

Mutual effects of single and combined CFTR modulators and bacterial infection in cystic fibrosis

Supplementary information

1 **Supplementary Methods**

2 **Minimum inhibitory concentration (MIC) measurement.** An aliquot of *P. aeruginosa* or *S. aureus*
3 strains from glycerol stocks was streaked for isolation on tryptic soy agar (TSA) and incubated at
4 37°C overnight (O/N). Bacterial glycerol stocks were not used more than three times to avoid
5 variability in the experiments. One colony was picked from the plate and used to inoculate 5 ml of
6 tryptic soy broth (TSB) (BD, Becton and Dickinson) and placed in a shaking incubator at 37°C 200
7 rpm O/N. The O/N bacterial suspension was diluted to 0.1 OD/ml in 20 ml of TSB / flask and grown
8 at 37°C at 200rpm up to reach the log phase (1). The bacteria were pelleted by centrifugation (2700
9 g, 15 min, 4°C) and resuspended in cation-adjusted Mueller-Hinton broth (MH-II broth). MICs of
10 CFTR modulators were determined using the broth microdilution susceptibility testing method,
11 following the Clinical and Laboratory Standards Institute guidelines (2, 3). CFTR modulators were
12 prepared using stock solutions in dimethyl sulfoxide (DMSO), and then tested in two-fold serial
13 diluted fashion in MH-II broth. MIC testing was run in sterile 96-well Microtiter plates. The MIC was
14 defined as the lowest compound concentration showing a reduction in the optical density at 620 nm
15 of approximately 90% (MIC₉₀) in comparison to the optical density of the bacteria grown with the
16 vehicle after 20 hrs.

17 **Checkerboard assay.** The synergistic activities of CFTR modulators combined with antibiotics were
18 determined by the checkerboard method in MH-II broth (4). Stock solutions of CFTR modulators
19 were prepared in DMSO, while those of antibiotics in sterile H₂O or DMSO, according to their
20 solubility. Serial dilutions of CFTR modulators and antibiotics were prepared based on the MIC value
21 of each drug. Then, a combination of different dilutions of each drug was prepared in 96-well
22 microtiter plates in a checkerboard formation and the bacterial strain was added. After 20 hrs
23 incubation at 37°C, bacterial growth was measured at the optical density of 620nm. Combinations
24 showing a decrease in the MIC of 90% in comparison to the MICs of the antibiotic and CFTR
25 modulator alone were recorded.. The fractional inhibitory concentration (FIC) index was calculated
26 as MIC₉₀ of CFTR modulator in combination/ MIC₉₀ CFTR alone + MIC₉₀ antibiotic in combination/
27 MIC₉₀ antibiotic alone (5). If no endpoint MIC value could be determined because of resistance to
28 compound or solubility limits, the next highest MIC value was chosen for the FIC calculation (i.e., if
29 the MIC>32, an MIC=64 was selected for the calculation) (6). The interaction between drugs was
30 interpreted as synergistic (FIC index <0.5), additive (FIC index \geq 0.50 and <1.0), indifferent (FIC index
31 \geq 1.0 and \leq 4) and antagonistic (FIC index >4) (5).

32 **Bacteria preparation for acute infection.** An aliquot of *P. aeruginosa* PAO1 reference strain from
33 glycerol stocks was streaked for isolation on TSA and incubated at 37°C O/N. One colony was picked
34 from the plate and used to inoculate 5 ml of TSB and placed in a shaking incubator at 37°C 200 rpm
35 O/N. The O/N bacterial suspension was diluted to 0.1 OD/ml in 20 ml of TSB / flask and grown for 3

36 hrs at 37°C at 200rpm, to reach the log phase (1). The bacteria were pelleted by centrifugation (2.700
37 g, 15 min, 4°C), resuspended in sterile phosphate-buffered saline (PBS) and diluted to give 1x10⁶
38 colony forming units (CFUs) / mouse in 60 µl.

39 **Mouse models.** C57BL/6NCrIBR male mice (8-10 weeks of age) were purchased from Charles
40 River (Calco, Italy), shipped in protective, filtered containers, transported in climate-controlled trucks,
41 and allowed to acclimatize for at least two days in the stabulary prior to use. Mice were maintained
42 in sterile ventilated cages in the biosafety level 3 (BSL3) facility at San Raffaele Scientific Institute
43 (Milano, Italia) where 3-5 mice per cage were housed. Mice were fed with standard rodent autoclaved
44 chow (VRFI, Special Diets Services, UK) and autoclaved tap water. Fluorescent lights were cycled
45 12hrs on, 12hrs off, and ambient temperature (23±1°C) and relative humidity (40-60%) were
46 regulated. For PK experiments, animals were separated into two groups, with IVA levels analyzed
47 in a model of acute *P. aeruginosa* lung infection or in uninfected mice. For infection experiments,
48 mice were anesthetized by an intraperitoneal injection of a solution of Avertin (2,2,2- tribromethanol,
49 97%) in 0.9% NaCl and administered at a volume of 0.015 ml/g body weight. Mice were placed in
50 supine position. The trachea was directly visualised by ventral midline, exposed and intubated with
51 a sterile, flexible 22-g cannula attached to a 1 ml syringe. An inoculum of 60 µl of planktonic bacterial
52 cells was implanted via the cannula into the lung. After inoculation, all incisions were closed by
53 suture. IVA was administered once, half an hour after the challenge, by intraperitoneal (i.p.) injection
54 at a dose of 3 mg/kg using as vehicle 10% PEG 400, 10% Tween 80, 80% saline. IVA dose used in
55 this model was calculated based on a single adult dose (150 mg) adjusted for mouse weight,
56 assuming that an adult with CF weighs 50 kg [28]. Another group of infected mice were treated with
57 the vehicle. Mice were euthanized with an overdose of carbon dioxide and the diaphragm opened.
58 Blood was collected in 0.5-ml K3-EDTA tubes through cardiac puncture. Time points for sampling
59 were before t = 0.1667, 1, 2, 6, and 24 hrs after administration. Blood samples were directly placed
60 on ice and centrifuged soon after in a precooled centrifuge, and plasma samples were stored at
61 -80°C until IVA plasma concentrations were determined. One sample was taken per mouse, and
62 every time point was sampled in triplicate for uninfected mice (3 mice) and quadruplicate for infected
63 ones (4 mice). Bronchoalveolar lavage (BAL) was performed immediately after blood collection. The
64 concentrations in ELF were determined by taking BAL fluid samples and a concomitant plasma
65 sample, as previously described [37]. In short, murine trachea was exposed by a 1-cm incision on
66 the ventral neck skin for insertion of the cannula. Lungs were instilled two times with 0.9 ml of sterile
67 saline, and the fluid was aspirated immediately. The aspirates recovered from each mouse were
68 pooled and directly placed on ice, then centrifuged in a precooled centrifuge and the supernatant
69 subsequently stored at -80°C. Lungs were excised aseptically, weighed and homogenized in 2 ml
70 of saline using the homogenizer gentleMACS™ Octo Dissociator, and then stored at -80°C. One-

71 hundred μ l of the lung homogenates and BAL fluids and 10-fold serial dilutions were spotted onto
72 TSA. CFUs were determined after overnight growth at 37°C.

73 **Protein binding and concentration determinations in plasma, lung and ELF.** Protein binding
74 was measured in mouse plasma and lung homogenate by Rapid Equilibrium Dialysis (RED). Stock
75 solutions of ivacaftor (IVA) and its main metabolite M1 were prepared in MeOH whereas M6 was
76 prepared in DMSO at a concentration of 1 mg/ml. Working solutions (WSs) were prepared by adding
77 10 μ l of each stock solution to 90 μ l of DMSO to have a concentration of 0.1 mg/ml. The standard
78 Diclofenac, used as control as highly binding standard (7), was prepared in DMSO (1 mM). 5 μ l of
79 WSs were diluted in 495 μ l plasma or lung homogenate to have a final compound incubation
80 concentrations of 1000 ng/ml and 10 μ M for Diclofenac. The concentration of 1000 ng/ml was chosen
81 because similar to IVA C_{max} *in vivo*. Spiked samples (200 μ l) were dialyzed in the Rapid Equilibrium
82 Dialysis (RED) inserts (Thermo Scientific) against 350 μ L of PBS buffer in the RED plate for 4 hrs at
83 37°C, under agitation (300 rpm orbital shaker). Samples were tested in duplicate. At the beginning
84 and the end of the incubation, 50 μ l of sample in the donor were added to 50 μ l of PBS buffer whereas
85 50 μ l of the blank matrix were added to 50 μ l of dialyzed PBS buffer. Samples were then mixed with
86 200 μ l of acetonitrile containing the internal standard (Lidocaine 10 ng/ml for IVA and metabolites,
87 and Verapamil 10 ng/ml for Diclofenac). Samples were centrifuged at 3500g for 5 minutes. Then,
88 supernatants were transferred into vials for ultra-performance liquid chromatography-tandem mass
89 spectrometry (UPLC-MS/MS) analysis. T_0 was used to check potential instability in the matrix. 5 μ l
90 supernatant was injected on a Waters Acquity Premiere Ultrahigh-Performance Liquid
91 Chromatography (UPLC) system equipped with a Kinetex C8 (2,6 μ m particle size, 100A 50*2.1mm
92 Phenomenex) column at 35 °C, running at a flow rate of 300 μ l/min and interfaced with an AB Sciex
93 API 3200 mass spectrometer with a TurbolonSpray source (positive mode) and the Analyst (version
94 1.6.3) software. The chromatographic gradient was in 0.1% formic acid in water or acetonitrile, going
95 from a 2% organic phase at 2 min to 98% at 2.5min. Isocratic condition at 98% was kept up to 4 min
96 and the system was reconditioned from 4.5 to 6 min at 2%B. Mass spectrometer was run in Positive
97 mode with a capillary Temperature 450 C, CUR 30, Gas 1 25 and Gas 2 45, IS was at 5500V.
98 Quantitation was performed in Multiple reaction Monitoring (MRM) using the transitions reported in
99 **Supplementary Table S6**. Protein binding was calculated as follows:

100 $\% \text{ Fb (fraction bound)} = [(\text{area donor} - \text{area acceptor}) / \text{area donor}] * 100$

101 Free fraction= 100- %Fb.

102 Concentrations of IVA and its metabolites in ELF were determined by using the ratio of the urea
103 concentration in BALF to that in plasma (8). Urea content was measured with a colorimetric Sigma
104 Aldrich MAK006 Kit. Absorbance was measured at 570 nm (A570) on a Tecan Sunrise
105 spectrophotometer. Urea concentrations are normally the same in plasma and ELF because of the
106 equilibrium across the capillary-alveolar membrane. Thus, the apparent ELF volume was estimated

107 by using urea as an endogenous marker of ELF dilution and was calculated as follows: drug
108 concentration in ELF = drug concentration in BALF × urea concentration in plasma/urea
109 concentration in BALF. Reference urea standards ranged from 20 to 100 nmol/ml and were linear
110 over the concentration range.

111 **Samples preparation.** Lung homogenates were prepared by homogenizing the whole tissue in 2 ml
112 of ammonium acetate buffer 20mM. An intermediate 50 µg/ml stock solution of all the three analytes
113 was prepared from the stock solutions 1 mg/ml. WSs of the three compounds, with concentrations
114 ranging from 5 to 10000 ng/ml, were prepared by sequential dilution of intermediate stock solution
115 in MeOH. Calibration curves in plasma and lung homogenate and BALF were prepared by spiking 5
116 µl of WSs into 45 µl of plasma, blank lung homogenate or blank BALF. QC samples were analogously
117 prepared at concentrations of 10, 50 and 500 ng/ml in the three matrices. Calibrants, QC samples
118 and samples (50 µl of matrices) were added to a 96 wells plate Phree Phospholipid Removal
119 Solutions (Phenomenex) with 200 ul of acetonitrile containing Lidocaine as internal standard (10
120 ng/ml) and 0.1 % of formic acid. The plate was shaken for 5 min on a 500g orbital shaker and
121 centrifuged for 15 min at 5°C with a collection plate. The eluates were transferred in a 96-well plate
122 for autosampler and analyzed. Calibration ranges were the following ones: for IVA 0.5-750 ng/ml in
123 plasma and BALF and 4-1000 ng/g in the lung homogenate; for M1 0.5-1000 in plasma, 0.5-500ng/ml
124 in BALF and 3.4-1000 ng/g in the lung homogenate; for M6 0.5-500ng/ml in plasma and BALF, and
125 3.4-2000ng/g in the lung homogenate. Concentration values reported as unquantifiable were
126 excluded from the evaluation.

127

Supplementary Tables

Supplementary Table S1. Phenotypic traits and antibiotics susceptibility of clinical *P. aeruginosa* isolates tested in this study. *P. aeruginosa* early isolates, collected from CF patients at the early stage of chronic colonization, and their pathoadaptive variants, collected after years of persistence, in the advanced stage of chronic colonization (late), were previously characterized for their clonality (all isolates collected from a single patient are clonal), phenotypic traits and susceptibility to antibiotics (3, 9, 10).

Patient	Isolate / type	Yrs (Δ)	Mucoidy μg uronic acid/ μg protein	Motility		Protease (\emptyset cm)	Pyocyanin (OD_{695} 26h)	Susceptibility to antibiotics			
				Twitching (\emptyset cm)	Swimming (\emptyset cm)			COL	CIP	MER	TOB
AA	2 / early		-	1.8	5.8	1.8	0.05	-	-	-	-
	43 / late	7.5	1.00	1.1	-	1.0	0.02	-	-	-	-
	44 / late	7.5	-	1.3	-	1.1	0.05	-	-	-	-
MF	1 / early		-	2.6	1.6	1.7	0.05	-	-	-	-
	51 / late	10.1	-	-	-	-	0.08	-	R	R	-
KK	1 / early		-	1.4	1.2	-	0.02	-	-	-	-
	2 / early		-	-	-	-	0.01	-	-	-	-
	71 / late	12.6	-	-	-	-	0.01	-	R	R	-
	72 / late	12.6	-	-	-	-	0.01	-	R	R	-
RP	73 / late	8.5	-	-	-	-	0.05			R	

Yrs (Δ): years between early and late in clonal sequential isolates; * Indicates swimming and twitching motility zone diameter. R: antibiotic resistance. COL: colistin, CIP: ciprofloxacin, MER: meropenem, TOB: tobramycin

Supplementary Table S2: Phenotypic traits of clinical *S. aureus* isolates tested in this study. *S. aureus* early isolates, collected from CF patients at the early stage of chronic colonization, and their pathoadaptive variants, collected after years of persistence, in the advanced stage of chronic colonization (late), were previously characterized for their clonality (all isolates collected from a single patient are clonal) and phenotypic traits (11).

Patient	Isolate name	Isolate type	Yrs (Δ)	Colony size	CP	β -HL
20	A10	early		n	+++	
	A12	late	13	i	++	
22	J6	early		n		
	J9	late	13.5	i		
25	F1	early		n	+	+
	F5	late	13	n		

Yrs (Δ): years between early and late in clonal sequential isolates; n: normally-sized colonies, i: intermediate-sized colonies; CP: capsule polysaccharide production, β -HL: presence of β -haemolysin activity.

Supplementary Table S3. Minimum inhibitory concentrations (MIC₉₀) of ivacaftor (IVA), lumacaftor (LUM), tezacaftor (TEZ), elexacaftor (ELX) and the triple combination elexacaftor/tezacaftor/ivacaftor (ELX/TEZ/IVA) against *P. aeruginosa* isolates collected from CF patients at different stages of colonization (early and late). *P. aeruginosa* isolates were grown for 20 hours in the presence of serial dilutions of IVA, LUM, TEZ, ELX and the triple combination ELX/TEZ/IVA. The concentrations tested ranged from 0.25 μ g/ml to 32 μ g/ml for IVA, LUM, TEZ and ELX, and from ELX 0.5 μ g/ml / TEZ 0.25 μ g/ml / IVA 0.375 μ g/ml to ELX 64 μ g/ml / TEZ 32 μ g/ml / IVA 48 μ g/ml for the triple combination. The MIC₉₀ was defined as the lowest compound concentration showing a reduction in the optical density at 620 nm of approximately 90% in comparison to the optical density of the bacteria grown with the vehicle after 20 hrs. Each experiment was performed at least two independent times (two technical replicates).

		<i>P. aeruginosa</i> isolates										
Isolate name		PAO1	AA2	AA43	AA44	MF1	MF51	KK1	KK2	KK71	KK72	RP73
Isolate type		ref	early	late	late	early	late	early	early	late	late	late
MIC ₉₀ (μ g/ml)	IVA	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32
	LUM	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32
	TEZ	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32
	ELX	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32	>32
	ELX / TEZ / IVA	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48	>64 / >32 / >48

ref, reference strain; early, isolate collected at the early stage of chronic colonization; late, isolate collected after years of persistence, in the advanced stage of chronic colonization.

Supplementary Table S4. Pharmacokinetic parameter estimates for M1 in plasma, epithelial lining fluid (ELF) and lungs of infected and non-infected mice. C57BL/6NCrIBR male mice (8 to 10 weeks of age) were infected with 1×10^6 colony forming units of *P. aeruginosa* PAO1 by intratracheal administration. A non-infected control group was also tested in parallel. After 30 min from the infection mice were treated with 3 mg/kg ivacaftor (IVA) in 10% PEG 400, 10% Tween 80, 80% saline by intraperitoneal administration. Mice were sacrificed at 10 min, 1, 2, 6 and 24 hrs after IVA administration. Blood was collected and processed to obtain plasma. Bronchoalveolar lavage fluid (BALF) was collected, centrifuged and the supernatant used to quantify IVA concentration. Lungs were excised, homogenized, centrifuged and the supernatants used to quantify IVA concentration. Plasma, lung homogenate and BALF (50 μ l) were added to a Phree Phospholipid Removal plate (Phenomenex) with acetonitrile and 0.1% formic acid in order to eliminate phospholipids, decreasing the matrix effect. Eluates were analyzed by UPLC-MS/MS method with a linear gradient in MRM positive mode. Calibration ranges were the following ones: M1 0.5-1000 in plasma, 0.5-500ng/ml in BALF and 3.4-1000 ng/g in the lung homogenate. Data are the geometric mean of values from 3-4 mice. Concentrations of M1 in ELF were determined using the ratio of the urea concentration in plasma to that in BALF. Concentration in ELF = drug concentration in BALF \times urea in plasma / urea in BALF. The data are the geometric means of values from 3-4 mice.

Sample	Mice group	Pharmacokinetic parameters							
		T _{max} (hrs)	C _{max} (ng/ml)	T _{last} (hrs)	C _{last} (hrs)	AUC _{last} (hrs ng/ml)	AUC _{INF} (hrs ng/ml)	T _{1/2} (hrs)	MRT (hrs)
plasma	non-infected	1	111.6	24	2.0	887.3	898.7	4,0	4.9
	infected	1	82.6	24	1.6	823.3	832.5	4,0	5.0
ELF	non-infected	6	10.8	6	10.8	55.1	nd	nd	3.3
	infected	1	11.6	6	9.9	59.9	406.6	24.3	3.1
lung	non-infected	2	7.0	6	6.4	36.3	nd	nd	3.2
	infected	6	18.4	6	1.7	239.8	nd	nd	6.7

T_{max}, time of maximum concentration; C_{max}, maximum concentration; T_{last}, time of last quantifiable concentration; C_{last}, last quantifiable concentration; AUC_{last}, AUC to the last quantifiable concentration level; AUC_{INF}, AUC to infinity; T_{1/2}, half-life; MRT, mean residence time; nd, not determined.

Supplementary Table S5. Penetration of M1 in lung and epithelial lining fluid (ELF). Plasma and lung concentrations were multiplied for the free fraction from the protein binding experiment (0.11% for plasma and 2.83% for lung; **Table 4**). ELF free fraction was calculated considering ELF protein binding equal to protein binding in the bronchoalveolar lavage fluid. Data are the geometric mean of values from 3-4 mice.

Mice group	Values by sample type						lung/plasma		ELF/plasma	
	plasma		lung		ELF		C _{max} [#]	AUC [#]	C _{max} [#]	AUC [#]
	fC _{max} (ng/ml)	fAUC (hrs ng/ml)	fC _{max} (ng/ml)	fAUC (hrs ng/ml)	C _{max} (ng/ml)	AUC (hrs ng/ml)				
non-infected	0.12	0.97	0.20	0.93	10.8	49.0	1.67	0.96	90.0	50.5
infected	0.09	1.10	0.95	1.57	11.6	52.0	10.6	1.43	129	47.3

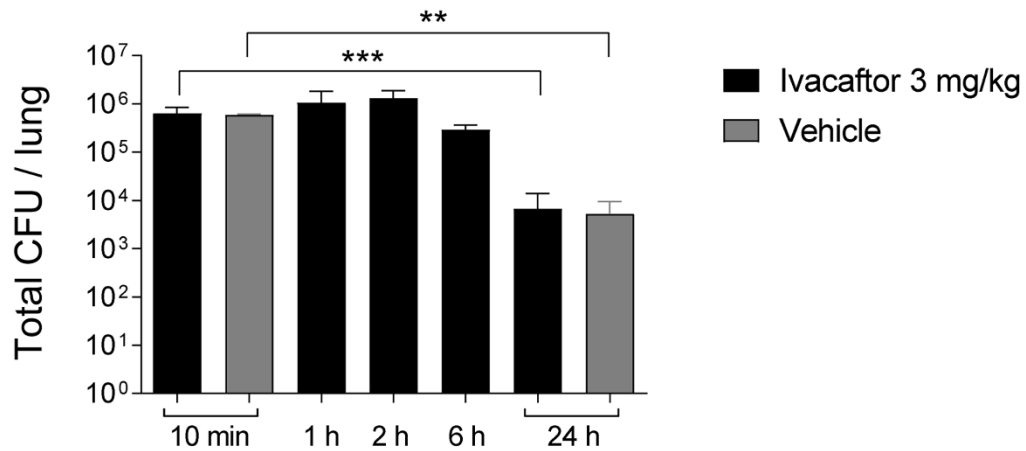
Values for lung/plasma (ratio of lung free maximum concentration or exposure to plasma free maximum concentration or exposure) and ELF/plasma (ratio of ELF maximum concentration or exposure to plasma free maximum concentration or exposure); fC_{max} and fAUC refer to unbound M1 in plasma and lung.

Supplementary Table S6. Multiple Reaction Monitoring (MRM) transitions and conditions

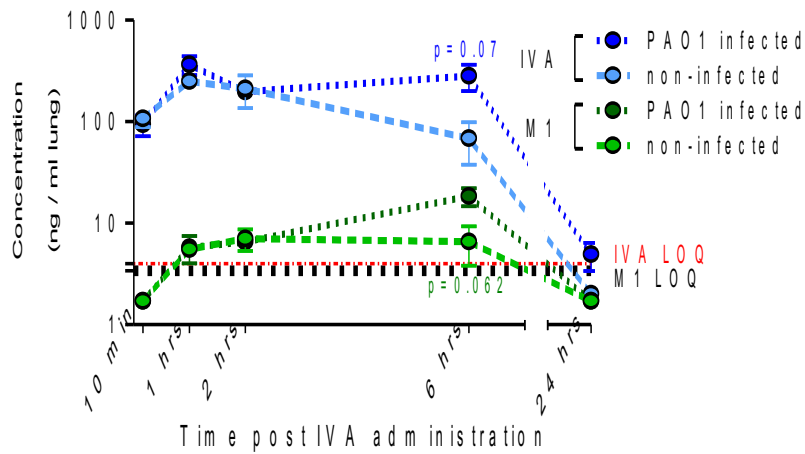
Analyte	Parent Ion MH ⁺ (Da)	Product Ion (Da)	Dwell Time (msec)	Declustering Potential (Volts)	Collision Energy(volts)
IVA_1*	393.2	172.2	100	38	38
IVA_2	393.2	319.2	100	43	41
M1_1*	409.1	172.3	100	44	48
M1_2	409.1	353.0	100	44	23
M6_1*	423.1	172.3	100	45	41
M6_2	423.1	367.2	100	45	23.5
Lidocaine 1*	235.3	86.1	100	49	26
Lidocaine 2	235.3	58.1	100	44	52

*MRM transitions used for quantitation

Supplementary Figures



Supplementary Figure S1. Ivacaftor (IVA) and M1 concentrations in murine lung. C57BL/6NCrIBR male mice (8 to 10 weeks of age) were infected with 1×10^6 colony forming units of *P. aeruginosa* PAO1 by intratracheal administration. After 30min from the infection mice were treated with 3 mg/kg IVA in the vehicle 10% PEG 400, 10% Tween 80, 80% saline or the vehicle by intraperitoneal administration. Mice were sacrificed at 10min, 1, 2, 6 and 24 hrs after IVA administration. After blood and bronchoalveolar lavage fluid collection, lungs were excised and homogenized. Both bronchoalveolar lavage fluids and lungs were plated on tryptic soy agar to determine the bacterial burden. Data are represented as mean values \pm standard errors of the means (SEMs). Statistical significance was calculated by two-way ANOVA with Bonferroni's multiple comparison test comparing mice treated with IVA and vehicle at 10 min and 24h. **: $p < 0.01$; ***: $p < 0.001$.



Supplementary Figure S2. Ivacaftor (IVA) and M1 concentrations in murine lung. C57BL/6NCrIBR male mice (8 to 10 weeks of age) were infected with 1×10^6 colony forming units of *P. aeruginosa* PAO1 by intratracheal administration. An uninfected control group was also tested in parallel. After 30min from the infection mice were treated with 3 mg/kg IVA in 10% PEG 400, 10% Tween 80, 80% saline by intraperitoneal administration. Mice were sacrificed at 10min, 1, 2, 6 and 24 hrs after IVA administration. After blood and bronchoalveolar lavage fluid collection, lungs were excised, homogenized and the supernatants used to quantify IVA and M1 by high-performance liquid chromatography–tandem mass spectrometry. Data, derived from 3-4 mice, are represented as mean values \pm standard errors of the means (SEMs). Limits of quantification (LOQ) for IVA and M1 are indicated. A value corresponding to LOQ/2 was assigned to undetectable samples at specific time-points (10 min and 24 hrs). Statistical significance was calculated by Mann-Whitney test comparing infected and uninfected mice at each time-point.

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