

AB521 (Casdatifan) Potently and Selectively Inhibits Hypoxia-Inducible Factor 2 Alpha (HIF-2 α) Dependent Pro-Tumorigenic Activity

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Abstract

Background: Hypoxia-Inducible Factor 2 α (HIF-2 α) is a transcription factor that regulates genes critical for cell adaptation and survival in low oxygen conditions, such as those present in the solid tumor microenvironment. HIF-2 α protein is stabilized and transcriptionally active in hypoxia but is targeted for proteasomal degradation in normoxic conditions by the von Hippel-Lindau (VHL) subunit of an E3 ubiquitin ligase complex. In clear cell renal cell carcinoma (ccRCC), VHL inactivating mutations are prevalent and cause a 'pseudohypoxic' state, wherein HIF-2 α protein is stabilized, even when oxygen levels are high. Notably, HIF-2 α is an established tumorigenic driver in ccRCC, and inhibition of HIF-2 α has been shown to be an effective strategy to improve clinical outcomes.

Materials and Methods: Applying a pharmacophore mapping and structure-based design approach, we identified a novel and potent small molecule HIF-2 α inhibitor, AB521 (casdatifan), that avidly binds to a hydrophobic pocket in the HIF-2 α PAS-B domain. The potency and pharmacology of AB521 were evaluated using *in vitro* and *in vivo* experimental systems.

Results: The *in vitro* impact of AB521 on ccRCC cells was assessed using the VHL and HIF1A mutant 786-O cell line. AB521 potently inhibited HIF-2 α -dependent reporter gene transcription in both low and high serum conditions with no off-target cytotoxicity. Additionally, VEGF secretion and soft agar colony formation were also inhibited by AB521. In hypoxic Hep3B hepatocellular carcinoma (HCC) cells, AB521 selectively inhibited HIF-2 α , but not HIF-1 α , target gene expression. AB521 also inhibited transcriptional activity of HIF-2 α in relevant human primary cells, including endothelial cells and pro-tumorigenic M2-polarized macrophages. When delivered orally to mice, AB521 caused significant regression of established ccRCC xenograft tumors and dose-dependent decreases in peripheral and tumor-specific HIF-2 α -associated pharmacodynamic markers. AB521 combined favorably with a clinical standard of care, the VEGF tyrosine kinase inhibitor cabozantinib, in a ccRCC xenograft model, as well as the clinical anti-PD-1 antibody, zimberelimab, in a humanized version of the ccRCC model with stably engrafted peripheral blood mononuclear cells. Outside of ccRCC, HIF-2 α inhibition also provided a survival benefit in a murine model of HCC.

Mechanism of Action

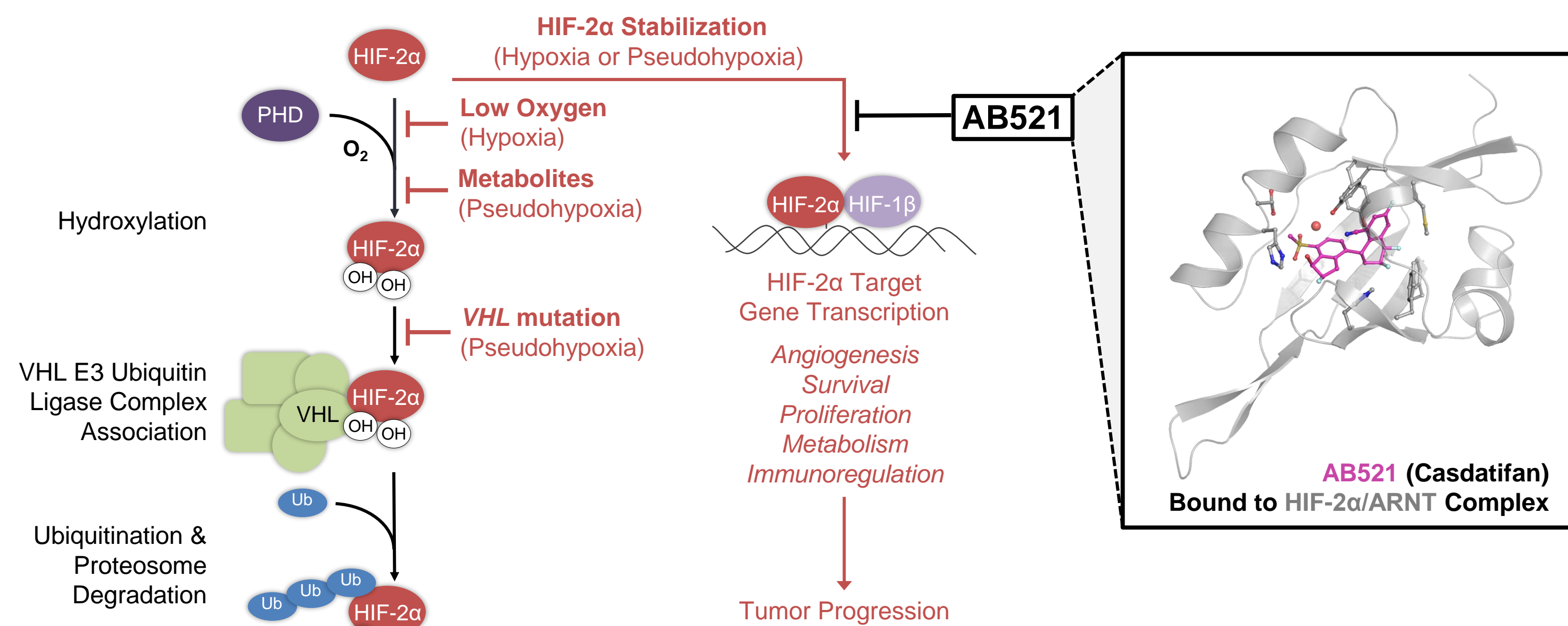


Figure 1. HIF-2 α drives transcriptional adaptations to low oxygen that can be hijacked to promote tumor progression. HIF-2 is a transcription factor comprised of two proteins: a stably expressed β subunit (HIF-1 β /ARNT) and an oxygen-sensitive α subunit (HIF-2 α). In hypoxic or pseudohypoxic conditions, HIF-2 α is stabilized and drives transcriptional programs that promote tumor progression. AB521 is a novel, potent, and selective allosteric small molecule inhibitor that can prevent HIF-2 α -dependent gene transcription and block tumor progression.

AB521 Selectively Inhibits HIF-2 α -Mediated Transcription and Colony Formation of Cancer Cells

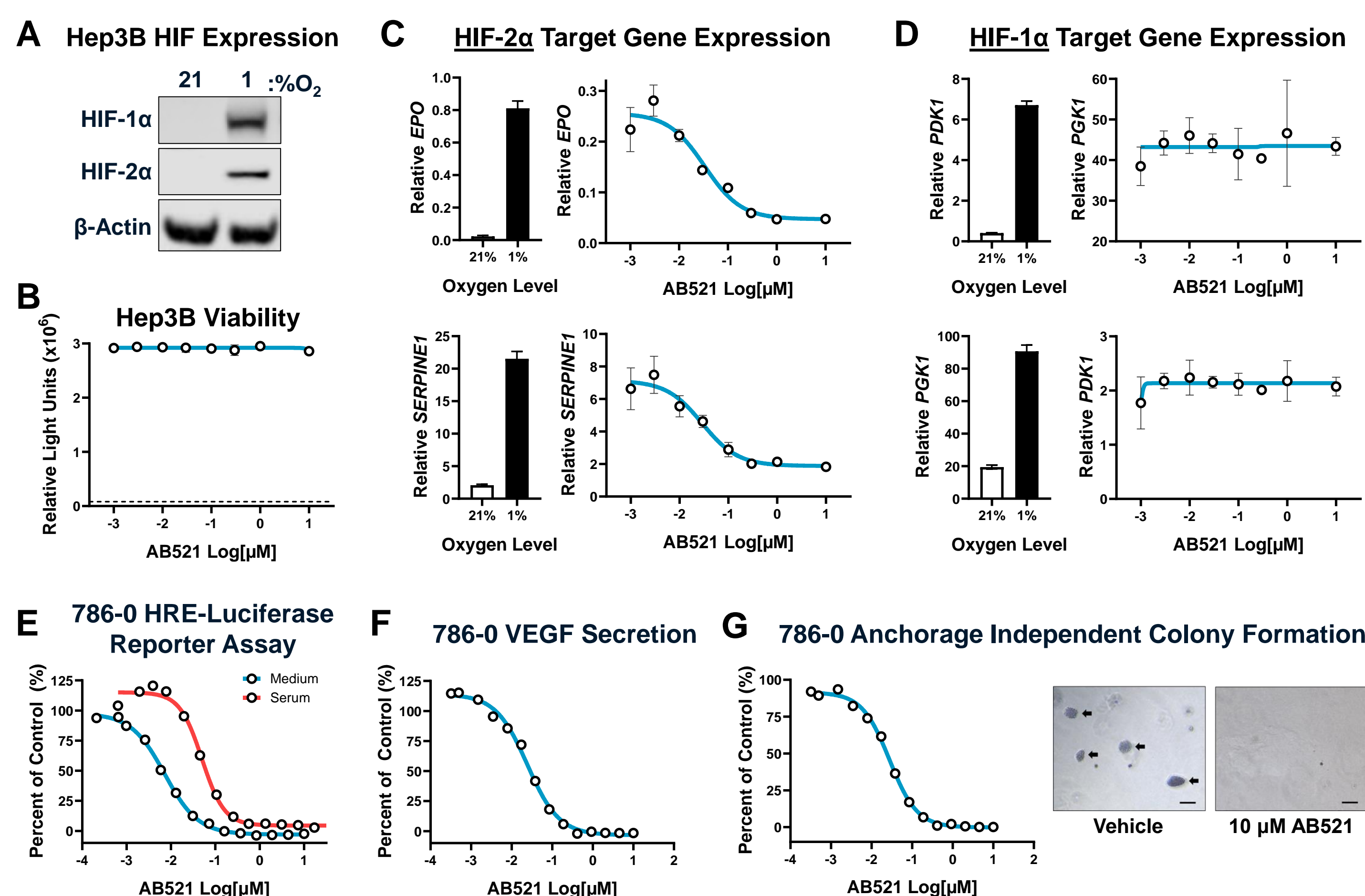


Figure 2. *In vitro* impacts of AB521 on cancer cell lines. A) HIF-1 α and HIF-2 α protein was stabilized under hypoxia (1% O₂) in Hep3B hepatocellular carcinoma cells. B) Viability of Hep3B cells was unaffected by AB521. C) AB521 selectively inhibited expression of HIF-2 α -regulated genes. D) HIF-1 α -regulated gene expression was unaltered by AB521 treatment. E) AB521 inhibited HIF-2 α -dependent hypoxia-response element (HRE) reporter expression in 786-O ccRCC cells in both medium and 100% human serum conditions. F) HIF-2 α -regulated endogenous VEGF secretion was inhibited by AB521. G) AB521 treatment inhibited 3D anchorage independent colony formation of 786-O cells.

AB521 Inhibits Transcription of HIF-2 α Regulated Genes in Macrophages and Endothelial Cells

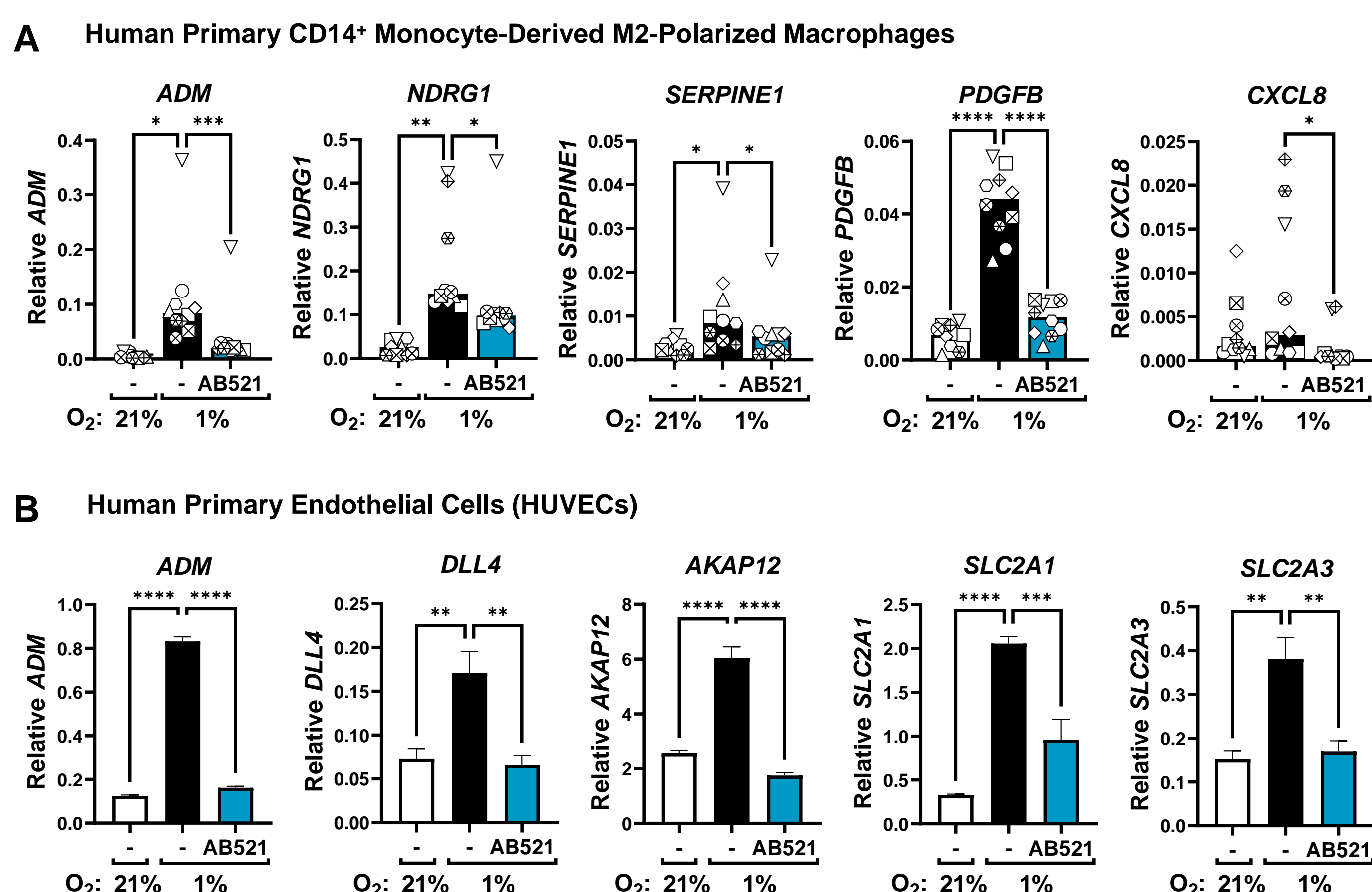


Figure 3. AB521 inhibits pro-tumorigenic gene expression in cell types relevant to the tumor microenvironment. A) Hypoxia-dependent upregulation of genes involved in angiogenesis (ADM, NDRG1), extracellular matrix remodeling (SERPINE1, PDGFB), and myeloid cell recruitment (CXCL8) was blocked with AB521 treatment in primary human M2-polarized macrophages. B) Hypoxia-dependent upregulation of genes involved in angiogenesis (ADM, DLL4, AKAP12), and glucose metabolism (SLC2A1, SLC2A3) was blocked by AB521 treatment in umbilical vein endothelial cells (HUVEC). AB521 used at 1 μ M, ANOVA with Dunnett's multiple comparisons * $p < 0.05$; ** $p < 0.01$; *** $p < 0.001$; **** $p < 0.0001$.

AB521 Shows Anti-Tumor Activity as a Single Agent and in Combination with TKI and I-O Therapies in Mice

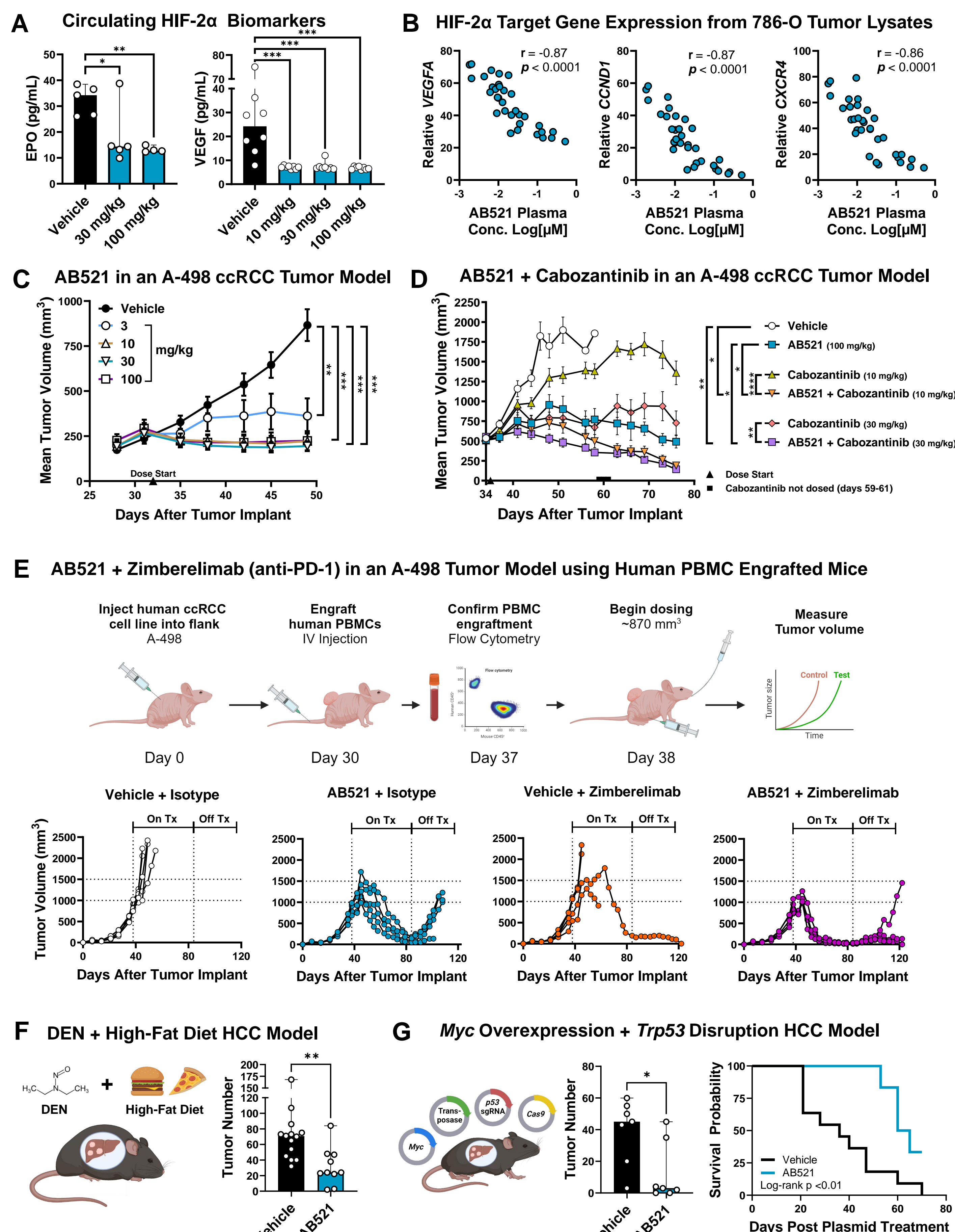


Figure 4. AB521 demonstrates anti-tumor activity in pre-clinical *in vivo* models. A) Mouse EPO and ccRCC tumor derived VEGF in plasma of mice after 8-hours (EPO) or 10-days (VEGF) of AB521 treatment. B) Relationship between AB521 plasma concentrations at trough and HIF-2 α target gene expression in 786-O xenograft lysates. C) Efficacy of AB521 in an A-498 ccRCC xenograft model. D) Combinatorial efficacy of AB521 with the tyrosine kinase inhibitor cabozantinib in an A-498 xenograft model. E) Evaluation of AB521 (30 mg/kg, PO QD) in combination with zimberelimab (anti-PD-1, 10 mg/kg IP Q3D) in an A-498 xenograft model using human peripheral blood mononuclear cell (PBMC) engrafted mice. F) AB521 therapeutic efficacy in a diethylnitrosamine (DEN) and high-fat diet driven hepatocellular carcinoma (HCC) model. G) AB521 therapeutic efficacy and impact on survival in a Myc and Trp53 driven HCC model.

Conclusions

- ❖ AB521 is a novel, potent, selective, and orally bioavailable HIF-2 α inhibitor
- ❖ Anti-tumor efficacy of AB521 as a single agent and in combination with clinically relevant therapies was observed in preclinical mouse tumor growth models
- ❖ AB521 (Casdatifan) is in clinical trials for ccRCC and other solid tumors
 - ARC-20: NCT05536141, STELLAR-009: NCT0691796
 - Upcoming trials in ccRCC: PEAK-1 Phase 3 in combination with cabozantinib and Phase 1 in combination with volrustomig, an investigational PD-1/CTLA-4 bispecific antibody