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An oral SARS-CoV-2 M^{pro} inhibitor clinical candidate for the treatment of COVID-19

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The worldwide outbreak of coronavirus disease 2019 (COVID-19) caused by severe acute respiratory syndrome coronavirus 2 (SARS-CoV-2) has become a global pandemic. Alongside vaccines, antiviral therapeutics are an important part of the healthcare response to counter the ongoing threat presented by COVID-19. Here, we report the discovery and characterization of PF-07321332, an orally bioavailable SARS-CoV-2 main protease inhibitor with in vitro pan-human coronavirus antiviral activity and excellent off-target selectivity and in vivo safety profiles. PF-07321332 has demonstrated oral activity in a mouse-adapted SARS-CoV-2 model and has achieved oral plasma concentrations exceeding the in vitro antiviral cell potency in a phase I clinical trial in healthy human participants.

Human coronavirus infections are common, with at least four examples (229E, NL63, OC43, HKU1) now considered endemic (*1*). However, the emergence within the last 20 years of SARS-CoV-1, Middle East Respiratory Syndrome (MERS-CoV) and SARS-CoV-2 as novel human coronaviruses has signaled the significant threat potential of this viral class. The catastrophic SARS-CoV-2 outbreak of 2019 has resulted in 235 million confirmed cases of COVID-19 causing over 4.8 million deaths globally as of October 2021. SARS-CoV-2 is a highly infectious, ribonucleic acid (RNA) beta coronavirus that can cause life-threatening viral pneumonia in the most serious cases. While effective COVID-19 vaccines have been developed within unprecedented timelines, a significant number of people are either unable, due to pre-existing medical conditions, or unwilling to be vaccinated, and global access challenges remain. Limited therapeutic options are available to those who are infected. Oral SARS-CoV-2 specific therapeutics are urgently needed to prevent more severe disease, hospitalization, and death. Treatment may also reduce the period of infectivity. Repurposing of approved drugs in the search for small molecule antiviral agents that target

SARS-CoV-2 has thus far been minimally effective (*2, 3*). Viral RNA-dependent RNA polymerase inhibitors such as molnupiravir and AT-527 are currently undergoing clinical trials for the treatment of COVID-19 (*4, 5*).

The SARS-CoV-2 genome encodes two polyproteins (pp1a and pp1ab) and four structural proteins (*6, 7*). The polyproteins are cleaved by the critical SARS-CoV-2 main protease (M^{pro}, also referred to as 3CL protease) at eleven different sites to yield shorter, non-structural proteins vital to viral replication (*8, 9*). The coronavirus M^{pro} is a three-domain cysteine protease, which features a Cys145-His41 catalytic dyad located in the cleft between domains I and II. Several common features are shared among M^{pro} substrates, including the presence of a Gln residue at P1 (using Schechter-Berger nomenclature (*10*)). No known human cysteine protease cleaves after Gln, thus offering potential selectivity for this viral target over the human proteome (*11–13*). Viral proteases are tractable targets for small molecule oral therapies in the treatment of human immunodeficiency (HIV) and hepatitis C (HCV) viruses (*14, 15*). Moreover, a recent report has demonstrated oral activity of an M^{pro} inhibitor in a transgenic

mouse model of SARS-CoV-2 infection (16). Given the pivotal role of SARS-CoV-2 M^{pro} in viral replication, its potential for mechanistic safety, and expected lack of spike protein variant resistance challenges, SARS-CoV-2 M^{pro} inhibition represents an attractive small molecule approach for an oral antiviral therapy to treat COVID-19.

An effort to identify inhibitors of the SARS-CoV-1 M^{pro} in response to the 2002 SARS outbreak led to the identification of PF-00835231 (**1**, Fig. 1) as a potent inhibitor of recombinant SARS-CoV-1 M^{pro} in a fluorescence resonance energy transfer (FRET)-based substrate cleavage assay (17). PF-00835231 also demonstrated potent inhibition (inhibition constant (K_i) = 0.271 nM) of recombinant SARS-CoV-2 M^{pro}, which is expected given that the SARS-CoV-1 and -CoV-2 M^{pro} share 100% sequence homology across their respective substrate binding sites (18). Antiviral activity against SARS-CoV-2 was also observed with PF-00835231 (half-maximal effective concentration (EC₅₀) of 231 nM) by monitoring cytopathic effect (CPE) in epithelial Vero E6 cells. As Vero E6 cells express high levels of the efflux transporter P-glycoprotein, for which **1** is a substrate (17, 19), antiviral assays in this cell line were conducted in the presence of the P-glycoprotein efflux inhibitor CP-100356 (20). TMPRSS2 expression was not detected in the Vero E6 cells by RTqPCR. The phosphate prodrug form (PF-07304814) of PF-00835231 is currently under investigation as an intravenous treatment option for COVID-19 in hospitalized patients (20).

In order to improve upon the low passive absorptive permeability (P_{app} < 0.207 × 10⁻⁶ cm/s) (21) and poor oral absorption of **1** in animals, we aimed to remove the hydrogen bond donor (HBD) of the P1' α-hydroxymethyl ketone moiety in **1** (Fig. 2A), as increased HBD count has been correlated with poor oral bioavailability (22). To this end, we pursued two functional groups preceded as covalent warheads for cysteine proteases in parallel: nitriles (23, 24) and benzothiazol-2-yl ketones (25, 26). The nitrile compound **2** demonstrated a significant increase in rat oral absorption (oral bioavailability (F) = 7.6% and fraction of oral dose absorbed from the gastrointestinal tract (F_a × F_g) = 38%) (27), while maintaining reasonable metabolic stability (intrinsic clearance (CL_{int}) against oxidative metabolism in human liver microsomes (HLM) (28) relative to **1** (Fig. 1). However, the in vitro FRET M^{pro} potency (K_i = 27.7 nM) and SARS-CoV-2 antiviral activity (EC₅₀ = 1364 nM) of **2** was inferior to **1**. Introduction of a 6,6-dimethyl-3-azabicyclo[3.1.0]hexane as a cyclic leucine mimetic at P2 (Fig. 2B) removed the HBD from the P2/P3 amide linkage. Analog **3**, resulting from the combination of this cyclic P2 fragment with a P1' benzothiazolyl ketone, displayed high passive absorptive permeability (P_{app} = 10.3 × 10⁻⁶ cm/s). The reduced biochemical SARS-CoV-2 M^{pro} inhibitory potency of **3** (K_i = 230 nM) relative to other reported benzothiazole-2-yl SARS-CoV-2 M^{pro} inhibitors (29) containing leucine

P2 groups can be rationalized from the binding mode observed for **3** (Fig. 2C). While the 6,6-dimethyl-3-azabicyclo[3.1.0]hexane effectively fills the lipophilic S2 pocket formed by Met49, Met169, His41, and Gln 189, productive hydrogen bonding of the ligand backbone to Gln189 is no longer possible (Fig. 2C). The inferior SARS-CoV-2 M^{pro} potency and the high CL_{int} (337 μl/min/mg) precluded further investments in compound **3**. Similar to **1**, the P3 indole of **3** does not protrude into the S3 pocket (Fig. 2, A and C). To better occupy the S3 pocket, we introduced branched, acyclic P3 groups. The methanesulfonamide in compound **4** extends underneath Gln189, productively engaging P3 pocket residues and achieving improved hydrogen bonding interactions with the Glu166 backbone (Fig. 2D) relative to **1** and **3**. Compound **4** demonstrated improved SARS-CoV-2 M^{pro} biochemical potency (K_i = 7.93 nM), Vero E6 antiviral activity (EC₅₀ = 909 nM), and HLM CL_{int} (127 μl/min/mg) relative to **3** (Fig. 1). Examination of the rat pharmacokinetics of **4** also revealed improvements in oral absorption (F = 10%, F_a × F_g = 84%) (Fig. 1). An effort to identify alternate P3 capping groups to sulfonamide led to trifluoroacetamide **5**. Compound **5** exhibited comparable biochemical potency (K_i = 12.1 nM) to **4**, but with greatly improved SARS-CoV-2 Vero E6 antiviral activity (EC₅₀ = 85.3 nM) and passive permeability (P_{app} = 13.1 × 10⁻⁶ cm/sec) as well as increased metabolic stability in HLM (CL_{int} = 30.3 μl/min/mg) (Fig. 1). **5** also shows greatly improved oral pharmacokinetics in both rats (F = 33%, F_a × F_g = 100%) (Fig. 1) and monkeys (F = 7.9%, F_a × F_g = 66%) (table S1). Introduction of the P1' nitrile to this scaffold led to the identification of the clinical candidate PF-07321332 (**6**, Fig. 1). Compound **6** is a potent inhibitor of SARS-CoV-2 M^{pro} biochemical activity (K_i = 3.11 nM) with improved Vero E6 antiviral activity (EC₅₀ = 74.5 nM) relative to compounds **2–4**. Compound **6** displays a similar binding mode (Fig. 2E) to compound **4**. The P1' nitrile of **6** forms a reversible covalent thioimidate adduct with the catalytic Cys145 (Fig. 2F). Reversibility of M^{pro} inhibition by **6** was demonstrated upon incubation of SARS-CoV-2 M^{pro} (2 mM) with 2 mM of either **6** or an irreversible M^{pro} inhibitor (compound **7**, Fig. 3A) (17) for 30 min and monitoring M^{pro} activity following 100-fold dilution of the incubation mixtures. No recovery of activity was observed following M^{pro} incubation with **7**. The recovery of >50% M^{pro} activity following incubation with **6** indicates that inhibition of SARS-CoV-2 M^{pro} is reversible (Fig. 3A). We selected the nitrile compound **6** (named PF-07321332) over compound **5** as the clinical candidate based on ease of synthetic scale-up, enhanced solubility that allowed for a simple formulation vehicle in support of pre-clinical toxicology, and reduced propensity for epimerization at the P1 stereocenter.

PF-07321332 demonstrated potent inhibition in FRET M^{pro} assays representing M^{pro} from all coronavirus types known to infect humans (6, 7, 30), including bet-

coronaviruses (SARS-CoV-2, SARS-CoV-1, HKU1, OC43, MERS) as well as alpha-coronaviruses (229E, NL63) (Fig. 3B and table S2). No inhibitory effects were noted against several mammalian cysteine (caspase 2, cathepsin B, cathepsin L), serine (chymotrypsin, elastase, thrombin) and aspartyl (cathepsin D) proteases at the highest concentration tested (100 μ M) of PF-07321332 (table S3). This was also the case for HIV-1 protease, a viral aspartyl protease (table S3).

The in vitro antiviral activity of PF-07321332 was also evaluated in two physiologically relevant cellular systems; human adenocarcinoma-derived alveolar basal epithelial (A549) cells constitutively expressing ACE2 (19) and differentiated normal human bronchial epithelial (dNHBE) cells (31). In contrast with Vero E6 cells, A549 and dNHBE cells do not overly express P-glycoprotein, and therefore, co-dosing with a P-glycoprotein inhibitor was not necessary to gauge antiviral activity in these cell lines (19). PF-07321332 inhibited SARS-CoV-2 replication as assessed using a nanoluciferase reporter virus in A549-ACE2 cells with EC₅₀ and EC₉₀ values of 77.9 nM and 215 nM respectively, with no cytotoxicity detected at concentrations up to 3 μ M (Fig. 3C and table S4). Treatment of dNHBE cells with varying concentrations of PF-07321332 for 3 days led to inhibition of SARS-CoV-2 viral replication with EC₅₀ and EC₉₀ values of 61.8 nM and 181 nM, respectively (Fig. 3C) as monitored by titration of virus harvested from the apical compartment using a 50% cell culture infective dose (CCID₅₀) assay in Vero76 cells. Increasing the duration of the dNHBE study to 5 days saw viral replication inhibited with EC₅₀ and EC₉₀ values of 32.6 nM and 56.1 nM (Fig. 3C). Since optimal therapeutic efficacy with marketed viral protease inhibitors and other antiviral agents is generally achieved when the minimum systemic unbound plasma concentrations of inhibitor (C_{min}) are maintained above cellular antiviral EC₉₀ (32), we selected the more conservative day 3 EC₉₀ value of 181 nM PF-07321332 in the dNHBE assay as the C_{min} to be maintained when predicting efficacy in animal models and in the clinic. We further evaluated the in vitro cellular antiviral activity of PF-07321332 against SARS-CoV-1, MERS and human coronavirus 229E using cytopathic effect (CPE) assays. PF-07321332 demonstrated potent antiviral activity against SARS-CoV-1 (EC₉₀ = 317 nM), MERS (EC₉₀ = 351 nM) and 229E (EC₉₀ = 620 nM) in their respective cellular assays (Fig. 3D and table S5).

We evaluated the in vivo antiviral activity of PF-07321332 in a mouse-adapted SARS-CoV-2 (SARS-CoV-2 MA10) model (33). Intranasal infection of BALB/c mice with SARS-CoV-2 MA10 leads to ~10% body weight loss and minimal mortality in 10-week-old mice. As shown in Fig. 4A, following infection with SARS-CoV-2 MA10, mice treated twice daily with PF-07321332 (at both 300 mg/kg and 1000 mg/kg doses), were protected from weight loss versus vehicle-treated mice. At 4 days post infection, mice were sacrificed, and lung viral titers

were evaluated in CCID₅₀ assays. Infected animals in the placebo group (n=12, two independent studies) had robust infection in the lungs (mean lung titer of log₁₀ 4.93 \pm 0.140 CCID₅₀/ml SARS-CoV-2 MA10) (Fig. 4B) whereas virus levels in mice treated with PF-07321332 were significantly reduced (mean lung titers of log₁₀ 3.53 \pm 0.187 and log₁₀ 3.02 \pm 0.423 CCID₅₀/ml for 300 and 1000 mg/kg PF-07321332-treated groups, respectively). In a satellite group of uninfected mice, the 300 mg/kg BID dose of PF-07321332 used in the mouse-adapted viral in vivo efficacy study maintained C_{min} unbound plasma concentrations above ~0.9 x EC₉₀ (Fig. 4C), as defined in the dNHBE primary cell assay. The 1000 mg/kg BID dose of PF-07321332 maintained a C_{min} unbound plasma concentration of ~4 x EC₉₀ (Fig. 4C). This confirms that PF-07321332 is effective at reducing SARS-CoV-2 MA10 viral load in mouse lungs at concentrations consistent with the observed in vitro anti-viral potency, and those being targeted clinically.

Disease in this model is manifest by weight loss and pathological changes in the lungs of the infected mice (33). Histopathological analysis and immunostaining of lungs from the SARS-CoV-2 MA10 infected mice shows that PF-07321332 limits cellular infiltration (Fig. 4D and fig. S2) and protects lung tissue from damage due to virus replication (Fig. 4E). Immunohistochemical analysis using a viral nucleocapsid antibody to detect viral antigen levels in the lungs revealed that PF-07321332 inhibits virus replication in a dose-dependent manner (Fig. 4E). Lungs from vehicle-treated, infected mice showed the strongest staining whereas mock-infected lungs were negative for nucleocapsid staining. Histopathological evaluation of lungs from the vehicle-treated mice demonstrated evidence of increased perivascular inflammation, bronchial or bronchiolar epithelial degeneration or necrosis, bronchial or bronchiolar inflammation, cellular debris in alveolar lumen, and alveolar inflammation and thickening of the alveolar septum compared to PF-07321332-treated mice and mock-infected mice (fig. S2). Most of the infected mice exhibited multifocal pulmonary lesions, which were significantly reduced in PF-07321332-treated mice.

PF-07321332 exhibited moderate plasma clearance (CL_p) in rats and monkeys, with elimination half-lives (t_{1/2}) of 5 hours and <1 hour, respectively, after intravenous dosing (table S6). Following oral administration to rats, crystalline PF-07321332 (10 mg/kg) demonstrated oral dose absorbed (F_a x F_g) and oral availability (F) values ranging from 65-95% and 34-50% respectively, depending on the crystalline form used (Fig. 1 and table S6). Oral administration of PF-07321332 (10 mg/kg) to monkeys led to a relatively poor F of 8.5% (F_a x F_g = 20%) (table S6), which is attributed to first-pass metabolism along the gastrointestinal tract by cytochrome P450 (CYP) enzymes, consistent with rapid (t_{1/2} = 20.5 min, CL_{int} = 33.8 μ l/min/mg) NADPH-dependent metabolic turnover of

PF-07321332 in monkey intestinal microsomes (table S7). PF-07321332 was resistant ($t_{1/2} > 240$ min, $CL_{int} < 2.89 \mu\text{l}/\text{min}/\text{mg}$) to CYP-mediated metabolism in intestinal microsomes from rat and human. PF-07321332 (0.3–10 μM) exhibited concentration-independent plasma protein binding in rat (mean plasma unbound fraction ($f_{u,p}$) = 0.478), monkey (mean $f_{u,p}$ = 0.434), and human (mean $f_{u,p}$ = 0.310) under equilibrium dialysis conditions (34).

Drug-metabolizing enzymes involved in the metabolism of PF-07321332 were also studied. In NADPH-supplemented HLM, PF-07321332 demonstrated moderate CL_{int} (24.5 $\mu\text{l}/\text{min}/\text{mg}$) (Fig. 1 and table S7), which was significantly inhibited ($\geq 82\%$) by the selective CYP3A4/5 inhibitor ketoconazole (28) (table S8). Moreover, the oxidative metabolic profile of PF-07321332 in HLM, which includes modifications on the P2 6,6-dimethyl-3-azabicyclo[3.1.0]hexane, the *tert*-butyl group at the P3 position, and the P1 pyrrolidinone ring, was reproduced in incubations of PF-07321332 with recombinant human CYP3A4 (fig. S1). These *in vitro* studies, which established a predominant role for CYP3A4 in the metabolism of PF-07321332, also presented an opportunity to boost therapeutic concentrations of PF-07321332 in the clinic via co-dosing with the potent CYP3A4 inactivator ritonavir (RTV), which is used as a pharmacokinetic enhancer of several marketed protease inhibitors (e.g., darunavir, lopinavir) that are subject to metabolic clearance via CYP3A4 (35, 36).

PF-07321332 demonstrated a favorable off-target selectivity profile in a broad panel of G protein-coupled receptors, kinases, transporters and phosphodiesterase enzyme inhibitor screens, and was devoid of activity against the cardiac ion channels $K_v1.1$, $Ca_v1.2$, and $Na_v1.5$ (tables S9 and S10). PF-07321332 was not mutagenic or clastogenic in *in vitro* genetic toxicity studies and was negative in an *in vivo* rat micronucleus assay (table S11). Repeat oral dosing of PF-07321332 in 2-week regulatory toxicity studies in monkeys (60–600 mg/kg) and rats (40–1000 mg/kg) led to dose-dependent increases in both maximal plasma concentrations (C_{max}) and area-under-the-plasma concentration versus time curves (AUC) (Fig. 5, A and B, and table S12). This resulted in unbound C_{max} and average concentrations (C_{av}) margins of 273-fold and 65-fold in rats (day 14, 1000 mg/kg) and 510-fold and 245-fold in monkeys (day 15, 600 mg/kg) over the measured unbound EC₉₀ value of PF-07321332 determined in the SARS-CoV-2 day three dNHBE cellular assay. PF-07321332 was well tolerated with no adverse findings in either species; the corresponding no observed adverse effect levels (NOAEL) were the highest doses tested (600 mg/kg/day in monkeys and 1000 mg/kg/day in rats).

The safety, tolerability, and pharmacokinetics of PF-07321332 as a single agent and in combination with RTV are under investigation in a randomized, double-blind, placebo-controlled, single ascending dose study in healthy adult

participants (table S13, ClinicalTrials.gov Identifier: NCT04756531). At each dose tested, 4 participants were randomized to receive active treatment while 2 participants received placebo. In the PF-07321332/RTV co-administration dosing paradigm, each subject (active and placebo) received one tablet (100 mg) of RTV at -12 hours, 0 hours and 12 hours. PF-07321332 was administered as an oral suspension under fasted conditions at 0 hours (minimum fast of ~10 hours prior to treatment). Preliminary plasma concentration versus time pharmacokinetic profiles achieved from two oral doses, PF-07321332 (150 mg) alone and PF-07321332 (250 mg) with RTV are presented in Fig. 5C. PF-07321332 was safe and well tolerated, and exhibited a significant boost in plasma concentrations when co-administered with RTV. Oral plasma concentrations of PF-07321332 (250 mg with RTV) were considerably above the SARS-CoV-2 antiviral EC₉₀ value (total EC₉₀ = 292 ng/ml, unbound EC₉₀ = 90.5 ng/ml, 181 nM) at 12 hours post-dose, thus increasing confidence in achieving robust pan-coronavirus anti-viral activity clinically (Fig. 5D). Based on these observations and the high safety margins at the NOAEL doses in animals, the efficacy of PF-07321332 in COVID-19 patients will be assessed with a twice daily dosing paradigm with the potential to increase the dose of PF-07321332 as a single agent and/or co-administration with RTV.

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Author contributions: Conceptualization: CMNA, ASA, DRO, MP Formal Analysis: LD, BB, SEG, QY, DKR, JJN, AD, RSO, RSPS Investigation: MA, SB, JCL, JL, KO, LW, RF, KSG, WL, RSO, DKR, SAG, LA, SN, HE Methodology: JBT, QY, MFS, PRV, MRR, MP, DRO, NCP, AC, EPK, KJC, RSPS Project Administration: DRO, MRR, JBT, ASK, CMS, LU, JGS, BLH, YZ, SWM, RDC, NCP Resources: AD, JJN, KJC Supervision: MRR, CES, AES, BLH, PRV, RDC Visualization: SWM, DKR, SN, BB, LA, HE Writing – original draft: DRO, ASK, MFS, CMS, RDC, MRR, JGS, CMS, YZ Writing – review and editing: DRO, ASK, MFS, ASA, CMNA **Competing interests:** DRO, CMNA, ASA, LA, MA, SB, BB, RDC, AC, KJC, AD, LD, HE, RF, KSG, SEG, EPK, ASK, JCL, JL, WL, SN, RSO, KO, NCP, DKR, MRR, MFS, JGS, RPS, CMS, AS, JBT, LU, PRV, LW, QY, YZ are employees of Pfizer and some of the authors are shareholders in Pfizer Inc. SWM, JJN, MP were employees of Pfizer Inc. during part of this study. **Data and materials availability:** All non-clinical data are available in the main text or the supplementary materials. X-ray coordinates and structure factors are deposited at the RCSB Protein Data Bank under accession codes 7RFR, 7RFU, 7RFS and 7RFW. Individualized Fig. 5 data, and statistical analysis plan are available on www.Figshare.com with the following DOI: 10.25454/pfizer.figshare.16699669. Pfizer compound use requests are processed via the 'Pure Compound Grants' program see <https://www.cybergrants.com/pfizer/Research>. Pfizer Inc. has applied for patent applications covering PF-07321332 as well as related compounds. A phase 1 clinical trial has been registered at clinicaltrials.gov (identifier NCT04756531).

SUPPLEMENTARY MATERIALS

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Materials and Methods

Figs. S1 and S2

Tables S1 to S13

References (37–70)

MDAR Reproducibility Checklist

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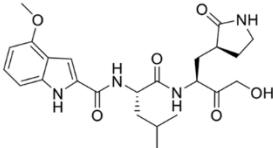
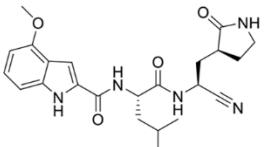
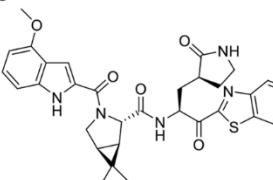
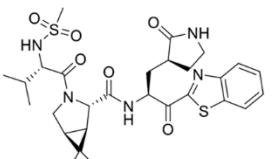
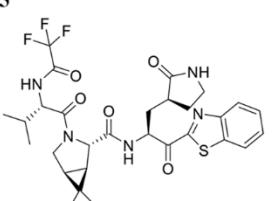
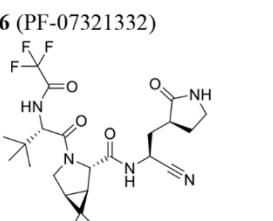
Number/Structure	SARS-CoV2 M ^{pro} Ki (nM) ^a	VeroE6-enACE2 CPE EC ₅₀ (nM) ^b	MDCK-LE P _{app} (x 10 ⁻⁶ cm/sec) ^c	HLM CL _{int} (μl/min/mg) ^d	Rat CL _p (mL/min/kg) ^e	Oral F (%) ^{e,f}	F _a x F _g (%) ^{e,h}
1 (PF-00835231) 	0.271 (0.155 – 0.471, n=6)	231 (158 – 338, n=8)	< 0.207 ± 0.048 (n=6)	7.47 ± 0.88	27.0 ± 3.1	1.4 ± 0.8	3.3
2 	27.7 (18.4 – 41.7, n=5)	1364 (860 – 2164, n=15)	0.945 ± 0.281 (n=6)	34.4 ± 0.7	39.3 (37.0, 41.5)	7.6 (7.4, 7.8)	38
3 	230 (181 – 292, n=4)	5593 (3457 – 9051, n=8)	10.3 ± 2.4 (n=6)	337 ± 9	N.D.	N.D.	N.D.
4 	7.93 (3.62 – 17.4, n=5)	909 (557 – 1482, n=14)	1.56 ± 0.38 (n=6)	127 ± 3	42.9 (38.2, 47.6)	10 (7.5, 13)	84
5 	12.1 (8.05 – 18.1, n=7)	85.3 (76.5 – 95.2, n=36)	13.1 ± 2.0 (n=8)	30.3 ± 0.6	31.0 (30.6, 31.4)	33 (33, 34)	100
6 (PF-07321332) 	3.11 (1.47 – 6.59, n=6)	74.5 (66.5 – 83.4, n=20)	1.71 ± 0.28 (n=4)	24.5 ± 0.2	27.2 (22.5, 31.9)	50 (30, 71), 34 ± 19 ^g	95, 65 ^g

Fig. 1. In vitro and in vivo parameters optimized in identifying oral SARS-CoV-2 M^{pro} inhibitors. ^aK_i values were fit to the Morrison equation with substrate, K_m and M^{pro} concentration parameters fixed to values described in the supplemental information. Data are geometric mean values with 95% confidence interval (CI) values and replicate numbers in parentheses. ^bEC₅₀ values were calculated using data normalized to controls within the assay and fit to a 4-parameter logistic curve fit (see supplemental information for details). Data are geometric mean values with 95% CI values and replicate numbers in parentheses. ^cApparent passive permeability (P_{app}) from apical to basolateral direction was determined in Madin-Darby canine kidney-low efflux (MDCK-LE) cells (21). ^dCL_{int} refers to total intrinsic clearance obtained from scaling of half-lives of test compounds in NADPH-supplemented HLM (28). Incubations were conducted on a single day in triplicate. ^ePharmacokinetic parameters were calculated from plasma concentration–time data and are reported as mean values (n= 2–3 male Wistar-Han rats/dosing route). See supplemental information for additional details. ^fOral pharmacokinetics studies were conducted in the fed state. Oral bioavailability (F) is defined as the dose-normalized AUC after oral administration divided by the dose-normalized AUC after intravenous administration. ^gCrystalline 6 was orally administered in anhydrous (Form 1) as well as anhydrous methyl-tert-butyl ether cosolvate form. ^hThe fraction of the oral dose absorbed from the gastrointestinal tract (F_a × F_g) was estimated using the equation F_a × F_g = F/(1 – CL_{blood}/Q) (27) (see supplemental section for additional details).

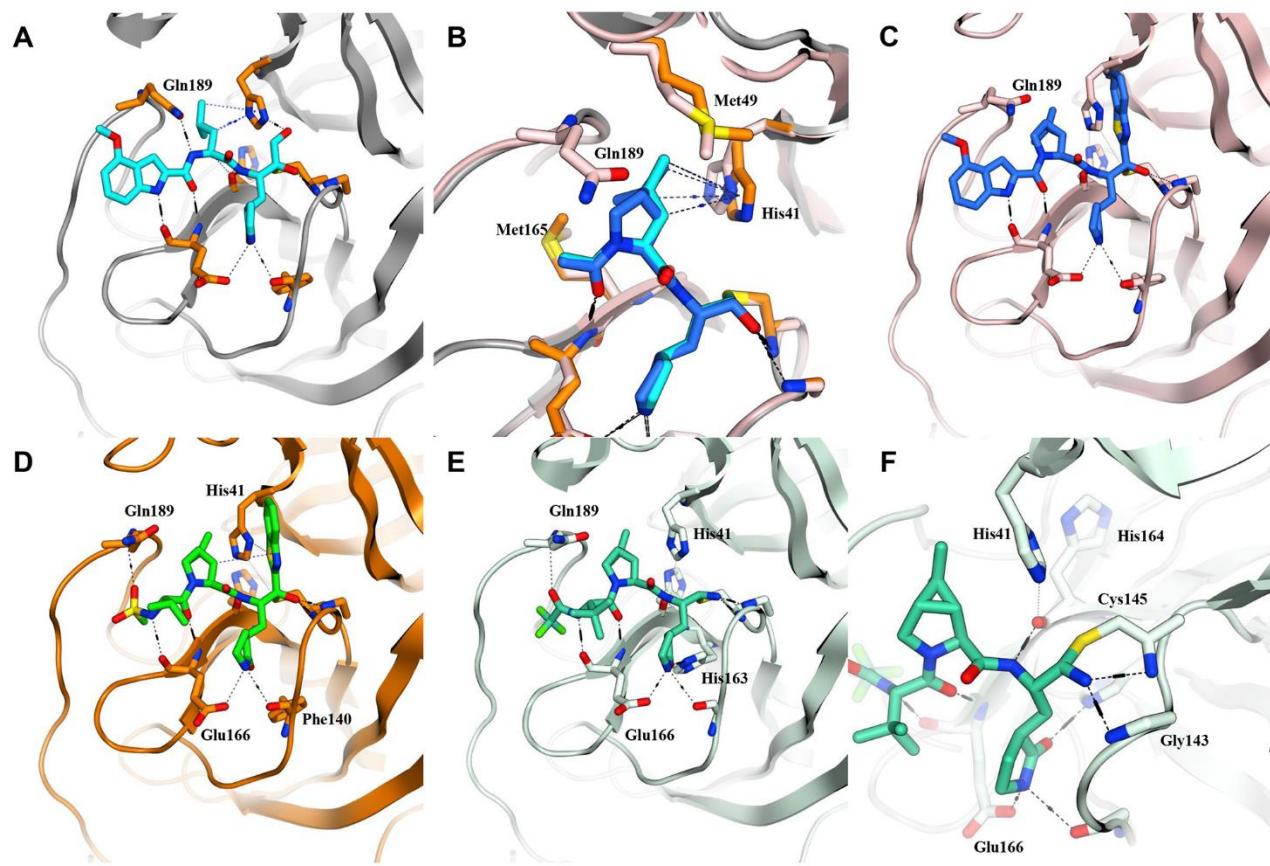


Fig. 2. SARS-CoV-2 M^{pro} structural biology. (A) Co-crystal structure of PF-00835231 (1) with SARS-CoV-2 M^{pro}. Key interactions are indicated. (B) Modeled overlap of dimethyl-bicyclo[3.1.0] proline from compound 3 (blue) as a mimic of P2 leucine residue (cyan) found in the viral polyprotein substrate and 1. This tolerated P2 change eliminates an H-bond donor from resulting inhibitors. (C) Compound 3 effectively fills the lipophilic S2 pocket formed by Met49, Met165, and His41 but productive hydrogen bonding to Gln189 is no longer possible. (D) Compound 4 with optimized acyclic P3 group and restored Gln189 interaction. (E) SARS-CoV-2 M^{pro} bound crystal structure of clinical candidate PF-07321332 (6). (F) A reversible covalent Cys145 adduct is formed with the nitrile substituent in compound 6.

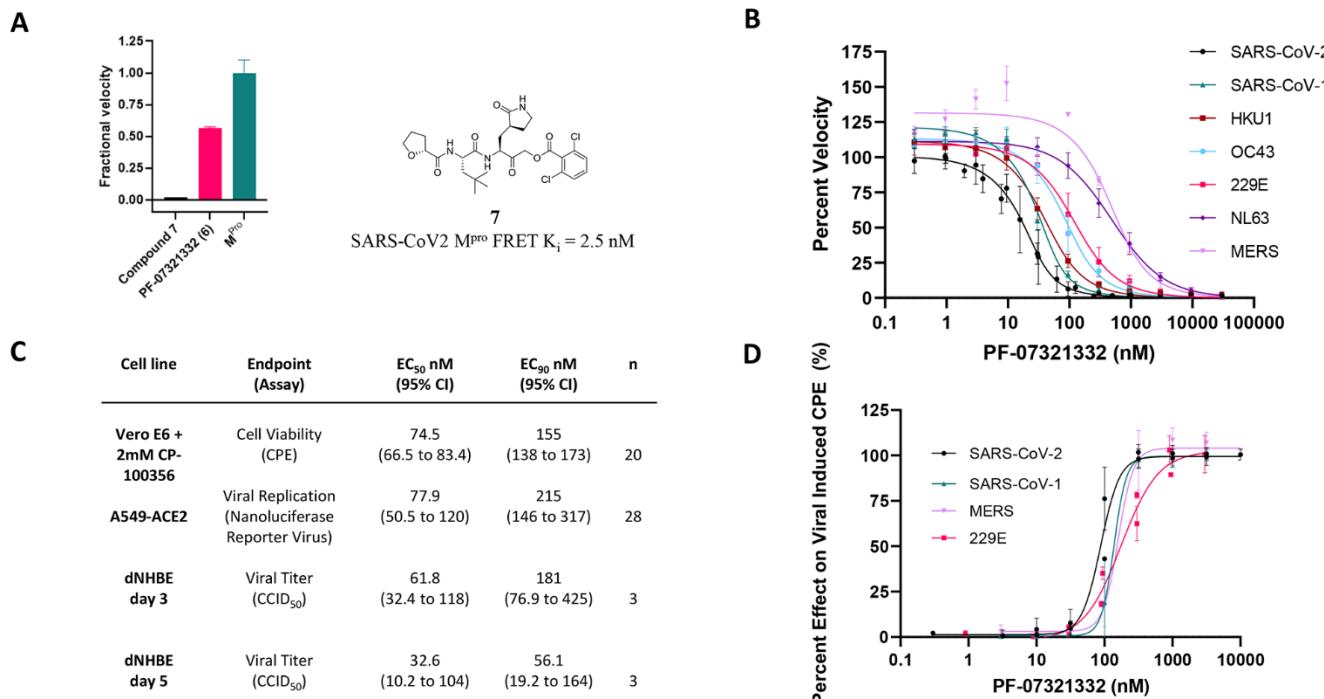


Fig. 3. PF-07321332 biochemical and antiviral activity. (A) PF-07321332 is a reversible inhibitor of SARS-CoV-2 M^{pro} as demonstrated by recovery of enzymatic activity following a 100-fold dilution of the enzyme inhibitor complex. Compound 7 (PF-00956378), an irreversible inhibitor, was included as a control. Data are representative of 3 independent experiments. (B) PF-07321332 is a potent inhibitor of the proteolytic activity of SARS-CoV-2 M^{pro} as well as related coronaviruses in FRET assays. Data shown are the mean \pm SD from 3 independent experiments. (C) PF-07321332 demonstrates potent SARS-CoV-2 antiviral cellular activity. PF-07321332 inhibited SARS-CoV-2 induced cytopathic effect (CPE) in Vero E6 cells enriched for ACE2. A P-glycoprotein inhibitor, CP-100356 (efflux inhibitor, EI) was added at 2 μM to inhibit the P-glycoprotein mediated efflux of PF-07321332. In Vero E6 cells with no EI, the EC_{50} of PF-07321332 (95% confidence interval (CI)) was 4.48 μM (3.55 to 5.65 μM) (N=8, n=20). Cytotoxicity of PF-07321332 was evaluated in non-infected cells and the CC_{50} was >100 μM (table S4). PF-07321332 inhibits SARS-CoV-2 replication in A549 cells expressing ACE2 in 4 independent experiments. The CC_{50} in non-infected A549 cells was >3 μM (table S4). In differentiated normal human bronchial epithelial (dNHBE) cells, PF-07321332 decreased SARS-CoV-2 viral replication (N=3). Data shown are the geometric mean and 95% confidence intervals (CI). (D) PF-07321332 demonstrates pan-coronavirus antiviral activity. PF-07321332 inhibition in viral-induced CPE assays: SARS-CoV-1 in Vero E6 cells (in the presence of 2 μM EI CP-100356), h-CoV-229E in MRC-5 cells and MERS in Vero 81 cells (in the presence of 1 μM EI CP-100356). Data shown are the mean and SD. CC_{50} values determined in all assays to be >100 μM (table S4).

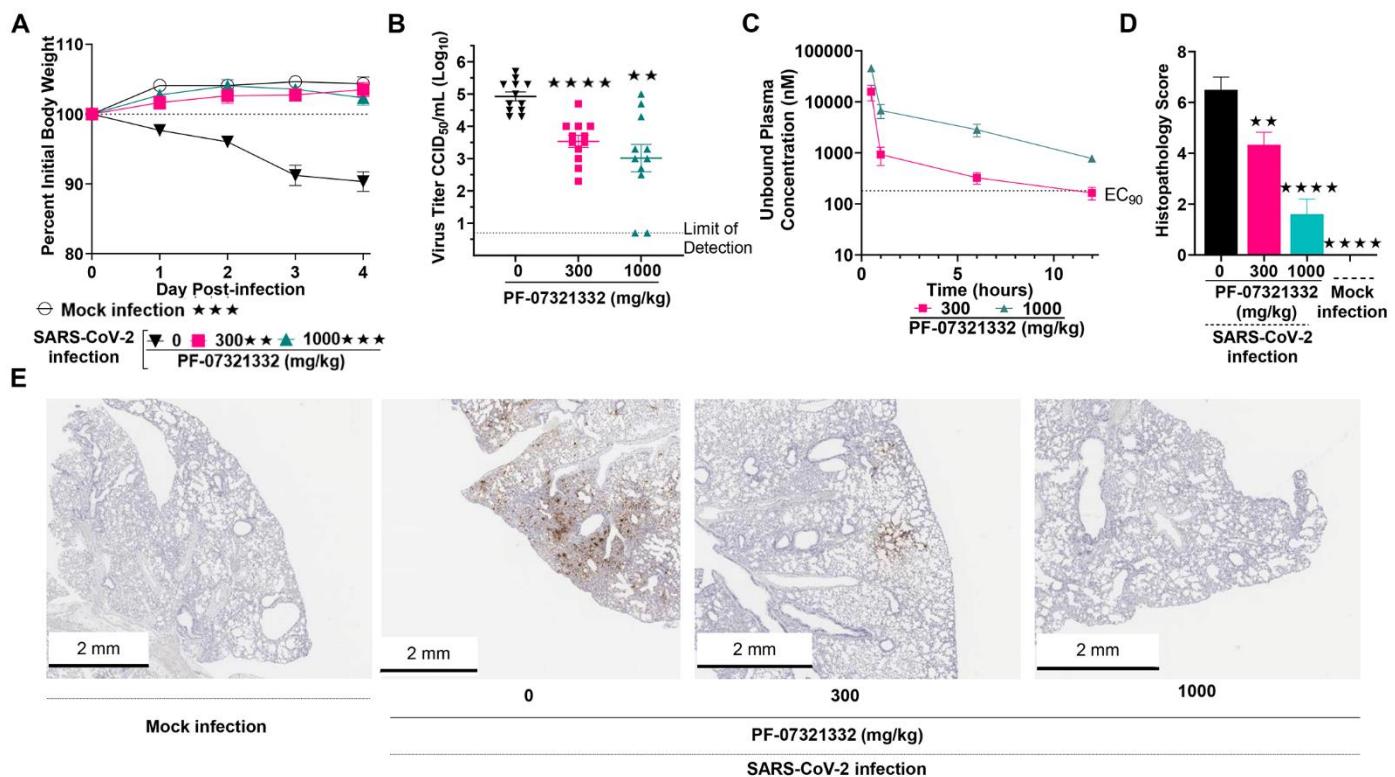


Fig. 4. In vivo efficacy of PF-07321332 against SARS-CoV-2 MA10 infection in mice. Six-mice/group were challenged intranasally with $1 \times 10^{5.0}$ 50% cell culture infectious doses of SARS-CoV-2 MA10 (CCID₅₀). Animals were orally administered 300 or 1000 mg/kg PF-07321332 or vehicle (placebo) twice daily (BID) at 4 hours post-infection. Animals were euthanized at 4 days post infection (dpi) and lungs collected for virus titers. Data for A-D are compiled from two independent studies (n=12, BALB/c mice). (A) Weight loss during infection. Mice were weighed daily. (B) Lung viral titer at 4 dpi. Lung titers are graphed as mean Log₁₀ CCID₅₀/ml +/- std error of mean. Dotted line represents the limit of detection for the CCID₅₀ assay. (C) 12-hour PF-07321332 exposure levels of 300 and 1000mg/kg doses in uninfected, orally treated mice. EC₉₀ represented as determined in the day 3 dNHBE primary cell assay (D) Histopathology scores on a scale of 0-5 where 0 is a normal healthy lung and 5 is severe coalescing areas of necrosis and confluent areas of inflammation. (E) SARS-CoV-2 Nucleocapsid Protein immunohistochemistry. Digital light microscopic scans of mouse lung tissue sections of mock-infected, placebo, 300 mg/kg PF-07321332, and 1000 mg/kg PF-07321332 treated mice stained with SARS-CoV-2 Nucleocapsid antibody. Data are scans from one study. The scale bars represent 2mm. Magnification shown is 1x.

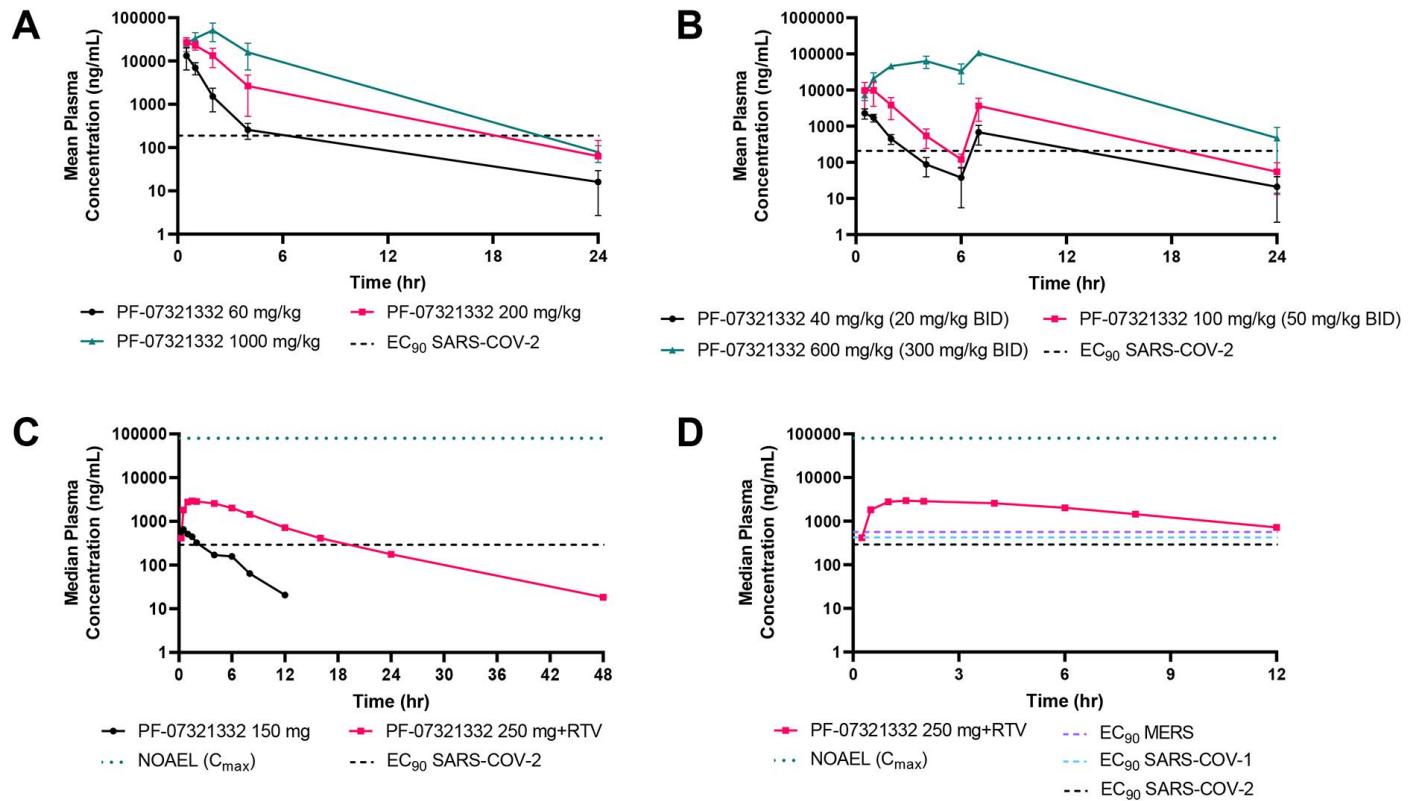


Fig. 5. Preclinical toxicology and healthy adult participant single ascending dose study exposures for PF-07321332. (A) Rat oral toxicokinetic exposures (day 14) of once-daily administered PF-07321332 compared with day 3 antiviral EC₉₀ values in dNHBE cells. (B) Monkey oral toxicokinetic exposures (day 15) of twice-daily administered PF-07321332 compared with day 3 antiviral EC₉₀ in dNHBE. (C) Human plasma concentrations (total) versus time profile after oral administration (fasted state) of PF-07321332 (150 mg) and PF-07321332 (250 mg) with ritonavir (RTV) (100 mg at t = -12 hours, 0 hours and 12 hours) compared with day 3 antiviral EC₉₀ in dNHBE. (D) Human antiviral target coverage >EC_{90,u} at 12 hours for SARS-CoV-1, SARS-CoV-2 and MERS post oral administration of PF-07321332 (250 mg) and ritonavir (100 mg at t = -12 hours, 0 hours and 12 hours). The in vitro unbound SARS-CoV-2 EC₉₀ of 181 nM was converted to ng/ml using a molecular weight of 499.5 g/mol for PF-07321332. Total EC₉₀ was calculated by dividing unbound EC₉₀ by f_{u,plasma} (rat 0.479, NHP 0.208, and human 0.310). This resulted in total EC₉₀ values of 189, 208, and 292 ng/ml for rat, NHP, and human, respectively. The calculated total human EC₉₀ for MERS and SARS-CoV-1 were 566 and 422 ng/ml, respectively. A human NOAEL of 79,700 ng/ml (for C_{max}) was estimated from the rat C_{max} value of 51500 ng/ml at the NOAEL dose of 1000 mg/kg, normalizing for plasma unbound fraction differences.

An oral SARS-CoV-2 M inhibitor clinical candidate for the treatment of COVID-19

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