

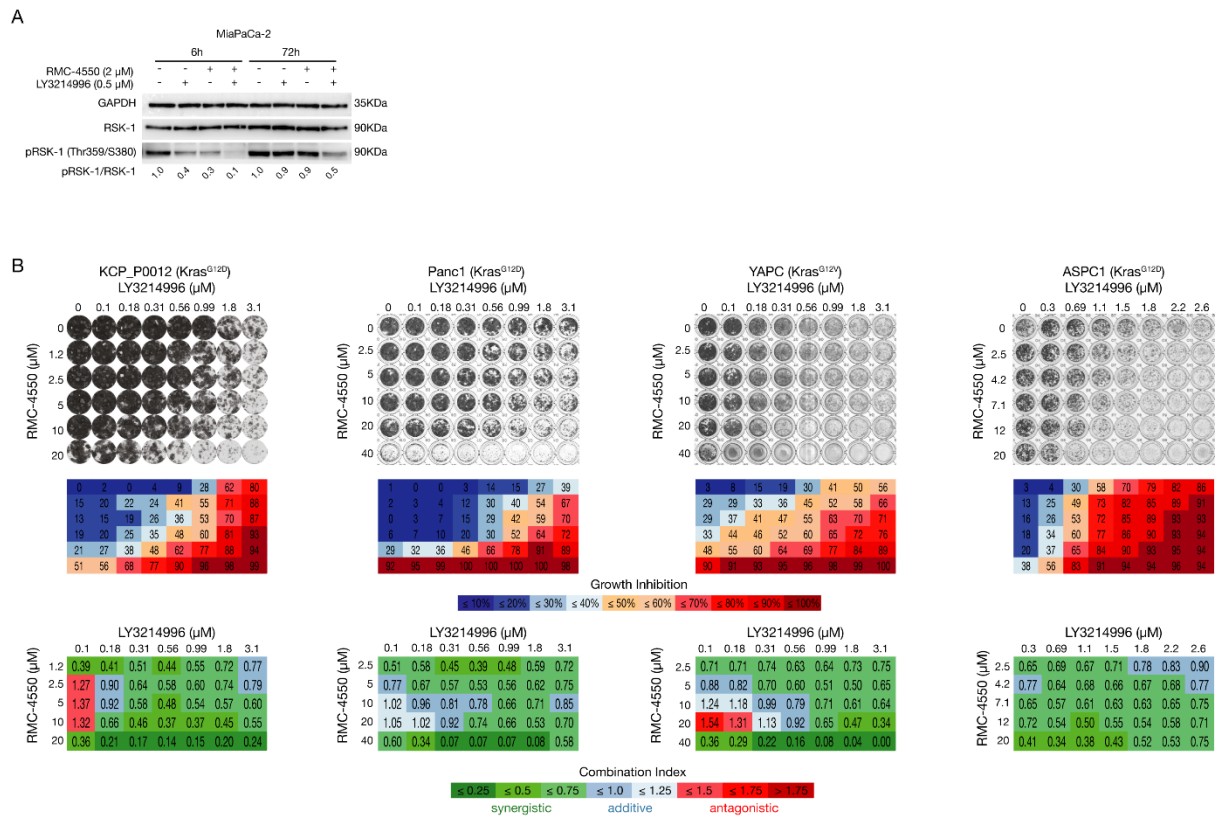
**Cell Reports Medicine, Volume 3**

**Supplemental information**

**Extensive preclinical validation of combined  
RMC-4550 and LY3214996 supports clinical  
investigation for KRAS mutant pancreatic cancer**

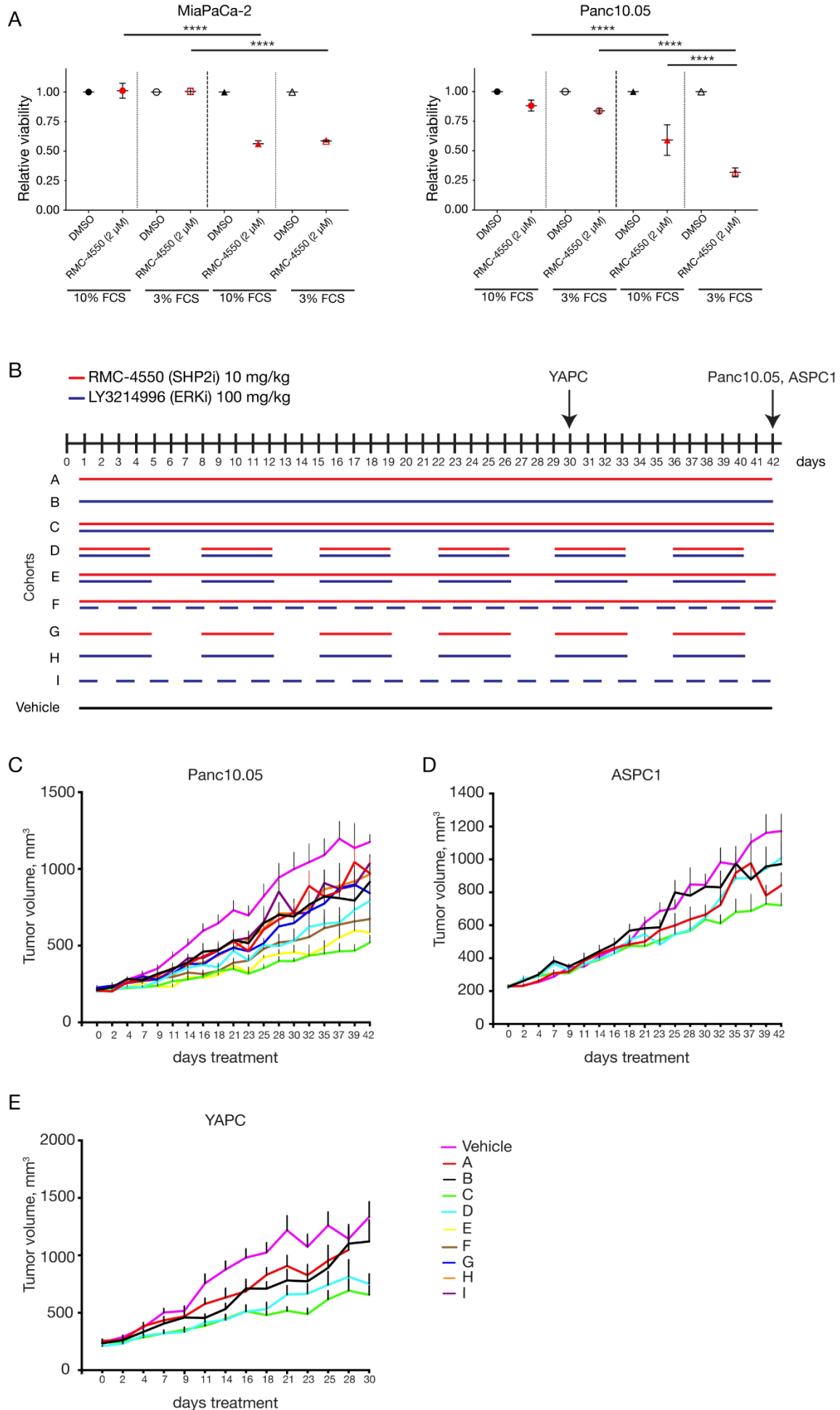
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## Supplemental material



### Supplementary Figure S1. Assessing the treatment response in murine and human *KRAS*-mutant pancreatic cancer cell lines, related to Figure 1

**A:** Western blot analysis with human cancer cell line MiaPaCa-2. Cells were treated as depicted and collected for lysis at the indicated time points. Protein extracts were probed with specific antibodies against total RSK-1, phosphorylated RSK-1 (pRSK-1), and GAPDH (as loading control). Numerical values indicate the pRSK-1/RSK-1 ratio quantified by densitometry. The blots are representative of at least three independent experiments. RSK-1 = Ribosomal s6 kinase 1. **B:** Synergistic effects of SHP2i and ERKi administration were evaluated by colony formation assay with murine cancer cell line KCP\_P0012 derived from KCP mouse model (Kras<sup>G12D</sup>) of spontaneous tumor formation and human cancer cell lines: Panc1 (KRAS<sup>G12D</sup>), YAPC (KRAS<sup>G12V</sup>), and ASPC1 (KRAS<sup>G12D</sup>). SHP2i and ERKi were combined at the concentrations indicated. Representative crystal violet staining of cells is shown (top panel). Box-matrices below the plate-scans depict quantification of growth inhibition in relation to control wells (middle panel). Bottom panel: Calculation of the Combination Index (CI) Scores from the growth inhibition values (shown above) via CompuSyn software demonstrating strong synergism between SHP2i and ERKi across a wide range of combinatorial concentrations. CI < 0.75 (shades of green) indicates synergism, CI = 0.75 – 1.25 (shades of blue) indicates additive effects and CI > 1.25 (shades of red) indicates antagonism. Experiments were repeated independently at least three times each, with similar results.

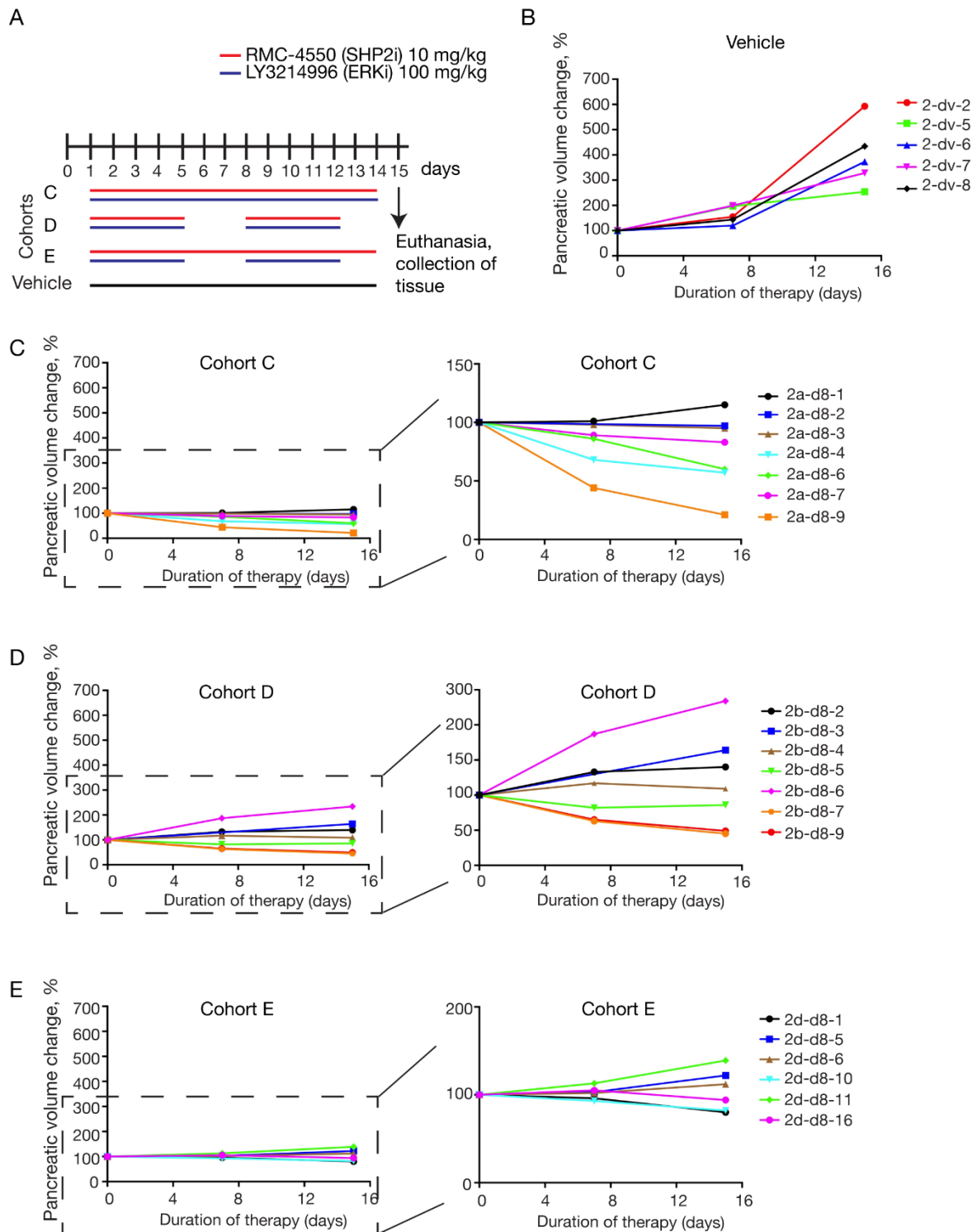


**Supplementary Figure S2. *In vivo* evaluation of the combined administration of RMC-4550 (SHP2i) and LY3214996 (ERKi) on tumor growth in different PDAC models, related to Figures 3 and 4**

**A:** Comparison of 2D vs. 3D growth and 10% serum vs. 3% serum conditions in the indicated human cancer cell lines treated with 2  $\mu$ M RMC-4550. Cell viability was measured after 4 days. Statistical analyses compared the effect of SHP2 inhibition between 2D and 3D growth both in 10% serum and 3% serum using ordinary one-way ANOVA test. Experiments were repeated independently at least three times each, with similar results.

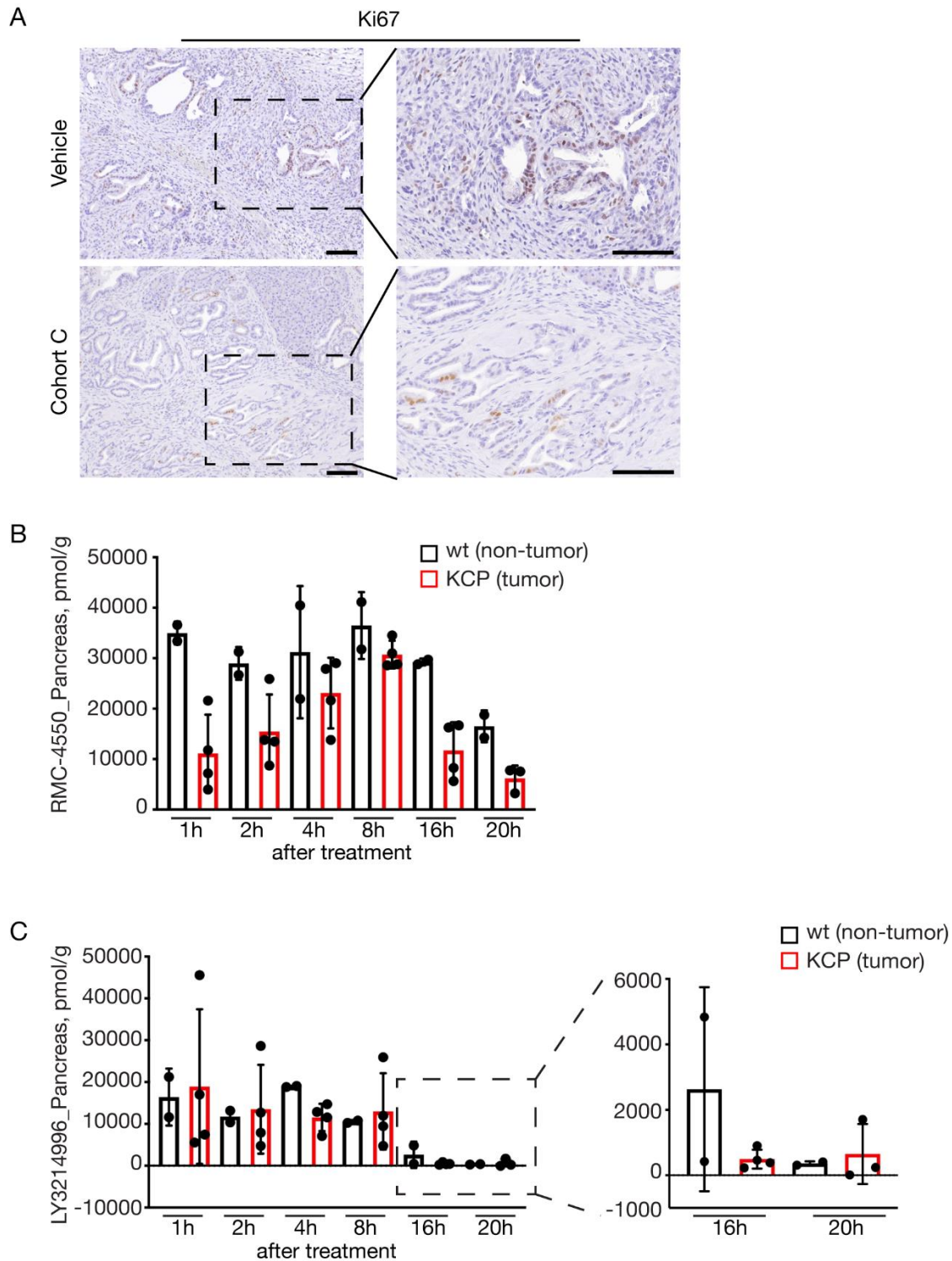
**B:** Schematic representation of the treatment schedule applied in MiaPaCa-2, ASPC1, and YAPC xenograft models. Cohort A: Continuous treatment with SHP2i alone daily; Cohort B: Continuous treatment with ERKi alone daily; Cohort C: Continuous treatment with the combination of SHP2i and ERKi daily; Cohort D: Intermittent treatment with the combination of SHP2i and ERKi 5 days on / 2 days off; Cohort E: Semi-continuous treatment schedule with daily dosing of SHP2i and intermittent dosing with ERKi 5 days on / 2 days off. Cohort F: Continuous treatment with SHP2i and on alternate days with ERKi. Cohort G: Intermittent dosing with SHP2i alone 5 days on / 2 days off. Cohort H: Intermittent dosing with ERKi alone 5 days on / 2 days off. Cohort I: Treatment with ERKi alone on alternate days. Control mice were continuously treated with vehicle. For all the xenograft experiments,  $5 \times 10^6$  cells were subcutaneously injected into the right flank of NOD scid gamma (NSG) mice, respectively. When tumors reached 200 - 250 mm<sup>3</sup>, mice were randomly assigned into cohorts and treated by oral gavage with inhibitors or vehicle according to treatment schedule. Mice were sacrificed after 1, 3 or 6 weeks of treatment (n = 8 mice per time point and per cohort) **C-**

**E:** Treatment response was assessed through tumor volume change using caliper measurements 3 times/week in different xenograft models: Panc10.05 (**C**), ASPC1 (**D**), and YAPC (**E**). Results represent mean  $\pm$  SD.



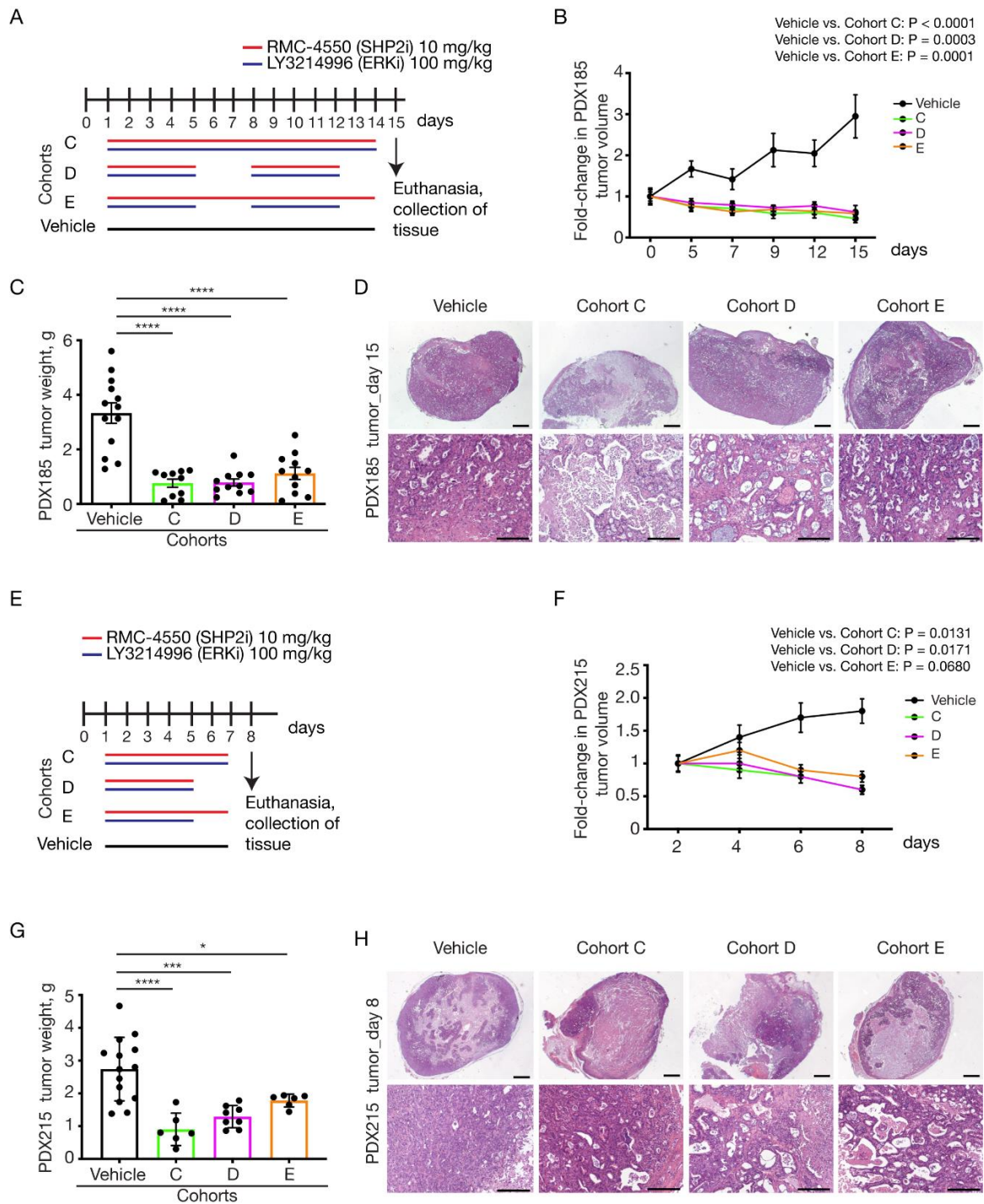
**Supplementary Figure S3. *In vivo* assessment of treatment response in an endogenous murine PDAC model, related to Figure 5**

**A:** Schematic representation of the treatment schedule applied in an endogenous (KCP) murine model of spontaneous tumor formation. Cohort C: Continuous treatment with the combination of SHP2i and ERKi daily (n = 9); Cohort D: Intermittent treatment with the combination of SHP2i and ERKi 5 days on / 2 days off (n = 7); Cohort E: Semi-continuous treatment schedule with daily dosing of SHP2i and intermittent dosing with ERKi 5 days on / 2 days off (n = 6). Control mice (n = 5) were treated with vehicle for 14 consecutive days. **B - E:** Volume-tracking curves for individual mice over the whole course of therapy. The y axis shows tumor volume change in %. Tumor volume at baseline before commencement of therapy is indicated by 100%.



**Supplementary Figure S4. Evaluation of drug distribution of RMC-4550 and LY3214996 and of tumor growth in an endogenous murine PDAC model, related to Figure 5**

**A:** Immunohistochemical analysis of paraffin-embedded pancreatic tissue samples of mice, treated with vehicle (n=5) or with the combination of SHP2i and ERKi daily (Cohort C, n = 9), using Ki67 antibody. Scale bars represent 100  $\mu$ m. **B, C:** Pharmacokinetic analysis regarding pancreatic tissue derived from KCP mouse model of spontaneous tumor formation and non-tumor bearing littermates over time. Mice were administered a single dose of 10 mg/kg RMC-4550 (SH2P2i) (**B**) or 100 mg/kg LY3214996 (ERKi) (**C**) and sacrificed after 1h (n = 4), 2h (n = 4), 4h (n = 4), 8h (n = 4), 16h (n = 4) and 20h (n = 3). Distribution for each drug in the tumor and healthy pancreas was determined by LC-MS/MS. Results represent mean  $\pm$  SD.



**Supplementary Figure S5. Treatment response in Patient-Derived Xenograft (PDX) models, related to Figure 6**

**A:** Treatment schedule for the PDX185 model. Mice were treated with the combination of RMC-4550 (SHP2i) and LY3214996 (ERKi) once per day via oral gavage for 14 consecutive days (Cohort C) or 5 days on / 2 days off (Cohort D) or with SHP2i continuous and ERKi 5 days on / 2 days off (Cohort E). For the PDX models, tumor pieces of 50 mm<sup>3</sup> were subcutaneously implanted into both flanks of NSG mice (n = 7 mice per cohort). When tumors reached 200 - 250 mm<sup>3</sup> (approximately 6 - 8 weeks after subcutaneous transplantation), mice were randomly assigned into cohorts and treated by oral gavage with inhibitors or vehicle according to treatment schedule for the indicated times. **B, C:** Treatment response of PDX185 was assessed

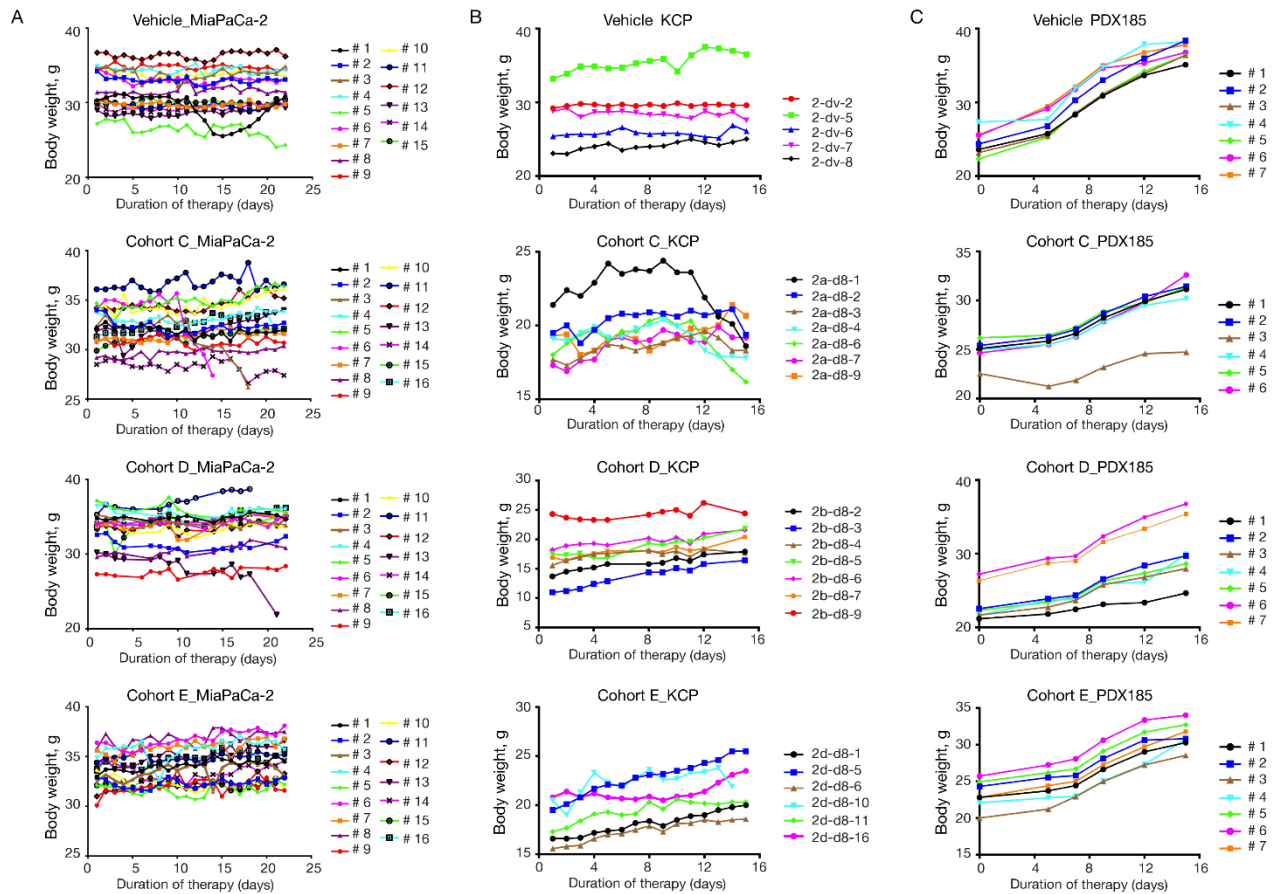
through tumor volume changes using daily caliper measurements (**B**) and tumor weight at endpoint (**C**). Results represent mean  $\pm$  SD. \*\*\*\* P <0.0001. Significance was determined by one-way ANOVA with Bonferroni's multiple comparison test. **D**: Representative H&E-stained sections of vehicle- and combination therapy-treated PDX185 tumors at day 15. Scale bars represent 1000  $\mu$ m (top) and 200  $\mu$ m (bottom). **E**: Treatment schedule for the PDX215 model. Mice were treated with the combination of RMC-4550 (SHP2i) and LY3214996 (ERKi) once per day via oral gavage for 7 consecutive days (Cohort C) or 5 days on / 2 days off (Cohort D) or with SHP2i continuous and ERKi 5 days on / 2 days off (Cohort E). **F, G**: Treatment response of PDX215 was assessed through tumor volume changes using daily caliper measurements (**F**) and tumor weight at endpoint (**G**). Results represent mean  $\pm$  SD. \* P <0.05, \*\*\* P <0.001, \*\*\*\* P <0.0001. Significance was determined by one-way ANOVA with Bonferroni's multiple comparison test. **H**: Representative H&E-stained sections of vehicle- and combination therapy-treated PDX215 tumors at day 8. Scale bars represent 1000  $\mu$ m (top) and 200  $\mu$ m (bottom).



**Supplementary Table S1:** Summary of Pathogenic and Likely Pathogenic Mutations in cells derived from PDX354 and PDX185, related to Figures 6 and S5

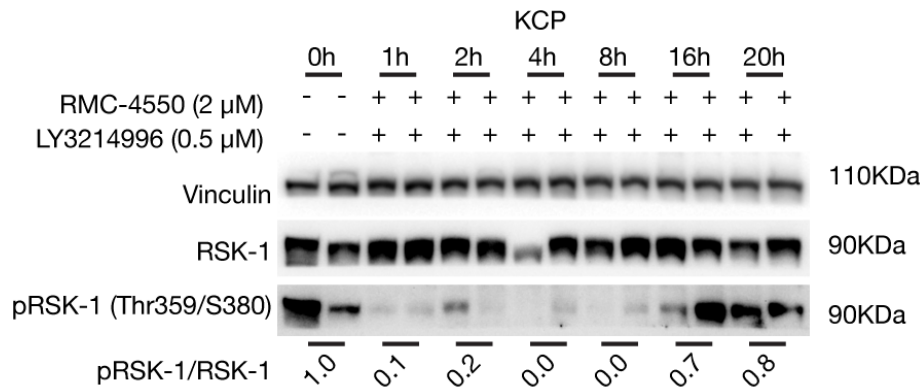
	<i>CDKN2A</i>	<i>EGFR</i>	<i>KRAS</i>	<i>SMAD4</i>	<i>TP53</i>
<b>PDX 354</b>	ENST00000498124.1:c.158T>G(p.Met53Arg)	NM_005228.5:c.2291A>G(p.Ty r764Cys)	NM_033360.4:c.35G>A(p.Gly 1 2Asp)	no pathogenic or likely pathogenic variants detected	ENST00000269305.4:c.1100+2 T>C
	ENST00000498124.1:c.221A>C(p.Asp74Ala)				
	ENST00000498124.1:c.206A>G(p.Glu69Gly)				
<b>PDX 185</b>	no pathogenic or likely pathogenic variants detected	NM_005228.5:c.868A>G(p.Thr 290Ala)	NM_033360.4:c.35G>A(p.Gly 1 2Asp)	NM_005359.6:c.1055del(p.Gly3 52AspfsTer32)	ENST00000269305.4:c.520A>G(p.Arg174Gly)
				NM_005359.6:c.1001A>G(p.Gln334Arg)	
				NM_005359.6:c.1022T>C(p.Val 341Ala)	

HGVS-nomenclature is used for each variant. NM = single nucleotide variant; ENST = splice variant



**Supplementary Figure S6. Comparison of body weight–time profiles between the different combination therapy cohorts in the endogenous murine PDAC model, the MiaPaCa-2 xenograft as well as the PDX model, related to Figures 3, 5 and S5.**

**A:** Individual body weight–time profile of the vehicle and combination therapy groups in MiaPaCa-2 xenograft bearing mice: Vehicle (n = 15), Cohort-C (n = 16), Cohort-D (n = 16), Cohort-E (n = 16). **B:** Individual body weight–time profile of the vehicle and combination therapy groups in the endogenous (KCP) murine model of spontaneous tumor formation: Vehicle (n = 5), Cohort-C (n = 7), Cohort-D (n = 7), Cohort-E (n = 6). **C:** Individual body weight–time profile of the vehicle and combination therapy groups in PDX185 bearing mice: Vehicle (n = 7), Cohort-C (n = 6), Cohort-D (n = 7), Cohort-E (n = 7).



**Supplementary Figure S7. Evaluation of pRSK-1 protein expression over time after administration of RMC-4550 (SHP2i) and LY3214996 (ERKi) in an endogenous murine PDAC model, related to Figures 1 and 7**

Western blot analysis with murine pancreatic tissue derived from KCP mouse model (KRAS<sup>G12D</sup>) of spontaneous tumor formation. Mice were treated with a single dose 8 of RMC-4550 (SHP2i) and LY3214996 (ERKi) and sacrificed after 1h (n = 2), 2h (n = 2), 4h (n = 2), 8h (n = 2), 16h (n = 2) and 20h (n = 2). Non-treated mice served as controls (0h, n = 2). Protein extracts of pancreatic tissue were probed with specific antibodies against total RSK-1, phosphorylated RSK-1 (pRSK-1), and alpha-tubulin (as loading control). Numerical values indicate the pRSK-1/RSK-1 ratio quantified by densitometry. The blots are representative of at least three independent experiments. RSK-1 = Ribosomal s6 kinase 1.