

Development of MK-8353, an orally administered ERK1/2 inhibitor, in patients with advanced solid tumors

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BACKGROUND. Constitutive activation of ERK1/2 occurs in various cancers, and its reactivation is a well-described resistance mechanism to MAPK inhibitors. ERK inhibitors may overcome the limitations of MAPK inhibitor blockade. The dual mechanism inhibitor SCH772984 has shown promising preclinical activity across various *BRAF*^{v500}/*RAS*-mutant cancer cell lines and human cancer xenografts.

METHODS. We have developed an orally bioavailable ERK inhibitor, MK-8353; conducted preclinical studies to demonstrate activity, pharmacodynamic endpoints, dosing, and schedule; completed a study in healthy volunteers (P07652); and subsequently performed a phase I clinical trial in patients with advanced solid tumors (MK-8353-001). In the P07652 study, MK-8353 was administered as a single dose in 10- to 400-mg dose cohorts, whereas in the MK-8353-001 study, MK-8353 was administered in 100- to 800-mg dose cohorts orally twice daily. Safety, tolerability, pharmacokinetics, pharmacodynamics, and antitumor activity were analyzed.

RESULTS. MK-8353 exhibited comparable potency with SCH772984 across various preclinical cancer models. Forty-eight patients were enrolled in the P07652 study, and twenty-six patients were enrolled in the MK-8353-001 study. Adverse events included diarrhea (44%), fatigue (40%), nausea (32%), and rash (28%). Dose-limiting toxicity was observed in the 400-mg and 800-mg dose cohorts. Sufficient exposure to MK-8353 was noted that correlated with biological activity in preclinical data. Three of fifteen patients evaluable for treatment response in the MK-8353-001 study had partial response, all with BRAF^{V600}-mutant melanomas.

CONCLUSION. MK-8353 was well tolerated up to 400 mg twice daily and exhibited antitumor activity in patients with *BRAF*^{v600}-mutant melanoma. However, antitumor activity was not particularly correlated with pharmacodynamic parameters.

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Introduction

ERK1/2 are members of the MAPK family of signaling proteins, a superfamily of cytoplasmic serine/threonine kinases that normally transmit signals from various extracellular stimuli to intracellular targets and regulate a number of cellular processes, including proliferation, adhesion, migration, survival, differentiation, and metabolism (1, 2). Activating mutations in upstream effectors that result in constitutive MAPK activation are present in almost 35% of all cancers (3). Reactivation of the ERK1/2 phosphorylation and signaling has been frequently associated with secondary resistance to BRAF and MEK inhibitors (4, 5), highlighting the importance of exploring targeted inhibition of more downstream components of the MAPK signaling pathway. In fact, in *BRAF*^{V600}-mutant melanoma cell lines that were established from patients who developed resistance to MAPK pathway inhibitors, pharmacologic inhibition of ERK1/2 was more potent than MEK inhibition in suppressing MAPK activity and cell proliferation and enhancing cell death (6).

While the development of small-molecule inhibitors of RAS currently remains pharmacologically challenging (reviewed in ref. 7), a number of ERK inhibitors (ERKi) have been successfully developed and are currently in early clinical development (8, 9). These compounds have been shown to either disrupt ERK dimerization (10) or to bind to the active conformation, inhibit intrinsic kinase activity, and, therefore, prevent phosphorylation of downstream kinases ("single mechanism," ATP-competitive, type I) (11–13). SCH772984 is a unique ERK1/2 inhibitor, because it does not only bind to and inhibit ERK1/2-intrinsic kinase activity, but it also induces a conformational change upon binding to ERK1/2 that prevents ERK1/2 phosphorylation by upstream kinases ("dual mechanism") (11, 14, 15). More specifically, SCH772984 potently and selectively inhibits ERK1 and ERK2 activity, with IC₅₀ values of 4 and 1 nM, respectively. Treatment of *BRAF/RAS*-mutant melanoma and colon cancer cell lines with SCH772984 led to sustained inhibition of the downstream effectors of ERK1/2 signaling, transient inhibition of MEK-mediated ERK phosphorylation (pERK), decreased cell proliferation, and increased apoptosis. In addition, SCH772984 suppressed proliferation of melanoma and colorectal cancer cell lines that had developed secondary resistance to BRAF/MEK inhibitors (16, 17). Finally, concurrent administration of vemurafenib and SCH772984 in various melanoma cell lines was significantly more potent than that with either agent alone (17).

Based on these results, we developed an orally bioavailable dual mechanism ERKi, MK-8353 (formerly SCH900353), with improved pharmacokinetic properties compared with the preclinical tool compound SCH772984. Preclinically, MK-8353 showed similar selectivity and potency to SCH772984. Antitumor activity was comparable in various preclinical models. Following an early study of MK-8353 administered as a single dose in healthy human subjects (P07652), we performed a phase I trial of MK-8353 in patients with advanced solid tumors (MK-8353-001). The first objective in the MK-8353-001 clinical trial was to define the dose-limiting toxicity (DLT) and maximum tolerated dose (MTD) of MK-8353 and to determine the recommended phase 2 dose (RP2D). The second objective was to evaluate the antitumor efficacy of MK-8353 in patients with metastatic melanoma and colorectal cancer bearing *BRAF* or *RAS* mutation(s).

Results

MK-8353 is a highly potent, orally bioavailable, dual-specificity ERK1/2 inhibitor. MK-8353 [HCl ($C_{37}H_{42}CI-N_9O_3S$), molecular weight, 728.31, Figure 1A] is an orally bioavailable, selective, and potent ERKi that inhibits activated ERK1 and ERK2 in vitro, with IC_{50} values of 23.0 nM and 8.8 nM, respectively (IMAP kinase assay), and nonactivated ERK2, with an IC_{50} of 0.5 nM (MEK1-ERK2-coupled assay). MK-8353 demonstrates kinase selectivity over a 227-human kinase panel; no additional kinase in the panel was inhibited by more than 35% at the 0.1 μM concentration, and only 3 kinases (CLK2, FLT4, and Aurora B) were inhibited >50% at the 1.0 μM concentration (data not shown). Similar to SCH772984, MK-8353 caused a dose-proportional decrease in the phosphorylated-activated forms of ERK1 (ERK1), ERK2 (ERK2), and ribosomal S6 kinase (ERK1) proteins, with complete suppression of ERK1 and ERK2 observed at 30 nM ERK1 mutant cancer cell lines (Supplemental Table 1; supplemental material available online with this article; https://doi.org/10.1172/jci.insight.92352DS1).

The in vivo antitumor activity of MK-8353 was evaluated in a panel of human xenograft models, including the BRAFV600 mutant Colo-205 colon cancer model and the BRAFV600 mutant SK-MEL-28 melanoma model (Supplemental Table 2). MK-8353 inhibited the growth of both models in a dose-dependent manner at dose levels of 30–60 mpk, by oral gavage, twice daily. The 30 mpk dose significantly inhibited the growth of both models, while the 60 mpk dose led to tumor regressions (Figure 2, A and B).



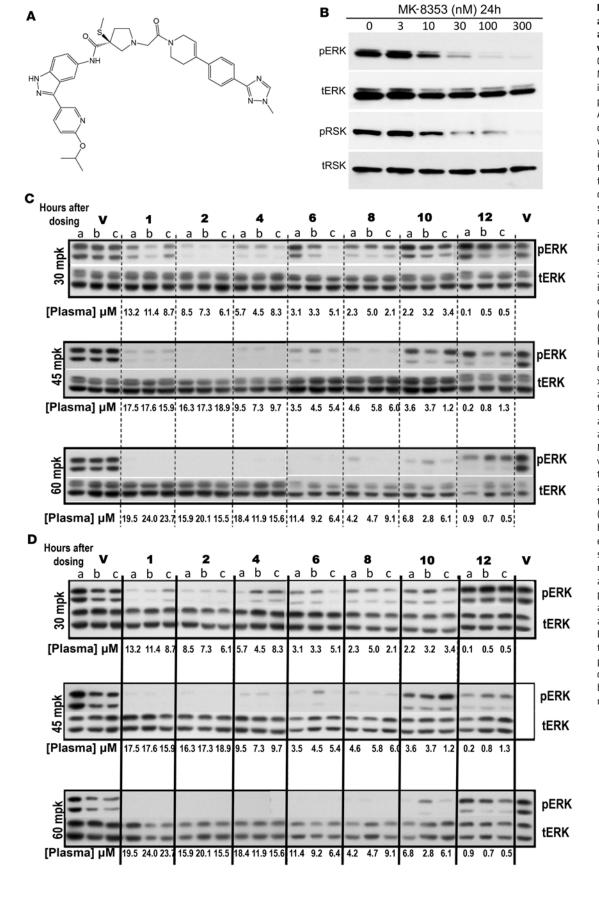


Figure 1. Structure and biochemical activity of MK-8353 in vitro and in vivo. (A) Chemical structure of MK-8353. (B) MK-8353 inhibits ERK and RSK phosphorylation. A2058 cells (1 \times 10 6 cells per 10-cm dish) were treated with increasing concentrations of MK-8353 (nM) for 24 hours. Whole cell lysates were subjected to immunoblot analysis using antibodies described in the Methods section. Image shows a representative immunoblot of a total of two experiments. (C and D) MK-8353 (30-60 mpk) inhibits ERK phosphorylation in Colo-205 human colon cancer tumor xenograft tissues (C) and paired normal skin tissues (**D**) as early as 1 hour following a single dose of MK-8353 versus drug vehicle (V). Xenograft tumors and same animal normal skin tissue were harvested (3 mice per group), and homogenates from excised tissues were subjected to immunoblot analysis using antibodies against phosphorylated/ activated ERK (pERK) and total ERK (tERK). Experiment in each tumor xenograft was performed once. See complete unedited blots in the supplemental material.



Overall, treatment of various other human cancer xenografts with MK-8353 administered by oral gavage at 60 mpk twice daily resulted in at least 50% tumor growth inhibition or regression in 83% of the animal models tested. In particular, treatment of Colo-205 colon cancer xenografts with MK-8353 at 30-60 mpk resulted in significant inhibition of pERK, as detected by immunoblot analysis of whole cell lysates obtained from tumor tissues as early as 1 hour following a single dose of MK-8353; however, activity of ERK1/2 rebounded after 10-12 hours (Figure 1C). Suppression of pERK in tumor tissue, as detected by Western blot analysis of corresponding tissue homogenates (Figure 1C versus Figure 1D) and by immunohistochemistry (Figure 2C versus Figure 2D; also Supplemental Figure 1), did correlate with suppression of pERK in matched skin samples from Colo-205 human colon cancer xenografts, especially as early as 2-4 hours after a single dose. In fact, modeling of the mouse preclinical data by quantitative pharmacology and pharmacometrics indicated that >65% suppression of mean pERK protein expression in skin biopsies from tumor-bearing nude mice was likely to correlate with >95% suppression of pERK in tumor samples and, therefore, result in meaningful clinical activity (data not shown). We therefore conclude that MK-8353 administered orally at doses ranging between 30 and 60 mpk results in dose-proportional tumor growth inhibition or regression across various human cancer models and that twice daily dosing is required to induce sustained dose-proportional suppression of pERK in tumor tissue. Finally, detection of pERK in skin biopsies by immunohistochemistry within 2–4 hours after MK-8353 dosing can be used as a surrogate tissue pharmacodynamic biomarker of MK-8353 activity within a given window from dosing in vivo.

The pharmacokinetic parameters of MK-8353 in female nude mice were determined across various dose levels (Supplemental Table 3). Exposure levels [AUC from 0 time to 24 hours, (AUC_{0-24hours})] for a dose range between 30 and 60 mpk, the doses corresponding with both significant inhibition of pERK in Colo-205 human xenografts and with tumor growth inhibition or regression across various human cancer xenograft animal models, were established to be approximately 15.3–55 µM·h (Supplemental Figure 1). Therefore, these exposure ranges were benchmarked as the approximate exposure levels that would result in antitumor activity in cancer patients. The no-observed-adverse-effect level (NOAEL) corresponding to repeated oral twice daily administration of MK-8353 across various species/strains was quite variable but within the dose range that resulted in both biologically effective dose and tumor growth inhibition or regression in mice (Supplemental Table 4). The most conservative NOAEL from the 4-week oral rat toxicity study (10 mg/kg) was used to calculate the human equivalent dose, which is 1.62 mg/kg. Based on a safety factor of 10, the lowest recommended safe single dose of oral MK-8353 in healthy volunteers (average weight was considered 60 kg) was 10 mg.

MK-8353 administration in healthy volunteers (P07652 study). A healthy volunteer study of MK-8353 was completed (P07652) in the Netherlands to collect toxicity, pharmacokinetic, and pharmacodynamic data. As part of this study, adult males received a single oral dose of MK-8353 in increasing (10-to 400-mg) dose cohorts (10 mg, n = 6; 40 mg, n = 6; 80 mg, n = 12; 150 mg, n = 6; 300 mg, n = 6; 400 mg, n = 12) in 2 alternative cohorts. In cohort 1, 6 subjects received MK-8353 and 2 subjects received placebo at each dose level of 10 mg, 40 mg, 150 mg, 400 mg, and 400 mg plus 5 days of pantoprazole. In cohort 2, 6 subjects received an active dose and 2 subjects received placebo at each dose level of 20 mg, 80 mg, 300 mg, 400 mg, and 80 mg plus food.

MK-8353 was generally safe and well tolerated across all doses, and no serious adverse events (SAEs) occurred. Adverse events (AEs) considered to be related to MK-8353 were diarrhea, pruritus, and rash. Following single-dose oral administration, peak plasma levels were attained after the median time had elapsed to reach maximum concentration ($t_{\rm max}$) of 1.5–3 hours, which was longer than the $t_{\rm max}$ observed in the corresponding single-dose animal studies; the $t_{\rm max}$ subsequently declined biexponentially, with a terminal estimated half-life of 4.2–8.9 hours. A dose-proportional increase in peak serum concentrations ($C_{\rm max}$) and the AUC_{0-24hours} was noted (Supplemental Table 5). The predicted steady-state exposure of MK-8353 administered at 100 mg orally twice daily suggested that the AUC_{0-24hours} in subjects treated with a single 100-mg dose of MK-8353 was predicted to be 16 μ M·h at steady state, which is below the NOAEL exposure level in the most sensitive species from the safety Good Laboratory Practice toxicology studies. MK-8353 administered at doses greater than or equal to 300 mg achieved the AUC_{0-24hours} range of 15.5–55 μ M·h; this range had been predicted to have a considerable biologic and antitumor efficacy in mice (Supplemental Table 3 and Supplemental Figure 1). This corresponds to an AUC_{0-12hours} of at least 12.6 μ M·h, which has a substantial biological effect. However, no significant change in pERK expression was seen in hair follicles and peripheral blood mononuclear cells in any of the different dose cohorts compared with baseline (data not shown).



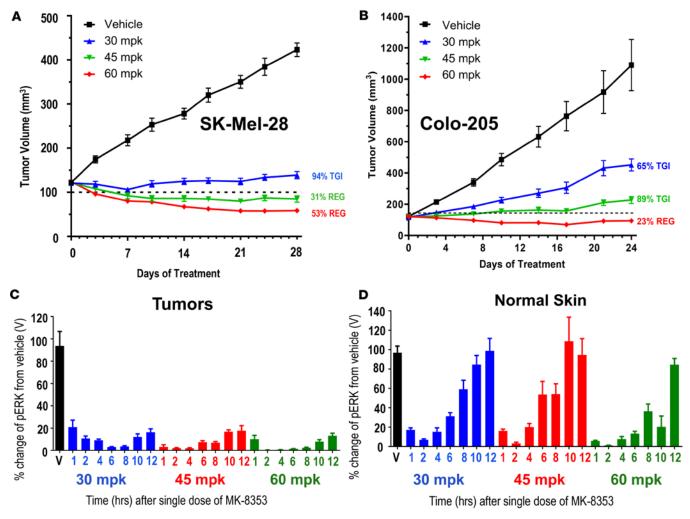


Figure 2. MK-8353 induces tumor growth inhibition or regression across various human cancer xenograft models. Female athymic nude mice (Colo-205) or SCID mice (SK-MEL-28) were injected with human Colo-205 colon cancer cells (B) and SK-MEL-28 melanoma cells (A) (both BRAF^{V600E}-mutant), as described in the Methods section. (A and B). Tumor growth curves show mean tumor volumes ± SEM (10 mice per group for each xenograft experiment). There was a statistically significant dose-dependent effect on tumor growth compared with the vehicle-treated group in both xenograft models that ranged from tumor growth inhibition (TGI) to regression (REG) and was present as early as day 3 from treatment initiation (*P* < 0.01, across all time points except for the day 3 time point in the Colo-205 xenograft model, in which the 30-mpk group was significantly different, *P* < 0.05; Student's *t* test). (C and D) Expression of pERK by tissue immunohistochemistry in Colo-205 tumor xenografts (C) and same-animal normal skin tissues (D) following a single dose of MK-8353 administered by oral gavage at various doses. Representative tissue sections from corresponding tissues that were harvested at various time points (0-12 hours) following a single dose of MK-8353 were stained with antibodies against pERK. See the Methods section for details and Supplemental Figure 1 for representative images of 3 animals, with tumors and normal skin imaged at each time point. Results are shown as percentage change of pERK from vehicle using the formula described in the Methods section. mpk, milligrams per kilogram; TGI, tumor growth inhibition; REG; regression; Veh, drug vehicle.

Safety and DLT in the MK-8353-001 study. Between November 2011 and February 2014, 26 patients were enrolled and 25 patients received MK-8353 (Figure 3 and Tables 1 and 2). Ten patients were treated in part 1a at doses ranging from 100 mg to 800 mg, and 15 patients were treated in part 1b at doses ranging from 200 mg to 400 mg. Dose escalation proceeded up to the 800-mg dose cohort, at which 2 of 6 patients experienced DLT, making us consider this dose to be beyond the MTD. Subject 1006 developed grade 3 diarrhea, hyperbilirubinemia, nausea, vomiting, and grade 2 vision changes, and subject 1010 developed grade 3 fatigue and grade 3 hyperbilirubinemia. Two of seven patients experienced DLT in the 400-mg dose cohort. Subject 2003 developed grade 2 diarrhea, which was considered to be a DLT due to several missed doses and grade 3 rash. The most frequent AEs of any grade included diarrhea (44%), fatigue (40%), nausea (32%), maculopapular rash (28%), and vomiting (28%) (Table 3). Increased serum bilirubin was determined not be a DLT, because both drug-related and patient-related factors may have contributed. More specifically, inhibition of UGT1A1-mediated bilirubin glucuronidation was most like-



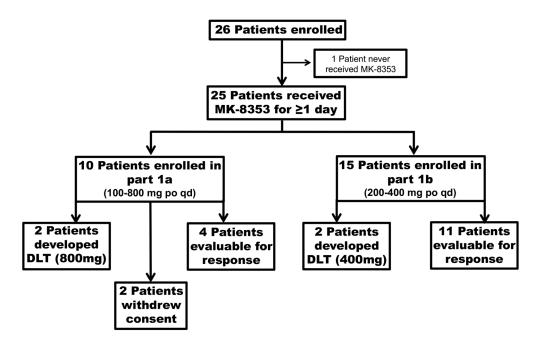


Figure 3. CONSORT diagram of the MK-8353-001 clinical trial. The flow chart shows the number of patients who were enrolled in parts 1a and 1b, received at least 1 dose of MK-8353, developed dose-limiting toxicity (DLT), and were evaluable for antitumor response by RECIST v1.1 criteria.

ly the cause of the hyperbilirubinemia observed, since some patients with elevated indirect bilirubin were found to have genetic polymorphisms in bilirubin metabolism-related genes. The most frequent grade 3/4 AEs were diarrhea (16%), isolated asymptomatic increased indirect serum bilirubin (16%), and maculopapular rash (8%). The 400-mg dose was identified as the preliminary MTD in part 1a. Due to early discontinuation of the study by the sponsor for strategic reasons and not due to toxicity, the RP2D was not determined. However, dosing up to 400 mg was safe and well tolerated during part 1b.

Pharmacokinetic analysis of drug exposure from the MK-8353-001 study. Plasma pharmacokinetic data from 24 patients at all 6 dose levels showed that the $t_{\rm max}$ ranged from 2 to 5 hours, mean terminal phase half-life ranged from 5 to 14 hours, and a biexponential decline occurred after peak. Individual AUC and $C_{\rm max}$ values were somewhat more variable within doses compared with the healthy volunteer study, but, nonetheless, near proportional increases in their median values were observed. Notable outliers included 3 patients with unexplained rises in MK-8353 concentration from 12 to 24 hours after a single dose (subjects 2006, 2007, and 2009) and subject 1004, who exhibited substantially reduced exposure, presumably due to incomplete absorption of MK-8353 from a Whipple procedure for pancreatic cancer (Figure 4 and Supplemental Table 6). Following continuous twice daily dosing, steady-state levels were achieved by day 8 in all patients (data not shown), with an accumulation ratio of approximately 2-fold in plasma on day 15. Interestingly, unexplained approximately 2-fold higher exposures in cancer patients compared with healthy subject volunteers without significant differences in mean $t_{\rm max}$ and $t_{\rm 1/2}$ values were noted.

Using the $AUC_{0-12hour}$ cutoff of 12.6 μ M·h as the minimum desired AUC exposure to achieve a biologically substantial effect, nearly all patients across all cohorts had sufficient dose-dependent exposure. Unfortunately, none of the 3 patients (1005, 1006, and 1009) who were dosed with 800 mg, the dose cohort with the highest $AUC_{0-12hours}$ exposure, were on the drug long enough to assess antitumor response.

At high doses, MK-8353 (400 mg and 800 mg orally twice daily) was found to induce unconjugated asymptomatic hyperbilirubinemia in 3 subjects (1006, 1010, and 2004). No significant differences in pharmacokinetic exposure (C_{max} , AUC_{0-12h}, $t_{1/2}$) or MK-8353 metabolism were seen in these 3 subjects compared to all other patients (Figure 4), suggesting that high pharmacokinetic exposure of the body to MK-8353 does not correlate with elevated bilirubin. Alternatively, hyperbilirubinemia may be the result of either altered metabolism by uridine 5'-di-glucuronosyltransferases (UGT) or altered hepatic uptake of unconjugated bilirubin. Although multiple MK-8353 glucuronides (A, B, and C) can be potentially formed via various UGT1 enzymes (UGT1A1, UGT1A3, UGT1A4), MK-8353 inhibits glucuronidation mediated by UGT1A1, a



Table 1. Patient demographics and characteristics for the MK-8353-001 study

Characteristic	Total (n = 25)				
Sex, n (%)					
Male	9 (36)				
Female	16 (64)				
Race, n (%)					
White	22 (88)				
Nonwhite	3 (12)				
Asian	1 (4)				
Black or African American	1 (4)				
Hispanic	1 (4)				
Age (yr), median (range)	59 (40-73)				
18-64	15 (60)				
≥65	10 (40)				
Tumor histology (%)					
Melanoma	12				
Colorectal	7				
Pancreas	2				
NSCLC	2				
Head and neck	1				
Papillary thyroid	1				
Prior systemic treatments, n (%)					
Median no. (range)	2 (0-6)				
ECOG performance status					
Median (range)	1 (0-1)				
NSCLC, non-small cell lung cancer; ECOG, Eastern Cooperative Oncology Group.					

UGT1 member that has been associated with reversible hyperbilirubinemia with HIV protease inhibitors (18), with high affinity (IC $_{50}$ 0.58 μ M based on an in vitro inhibition assay). Bilirubin transport via the organic anion-transporting polypeptides, OATP1B1 and OATP1B3 (IC $_{50}$ 0.31 μ M and 0.97 M, respectively, based on in vitro inhibition study), is a less likely possibility, since this mechanism would result in both conjugated and unconjugated bilirubin increase in serum.

Analysis of inhibition of pERK in skin biopsies from the MK-8353-001 study. Normal skin biopsies from baseline, the day 8 on-treatment time point (3–5 hours after dose, corresponding to the approximate C_{max}), and the day 15 on-treatment time point (8-10 hours after dose corresponding to the approximate C_{\min}) were obtained. Baseline and day 8 biopsies were available from 18 patients, whereas 13 patients had all 3 biopsies available. Development of AEs before day 15 (1007), withdrawal of consent (2007 and 2013), and lack of submission of specimen from the study site (2003 and 2016) prevented collection from 5 subjects on the day 15 time point. With the exception of 2 subjects from whom no skin biopsies were available from any time point (1001 and 2006) due to shipping issues, 5 subjects who were enrolled in the 800-mg dose cohort did not have any on-treatment biopsies because they remained on study treatment for less than 8 days. Figure 5 shows corrected histoscore (H-score) for pERK for each patient for a given time point, and Supplemental Table 7 shows actual data values. The average baseline pERK expression in normal epidermis was variable among patients (mean ± SD, 113 ± 38). Sustained inhibition of pERK protein expression in both on-treatment biopsies was seen in 5 subjects, and optimal (>65%) suppression of pERK protein expression in any of the on-treatment skin biopsies was observed in 3 of the 18 evaluable subjects. Eleven of eighteen patients showed at least 20% and seven of eighteen patients showed at least 40% suppression of mean pERK protein expression in one or more of the two on-treatment normal skin biopsies. All patients with an objective response to

therapy showed >20% suppression of mean pERK protein expression in this analysis. Finally, more than 40% suppression of mean pERK protein expression in any of the two on-treatment time points was seen in the 200-mg (1 of 2 evaluable), 350-mg (1 of 3 evaluable), and 400-mg (2 of 7 evaluable) dose cohorts.

Antitumor activity of MK-8353 in the MK-8353-001 study. Ten patients were inevaluable for radiographic response on day 28: four patients due to DLT, two patients due to MK-8353-related SAEs, one patient due to asymptomatic hyperbilirubinemia related to MK-8353, one patient due a SAE related to cancer progression and, additionally, two patients who withdrew consent with symptoms (blurry vision and grade 1 nausea and vomiting). Figure 6 shows waterfall plots corresponding to the best antitumor response from baseline for the 15 evaluable patients based on investigators' assessment. Three partial responses (PRs) were seen (2008, 2011, 2014; 2008 and 2011 were confirmed) enduring for 167, 95, and 57 days, respectively; therefore, the overall response rate was 20% (evaluable patients) or 12% when both evaluable and inevaluable patients were considered. In addition, subject 2007, who had a history of BRAF^{V600E}-mutant melanoma (no other gene mutation was tested) and a prior PR to single-agent vemurafenib, had confirmed stable disease enduring for 114 days while on 300 mg MK-8353. All responders were males of European descent bearing metastatic (AJCC M1b) BRAF^{V600}-mutant melanoma, irrespective of prior treatment with MAPK pathway inhibitor. In fact, 3 of 8 patients with BRAF^{v600}-mutant melanoma achieved PR, including 4 patients who have previously received MAPK pathway inhibitor-based treatments. Six patients (2010, 2005, 1002, 1003, 2012, and 2016) were found to have progressive disease after 2 cycles, and three patients (2002, 2013, and 2004) progressed after 1 cycle. Subject 2008 (300-mg dose cohort) was 65 years old with a history of BRAF^{V600E}-mutant melanoma without other coexisting mutations based on a 46-gene panel and no prior MAPK pathway inhibitor-based treatments. Upon treatment with MK-8353, he developed a PR after 2 cycles and resolution of all subcentimeter lung nodules following 3 cycles. Following progression on MK-8353, he was subsequently treated with concurrent dabrafenib plus trametinib (D+T);



Table 2. Individual patient data of MK-8353-001 study subjects

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Subject no.	Cancer diagnosis	Mutation	Prior systemic treatments (BOR)	Study part	Dose cohort (mg BID)	Evaluable for efficacy	Reason for D/C	No. days on MK-8353
1001	Pancreas	KRAS ^{12V}	1 (SD)	1a	100	No	AE	13
1002	Colorectal	$KRAS^{WT}$	5 (adj/SD/unk)	1a	100	Yes (PD)	PD	56
1003	Head and neck	Nontest	3 (PD/unk)	1a	200	Yes (PD)	PD	56
1004	Pancreas	KRAS ^{12R}	1 (PD)	1a	400	Yes (SD) ^A	PD	43
1005	Papill thyroid	Nontest	2 (PD)	1a	800	No	WC(AE)	4
1006	NSCLC	No mutat	3 (SD/PD/unk)	1a	800	No	AE	7
1007	NSCLC	KRAS ^{12V}	3 (SD/PD)	1a	800	No	AE	12
1008	Colorectal	KRAS ^{12D}	3 (PD)	1a	800	No	PD/AE	4
1009	Colorectal	KRAS ^{12V}	3 (SD/PD/unk)	1a	800	No	AE	4
1010	Colorectal	KRAS ¹²⁵	1 (PD)	1a	800	No	AE	5
2002	Colorectal	KRAS ^{12D}	2 (PR/PD)	1b	400	Yes (PD) ^A	PD	14
2003	Colorectal	KRAS	4 (adj/SD/unk)	1b	400	No	WC(AE)	22
2004	Melanoma	BRAF ^{600E}	2 ^v (PD)	1b	400	Yes (PD)	AE	16
2005	Melanoma	BRAF ^{600E}	2 ^N (PR/PD)	1b	200	Yes (PD)	PD	54
2006	Melanoma	BRAF ^{600E}	3 ^v (SD/PD)	1b	200	No	PD/AE	26
2007	Melanoma	BRAF ^{600E}	1 ^y (PR)	1b	300	Yes (SD)	PD	168
2008	Melanoma	BRAF ^{600E}	1 ^N (adj)	1b	300	Yes (PR)	PD	252
2009	Melanoma	NRAS ^{61L}	5 (SD/PD)	1b	300	No	PD/AE	29
2010	Melanoma	BRAF ^{600K}	3 ^N (SD/PD)	1b	300	Yes (PD)	PD	55
2011	Melanoma	BRAF ^{600K}	O _N	1b	350	Yes (PR)	PD	149
2012	Melanoma	NRAS ^{61K}	2 (SD/PD)	1b	350	Yes (PD)	PD	28
2013	Colorectal	NRAS ^{61R}	6 (SD/PD)	1b	400	Yes (PD) ^A	WC (AE)	28
2014	Melanoma	BRAF ^{600E}	3 ^v (CR/PR/PD)	1b	400	Yes (PR)	PD	112
2015	Melanoma	NRAS ^{61R}	1 (PD)	1b	350	Yes (SD)	WC (other)	61
2016	Melanoma	NRAS ^{61K}	2 (adj/PD)	1b	400	Yes (PD) ^A	PD/AE	25

NSCLC, non-small cell lung cancer; BOR, best overall therapeutic response by RECIST v1.1 criteria; adj, adjuvant therapy; SD, stable disease; unk, unknown; PD, progressive disease; PR, partial response; CR, complete response; D/C, discontinuation; WC, consent withdrawal. Prior to this study enrollment, subjects received (Y) MAPK pathway inhibitor or did not (N). Antitumor response by radiographic measurements was assessed on day 28 after MK-8353 treatment.

however, his antitumor response could not be determined. Subject 2011 was 65 years old, with a history of *BRAF*^{V600K}-mutant melanoma without other coexisting mutations based on a 46-gene panel and no prior MAPK pathway inhibitor-based treatments. He was treated with 350 mg MK-8353 orally twice daily and developed a PR after 2 cycles. Ultimately, he was taken off the study after he was found to have brain metastases. He developed a PR when he was subsequently treated with concurrent D+T. Subject 2014 was 58 years old with a history of *BRAF*^{V600E}-mutant melanoma and concurrent *CDKN2A*^{R58*} nonsense mutation based on a 46-gene panel; he had previously developed complete response to single-agent vemurafenib and PR to concurrent administration of D+T. He was treated with 400 mg MK-8353 orally twice daily following progression on D+T and developed a PR following 2 cycles of MK-8353. However, he developed disease progression after 4 cycles and was taken off the study. Following progression on MK-8353, he developed an antitumor response to dacarbazine plus ipilimumab, no response to pembrolizumab, and, more recently, antitumor response to dabrafenib plus nivolumab.

Six patients (2010, 2005, 1002, 1003, 2012, and 2016) were found to have progressive disease after 2 cycles, and 3 patients (2002, 2013, and 2004) progressed after 1 cycle. Interestingly, subject 2010, who had metastatic melanoma bearing *BRAF*^{V600E} plus other mutations (e.g., *GNAS*^{S201H} and *RB1*^{E137D} mutations), and had not received any MAPK pathway inhibitor–based therapy prior to enrollment in this trial, did not respond to subsequent challenge with concurrent D+T. In contrast, the 3 *BRAF*^{V600}-mutant subjects (2005, 2008, and 2011) that were naive to MAPK pathway inhibitors, who progressed from MK-8353 treatment did respond to subsequent MAPK pathway inhibitor–based treatments. Of note, no other mutations other than *BRAF*^{V600} were identified for subjects 2008 and 2011, whereas subject 2005 was found to have *CDK4*^{R24L} missense mutation based on the TruSight Tumor 26-gene panel (Illumina). No other mutations other than the *BRAF*^{V600} mutation were screened in pretreatment tumors from subjects 2004 and 2007.



Table 3. Summary of AE possibly or probably related to MK-8353 (CTCAE v4.03) in the MK-8353-001 study

AE	No. of patients having AE (%)					
	All grades	Grade 3	Grade 4			
Diarrhea	11 (44)	4 (16)	0			
Fatigue	9 (36)	1 (4)	0			
Maculopapular rash	7 (28)	2 (8)	0			
Nausea	7 (28)	1 (4)	0			
Vomiting	6 (24)	1 (4)	0			
Bilirubin serum, increased	5 (20)	4 (16)	0			
Creatinine serum, increased	4 (16)	0	0			
Abdominal pain	4 (16)	0	0			
Anemia	3 (12)	1 (4)	0			
Pruritus	3 (12)	1 (4)	0			
Decreased appetite	3 (12)	0	0			
Chills	3 (12)	0	0			
BUN serum, increased	3 (12)	0	0			
Dysgeusia	3 (12)	0	0			
Impaired gastric emptying	3 (12)	0	0			
Hyponatremia	2 (8)	0	0			
Oropharyngeal pain	2 (8)	0	0			
Pyrexia	2 (8)	0	0			
Vision blurred	2 (8)	0	0			
Dermatitis acneiform	2 (8)	0	0			
Dyspnea	2 (8)	0	0			
Muscular weakness	2 (8)	0	0			
Dehydration	2 (8)	0	0			
Headache	2 (8)	0	0			
Pain	1 (4)	0	0			
Increased serum AST	2 (8)	0	0			
Muscular weakness	2 (8)	0	0			
Increased serum Alk Phos	1 (4)	0	0			
Increase CPK	1 (4)	0	1(4)			
Increased γGT	1 (4)	0	0			
Dyspepsia	1 (4)	0	0			
Hypoalbuminemia	1 (4)	0	0			
Hypocalcemia	1 (4)	0	0			
 Hypomagnesemia	1 (4)	0	0			
Hypermagnesemia	1 (4)	0	0			
Hypokalemia	1 (4)	0	0			
Hyponatremia	1 (4)	0	0			
Acute renal failure	1 (4)	1 (4)	0			
Amnesia	1 (4)	0	0			
Depressed mood	1 (4)	0	0			
Disorientation	1 (4)	0	0			
Incontinence	1 (4)	0	0			
Cough	1 (4)	0	0			
Urticaria	1 (4)	0	0			
No MK-8353-related deaths (grade 5) occurred. AST, aspartate aminotransferase; Alk Phos,						

No MK-8353-related deaths (grade 5) occurred. AST, aspartate aminotransferase; Alk Phos, alkaline phosphatase; CPK, creatine phosphokinase; YGT, gamma-glutamyl transferase.

Among the 10 nonevaluable patients, subject 2009, who was diagnosed with NRASQ61L-mutant cutaneous melanoma from the lower extremity, was noteworthy. Prior to study enrollment, she received multiple systemic treatments, including ipilimumab and pembrolizumab, without any significant antitumor response. She was enrolled into the MK-8353 trial based on in vitro drug sensitivity data from 2 melanoma cell lines that were established from her own tumors, M435-1 and M428, which showed high sensitivity to SCH772984 (Supplemental Figure 2A). Despite the initial clinical response in bulky subcutaneous tumors (Supplemental Figure 2B), she unfortunately developed symptomatic metastases to the brain, from which she subsequently died.

Discussion

We have developed an orally bioavailable, dual mechanism, direct ERK1/2 inhibitor, MK-8353, which appears to have comparable potency and selectivity with the preclinical tool compound, SCH772984 (16). Both compounds have higher potency in cellbased assays than ulixertinib (BVD-523) and CC-90003, which are currently in clinical development (9, 11). Comparison of the doses required for substantial biological activity (i.e., suppression of pERK in tissues), tumor growth inhibition or even regression, and toxicology suggest that MK-8353 administered orally twice daily may have a narrow therapeutic index, given that the drug target, ERK, is important for physiologic signaling in normal tissues (19-21) and the drug, MK-8353, binds to and inhibits both activated and nonactivated ERK.

Following an earlier study in healthy volunteers to establish preliminary toxicity, pharmacokinetic, and pharmacodynamic data (P07652), we performed a phase I study of MK-8353 in patients with advanced cancers (MK-8353-001). Pharmacokinetic analysis of the P07652 study suggested that doses between 300 and 400 mg corresponded to drug exposure (AUC_{0-24hours}) that resulted in both a biologic and antitumor effect (e.g., tumor growth inhi-

bition or regression) in preclinical models. However, in the P07652 study, we did not observe any changes in pERK expression in either the normal skin tissue or peripheral blood mononuclear cells in the respective tissue between subjects who received either placebo or a single dose of MK-8353. We have attributed the lack of significant pharmacodynamic changes to the great variability in pERK expression among normal human subjects in different normal tissues.



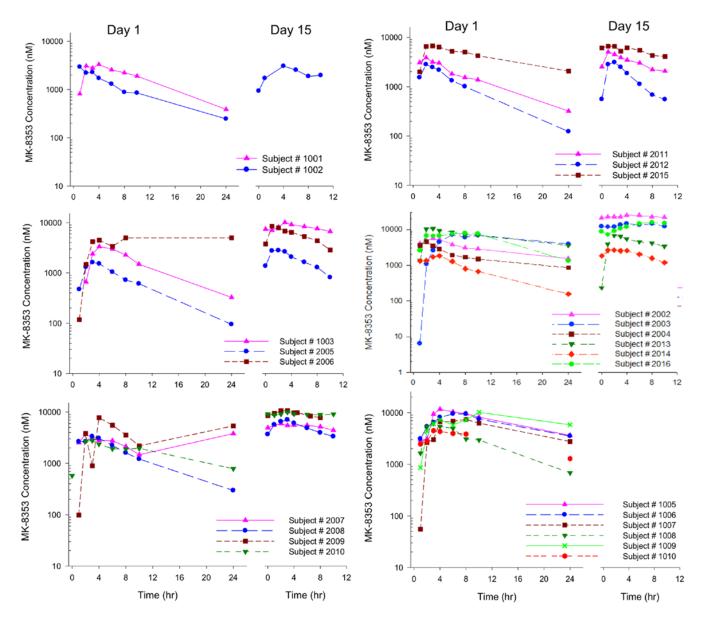


Figure 4. Mean plasma concentration-time profiles (semi-log scale) of MK-8353 in the MK-8353-001 study.

In the MK-8353-001 study, we found that MK-8353 was well tolerated at doses up to 400 mg twice daily and exhibited a side effect profile that is similar to that of MEK inhibitors (22), with diarrhea, nausea, vomiting, and rash being the most frequently seen. Although we did not establish the RP2D for MK-8353 in this study, pharmacokinetic analysis suggested adequate dose-dependent exposure across all dose cohorts according to the optimal AUC that was considered biologically effective in mouse studies. Despite the fact that 2 DLTs occurred in each of the 400-mg and 800-mg dose cohorts, no patient in the 800-mg dose cohort remained on treatment longer than 12 days; in addition, all patients in the 800-mg dose cohort stopped due to AEs (Table 2). Our inability to escalate above the 400-mg twice daily dose due to toxicity manifested as DLT or as compliance or early withdrawal from the study, which may have to do with the narrow therapeutic index of MK-8353. To compensate for the inherent high potency of this dual mechanism inhibitor that may account for patient intolerance at doses higher than 400 mg, future studies may consider continuous once daily or intermittent twice daily schedules. This alternative approach may balance higher, though more temporal suppression of pERK at the tumor tissue in favor of sparing sustained suppression of ERK signaling in normal tissues.



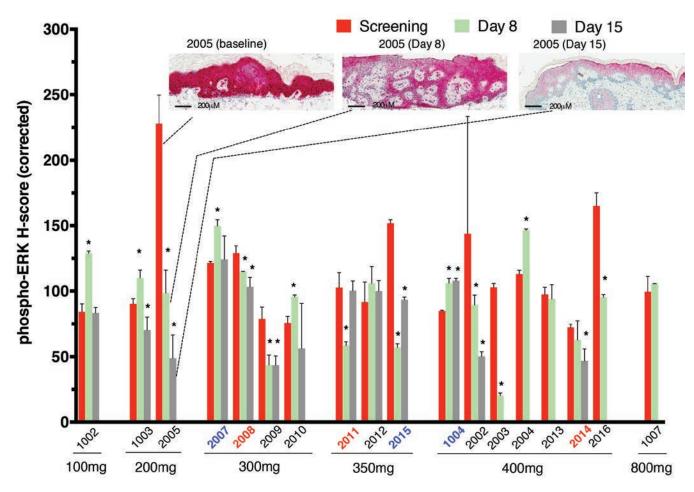


Figure 5. Effect of MK-8353 on pERK protein levels in normal epidermis in the MK-8353-001 study. Patients with the baseline (red bar) and at least 1 on-treatment biopsy (day 8, green bar; day 15, gray bar) are shown. Corrective pERK levels (H-score mean ± SD) are shown. Asterisks show significant differences compared with baseline (*P < 0.05, 1-way ANOVA). Stable disease (SD) is shown in blue font; partial response (PR) is shown in red font. All other patients were nonevaluable for the efficacy endpoint. Scale bar: 200 μm.

Reversible unconjugated hyperbilirubinemia was seen in 3 patients in the MK-8353-001 study who received MK-8353 ≥400 mg twice daily. None of the 3 subjects had history of or suspected Gilbert's syndrome, which was an exclusion criterion for our study, given the possibility of MK-8353-mediated inhibition of the UGT1A1 enzyme. Development of asymptomatic hyperbilirubinemia was not eventually determined to be a DLT and is not expected to be an issue in further clinical development of MK-8353, although close monitoring of bilirubin levels in ongoing clinical trials using MK-8353 is recommended.

Administration of MK-8353 led to antitumor responses in 3 of 8 patients with *BRAF*^{V600}-mutant metastatic melanoma, regardless of prior therapy with MAPK pathway inhibitor. The fact that antitumor responses were seen in doses greater or equal to 300 mg twice daily is in line with the dose-proportional increased exposure (AUC_{0-12hours}) to MK-8353. Although the number of patients with *BRAF*^{V600}-mutant melanoma that were treated with MK-8353 as part of this study is small, the response rate was similar to what was seen in early, but much larger than the current, clinical trials testing single-agent MEK inhibitors (23). In addition, there is early evidence of temporary disease stability in patients with *NRAS*^{Q61}-mutant melanoma, although no antitumor responses were observed in any MK-8353-001 study patients bearing *NRAS* or *KRAS*-mutant cancers.

Similar to the healthy volunteer P07652 study, the advanced solid tumor MK-8353-001 study failed to show consistently significant dose-dependent suppression of pERK expression in serial normal skin biopsies in any of the doses explored. Variability in expression levels of pERK in normal skin biopsies by immuno-histochemistry has been previously reported as similar or slightly lower compared with the corresponding pERK expression levels seen in melanoma biopsies (4). Normal tissues have been previously successfully



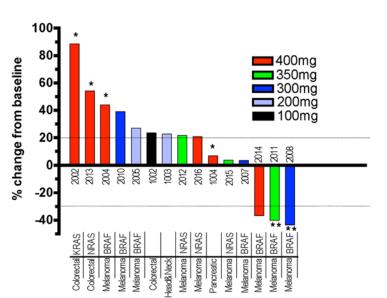


Figure 6. Investigators' assessment of objective tumor responses in evaluable patients (n = 15) in the MK-8353-001 study. With the exception of subjects 1004, 2002, 2013, and 2004, whose tumor assessments were performed prior to completion of 2 cycles (*), as well as subjects 2011 and 2008, whose best antitumor response occurred after 4 cycles (**), tumor assessments were completed following completion of 2 cycles. Three patients achieved partial response (PR).

used as pharmacodynamic markers for other targeted therapies (24), including other ERKi currently in clinical development (9). In the early clinical studies of single-agent vemurafenib in patients with BRAF^{v600}-mutated metastatic melanoma, greater than 80% inhibition of cytoplasmic pERK was seen in patients with tumor regressions (25). Inhibition of pERK by more than 65%, the presumed cut point for biologically substantial suppression of pERK based on preclinical data in tumor-bearing mice, occurred in on-treatment tissue specimens in only the 3 subjects in the 400-mg and 200-mg dose cohorts. Nevertheless, no antitumor responses were observed in those 3 patients who indeed had substantial suppression of pERK. The comparison of the pERK immunohistochemical data between normal skin biopsies and tumor tissue in mouse studies suggests that the duration of pERK suppression following a single dose of MK-8353 is much shorter in normal skin compared with tumor tissue (Figure 1, C and D). This paired with the fact that the t_{max} for MK-8353 is longer in humans compared with mice may explain why, despite our best efforts to obtain normal skin biopsies 5-7 hours after the MK-8353 dose, we have not been able to consistently capture true pERK suppression that may have been truly achieved in normal skin biopsies before or after the actual timing of the biopsy. Since tumor tissues express more pERK than normal tissues and the duration of pERK suppression is longer in tumor tissues, analysis of pERK expression in tumor tissues is obviously expected to correlate more closely with antitumor response to MK-8353. Lack of collection of tumor tissue at baseline and at disease progression and antitumor response are all inherent weaknesses of the MK-8353-001 study. Alternatively, the lack of pERK suppression in normal skin biopsies may not be as clinically relevant, as these normal tissues may not consistently express activated-phosphorylated ERK under physiologic conditions in humans. Another important factor to consider is related with the dual mechanism of action of MK-8353; namely, any assay that solely relies on pERK expression levels does not take into account the fact that MK-8353 can also bind unphosphorylated ERK1/2, which subsequently prevents ERK1/2 phosphorylation by upstream kinases.

Molecular testing for all 3 partial responders using a 46-gene panel suggests a lack of clinically important coexisting mutations other than *BRAF*^{V600} that have previously shown to impair response to MAPK pathway inhibitor–based treatments (26). While coexisting mutations in clinically important genes in *BRAF*^{V600E}-mutant melanoma subjects 2005 and 2010 could potentially account for the lack of response to any MAPK pathway inhibitor, including MK-8353, we have no additional molecular information about coexisting mutations in the other 2 nonresponding *BRAF*^{V600}-mutant subjects, 2004 and 2007. It is likely that additional genetic, or even epigenetic, changes may account for the lack of response to MK-8353, which were beyond the scope of the MK-8353-001 trial. For example, a detailed analysis of pERK staining over a large number of melanoma samples by immunohistochemistry showed that pERK abundance was largely heterogeneous, not only among different melanoma tumor tissues, but also within the same melanoma specimen, and did not correlate with *BRAF* and *NRAS* mutation status (27). Among patients with *BRAF*^{V600}-mutant melanoma, baseline expression of pERK was also variable in melanoma biopsies (4). Similar results were seen in colorectal adenocarcinomas, although that study did not provide *RAS* mutational status data (28). Other possibilities are



that the ERK-dependent negative feedback on the RAF/MEK/ERK pathway is lost and/or other non-RAF/MEK/ERK signaling pathways may alternatively contribute to oncogenic growth. In line with this, a preclinical study of the ERKi SCH772984 in *KRAS*-mutant pancreatic cancer suggests that both such processes may be clinically important (29). Therefore, future directions in the clinical development of ERKi might include optimizing drug formulation for better bioavailability and longer half-life, which may improve drug exposure to tumors and result in significant inhibition of pERK. However, even if such significant inhibition of pERK in tissues is achieved, combination strategies with other inhibitors of the RAF or PI3K pathway may be clinically more effective, albeit more toxic. Our preclinical data on the superiority of the combination of trametinib with SCH772984 in melanoma cell lines in vitro as well as a recent in vitro report on the synergy between trametinib and a peptide that blocks nuclear translocation of ERK1/2 (30) support possible future evaluations of concurrent MEK and ERK inhibition for several *BRAF*^{v600} wild-type and *BRAF*^{v600}-mutant melanomas. Whether ERK inhibition may be more effective than MEK inhibition when each of these targets are targeted in combination with BRAF inhibitors in melanoma is currently unknown.

In conclusion, we tested the first dual-mechanism direct ERKi to our knowledge, MK-8353, in patients with advanced solid tumors, and we found that it was well tolerated at doses ≤400 mg orally twice per day. PK/PD data suggest that the dosing of MK-8353 between 400 mg and 800 mg orally twice per day, or, even better, once daily, may be further explored to balance exposure, biological activity, and toxicity. The AE profile (e.g., diarrhea, fatigue, skin rash) and the antitumor activity (12% response rate) were comparable to those seen in a larger phase I trial of ulixertinib in patients with advanced solid tumors (9). Given the promising clinical activity and the lack of SAEs, further clinical development of MK-8353 in combination with immunotherapies and for select cancer types is currently under consideration (NCT02972034). This phase I clinical study of MK-8353 provided valuable information regarding clinical activity, toxicity profile, and tolerability of MK-8353 administered twice daily; lessons about optimal pharmacodynamic endpoints; and solid tumor cancer types in which single-agent ERKi may have clinical benefit. It is likely that more ERKi will be clinically developed in the future, perhaps in combination with other targeted therapies of the MAPK or PI3K pathways for treatment of ERK-dependent cancers or even immunotherapies.

Methods

Cell lines. All cell lines used for the preclinical development of MK-8353 (Supplemental Tables 1 and 2) were purchased from ATCC. Exceptions were the UACC-62, M14, LOX, OVCAR-5, IGGROV-1, and A2780 cell lines, which were provided by the National Cancer Institute (NCI) Tumor Repository (Frederick, Maryland, USA), and the 8505C, IPC-298, SK-MEL-30, and BHT-101 cell lines, which were provided by the Leibniz-Institut DSMZ (Deutsche Sammlung von Mikroorganismen und ZellKulturen GmbH; Braunscheweig, Germany). Cell lines were maintained in culture medium as per the manufacturer's recommendations. The identities of all cell lines were confirmed using short-tandem repeat DNA analysis.

M428 and M435 are human melanoma cell lines established from baseline biopsies from MK-8353-001 study patient 2009, as previously described (31). M428 and M435 cells were cultured in RPMI 1640 with L-glutamine (Mediatech Inc.), 10% fetal bovine serum (Omega Scientific), and 1% penicillin, streptomycin, and fungizone (Omega Scientific) and were screened for mycoplasma contamination. Cultures were incubated in a water-saturated incubator at 37°C with 5% CO₂.

Drug compounds used in in vitro cell viability assays. For in vitro experiments, MK-8353 was synthesized by Merck Sharp & Dohme Corp. SCH772984 was provided to the University of California Los Angeles (AR) through a material transfer agreement with Merck Sharp & Dohme Corp. Trametinib was purchased from Selleck Chemicals. Stocks and solutions for all compounds were prepared in DMSO solvent to a final concentration of 10 mM. Tumor cell growth assays using MK-8353 were performed in 96-well format by plating 4,000 cells per well. On the next day cells were treated with DMSO or a 9-point IC_{50} dilution (0.001–10 μ M) to a final concentration of 1% DMSO. Six replicates were assigned to each dose, and cells were treated with MK-8353 for 3 days. Cell viability was assayed using the CellTiterGlo Viability Assay luminescent kit (Promega), as per the manufacturer's instructions (17). All cell growth assays were performed 3 time for each cell line. Half-maximal effective concentration (EC₅₀) values were calculated using Microsoft Excel.

M428 and M435 cells were treated in duplicate with 0–10 M MK-8353 and 0–1 μ M trametinib alone or in combination. A constant amount of DMSO was kept for all conditions. After incubation for 120 hours, the cell viability was determined using the CellTiterGlo Assay. Each experiment was repeated independently twice, always in duplicate.



Animal studies. Female athymic nude mice and C57/SCID mice, both 4–6 weeks of age, were purchased from Charles River Laboratories and were maintained in a viral antibody-free barrier facility. All animals received food and water ad libitum. Regardless of the cancer cell line used, animals were inoculated subcutaneously with 5×10^6 cancer cells in 50% Matrigel. Resultant tumors were grown to approximately 100 mm^3 in size before initiating treatment. Mice were then randomized to treatment groups (10 mice/group) and treated by oral gavage with either MK-8353 (30–60 mpk twice daily) or 20% HPBCD vehicle twice daily. Tumor length (L), width (W), and height (H) were measured using a caliper twice weekly on each mouse. Animal body weight was measured twice weekly for the duration of treatment.

Antibodies and protein detection. For immunoblot analysis MK-8353–treated cells were lysed in radioimmunoprecipitation assay buffer with protease inhibitor cocktail (Roche Diagnostics GmbH) and cleared by centrifugation. Protein concentration was determined using bicinchoninic acid reagent (Pierce Chemical). Protein samples were separated by 10% SDS-PAGE and transferred onto a nitrocellulose membrane. Blots were incubated with primary antibodies against phosphorylated ERK1/2 T185/Y187 (rabbit polyclonal, BioSource/Thermo Fisher Scientific, catalog no. 44-680G, dilution 1:1,000), ERK1/2 (rabbit polyclonal, BioSource/Thermo Fisher Scientific, catalog no. 44-654G, dilution 1:1,000), phosphorylated p90 RSK T359/S363 (rabbit monoclonal, clone E238, Cell Signaling Technology, catalog no. 9344, dilution 1:500), total RSK (mouse monoclonal, clone 78/RSK, BD Transduction Laboratories, catalog no. 610226, dilution 1:5,000), and cleaved poly (ADP-ribose) polymerase p85 fragment (rabbit polyclonal, Promega, catalog no. G7341, dilution 1:2,000). Following incubation with primary and secondary antibodies, band intensity was detected by chemiluminescence using enhanced chemiluminescence detection reagents (Amersham). Densitometric quantification of each band was determined using Quantity One software (PDI Inc.) and was normalized in comparison with the relevant expression of β-actin in the reprobed blot.

For immunohistochemical detection of phospho-ERK, 5-µM-thick formalin-fixed paraffin-embedded tissue sections (tumor tissue and normal skin) were deparaffinized and rehydrated by serial passages through Clear-Rite (Thermo Fisher Scientific) (5 minutes, 3 times), 100% Flex (5 minutes, twice), 95% Flex (Thermo Fisher Scientific) (5 minutes, once), 80% Flex (5 minutes, once), 50% Flex (3 minutes, once), and dH₂0 (5 minutes) solutions. Antigen retrieval was performed in decloaking chamber at 98°C for 35 minutes using the DAKO Target Retrieval Solution (Dako/Agilent Pathology Solutions, catalog no. S1700). Tissue slides were then cooled for 5 minutes, rinsed with dH₂O for 20 minutes, and subsequently incubated for 5 minutes with 0.03% H₂O₂. Tissue slides were then incubated with 200 µl diluted (1:100) primary antibody against phospho-ERK (T202/Y204) (rabbit monoclonal, clone D13.14.4E, Cell Signaling, catalog no. 4370) in DAKO antibody diluent (Dako/ Agilent Pathology Solutions, catalog no. S0809) for 30 minutes followed by incubation with peroxidase-labeled polymer goat anti-rabbit for 30 minutes. HRP-labeled polymer anti-rabbit (DAKO EnVIsion+ System, Dako/ Agilent Pathology Solutions, catalog no. K4003) solution (200 µl per tissue slide) was then added for 5 minutes. Slides were then rinsed and counterstained with hematoxylin OS (Vector Labs, catalog no. H3404) for 1 minute. Tissue slides were finally rinsed with dH,O, dehydrated by serial passages through 70% Flex (20 seconds, once), 95% Flex (20 seconds, once), 100% Flex (20 seconds, twice), and Clear-Rite (1 minute, twice) and coverslipped. Five digital images were obtained from each immunohistochemically stained tissue section (tumor tissue and normal skin; 3 mice per each time point) using Zeiss Axioskop light microscope equipment. Signal was quantified using AxioVision software (Carl Zeiss Microscopy LLC).

Patients (MK-8353-001 study). Patients were recruited either via self-referral, referral by community medical oncologists, or as established patients across 9 academic institutions on 3 continents. Eligibility required a histologic diagnosis of advanced solid tumors, adequate organ function (hemoglobin ≥9.0 gm/l, absolute neutrophil count ≥1,500/µl, platelet count ≥100,000/l, serum cystatin C level ≤1.5× upper limits of normal [ULN], aspartate aminotransferase and alanine aminotransferase ≤5× ULN if liver metastases were present or ≤3× ULN if absent, and bilirubin 2× ULN, unless Gilbert's syndrome was suspected), ECOG performance status of 0 or 1, and the ability to swallow and absorb oral medications. Subjects with unstable or progressing brain metastases, gastrointestinal disorder that may alter drug absorption, QTcF interval >450 ms, and history of retinal vein occlusion, central serous retinopathy, or glaucoma or female subjects who were breastfeeding, pregnant, or intending to become pregnant were excluded. Investigators from each study site were required to provide subject data to the study sponsor, Merck Sharp & Dohme Corp., according to Sponsor's instructions and compliant with Good Clinical Practice guidelines.

Study design (MK-8353-001 study). The trial (NCT01358331) was conducted in 2 parts. Part 1a enrolled subjects with any incurable advanced solid tumor. In part 1a, the MK-8353 dose was increased, and



MTD range was preliminarily determined based on safety, tolerability, and pharmacokinetic/pharmacodynamics evaluations. Part 1b enrolled patients with melanoma and colon cancer bearing oncogenic hotspot RAS and/or BRAF mutations. Patients in part 1b must have had at least one measurable lesion as defined by RECIST v1.1 criteria and should have not received greater than 3 prior systemic treatments, excluding MAPK pathway inhibitor. Assessment of $BRAF^{V600}$ or RAS mutation status was performed in a Clinical Laboratory Improvement Amendments–certified laboratory or by centralized testing. The previously defined MTD range from part 1a would be further refined and defined in part 1b for subsequent phase II studies. Patients without any prior systemic treatment for their cancer were allowed to participate at any dose cohort.

Study drug administration (07652 and MK-8353-001 studies). MK-8353 was supplied in capsules by Merck Sharp & Dohme Corp., a subsidiary of Merck & Co. Inc., and was administered orally on an empty stomach 1 hour before or approximately 2 hours after a meal. In the 07652 study, MK-8353 was administered as a single dose to healthy adult males in the outpatient clinic, whereas in the MK-8353-001 study, MK-8353 was administered twice daily at home or in the outpatient clinic continuously in consecutive 28-day cycles until disease progression or unacceptable toxicity. In part 1a of the MK-8353-001 study, dose escalation followed an accelerated titration design. As long as no grade 2 toxicities occurred, the next dose level would be double that of the previous dose level. If a single ≥grade 2 toxicity (non-dose-limiting) occurred in any given dose cohort, the dose level would expand to 3 subjects. If no additional ≥grade 2 toxicity (non-dose-limiting) occurred, dose escalation would continue by doubling of the previous dose level. If one or more additional ≥grade 2 toxicity(-ies) (in 2 or 3 of the 3 patients) occurred, dose escalation would continue to increase the dose by approximately 67%, 50%, 33%, and 25%, respectively. In part 1b of the MK-8353-001 study, dose level started at the preliminary MTD with potential for dose escalation, depending on the number of subjects evaluable for analysis for DLT. DLT was counted toward RPTD decision in dose escalation.

Safety and response assessment (MK-8353-001 study). The prespecified safety endpoint was the DLT rate occurring during the first cycle. For determination of AE severity, the NCI's Common Terminology Criteria for Adverse Event (NCI-CTCAE) v4.03 was used. The preliminary MTD was defined as the highest dose level for which DLT is <33%. DLT was defined as any ≥grade 3 hematologic toxicity lasting ≥1 week; any ≥grade 3 nonhematologic toxicity not clearly attributable to other causes; grade 3 nausea, vomiting, and anorexia, if maximal supportive care had been delivered; any grade 4 nonhematological laboratory value not related to underlying disease; >20% missed doses due to toxicity in the first cycle. Response rate was a secondary endpoint for this trial and was defined by RECIST v1.1. Imaging was performed every other cycle, unless the investigator felt clinical need for earlier assessments.

Pharmacokinetic assessments (tumor growth xenograft, 07652, and MK-8353-001 studies). For pharmacokinetic analysis, plasma was collected on cycle 1, day 1 (0, 1, 3, 4, 6, 8, 10, and 24 hours; 07652 and MK-8353-001 studies), and day 15 (immediately before the first dose on that day and 1, 2, 3, 4, 6, 8, and 10 hours after the first dose; MK-8353-001 study) of MK-8353 treatment. Plasma MK-8353 concentrations were used to estimate the following pharmacokinetic parameters of MK-8353 treatment: maximum observed plasma concentration (C_{max}), time-to-maximum observed plasma concentration (t_{max}), area-under-the-plasma concentration-time curve, apparent total body clearance (CL/F), terminal phase half-life ($t_{1/2}$), accumulation ratio, and apparent volume of distribution (Vd/F).

The analytical method for the analysis of MK-8353 was based on an automated 96-well format liquid-liquid extraction of drug from human plasma. Briefly, drug and internal standard were extracted from plasma with methyl tert-butyl ether. Samples were mixed-rotated and centrifuged in a 96-well plate. The organic supernatant layer was removed, evaporated under nitrogen, and reconstituted in 50:50 acetonitrile/water. The plate was sealed, and 2–10 μ l of the reconstituted sample was injected onto a high-performance liquid chromatography tandem mass spectrometry system for analysis. Drug and internal standard were chromatographed using reverse-phase liquid chromatography and detected with tandem mass spectrometry. The multiple reaction monitoring transitions were monitored with mass-to-charge ratio (m/z) 692.2 \rightarrow 424.1 for MK-8353. The lower limit of quantitation for this method was 2 ng/ml.

Pharmacodynamic assessments in clinical studies (07652 and MK-8353-001). The degree of pERK suppression, as detected by immunohistochemistry in normal 4-mm skin punch biopsies, was used as a surrogate biomarker of ERK activation at the following time points: at baseline, on cycle 1, and on days 8 and 15. Immunohistochemistry was performed on a Ventana Discovery XT immunostainer



(Ventana Medical Systems Inc.) using the DISCOVERY ChromoMap RED kit Alkaline Phosphatase (AP) staining kit (Ventana). Slides were deparaffinized, underwent heat-induced epitope retrieval using the RiboCC buffer (Ventana), and were incubated in 7% goat serum (MilliporeSigma) for 20 minutes to inhibit nonspecific staining. Slides were subsequently incubated with rabbit monoclonal antibody against phospho-ERK (clone D13.14.4E, Cell Signaling; 1:600 dilution) or rabbit monoclonal antibody IgG isotype control (DA1E, Cell Signaling) for 60 minutes. Antibody detection was performed using the DISCOVERY UltraMap anti-Rabbit AP detection system (Ventana). Tissues were counterstained with Hematoxylin II (Ventana) and developed using bluing reagent (Ventana). Stained slides were transferred to a Leica Multistainer/Coverslipper (ST5020/CV5030, Leica Microsystems Inc.), dehydrated in graded ethanol and xylene, and coverslipped.

Immunohistochemically stained slides for pERK1/2 from both xenograft and clinical studies were scanned into Aperio ScanScope XT. Digital images were analyzed using Aperio ImageScope software (Leica Microsystems Inc.). Analysis algorithm was developed using a panel of human skin tissues. Staining was evaluated using computer imaging analysis (color deconvolution algorithm) in which electronic gates were manually set to select for epidermis and exclude dermal and subdermal regions and structures. From each skin biopsy, two slides were used to stain pERK and one slide was used to stain for IgG negative control. A single lot of the pERK antibody was used for staining.

Statistics. Descriptive analysis of preclinical data was performed using GraphPad Prism 5.0 (GraphPad Software Inc.). EC_{50} were calculated using Microsoft Excel. Tumor volumes in animal experiments were calculated using the formula $(L \times W \times H)/2$. Tumor volume data are expressed as mean \pm SEM. Statistically significant differences in tumor growth curves and immunohistochemical expression of pERK among the different treatment groups in xenograft studies were determined by the Student's t test. An unadjusted t value of 0.05 was considered as statistically significant.

Descriptive analysis of clinical data was performed using the Statistical Analysis System (SAS) software suite release 9.1 (SAS Institute). Analysis of pharmacokinetic data was performed using Phoenix WinNonLin (Certara). Graphs were developed using Prism 6.0 (GraphPad Software Inc.). Subjects who received ≥ 1 dose of MK-8353 and had ≥ 1 efficacy assessment after baseline were included in the efficacy endpoint analysis, whereas subjects who received ≥ 1 dose of MK-8353 and completed cycle 1 or discontinued due to toxicity were evaluable for DLT. The DLT rates along with the 80% Bayesian credible intervals for the DLT rate in cycle 1 were provided. Plasma MK-8353 concentrations and pharmacokinetic parameters were listed and summarized by dose using descriptive statistics. The pharmacokinetic parameters AUC and C_{max} were log-transformed and analyzed based on a linear-mixed effects model containing a fixed effect for treatment and a random effect for subject.

For pharmacodynamic analysis of pERK inhibition by immunohistochemistry in normal human skin biopsies, the color deconvolution algorithm was used to assess staining on a "per pixel" basis and is reported as a score calculated in a manner similar to an H-score:

Score = $[(\% 1 + pixels) \times 1] + [(\% 2 + pixels) \times 2] + [(\% 3 + pixels) \times 3]$

Corrected score was defined as follows: (pERK score) – (IgG-negative control score). Percentage inhibition was then calculated using the corrected scores.

Study approval. The MK-8353-001 study protocol was approved by the IRBs of the University of North Carolina at Chapel Hill, Massachusetts General Hospital Cancer Center, The University of Texas MD Anderson Cancer Center, the Translational Genomics Research Institute, the Virginia G. Piper Cancer Center, the Roswell Park Cancer Institute, The University of Auckland and Auckland City Hospital, the Chaim Sheba Medical Center, the University of Pittsburgh School of Medicine, Merck Sharp & Dohme Corp., and the Jonsson Comprehensive Cancer Center at UCLA. M428 and M435 human melanoma cell lines were established under the University of California Los Angeles IRB (02–08-067)). Written informed consent was received from participants prior to inclusion in the study. Written informed consent was also provided for patient pictures appearing in the manuscript. Animal procedures were performed according to the NIH Guide For The Care And Use of Laboratory Animals (National Academies Press, 2011).

Author contributions

TSR, DZ, NRM, AAS, EHR, JS, BJL, PD, DC, PK, WRB, YD, AC, and GWS were involved in preclinical development of MK-8353. JR, AAS, PC, and EHR were involved in the healthy volunteer study plan and conduct. JR, PC, AAS, EHR, and KTF were involved in the MK-8353-001 study planning and conduct.



SJM, RJS, WJH, RKR, AAA, PCF, RSF, HAT, AR, and KTF were involved in patient enrolling, care, and assessment, as well as reporting of antitumor response and toxicity to MK-8353. TSR, DZ, NRM, AAS, BHM, LR, and AR were involved in the correlative analyses. SJM, RJS, WJH, JR, NRM, AAS, BJL, AR, and KTF were involved in manuscript writing, editing, and submission. SJM, RJS, BJL, AAS, EHR, AR, and KTF are responsible for the overall content.

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