inhibitor, as an effective combination agent in RAS mutated malignancies

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3. Combination of ulixertinib and palbociclib demonstrate significant efficacy in vivo

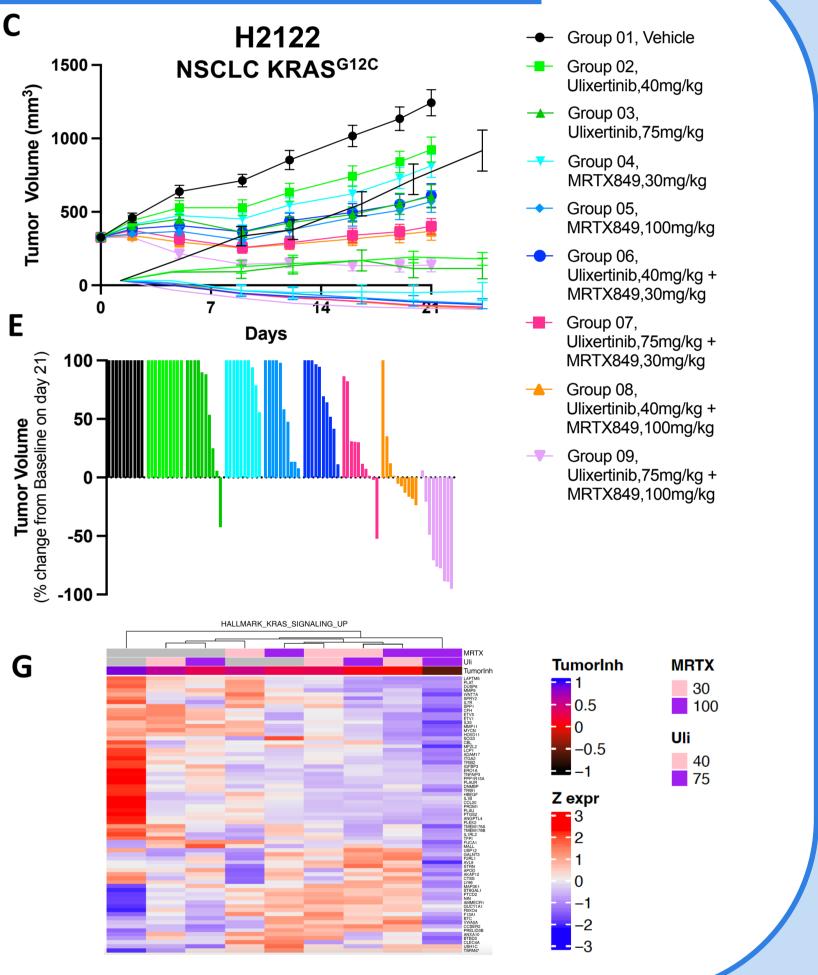




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Conclusions

• Ulixertinib demonstrates tumor regression in multiple RAS driven models as monotherapy, in combination with CDK4/6 inhibitors, and in combination with KRAS^{G12C} inhibitors.

• By targeting the last node in the MAPK pathway with ulixertinib, we believe that more durable MAPK inhibition will be possible and that the pathway reactivation resistance mechanisms will

• Ulixertinib has established tolerable BID doses in both adult and pediatric patient populations, with 500+ patients treated to date. Multiple clinical efforts are underway in RAS/MAPK

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