



inhibition leads to a reduction in MAPK and AKT signaling, but residual MAPK signaling due to feedback or signaling from other pathways can allow cells to escape cell death. (C) Combined inhibition of the KIT and MAPK pathways leads to cell death.

¹Ran L. *et al.* (2015) Combined inhibition of MAP Kinase and KIT Signaling Synergistically Destabilizes ETV1 and Suppresses GIST Tumor Growth. *Cancer Disc.* 5: 304-315

Combined inhibition of MAPK and KIT signaling induces pro-apoptotic proteins and apoptosis



Figure 2: DCC-2618 and trametinib combinations induce apoptosis in GIST cell lines. (A) Immunoblot of pKIT, pERK, BIM and pH2AX levels in imatinib-sensitive (GIST-T1) and imatinib-resistant (GIST-T1/D816E and GIST-T1/T670I) cells treated with imatinib (IM, 100 nM) or DCC-2618 (DCC, 100 nM) in the absence or presence of trametinib (Tram, 10 nM) for 48 hr. (B) Apoptosis induction in GIST-T1 and GIST-T1-T670I cells as measured by AnnexinV/PI staining. Cells were treated with compounds at indicated concentrations for 24, 48 and 72h and stained with Annexin V and PI.

DCC-2618, a broad-spectrum inhibitor of KIT and PDGFRA mutants, synergizes with inhibitors of the MAPK pathway

Anu Gupta¹, Cynthia B. Leary¹, Alfonso Garcia-Valverde², Joaquin Arribus², Cesar Serrano², Daniel L. Flynn¹, Bryan D. Smith¹ ¹Deciphera Pharmaceuticals, LLC, Waltham, MA; ²Preclinical Research program, Vall d'Hebron Institute of Oncology (VHIO), Vall d'Hebron University Hospital, Barcelona, Spain EORTC-NCI-AACR Annual Meeting, Nov 13-16, 2018, Dublin, Ireland

> Figure 3: DCC-2618 synergizes with MEK inhibitors to induce caspase activity in GIST cell lines. (A) Caspase activity in GIST cell lines with DCC-2618 and trametinib combination after 24 h of treatment. Values represent average of three replicates. P values for drug combination vs each drug alone were <0.001. (B) Surface mapping of combination dose-responses demonstrate synergy in GIST cell lines with DCC-2618 and trametinib. A matrix of various concentrations of drugs was used to measure caspase activity and the BLISS synergy score was calculated with Combenefit software. (C) Caspase 3/7 activity in GIST cell lines with DCC-2618 and binimetinib combination. (D) Surface maps of combination treatments to demonstrate synergy.





Figure 4: DCC-2618 synergizes with MEKi in HMC1.2 cells (V560G/D816V KIT). (A) Caspase 3/7 activity in HMC1.2 cells treated with DCC-2618 alone or in combination with trametinib for 24 h. (B) Surface map of combination treatments with trametinib, cobimetinib and binimetinib demonstrate synergy.



Figure 5: 100 cells for GIST-T1 and GIST-T1-D816E and 500 cells for GIST-T1-T670I were plated in 6 well plates and treated with compounds for 2 weeks. Drugs were washed off and cells were grown in complete media for an additional 10 days. Colonies were stained with crystal violet and counted. (A) Colony outgrowth in GIST cell lines treated with DCC-2618 and trametinib. (B) Average colony counts from three wells. (C) Average colony counts of GIST cell lines treated with a combination of DCC-2618 with binimetinib. Arrows indicate no visually determined colony outgrowth.

DCC-2618/MEKi combination induces apoptosis in N-Ras G12D expressing mastocytosis cells



Figure 6: Combination treatment inhibits colony outgrowth in N-Ras G12D transfected HMC1.2 cells. (A) HMC1.2 empty vector (EV) control and N ras mutant G12D transfected cells were grown in soft agar for 10 days in the presence of DCC-2618 and trametinib. Drug was washed off and cells were grown for 5 days in complete media. Colonies were stained and photographed. (B) Average colony counts from 3 wells. (C) Caspase activity in EV and N-Ras G12D transfected cells in combination with DCC-2618 and trametinib.



Saturation Mutagenesis



Figure 7. Combination of DCC-2618 and trametinib prevents emergence of drug resistance mediated by non-KIT mechanisms. Ba/F3 cells stably expressing V560D KIT were treated with ENU and grown in the presence of DCC-2618 +/- trametinib for 28 days. Wells exhibiting outgrowth were sequenced for KIT mutations. "V560D (Native)" indicates that no secondary mutations were found within the KIT gene, and mutations may have arisen in other pathways, such as the MAPK pathway, that allow for Ba/F3 cell growth. Arrows indicate no outgrowth.

DCC-2618 in combination with trametinib inhibits GIST xenograft tumor growth



- Vehicle				
DCC-2618 ~ 100 mg/kg chow				
trametinib 0.5 mg/kg PO BID				
→ DCC-2618 ~100 mg/kg + trametinib				
	Day 27		Day 67	
	PR		DD	
		UK	PR	CR
DCC-2618	4	6	2	CR 0
DCC-2618 Trametinib	4 0	6 0	2 0	CR 0 0

Figure 8. DCC-2618 and trametinib treatment results in tumor regressions in a GIST-T1 xenograft model. Mice were treated with DCC-2618 formulated into the diet to achieve ~100 mg/kg/day (red), trametinib at 0.5 mg/kg BID PO (green), or with the combination of drugs (orange). Dosing started on day 10 and ended on day 27. Tumor growth was monitored for additional 40 days. Partial and complete tumor regressions (PR and CR) are shown at Days 27 and 67.

Summary

- KIT inhibitors are cytostatic in GIST cells, and cells start to grow back when drug is removed. The combination of DCC-2618 with a MEK inhibitor induces apoptosis and prevents the outgrowth of colonies even after drug is removed.
- DCC-2618 shows strong synergy in inducing apoptosis in combination with MEK inhibitors trametinib and binimetinib in imatinib-sensitive and, importantly, in imatinib-resistant GIST and mastocytosis cell lines.
- Ras is frequently mutated in advanced forms of systemic mastocytosis and Ras mutant expressing HMC1.2 cells were more sensitive to the combination of DCC-2618 and trametinib.
- In saturation mutagenesis studies, treatment with DCC-2618 prevents the outgrowth of cells with drug-resistant KIT mutations. A few wells did exhibit outgrowth, in which cells presumably had mutations in other signaling proteins or pathways, such as the MAPK pathway. The combination of DCC-2618 with trametinib completely suppressed clonal growth.
- The combination of DCC-2618 and trametinib in a GIST-T1 xenograft model leads to more partial and complete tumor regressions than single agent treatment, even with a short treatment period of 18 days and 40 days of allowing tumors to regrow after dosing was stopped.