

1 **An assay toolkit for anti-tuberculosis drugs in the hollow fiber system**

2

3 Fernando Sanz-García<sup>1,6\*</sup>, Dominique Ndjogou<sup>2</sup>, Diana A. Aguilar-Ayala<sup>1</sup>, Maxime R.  
4 Eveque-Mourroux<sup>2</sup>, Ana Benítez<sup>1</sup>, Sergio Aznar<sup>1</sup>, Nuria Isach-Traver<sup>1</sup>, Alfonso  
5 Mendoza-Losana<sup>3</sup>, Nicolas Willand<sup>2</sup>, Ainhoa Lucía<sup>1,4,#</sup>, Santiago Ramón-García<sup>1,4,5,\*,#</sup> on  
6 behalf of the ERA4TB consortium.

7

8 <sup>1</sup>Department of Microbiology, University of Zaragoza, C/ Domingo Miral s/n 50009-  
9 Zaragoza, Spain

10 <sup>2</sup>Univ. Lille, Inserm, Institut Pasteur de Lille, U1177 - Drugs and Molecules for Living  
11 Systems, F-59000 Lille, France.

12 <sup>3</sup>Biomedical Sciences and Engineering Laboratory, Bioengineering Department,  
13 Universidad Carlos III de Madrid, 28903, Madrid, Spain.

14 <sup>4</sup>Spanish Network for Research on Respiratory Diseases (CIBERES), Carlos III Health  
15 Institute, 28029 Madrid, Spain

16 <sup>5</sup>Research & Development Agency of Aragón (ARAID) Foundation, Zaragoza, Spain.

17 <sup>6</sup>Lead contact

18

19 #: Equal contribution

20 \*Correspondence: [f.sanzg@unizar.es](mailto:f.sanzg@unizar.es) and [santiramon@unizar.es](mailto:santiramon@unizar.es)

21

22 **SUMMARY**

23 The hollow-fiber system for tuberculosis (HFS-TB) is an *in vitro*  
24 pharmacokinetic/pharmacodynamic (PKPD) tool qualified by the European Medicines  
25 Agency to support anti-TB drug development. It can simulate PK parameters of  
26 antimicrobials and feed *in silico* models to inform Phase II/III clinical trials. Yet, its  
27 implementation is challenging due to lack of consolidated technical guidelines, such as  
28 the compatibility of drugs with the most commonly used types of HFS-TB cartridges.  
29 Herein, we uncovered the compatibility of 10 anti-TB drugs alone: bedaquiline,  
30 clarithromycin, delamanid, ethambutol, isoniazid, linezolid, moxifloxacin, pretomanid,  
31 rifampicin and rifapentine; and a combination: bedaquiline-pretomanid-linezolid. The  
32 most hydrophilic compounds were within the compatibility range with all fibers; whereas  
33 the lipophilic ones required adjustments to be used in the system. Also, polysulfone and  
34 cellulose fibers were the most suitable for the tested drugs. Our data strengthen the  
35 importance of these preliminary studies and provide a useful toolkit towards wider HFS-  
36 TB implementation.

37

38 **Keywords:** hollow-fiber system, tuberculosis, pharmacokinetics, standardization,  
39 cartridge compatibility, lipophilicity, BPaL.

40

## 41 INTRODUCTION

42 Upon the first report in 2004 <sup>1</sup>, the hollow-fiber system for tuberculosis (HFS-TB) was  
43 subjected to formal evaluation by the European Medicines Agency (EMA), leading to  
44 endorsement for TB drug development in 2015 <sup>2,3</sup> and, subsequently, by the Food and  
45 Drug Administration (FDA) <sup>4</sup>. This preclinical model permits exposing a growing culture  
46 of *Mycobacterium tuberculosis* to any simulated antibiotic dose by mimicking human  
47 plasma or site of infection *in vitro* pharmacokinetic (PK) <sup>5,6</sup>. Bacteria grow in an enclosed  
48 bioreactor threaded with semi-permeable hollow fibers (i.e., a cartridge), which separate  
49 the system into two compartments: an extra-capillary space (ECS) where bacteria are  
50 harboured, and an intra-capillary space (ICS) where drugs are infused. The majority of  
51 nutrients, bacterial metabolites and compounds diffuse across the hollow fibers' pores,  
52 with different molecular weight cut-offs (MWCO), thereby equilibrating the content of  
53 both compartments. These pores are too small for bacteria to leave the ECS into the  
54 primary flow path inside the fibers, but they enable media to be constantly refreshed at a  
55 specific flow rate, as well as the addition and gradual clearance of drugs to and from the  
56 ECS <sup>7-9</sup>. Altogether, these traits allow users to mimic the PK profile of antimicrobial  
57 agents in the HFS-TB, and expose bacteria to a predefined posology to infer  
58 pharmacodynamic (PD) responses linked to said PK profiles. In combination with PKPD  
59 modelling and simulation, the HFS-TB may forecast clinical outcomes with a predictive  
60 accuracy of ca. 94.4% <sup>10</sup>. For instance, this technique can emulate dissimilar posology  
61 (e.g., once daily, QD vs. every other day, QAD). Further, HFS-TB readouts can unveil  
62 PKPD drivers for antibacterial activity of the unbound drug fraction, namely: area under  
63 the concentration-time curve (AUC), maximal concentration ( $C_{max}$ ) or percentage of time  
64 (%fT), over the minimal inhibitory concentration (MIC), i.e., AUC/MIC,  $C_{max}$ /MIC or  
65 %fT/MIC, respectively <sup>11-14</sup>. In line with this, the HFS-TB can inform the design of

66 clinical trials including novel or repurposed anti-TB compounds <sup>15-17</sup>, as well as drug  
67 combinations <sup>18-20</sup>, among other possibilities. In addition, this methodology allows  
68 repetitive sample collection, is not time-limited, enables bacteria to reach high densities  
69 (similar to the ones found at the sites of infection) and lacks the ethical concerns  
70 associated with animal models, besides other advantages over alternative *in vitro* and *in*  
71 *vivo* preclinical tools <sup>21</sup>.

72 Nonetheless, the HFS-TB is a relatively novel technology, with few research groups  
73 actively using it. This situation is, in part, due to the fact that existing literature lacks  
74 standard procedures, thorough recommendations, quality control tests and/or  
75 consolidated experimental technical guidelines <sup>8,22</sup>, a predicament that, along with the  
76 high cost and infrastructure requisites needed to implement the HFS-TB, hinders further  
77 spreading to other laboratories, thereby curbing technology adoption <sup>23</sup>. One of the most  
78 relevant procedures traditionally dismissed in the field is the report of the compatibility  
79 between the drug(s) of interest and the materials throughout the system; particularly, the  
80 hollow fibers within the cartridge. The physicochemical properties of the antimicrobials  
81 (or new molecules in development) may dictate non-specific binding to said materials,  
82 and therefore they might not be able to penetrate the semi-permeable membrane and  
83 diffuse between compartments. In particular, lipophilicity could play a substantial role in  
84 this matter, since as logP increases, there is an increased probability of binding to  
85 unspecific hydrophobic targets <sup>24,25</sup>. Hence, the compatibility of the compounds with the  
86 constituents of the HFS-TB must be assessed prior to launching PKPD experiments to  
87 ensure proper reproduction of the desired PK profile at the ECS, where bacteria reside.  
88 In spite of its utmost relevancy, validation of the compatibility of the drug(s) under study  
89 with the HFS-TB and confirmation of achievement of the desired PK profile within the  
90 ECS are hardly addressed in works published on HFS-TB <sup>8</sup>.

91 In this study, we inform on the compatibility of 10 anti-TB compounds: bedaquiline  
92 (BDQ, B), clarithromycin (CLA), delamanid (DLM), ethambutol (EMB), isoniazid  
93 (INH), linezolid (LZD, L), moxifloxacin (MXF), pretomanid (PTD, Pa), rifampicin  
94 (RMP) and rifapentine (RPT) <sup>26,27</sup>, with the three most common hollow-fiber cartridges  
95 materials: polysulfone (PS), polyvinylidene fluoride (PVDF) and cellulose. We carried  
96 out multiple dosing to check drugs' compatibility, which yields a more complete  
97 summary of the molecules' behaviour in the HFS-TB than previously reported.  
98 Furthermore, we designed a robust strategy to simulate the PK of BDQ, PTD and LZD in  
99 combination (i.e., BPAL therapy for drug-resistant TB <sup>28</sup>), a challenging combination to  
100 work with in the HFS-TB due to the highly lipophilic nature of BDQ and PTD. With this  
101 information, we provide an open toolkit to ease the standardization of HFS-TB  
102 experiments and its wider implementation.

## 103 RESULTS

104 **Ethambutol and linezolid fit within the compatibility range for tested HFS-TB**  
105 **fibers.** EMB was compatible with PS, PVDF and cellulose cartridges: the  $AUC_{0-24h, ECS, ratio}$   
106 (ca. 1.2) and the  $C_{max, ss, ECS, ratio}$  (1.3, 0.8 and 1.2 for PS, PVDF and cellulose fibers,  
107 respectively) fit the required compatibility range. The drug was also cleared from the  
108 cartridge as expected, i.e., actual  $T_{1/2}$  in the ECS met established criteria (**Figure 1a,**  
109 **Table 1**). It is worth outlining that PVDF fibers, though compatible with EMB, may not  
110 be the most suitable ones to work with this anti-TB drug since  $C_{max, ss, ECS}$  was not reached  
111 until administration of the second dose (**Figure 1a, Table S1**).

112 The PK profile of LZD was also accurately simulated within the compatibility range in  
113 either type of HFS-TB cartridge, i.e.,  $AUC_{0-24h, ECS, ratio}$  (0.9, 0.9 and 0.8),  $C_{max, ss, ECS, ratio}$   
114 (1, 0.8 and 0.8) and  $T_{1/2, ECS, ratio}$  (1.3, 1.4 and 1.1) for PS, PVDF and cellulose fibers,

115 respectively (**Figure 1b, Table 1**). Although no major differences in the AUC were  
116 observed between these materials, measured  $C_{\max, ss, ECS}$  were closer to expected when  
117 using PS fibers (**Figure 1b, Table S1**). Therefore, PS cartridges stood out as the most  
118 adequate to perform HFS-TB assays with LZD.

119 Drug concentrations in the ECS for both drugs were slightly lower than those measured  
120 in the ICS during the first samplings, indicating that a short time lapse may be needed to  
121 reach equilibrium between both compartments. In contrast, these concentrations were  
122 slightly higher than those detected in the ICS during days 2 and 3, thereby suggesting  
123 some degree of accumulation (**Figure 1, Table S1**).

124 **Several anti-TB drugs fit within the compatibility range only for specific types of**  
125 **HFS-TB fibers.** INH was compatible with PS and cellulose cartridges, as endorsed by  
126 the observed PK parameters within the compatibility range:  $AUC_{0-24h, ECS, ratio}$  (1.1 and  
127 0.9),  $C_{\max, ss, ECS, ratio}$  (0.7 and 0.9) and  $T_{1/2, ECS, ratio}$  (1.3 and 1.3) for PS and cellulose,  
128 respectively. To the contrary, INH measured  $AUC_{0-24h, ECS, ratio}$  and  $C_{\max, ss, ECS}$  in PVDF  
129 cartridges did not match the criteria (0.4 and 0.4, respectively) (**Figure 2a, Table 2**).

130 The observed PK parameters for CLA and MXF fit compatibility requirements when  
131 tested with PS fibers:  $AUC_{0-24h, ECS, ratio}$  (1.1 and 1.1),  $C_{\max, ss, ECS, ratio}$  (0.9 and 1) and  $T_{1/2,$   
132  $ECS, ratio}$  (1.3 and 0.9), respectively; but not with PVDF fibers, for which their calculated  
133  $C_{\max, ss, ECS, ratio}$  (0.5 and 0.6) and  $T_{1/2, ECS, ratio}$  (3.9 and 2.6) notably differed from targeted  
134 compatibility ranges for CLA and MXF, respectively, and  $AUC_{0-24h, ECS, ratio}$  (0.7) for  
135 MXF. Likewise, cellulose fibers were outside the compatibility range with both  
136 compounds, because of divergences in  $T_{1/2, ECS, ratio}$  (1.7 and 1.6 for CLA and MXF,  
137 respectively). Given that the other PK traits were adequately imitated, these fibers might

138 still be suitable for these antimicrobial agents after proper experimental adjustments  
139 (**Figure 2b-c, Table 2**).

140 RMP was recently described not to be used with PS<sup>22</sup>. Similarly, the PK profile of RMP  
141 was not adequately mimicked in the HFS-TB when using PVDF cartridges (**Figure 2d**  
142 and **Table 2**). Most concentration values from the ECS were below the expected profile,  
143 indicating unspecific binding of the drug to the components of the system (**Table S1**).  
144 Thence,  $AUC_{0-24h, ECS, ratio}$  and  $C_{max, ss, ECS, ratio}$  were much lower than required: 0.4 and 0.2,  
145 respectively (**Table 2**). Moreover,  $T_{1/2}$  could not be calculated because no exponential  
146 decline in RMP concentrations was registered along the time course of the experiment  
147 (**Figure 2d**). Concerning cellulose fibers, RMP's  $AUC_{0-24h, ECS, ratio}$ ,  $C_{max, ss, ECS, ratio}$  and  
148  $T_{1/2, ECS, ratio}$  did not meet the compatibility range requirements either; however, their  
149 values were the nearest to fit within the ranges among all tested types of fibers: 1.6 (vs  
150 0.8-1.25), 0.6 (vs 0.7-1.3) and 2.5 (vs 0.6-1.4), respectively (**Table 2**). To note that the  
151 first RMP dose rendered a slightly higher  $C_{max, ss, ICS}$  than expected (**Figure 2d**), which  
152 could be due to minor discrepancies between the targeted and actual syringe stock. This  
153 might have counterbalanced the decline in concentration of RMP that diffusion between  
154 ICS and ECS entailed in PVDF fibers, thus rising  $C_{max, ss, ECS}$ . Overall, these results  
155 indicate that, upon certain adjustments -i.e., adding extra amount of compound and/or  
156 using top-up compensation dosing-, cellulose fibers should be the ones of choice when  
157 working with RMP in the HFS-TB. In light of this knowledge and taking into account  
158 RPT's chemical resemblance with RMP<sup>29</sup>, RPT's compatibility was only checked with  
159 cellulose fibers. As shown in **Figure 2e** and **Table 2**,  $AUC_{0-24h, ECS, ratio}$  sided with the  
160 expected: 0.8 (range 0.8-1.25), while  $C_{max, ss, ECS, ratio} = 0.6$  (range 0.7-1.3) and  $T_{1/2, ECS, ratio}$   
161  $= 1.6$  (range 0.6-1.4) were near to meet compatibility requirements, supporting the use of

162 cellulose cartridges to work with RPT in the HFS-TB. Thence, results from these assays  
163 suggest the sole validity of cellulose to test rifamycins in the HFS-TB.

164 For most of these molecules, a general tendency was discerned: concentrations in the ECS  
165 were lower than those detected in the ICS, especially in the case of drug-fiber duos outside  
166 the compatibility range -e.g., RMP-PVDF-. This may indicate that a small fraction of the  
167 compounds is unable to traverse the membrane between compartments due to binding  
168 issues. Nevertheless, there were still a few cases in which a minor degree of compound  
169 accumulation in the ECS was observed during the assay, particularly during the samplings  
170 of the third dose -e.g., CLA-cellulose- (**Figure 2, Table S1**).

171 **Sub-optimal drug-fiber compatibilities might be overcome with methodological**  
172 **adjustments.** PTD and DLM were outside the compatibility range with all fibers (**Figure**  
173 **3 and Table 3**). Most PK parameters for PTD largely differed from the targeted  
174 compatibility ranges:  $AUC_{0-24h, ECS, ratio}$  (0.3, 0.006 and 0.4),  $C_{max, ss, ECS, ratio}$  (0.2, 0.005  
175 and 0.2) and  $T_{1/2, ECS, ratio}$  (2.5, 2.5 and 1.2) for PS, PVDF and cellulose fibers, respectively.  
176 Nearly all DLM calculated ratios of PK parameters were below the expected ones:  $AUC_{0-}$   
177  $24h, ECS, ratio$  (0.001, 0.001 and 0.2) and  $C_{max, ss, ECS, ratio}$  (0.002, 0.002 and 0.1) for PS, PVDF  
178 and cellulose fibers, respectively; while  $T_{1/2, ECS, ratio}$  could only be calculated for cellulose  
179 fibers (0.9), since no exponential decline in DLM concentrations was discerned in the  
180 other two materials (**Figure 3a-b, Table 3, Table S1**). Altogether, these results placed  
181 PTD and DLM outside the compatibility range with all tested HFS-TB fibers.  
182 Nonetheless, it could be ascertained that cellulose fibers were the ones that gave rise to  
183 PK profiles with the highest similarity to the targeted ones. In fact, their  $T_{1/2, ECS, ratio}$   
184 entered the required range for both compounds, which suggests that the unbound drug  
185 fraction was being accurately cleared. Ergo, upon further optimization, cellulose might  
186 be the fibers of choice to work with PTD and DLM in the HFS-TB.

187 As a first approach, the fiber's compatibility of BDQ was tested with a non-clinical single  
188 dose with a  $C_{max}$  holding step in PS and PVDF fibers, being outside the compatibility  
189 range with both fibers (**Figure 3c** and **Table 3**).  $AUC_{0-24h, ECS, ratio}$  (0.002) and  $C_{max, ss, ECS, ratio}$   
190 (0.006) were below the compatibility range in PVDF fibers, whereas  $T_{1/2, ECS, ratio}$  in  
191 this cartridge and all PK parameters in PS fibers could not be calculated, because most  
192 samplings, especially in the ECS, were below the limit of quantification of the LC-  
193 MS/MS (**Table S1**). These results posed BDQ as the most challenging compound in our  
194 compilation to work with in the HFS-TB.

195 In an attempt to overcome these challenges, and considering the extended terminal half-  
196 life of BDQ in the clinic<sup>30</sup>, a modified experimental approach was designed to detect  
197 BDQ in the system and optimize its PK profile. First, the use of polypropylene and  
198 silicone materials was minimized to avoid unspecific binding<sup>31,32</sup>. Second, data from the  
199 compatibility tests and the TB-Platform for the Aggregation of Preclinical Experiments  
200 Data (TB-APEX)<sup>33</sup> were used to calculate the amount of drug and number of infusions  
201 needed to reach the targeted PK. In that sense, BDQ median concentration at protein-  
202 unbound steady state in the lung ( $C_{med, ss} = 0.061 \mu\text{g/mL}$ ) was aimed instead of the usual  
203 PK parameters ( $AUC_{0-24h}$ ,  $C_{max, ss}$  and  $T_{1/2}$ )<sup>30</sup>. Third, BDQ was directly infused in the  
204 ECS, in order to circumvent drug loss through the barrier between compartments. With  
205 this approach, the  $C_{med, ss}$  was accurately imitated in PVDF fibers (0.082  $\mu\text{g/mL}$ ), unlike  
206 in PS fibers (0.008  $\mu\text{g/mL}$ ) and cellulose ones (0.571  $\mu\text{g/mL}$ ) (**Figure 4** and **Table S2**).  
207 Nevertheless, cellulose showed higher BDQ concentrations than targeted ones. This  
208 suggests that the binding that was taken into consideration to program drug infusions was  
209 overestimated in cellulose, which is in agreement with the more promising behaviour of  
210 PTD and DLM within these fibers (**Figure 3a-b**). Fluctuations in BDQ levels were  
211 registered in all fibers, but most measurements in PVDF and cellulose fit within a close

212 range: 0.036 to 0.090  $\mu\text{g/mL}$  and 0.200 to 1.287  $\mu\text{g/mL}$ , respectively. Hence, though  
213 further adjustments would be needed to refine the PK profile of BDQ in the HFS-TB, this  
214 approach could help to pave the way towards the use of lipophilic compounds in the HFS-  
215 TB.

216 **Pharmacokinetic profile mimicking of the BPaL combination in the HFS-TB.** We  
217 took advantage of our assay toolkit for anti-TB drugs in the HFS-TB to design an optimal  
218 setup for BPaL testing. As a proof-of-concept, we targeted a 3-days PK profile of BDQ,  
219 PTD and LZD in a cellulose cartridge, based on our previously generated monotherapy  
220 data. An unbound  $C_{\text{med, ss}}$  in the lung was aimed for BDQ, whilst PTD's PK profile was  
221 targeted by infusing higher amounts of drug and top-up compensations, in order to  
222 overcome unspecific binding and countervail LZD's clearance rate, respectively.

223 As displayed in **Figure 5**,  $C_{\text{med, ss}}$  of BDQ was correctly reproduced: 0.050  $\mu\text{g/mL}$  (*vs*  
224 0.061  $\mu\text{g/mL}$ ), and most measurements fit within a small range: 0.014 to 0.099  $\mu\text{g/mL}$   
225 (**Table S3**). Likewise, LZD's PK parameters fit compatibility requirements:  $\text{AUC}_{0-24\text{h}}$ ,  
226  $\text{ECS}_{\text{ratio}} = 1.2$  (0.8-1.25),  $C_{\text{max, ss, ECS, ratio}} = 0.8$  (0.7-1.3) and  $T_{1/2, \text{ECS, ratio}} = 1.2$  (0.6-1.4)  
227 (**Table 1**). Conversely, PTD's  $T_{1/2, \text{ECS, ratio}}$  sided with the expected (0.8 *vs* 0.6-1.4), but  
228  $\text{AUC}_{0-24\text{h, ECS, ratio}} = 0.5$  (*vs* 0.8-1.25) and  $C_{\text{max, ss, ECS, ratio}} = 0.4$  (*vs* 0.7-1.3) were outside  
229 the compatibility range. However, these values were near our requirements and notably  
230 improved the results from PTD's compatibility test alone (**Table 3**). Given that  $T_{1/2, \text{ECS}}$   
231 was accurately mimicked,  $\text{AUC}_{0-24\text{h, ECS}}$  and  $C_{\text{max, ss, ECS}}$  could be easily boosted by  
232 increasing PTD's concentration in each infusion.

## 233 **DISCUSSION**

234 The HFS-TB has become a useful *in vitro* preclinical tool to characterize PKPD profiles  
235 of new and repurposed antibiotics and regimens for TB treatment, as well as informing

236 the design of prospective clinical trials<sup>6</sup>. However, access to essential information needed  
237 for inter-laboratory experimental reproduction of results has been elusive. Indeed, recent  
238 publications in the field have emphasized the urgent need for standard procedures and  
239 experimental specifications in HFS-TB reports, to facilitate laboratories to *de novo*  
240 implement this technology<sup>8,22,34</sup>. Delivering this information will increase capacity and  
241 provide confidence in published data, allowing researchers to reproduce them.

242 The lack of information on the compatibility between drugs and the different types of  
243 hollow fibers stands out as a major limitation in the use of the HFS-TB. In this sense, it  
244 is relevant to stress that not all molecules are compatible with the materials the cartridges  
245 are made of: they may bind to the hollow fibers, plastics or other components of the  
246 system; and/or they may not be able to penetrate the semi-permeable membrane of said  
247 fibers. Therefore, as advised by Aguilar-Ayala *et al.*<sup>22</sup>, compatibility tests for the drug(s)  
248 of interest in the absence of bacteria should be routinely assessed and reported prior to  
249 conducting PKPD experiments in the HFS-TB. To shorten this gap, in this work, we  
250 carried out compatibility tests with the three most common hollow-fiber materials, i.e.,  
251 PS, PVDF and cellulose, and a catalogue of 10 drugs including representatives of distinct  
252 anti-TB families, so as to span a wide window of chemical features: ethanolamines  
253 (EMB), diarylquinolines (BDQ), INH, macrolides (CLA), nitroimidazoles (PTD, DLM),  
254 oxazolidinones (LZD), quinolones (MXF) and rifamycins (RMP, RPT) (**Figure 6**).  
255 Further, a currently prominent three-drug combination in the anti-TB field (BPaL,  
256 featuring BDQ, PTD and LZD) was assayed in the system upon the information gathered  
257 from the drug-fiber compatibility tests of each antibiotic alone. A specific PK profile of  
258 each antimicrobial was targeted in the system in absence of bacteria, and its accuracy was  
259 verified by measuring drug levels in the ICS and ECS over time. Compatibility ranges  
260 were established based on obtained  $AUC_{0-24h}$ , ECS,  $C_{max, ss, ECS}$  and  $T_{1/2, ECS}$  compared to

261 the expected parameters <sup>35</sup>. It is of note that drug concentrations in the ECS have been  
262 scarcely reported in the field, despite being the compartment where bacteria actually face  
263 the compound(s) and the actual pharmacodynamic interaction occurs.

264 Our results indicate the following: (i) EMB and LZD fit within the compatibility range  
265 with the three types of fibers; (ii) INH with PS and cellulose, but not with PVDF; (iii)  
266 CLA and MXF only with PS; (iv) rifamycins (RMP, RPT) could only be used with  
267 cellulose; (v) PTD and DLM were outside the compatibility range with all fibers, but their  
268 PK profiles in cellulose cartridges appeared more promising upon further modifications;  
269 (vi) BDQ was outside the compatibility range with PS and PVDF fibers, but alternative  
270 approaches could be implemented to overcome this issue (**Figures 1-4 and 6, Tables 1-**  
271 **3**). These results accentuate the importance of deciphering the suitability of each  
272 compound with the HFS-TB cartridges before performing PKPD studies, as many of them  
273 were unfit to be used with at least some of the fibers' materials. We could also infer that  
274 PVDF cartridges are the least user-friendly, since they were outside the compatibility  
275 range with 7 out of 9 compounds (BDQ, CLA, DLM, INH, MXF, PTD and RMP). PS  
276 fibers were the most suitable for 5 out of 9 compounds tested (CLA, EMB, INH, LZD  
277 and MXF). In contrast, cellulose cartridges seem to be the most versatile ones; although  
278 only 3 out of 10 compounds were within the compatibility range, they would allow  
279 working with of all the compounds in our panel -upon infusions' adjustments- with  
280 different degrees of lipophilicity (**Figure 6**).

281 HFS-TB studies have traditionally only reported drug levels at the ICS, thereby  
282 disregarding the ECS, where bacteria are effectively exposed to the drug <sup>34,36,37</sup>. This  
283 work, along with Aguilar-Ayala *et al.* <sup>22</sup>, reinforces the fact that there are usually non-  
284 negligible discrepancies between the drug concentrations in the ICS and the lumen of the  
285 cartridges (i.e., ECS), being the latter the one that should be rigorously reported. These

286 permeability issues might be due to the pore's and fiber's inner diameter, the membrane  
287 surface area or the effective length of the tubing, among other factors related to the system  
288 itself<sup>38</sup>; but also to the hydrophilicity/lipophilicity attributes of the compounds and the  
289 components of the system.

290 Regarding this hydrophilicity/lipophilicity dichotomy, the general tendency we observed  
291 is that molecules with the lowest logP and highest water solubility showed less binding  
292 to the fibers and free diffusion between the ICS and the ECS (**Figure 6**). This applied to  
293 EMB and LZD, with low experimental and/or predicted logP (0.4 and 0.9 for EMB and  
294 LZD, respectively), which is usually associated with low lipophilicity<sup>39</sup>. Their water  
295 solubility is high as well: 1,000 mg/mL and 3 mg/mL, respectively (**Table S4**). INH also  
296 shows high hydrophilicity (logP = -0.7) and water solubility (140 mg/mL), but did not fit  
297 within the compatibility range with PVDF fibers. This is likely because the concentrations  
298 quantified in the first samplings were below the expected ones, thus rendering an overall  
299  $AUC_{0-24h, ECS, ratio}$  and  $C_{max, ss, ECS, ratio}$  that deviated from the required criteria (**Figure 2a**,  
300 **Table 2**). By adjusting the amount of INH in the first dosing to compensate for this delay  
301 in the equilibrium between compartments, the INH-PVDF drug-fiber duo might be turned  
302 into a compatible pair for HFS-TB experiments. Nonetheless, only INH-PS or INH-  
303 cellulose duos have been published to date<sup>40,41</sup>, so this hypothesis should be further  
304 confirmed. In contrast, drugs with medium-high lipophilicity (e.g., PTD and DLM:  
305 predicted logP = 4.14 and 6.14 and water solubility =  $<1$  mg/mL and 0.0022 mg/mL,  
306 respectively; **Table S4**) did not meet the compatibility ranges for most PK parameters  
307 and hardly passed through the fibers' membrane, probably due to unspecific binding.  
308 Interestingly, they were prone to behave better in cellulose (or PS, in the case of PTD)  
309 than in PVDF, for which a sizable degree of binding was ascertained (**Figure 3**, **Table 3**,  
310 **Table S1**). This may be justified by the lipophilicity of the fibers themselves: PVDF

311 appears to be the most hydrophobic, whereas cellulose would be the most hydrophilic,  
312 followed by PS<sup>42,43</sup>. As a consequence, it is not unforeseen that cellulose worked better  
313 with BDQ, DLM, PTD, RMP or RPT (which are among those with the highest logP)  
314 because a hydrophilic ambience may prevent hydrophobic compounds from sticking to  
315 inner surfaces<sup>44</sup>. In brief, our work supports these observations and suggest that  
316 deepening into the chemical properties -mainly logP and water solubility- of the drug(s)  
317 under study prior to performing HFS-TB assays could aid the selection of the most  
318 accurate cartridge to work with. Nevertheless, cellulose fibers outstand as the most  
319 versatile choice when dealing with highly lipophilic molecules.

320 It is worth highlighting that some PK parameters of RMP, RPT and, specially, BDQ,  
321 DLM and PTD were not successfully mimicked in any of the cartridges tested (**Tables 2-**  
322 **3**), which according to our criteria placed them outside the compatibility range. In fact,  
323 as depicted in **Figure 3**, a sizable amount of compound was lost when crossing the semi-  
324 permeable hindrance between the ICS and the ECS. Several strategies could be adopted  
325 to overcome this hurdle, as the ones we applied to BDQ: (i) rising the amount of  
326 compound in each infusion, (ii) using compensating infusions to keep a median  
327 physiological concentration and (iii) adding the drug directly into the ECS to avoid the  
328 barrier between compartments. In addition, hydrophilicity performance of the fibers can  
329 be enhanced, which might reduce the unspecific binding of hydrophobic molecules. This  
330 has been previously addressed with PS hollow-fiber membranes, using calcium alginate,  
331 zinc oxide or graft polymerization with 2-hydroxyethylacrylate, among other possibilities  
332<sup>45-48</sup>; albeit, extrapolation to hollow-fiber cartridges would require further research.  
333 Oppositely, a hydrophobic coating may saturate the unspecific binding sites of the HFS-  
334 TB materials and, as a consequence, prevent drug levels to decline. This strategy was  
335 employed by Raaijmakers J. *et al*<sup>31</sup> using clofazimine to block the binding sites of BDQ.

336 However, this alternative was ruled out in this study since clofazimine has antibacterial  
337 activity against *M. tuberculosis* <sup>49</sup>.

338 In this work, we propose criteria to decide whether a drug is within the compatibility  
339 range or not with a certain type of hollow-fiber cartridge, based on three primary PK  
340 parameters in the ECS ( $AUC_{0-24\text{ h}}$ ,  $C_{\text{max, ss}}$  and  $T_{1/2}$ ). This classification lies within a  
341 methodological perspective that aims to inform on the most suitable cartridges when  
342 working with certain compounds in the system. Albeit, it should be remarked that HFS-  
343 TB experiments are often used to feed *in silico* models to inform the design of clinical  
344 trials <sup>10</sup>. This means that modelling might still benefit from data derived from a duo  
345 outside the compatibility range, e.g., INH-PVDF, CLA-Cellulose or MXF-Cellulose.  
346 Many drugs included in this study had AUC/MIC as their PKPD driver for antibacterial  
347 efficacy <sup>50-53</sup>, being thus  $AUC_{0-24\text{ h}}$ , ECS, ratio a credible parameter to validate their PK;  
348 however, murine models have described PTD activity as time-dependent and its  
349 bactericidal activity best associated with the percentage of the dosing interval in which  
350 free drug concentration surpasses the MIC (%fT/MIC) <sup>54</sup>. Being 0.125-0.25  $\mu\text{g/mL}$  the  
351 MIC to PTD of *M. tuberculosis* H37Rv strain <sup>55</sup>, we were able to kept a 100% fT/MIC to  
352 PTD in our compatibility tests with PS and cellulose fibers. This indicates that, depending  
353 on the research goal, PTD may be used in the HFS-TB to provide useful PKPD outputs  
354 despite the technical quandary this drug entails. As a final note, we ought to state that  
355 preliminary compatibility tests do not replace the need to report further PK measurements  
356 during actual HFS-TB PKPD experiments with bacteria <sup>56</sup>.

357 BPAL therapy has dramatically changed the management of drug resistant TB <sup>28</sup>.  
358 However, to our knowledge, only a few publications have reported BPAL results in the  
359 HFS-TB <sup>16,20</sup>. In fact, an accurate reproduction of its combined PK profile poses a  
360 significant challenge, due to the highly lipophilic nature of BDQ and PTD. By leveraging

361 on our drug-fiber compatibility studies, we obtained information on their behavior within  
362 each cartridge, defined adjustments to refine their PK profiles and developed a robust  
363 strategy to simulate a combined PK profile of BDQ, PTD and LZD (**Figure 4**). In spite  
364 of PTD not fitting the compatibility range for 2 out of 3 PK parameters (**Table 3**), our  
365 results proved the importance of previous drug-alone tests, being accurate enough to  
366 facilitate the implementation of this combination in the HFS-TB. Since BPaL is one of  
367 the most important TB treatments against drug-resistant TB, and with the emerging  
368 concern of BDQ-resistant strains presenting mutations in the *rv0678c* gene <sup>57</sup>, the  
369 methodological improvements here reported hold major significance to aid in the  
370 optimization of the treatment against BDQ-resistant strains.

371 In summary, in this work we provide a comprehensive analysis of the suitability of the  
372 most common hollow-fiber materials to be used in the HFS-TB with anti-TB drugs to  
373 date. Our data show that several types of fibers are not suitable to work with certain  
374 compounds, and that measuring the actual drug concentrations that bacteria face inside  
375 the ECS (which are usually lower than expected) is crucial, underpinning that drugs  
376 should be tested for compatibility with the HFS-TB before carrying out a PKPD assay,  
377 either in monotherapy or combinations. We also provide a theoretical rationale to pre-  
378 select the cartridge material to be used for each new drug, which is based on a relationship  
379 between the hydrophilicity/lipophilicity of the molecules and the compatibility with  
380 fibers' materials. We hope that this report is of use to other laboratories trying to  
381 implement and optimize HFS-TB PKPD experiments. We also encourage other  
382 laboratories working with the HFS-TB to readily disclose this type of information for  
383 additional compounds not here tested. This collaborative framework would make a  
384 noteworthy step towards reaching best-practice principles that this preclinical technique  
385 should own to maximize its role in informing the design of Phase II/III clinical trials.

**386 RESOURCE AVAILABILITY****387 Lead contact**

388 Requests for further information and resources should be directed to and will be fulfilled  
389 by the lead contact, Fernando Sanz-García ([f.sanzg@unizar.es](mailto:f.sanzg@unizar.es)).

**390 Materials availability**

391 This study did not generate new unique reagents.

**392 Data and code availability**

- 393 • Data reported in this paper will be shared by the lead contact upon request.
- 394 • This paper does not report original code.
- 395 • Any additional information required to reanalyze the data reported in this paper is  
396 available from the lead contact upon request.

**397 Limitations of the study**

398 Our compatibility results were mostly based on PK profiles in the lung, which were  
399 derived from specific dosing chosen among the most commonly used for each antibiotic.  
400 However, depending on the research question, special situations and patient populations  
401 -e.g., children- might require alternative dosing strategies or mimicking PK profiles in  
402 different compartments -e.g., plasma *vs.* lung-. If these differences entailed major changes  
403 in expected drug concentrations, the compatibility of the molecule with the HFS-TB  
404 fibers might differ, assuming its binding is concentration-dependent. Thus, we advise  
405 users to ensure desired PK profiles are properly mimicked in their experimental setup.  
406 Although PS, PVDF and cellulose are by far the main fibers' materials used in the HFS-  
407 TB field, other options could have been examined to expand our assay toolkit, -e.g.,  
408 polyethersulfone cartridges<sup>58</sup>-, especially when facing highly lipophilic compounds.

409 Similarly, we have limited our study to some representative anti-TB drugs; other less  
410 commonly used or currently under development were not included in our study because  
411 of lack of access to the compound or capacity limitations. However, we hope that, by  
412 assaying this initial “toolkit”, we contribute to the creation of a working framework that  
413 could enhance essential data accessibility to robustly performed HFS-TB studies.  
414 Additional contributions from the field would enlarge this assay toolkit for HFS-TB with  
415 more antibiotics.

#### 416 **ACKNOWLEDGEMENTS**

417 We would like to thank the technical support of Ana Picó Marco from the University of  
418 Zaragoza, and the recommendations on targeted PK parameters for BPaL combination of  
419 Evangelos Karakitsios, from Institute for Applied Computing “Mauro Picone”, Consiglio  
420 Nazionale delle Ricerche.

421 This work has received support from the Innovative Medicines Initiatives 2 Joint  
422 Undertaking (grant No 853989) to NW and SRG. The JU receives support from the  
423 European Union’s Horizon 2020 Research and Innovation Programme and EFPIA and  
424 Global Alliance for TB Drug Development Non-Profit Organisation, Bill & Melinda  
425 Gates Foundation, University of Dundee.

#### 426 **AUTHOR’S CONTRIBUTIONS**

427 CRediT (Contributor Roles Taxonomy) has been applied for author contribution.

428 Conceptualization: FSG, AL, SRG; Data curation: FSG; Formal Analysis: FSG, MREM  
429 DN; Funding acquisition: NW, SRG; Investigation: FSG, DAAA, MREM, DN, AB, SA,  
430 NI; Methodology: FSG, DAAA, AML, AL, SRG; Project administration: NW, AL, SRG;  
431 Supervision: NW, AL, SRG; Visualization: FSG, DAAA; Writing – original draft: FSG,

432 AL, SRG; Writing – review & editing: FSG, DAAA, MREM, DN, AL, SRG. All authors  
433 read and approved the final version of the document.

434 **DECLARATION OF INTERESTS**

435 The authors declare no competing interests.

436

437

438

439

440

441

442

443

444

445

446

447

448

449

450

451

452

Journal Pre-proof

453 **Figure 1 | Drug-fiber compatibility tests of ethambutol and linezolid in the HFS-TB.**  
454 Expected (dashed line) and observed (solid lines) pharmacokinetic (PK) profiles of (a)  
455 ethambutol and (b) linezolid, sampled from the intra-capillary space (ICS; green line) and  
456 extra-capillary space (ECS; purple line) of polysulfone (PS), polyvinylidene fluoride  
457 (PVDF) and cellulose HFS-TB cartridges. Targeted PK parameters for each antibiotic are  
458 described in **Table 4**. Drug concentration values are provided in **Table S1**. EMB:  
459 ethambutol, LZD: linezolid.

460 **Figure 2 | Drug-fiber compatibility tests of isoniazid, clarithromycin, moxifloxacin,**  
461 **rifampicin and rifapentine in the HFS-TB.** Expected (dashed line) and observed (solid  
462 lines) pharmacokinetic (PK) profiles of (a) isoniazid, (b) clarithromycin, (c)  
463 moxifloxacin, (d) rifampicin and (e) rifapentine, sampled from the intra-capillary space  
464 (ICS; green line) and extra-capillary space (ECS; purple line) of polysulfone (PS),  
465 polyvinylidene fluoride (PVDF) and cellulose HFS-TB cartridges. Targeted PK  
466 parameters for each antibiotic are described in **Table 4**. Drug concentration values are  
467 provided in **Table S1**. CLA: clarithromycin; INH: isoniazid; MXF: moxifloxacin; RMP:  
468 rifampicin; RPT: rifapentine.

469 **Figure 3 | Drug-fiber compatibility tests of pretomanid, delamanid and bedaquiline**  
470 **in the HFS-TB.** Expected (dashed line) and observed (solid lines) pharmacokinetic (PK)  
471 profiles of (a) pretomanid, (b) delamanid and (c) bedaquiline, sampled from the intra-  
472 capillary space (ICS; green line) and extra-capillary space (ECS; purple line) of  
473 polysulfone (PS), polyvinylidene fluoride (PVDF) and cellulose HFS-TB cartridges.  
474 Targeted PK parameters for each antibiotic are described in **Table 4**. Drug concentration  
475 values are provided in **Table S1**. BDQ: bedaquiline; DLM: delamanid; PTD: pretomanid.

476 **Figure 4 | Optimization approach to use bedaquiline in the HFS-TB.** Observed  
477 pharmacokinetic (PK) profiles of bedaquiline, sampled from the extra-capillary space

478 (ECS) of polysulfone (PS), polyvinylidene fluoride (PVDF) and cellulose HFS-TB  
479 cartridges. The targeted median concentration at protein-unbound steady state in the lung  
480 ( $C_{med, ss} = 0.061 \mu\text{g/mL}$ ) is indicated in **Table 4** and represented in the graphs by a dashed  
481 line. Drug concentration values are provided in **Table S2**. BDQ: bedaquiline.

482 **Figure 5 | Optimization approach to use bedaquiline-pretomanid-linezolid (BPaL)**  
483 **combination in the HFS-TB.** Expected (dashed lines) and observed (solid purple lines)  
484 pharmacokinetic (PK) profiles of a combination of bedaquiline, pretomanid and linezolid  
485 (i.e., BPaL) sampled from the extra-capillary space (ECS) of a cellulose HFS-TB  
486 cartridge. The targeted median concentration at protein-unbound steady state in the lung  
487 of BDQ ( $C_{med, ss} = 0.061 \mu\text{g/mL}$ ) is indicated in **Table 4** and represented in the graph by  
488 a straight dashed line. Likewise, targeted PK parameters for pretomanid and linezolid are  
489 described in **Table 4**. Drug concentration values are provided in **Table S3**. BDQ:  
490 bedaquiline, PTD: pretomanid, LZD: linezolid.

491 **Figure 6 | Schematic drug-fiber compatibility status of anti-TB drugs in the HFS-**  
492 **TB.** Anti-TB drugs are ordered based on their lipophilicity. Experimental and predicted  
493 logP values are encompassed in **Table S4**. PS: polysulfone; ND: not determined; BDQ:  
494 bedaquiline; CLA: clarithromycin; DLM: delamanid; EMB: ethambutol; INH: isoniazid;  
495 LZD: linezolid; MXF: moxifloxacin; PTD: pretomanid; RMP: rifampicin; RPT:  
496 rifapentine.

497 \*Optimization approach showed in **Figure 4** rendered cellulose as the fibers with less  
498 binding to BDQ.

499

500

501

502 **STAR★METHODS**503 • **Key resources table**504 • **Method details**

505 - Culture media and drugs

506 - HFS experimental set up

507 - Mimicking a pharmacokinetic profile in the HFS for drug-fiber compatibility tests

508 - Optimization approach to mimic a pharmacokinetic profile of bedaquiline in the  
509 HFS-TB510 - Optimization approach to mimic a pharmacokinetic profile of bedaquiline-  
511 pretomanid-linezolid (BPaL) combination in the HFS-TB

512 - Preparation of drug samples and drug quantification

513 • **Quantification and statistical analysis**514 **SUPPLEMENTAL INFORMATION**515 **Document S1.** Figure S1, Tables S2-S6 and supplemental references.516 **Table S1.**

517 **Table 1. Comparison of pharmacokinetic parameters of different drugs within the compatibility range for three types of HFS-TB fibers.**

Drug	Expected PK parameters			PS						PVDF						Cellulose					
	AUC <sub>0-24h</sub> ( $\mu\text{g}\cdot\text{h}/\text{mL}$ )	C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ )	T <sub>1/2</sub> (h)	AUC <sub>0-24h</sub> ( $\mu\text{g}\cdot\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)		AUC <sub>0-24h</sub> ( $\mu\text{g}\cdot\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)		AUC <sub>0-24h</sub> ( $\mu\text{g}\cdot\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)	
				ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS
EMB	24	3	3 <sup>a</sup>	32	31	4	4	3.3	3.6	22.5	29.6	3.1	2.5	3.4	3.9	25	28.8	3.7	3.6	3.3	4.1
			(1.2)	(1.2)	(1.3)	(1.3)	(1.1)	(1.2)	(0.9)	(1.2)	(1)	(0.8)	(1.1)	(1.3)	(1.0)	(1.2)	(1.2)	(1.2)	(1.1)	(1.4)	
			12 <sup>a</sup>					13.9	8.7					9.2	9					10.5	9
				(1.2)	(1.2)	(1.3)	(1.3)	(1.2)	(0.7)					(0.8)	(0.8)					(0.9)	(0.8)
LZD (alone)	57.8	8.6	3.5	55.3	54.7	8.5	8.6	4.6	4.7	53.9	52.8	7.6	6.6	5	4.9	52	46.2	8.5	6.7	4	3.8
				(0.9)	(0.9)	(1)	(1)	(1.3)	(1.3)	(0.9)	(0.9)	(0.9)	(0.8)	(1.4)	(1.4)	(0.9)	(0.8)	(1)	(0.8)	(1.1)	(1.1)
LZD (BPaL)				NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	13.7 <sup>b</sup>	NA	1.3 <sup>b</sup>	NA	4.5
																	(1.2)	(0.8)		(1.2)	

518

519 EMB: ethambutol, LZD: linezolid, BPaL: bedaquiline-pretomanid-linezolid. PS: polysulfone, PVDF: polyvinylidene fluoride, SS: steady state,

520 ICS: intra-capillary space, ECS: extra-capillary space. AUC ratio is calculated as  $AUC_{0-24h, \text{real}}/AUC_{0-24h, \text{expected}}$ . C<sub>max, ss</sub> ratio is calculated as C<sub>max,</sub>521 <sub>ss, real</sub>/ C<sub>max, ss, expected</sub>, being C<sub>max, ss, real</sub> an average of the three C<sub>max</sub> reached during the dosing. T<sub>1/2</sub> ratio is calculated as T<sub>max, 1/2, real</sub>/ T<sub>1/2 expected</sub>, being

522  $T_{\max, 1/2, \text{real}}$  an average of the three  $T_{1/2}$  calculated for each clearance after dosing. Drug concentration values are provided in **Table S1** for drugs  
523 alone and **Table S3** for LZD in the BPaL combination.

524 <sup>a</sup>EMB's PK is best described by a two-compartment model:  $T_{1/2,\alpha, 0-12 \text{ h}} = 3 \text{ h}$  and  $T_{1/2,\beta, 12-24 \text{ h}} = 12 \text{ h}$ .

525 <sup>b</sup>Due to changes detected in the syringe stock, LZD's expected PK parameters in BPaL were re-calculated as:  $AUC_{0-24\text{h}} = 11 \mu\text{g}\cdot\text{h}/\text{mL}$  and  $C_{\max, \text{ss}}$   
526  $= 1.6 \mu\text{g}/\text{mL}$ .

527

Journal Pre-proof

528 **Table 2. Comparison of pharmacokinetic parameters of different drugs within the compatibility range with specific types of HFS-TB**  
 529 **fibers.**

Drug	Expected PK parameters			PS						PVDF						Cellulose					
	AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ )	C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ )	T <sub>1/2</sub> (h)	AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)		AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)		AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)	
				ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS
CLA	30.7	3.5	4	37.1 (1.2)	34.3 (1.1)	3.5 (1)	3.1 (0.9)	5.1 (1.3)	5.4 (1.3)	26.6 (0.8)	27.3 (0.9)	2.7 (0.8)	1.6 (0.5)	5.8 (1.5)	15.5 (3.9)	33.7 (1.1)	35.9 (1.2)	3.5 (1)	2.8 (0.8)	4.7 (1.2)	6.9 (1.7)
INH	39.6	6.8	3	43.6 (1.1)	42.6 (1.1)	6.8 (1)	4.8 (0.7)	4.2 (1.4)	3.9 (1.3)	23.7 (0.6)	17.4 (0.4)	5.6 (0.8)	2.7 (0.4)	3.5 (1.2)	3.3 (1.1)	34.9 (0.8)	38.2 (0.9)	5.5 (0.8)	5.8 (0.9)	3.3 (1.1)	3.8 (1.3)
MXF	45.4	3.1	13	50 (1.1)	49.9 (1.1)	3.3 (1.1)	3.2 (1)	12 (0.9)	11.5 (0.9)	36.2 (0.8)	32.5 (0.7)	2.4 (0.8)	1.8 (0.6)	32.6 (2.5)	33.7 (2.6)	42.1 (0.9)	41.3 (0.9)	2.8 (0.9)	2.3 (0.7)	18.7 (1.4)	20.4 (1.6)
RMP	12.5	2.05	2.5	NA	NA	NA	NA	NA	NA	7.6 (0.6)	5 (0.4)	1.3 (0.6)	0.5 (0.2)	4.6 (1.8)	- <sup>a</sup>	22.5 (1.8)	19.8 (1.6)	2.7 (1.3)	1.2 (0.6)	3.6 (1.4)	6.3 (2.5)
RPT	12.5	2	2.5	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	10 (0.9)	9.7 (0.8)	1.6 (0.8)	1.2 (0.6)	3.2 (1.3)	4 (1.6)

530

531 CLA: clarithromycin, INH: isoniazid, MXF: moxifloxacin, RMP: rifampicin, RPT: rifapentine, BPAL: bedaquiline-pretomanid-linezolid. PS:  
532 polysulfone, PVDF: polyvinylidene fluoride, SS: steady state, ICS: intra-capillary space, ECS: extra-capillary space, NA: not applicable. AUC  
533 ratio is calculated as  $AUC_{0-24h, real}/AUC_{0-24h, expected}$ .  $C_{max, ss}$  ratio is calculated as  $C_{max, ss, real}/C_{max, ss, expected}$ , being  $C_{max, ss, real}$  an average of the three  
534  $C_{max}$  reached during the dosing.  $T_{1/2}$  ratio is calculated as  $T_{max, 1/2, real}/T_{1/2, expected}$ , being  $T_{max, 1/2, real}$  an average of the three  $T_{1/2}$  calculated for each  
535 clearance after dosing. Drug concentration values are provided in **Table S1**.

536 <sup>a</sup>  $T_{1/2}$  could not be calculated because the actual PK profile did not follow the expected exponential decline.

537

538 **Table 3. Comparison of pharmacokinetic parameters of drugs outside the compatibility range with tested HFS-TB fibers.**

Drug	Expected PK parameters			PS						PVDF						Cellulose					
	AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ )	C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ )	T <sub>1/2</sub> (h)	AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)		AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)		AUC <sub>0-24h</sub> ( $\mu\text{g}^*\text{h}/\text{mL}$ ) (ratio)		C <sub>max, ss</sub> ( $\mu\text{g}/\text{mL}$ ) (ratio)		T <sub>1/2</sub> (h) (ratio)	
				ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS	ICS	ECS
BDQ	8.9	0.5	17.8	- <sup>a</sup>	- <sup>a</sup>	- <sup>a</sup>	- <sup>a</sup>	- <sup>b</sup>	- <sup>a</sup>	0.3 (0.03)	0.02 (0.002)	0.06 (0.1)	0.003 (0.006)	8.1 (0.4)	- <sup>b</sup>	NA	NA	NA	NA	NA	NA
DLM	7.9	0.5	30	0.2 (0.02)	0.01 (0.001)	0.02 (0.04)	0.0009 (0.002)	- <sup>b</sup>	- <sup>b</sup>	0.6 (0.08)	0.01 (0.001)	0.08 (0.2)	0.001 (0.002)	- <sup>b</sup>	- <sup>b</sup>	5.7 (0.7)	1.7 (0.2)	0.3 (0.6)	0.05 (0.1)	20.4 (0.7)	26.5 (0.9)
PTD (alone)	32.7	2	18	21.2 <sup>c</sup> (0.5)	11.3 <sup>c</sup> (0.3)	1 <sup>c</sup> (0.5)	0.4 <sup>c</sup> (0.2)	39.7 (2.2)	44.5 (2.5)	1.4 (0.04)	0.2 (0.006)	0.2 (0.1)	0.01 (0.005)	16.3 (0.9)	45.4 (2.5)	27.6 <sup>c</sup> (0.5)	15.9 <sup>c</sup> (0.3)	1.7 <sup>c</sup> (0.9)	0.7 <sup>c</sup> (0.2)	20.7 (1.2)	21.9 (1.2)
PTD (BPAL)	70.5	4.3		NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	NA	30.3 (0.4)	NA	2.2 (0.5)	NA

539

540 BDQ: bedaquiline, DLM: delamanid, PTD: pretomanid, BPAL: bedaquiline-pretomanid-linezolid. PS: polysulfone, PVDF: polyvinylidene fluoride,

541 SS: steady state, ICS: intra-capillary space, ECS: extra-capillary space, NA: not applicable. AUC ratio is calculated as  $\text{AUC}_{0-24h, \text{real}}/\text{AUC}_{0-24h, \text{expected}}$ .542  $\text{C}_{\text{max, ss}}$  ratio is calculated as  $\text{C}_{\text{max, ss, real}}/\text{C}_{\text{max, ss, expected}}$ , being  $\text{C}_{\text{max, ss, real}}$  an average of the three (PTD) and two (DLM)  $\text{C}_{\text{max}}$  reached during

543 the dosing, or the five samplings taken during the  $C_{\max}$  holding step (BDQ).  $T_{1/2}$  ratio is calculated as  $T_{\max, 1/2, \text{real}} / T_{1/2 \text{ expected}}$ , being  $T_{\max, 1/2, \text{real}}$  an  
544 average of the three (PTD), two (DLM) or one (BDQ)  $T_{1/2}$  calculated for each clearance after dosing. Drug concentration values are provided in  
545 **Table S1** for drugs alone and **Table S3** for PTD in the BPaL combination.

546 <sup>a</sup>PK parameters could not be calculated because drug concentrations were below the limit of quantification

547 <sup>b</sup> $T_{1/2}$  could not be calculated because the actual PK profile did not follow the expected exponential decline.

548 <sup>c</sup>Due to changes detected in the syringe stocks, PTD-PS and PTD-cellulose compatibility test's expected PK parameters were re-calculated as:

549  $AUC_{0-24h} = 40.6 \mu\text{g}\cdot\text{h}/\text{mL}$ ,  $C_{\max,ss} = 2.5 \mu\text{g}/\text{mL}$ ; and  $AUC_{0-24h} = 53.2 \mu\text{g}\cdot\text{h}/\text{mL}$ ,  $C_{\max,ss} = 3.1 \mu\text{g}/\text{mL}$ ; respectively.

550

551 **Table 4. Summary of pharmacokinetic parameters for drug-fibers compatibility tests in the HFS-TB.**

Drug	Mimicking dosing	C <sub>max, ss</sub> (µg/mL)	T <sub>max</sub> (h)	T <sub>1/2</sub> (h)	Total/free fraction	Site of infection	Reference
BDQ	NA	0.5	1	17.8	NA	NA	NA
	200 mg TIW (3 days)	C <sub>med, ss</sub> = 0.061 µg/mL			Free	Lung (ELF)	Raaijmakers et al. <sup>31</sup> , van Heeswijk et al. <sup>59</sup> , Alghamdi et al. <sup>60</sup> , Moodliar et al. <sup>61</sup> , Svensson et al. <sup>62</sup> , Stephens et al. <sup>63</sup> , Ngwalero et al. <sup>64</sup>
CLA	500 mg BID (3 days)	3.5	2	4	Free	Plasma	Guay et al. <sup>65</sup>
DLM	10 mg/kg QAD (4 days)	0.5	3.4	30	Free	Plasma	Ramirez et al. <sup>66</sup> , TB-APEX <sup>33</sup>
EMB	25 mg/kg QD (3 days)	3	2	3; 12 <sup>a</sup>	Free	Lung (ELF)	Bekker et al. <sup>67</sup> , Tikiso et al. <sup>68</sup> , Lee et al. <sup>69</sup> , Srivastava et al. <sup>70</sup>
INH	600 mg QD (3 days)	6.8	1	3	Free	Lung (ELF)	Gumbo et al. <sup>20</sup>
LZD	600 mg QD (3 days)	8.6	1	3.5	Free	Lung (lesion)	Heinrichs et al. <sup>14</sup>
MXF	400 mg QD (3 days)	3.1	1.5	13	Total	Plasma	Ginsburg et al. <sup>71</sup> , Kim et al. <sup>72</sup>
PTD (alone)	200 mg QD (3 days)	2	1	18	Free	Plasma	Lyons et al. <sup>73</sup> , Liu et al. <sup>74</sup> , Gumbo et al. <sup>20</sup>
PTD (BPaL)	200 mg QD (3 days)	4.3	1	18	Free	Lung (lesion)	Lyons et al. <sup>73</sup> , Liu et al. <sup>74</sup> , Mehta et al. <sup>75</sup> , Karakitsios et al. <sup>76</sup> , Mehta et al. <sup>77</sup>
RMP	600 mg QD (3 days)	9	2	2.5	Free	Lung (ELF)	Drusano et al. <sup>36</sup> , Stott et al. <sup>78</sup>

RPT	600 mg QD (3 days)	2.05	2	2.5	Free	Lung (ELF)	Clewe et al. <sup>79</sup> , Stott et al. <sup>78</sup>
-----	--------------------	------	---	-----	------	------------	---------------------------------------------------------

552

553 BDQ: bedaquiline, CLA: clarithromycin, DLM: delamanid, EMB: ethambutol, INH: isoniazid, LZD: linezolid, MXF: moxifloxacin, PTD:  
554 pretomanid, RMP: rifampicin, RPT: rifapentine, BPAL: bedaquiline-pretomanid-linezolid.  $C_{med, ss}$ : median concentration at protein-unbound steady  
555 state in the lung, TIW: three times a week; BID: *bis in die*: twice a day; QAD: *quaque altera die*: every other day; QD: *quaque die*: once a day.  
556 ELF: epithelial lining fluid. NA: not applicable.

557 <sup>a</sup>EMB's PK is best described by a two-compartment model:  $T_{1/2, \alpha, 0-12 h} = 3 h$  and  $T_{1/2, \beta, 12-24 h} = 12 h$ .

558

## 559 REFERENCES

- 560 1. Gumbo, T., Louie, A., Deziel, M.R., Parsons, L.M., Salfinger, M., and Drusano, G.L.  
561 (2004). Selection of a moxifloxacin dose that suppresses drug resistance in  
562 Mycobacterium tuberculosis, by use of an in vitro pharmacodynamic infection model and  
563 mathematical modeling. *J. Infect. Dis.* *190*, 1642-1651. 10.1086/424849.
- 564 2. Cavaleri, M., and Manolis, E. (2015). Hollow Fiber System Model for Tuberculosis: The  
565 European Medicines Agency Experience. *Clinical infectious diseases : an official  
566 publication of the Infectious Diseases Society of America* *61 Suppl 1*, S1-4.  
567 10.1093/cid/civ484.
- 568 3. European Medicines Agency Qualification Opinion In-vitro Hollow Fiber System Model  
569 of Tuberculosis (HSF-TB). (2015) [https://www.fibercellsystems.com/wp-](https://www.fibercellsystems.com/wp-content/uploads/2022/11/Article-EMAEndorsement.pdf)  
570 [content/uploads/2022/11/Article-EMAEndorsement.pdf](https://www.fibercellsystems.com/wp-content/uploads/2022/11/Article-EMAEndorsement.pdf).
- 571 4. U.S. Food and Drug Administration (FDA). Pulmonary Tuberculosis: Developing Drugs  
572 for Treatment. Guidance Document (2022)  
573 <https://www.fda.gov/media/164606/download>.
- 574 5. Gloede, J., Scheerans, C., Derendorf, H., and Kloft, C. (2010). In vitro pharmacodynamic  
575 models to determine the effect of antibacterial drugs. *J. Antimicrob. Chemother.* *65*, 186-  
576 201. 10.1093/jac/dkp434.
- 577 6. Maitra, A., Solanki, P., Sadouki, Z., McHugh, T.D., and Kloprogge, F. (2021). Improving  
578 the Drug Development Pipeline for Mycobacteria: Modelling Antibiotic Exposure in the  
579 Hollow Fibre Infection Model. *Antibiotics* *10*. 10.3390/antibiotics10121515.
- 580 7. Drusano, G.L. (2017). Pre-clinical in vitro infection models. *Curr. Opin. Pharmacol.* *36*,  
581 100-106. 10.1016/j.coph.2017.09.011.
- 582 8. Sadouki, Z., McHugh, T.D., Aarnoutse, R., Ortiz Canseco, J., Darlow, C., Hope, W., van  
583 Ingen, J., Longshaw, C., Manissero, D., Mead, A., et al. (2021). Application of the hollow  
584 fibre infection model (HFIM) in antimicrobial development: a systematic review and  
585 recommendations of reporting. *J. Antimicrob. Chemother.* *76*, 2252-2259.  
586 10.1093/jac/dkab160.
- 587 9. Mason, A.B., and Dartois, V. (2021). Drug Sensitivity Testing of Mycobacterium  
588 tuberculosis Growing in a Hollow Fiber Bioreactor. *Methods. Mol. Biol.* *2314*, 715-731.  
589 10.1007/978-1-0716-1460-0\_31.
- 590 10. Gumbo, T., Pasipanodya, J.G., Romero, K., Hanna, D., and Nuermberger, E. (2015).  
591 Forecasting Accuracy of the Hollow Fiber Model of Tuberculosis for Clinical  
592 Therapeutic Outcomes. *Clinical infectious diseases : an official publication of the  
593 Infectious Diseases Society of America* *61 Suppl 1*, S25-31. 10.1093/cid/civ427.
- 594 11. Gumbo, T., Angulo-Barturen, I., and Ferrer-Bazaga, S. (2015). Pharmacokinetic-  
595 pharmacodynamic and dose-response relationships of antituberculosis drugs:  
596 recommendations and standards for industry and academia. *J. Infect. Dis.* *211 Suppl 3*,  
597 S96-S106. 10.1093/infdis/jiu610.
- 598 12. Kloprogge, F., Hammond, R., Kipper, K., Gillespie, S.H., and Della Pasqua, O. (2019).  
599 Mimicking in-vivo exposures to drug combinations in-vitro: anti-tuberculosis drugs in  
600 lung lesions and the hollow fiber model of infection. *Sci. Rep.* *9*, 13228. 10.1038/s41598-  
601 019-49556-5.
- 602 13. Wale, Y.M., Roberts, J.A., and Sime, F.B. (2024). Dynamic In Vitro PK/PD Infection  
603 Models for the Development and Optimisation of Antimicrobial Regimens: A Narrative  
604 Review. *Antibiotics* *13*. 10.3390/antibiotics13121201.
- 605 14. Heinrichs, M.T., Drusano, G.L., Brown, D.L., Maynard, M.S., Sy, S.K.B., Rand, K.H.,  
606 Peloquin, C.A., Louie, A., and Derendorf, H. (2019). Dose optimization of moxifloxacin  
607 and linezolid against tuberculosis using mathematical modeling and simulation. *Int. J.*  
608 *Antimicrob. Agents* *53*, 275-283. 10.1016/j.ijantimicag.2018.10.012.
- 609 15. Singh, S., Gumbo, T., Boorgula, G.D., Thomas, T.A., Philley, J.V., and Srivastava, S.  
610 (2024). Omadacycline pharmacokinetics/pharmacodynamics and efficacy against

- 611 multidrug-resistant Mycobacterium tuberculosis in the hollow fiber system model.  
 612 Antimicrob. Agents Chemother. *68*, e0108023. 10.1128/aac.01080-23.
- 613 16. Singh, S., Gumbo, T., Alffenaar, J.W., Boorgula, G.D., Shankar, P., Thomas, T.A.,  
 614 Dheda, K., Malinga, L., Raj, P., Aryal, S., and Srivastava, S. (2023). Meropenem-  
 615 vaborbactam restoration of first-line drug efficacy and comparison of meropenem-  
 616 vaborbactam-moxifloxacin versus BPaL MDR-TB regimen. *Int. J. Antimicrob. Agents*  
 617 *62*, 106968. 10.1016/j.ijantimicag.2023.106968.
- 618 17. van Rijn, S.P., Srivastava, S., Wessels, M.A., van Soolingen, D., Alffenaar, J.C., and  
 619 Gumbo, T. (2017). Sterilizing Effect of Ertapenem-Clavulanate in a Hollow-Fiber Model  
 620 of Tuberculosis and Implications on Clinical Dosing. *Antimicrob. Agents Chemother.* *61*.  
 621 10.1128/AAC.02039-16.
- 622 18. Deshpande, D., Srivastava, S., Nuermberger, E., Pasipanodya, J.G., Swaminathan, S., and  
 623 Gumbo, T. (2016). A Faropenem, Linezolid, and Moxifloxacin Regimen for Both Drug-  
 624 Susceptible and Multidrug-Resistant Tuberculosis in Children: FLAME Path on the  
 625 Milky Way. *Clinical infectious diseases : an official publication of the Infectious Diseases*  
 626 *Society of America* *63*, S95-S101. 10.1093/cid/ciw474.
- 627 19. Srivastava, S., Deshpande, D., Magombedze, G., van Zyl, J., Cirrincione, K., Martin, K.,  
 628 Bendet, P., Berg, A., Hanna, D., Romero, K., et al. (2020). Duration of  
 629 pretomanid/moxifloxacin/pyrazinamide therapy compared with standard therapy based  
 630 on time-to-extinction mathematics. *J. Antimicrob. Chemother.* *75*, 392-399.  
 631 10.1093/jac/dkz460.
- 632 20. Gumbo, T., Chapagain, M., Magombedze, G., Srivastava, S., Deshpande, D.,  
 633 Pasipanodya, J.G., Gumbo, R., Dheda, K., Diacon, A., Hermann, D., and Hanna, D.  
 634 (2022). Novel tuberculosis combination regimens of two and three-months therapy  
 635 duration. *Biorxiv*. <https://doi.org/10.1101/2022.03.13.484155>.
- 636 21. Velkov, T., Bergen, P.J., Lora-Tamayo, J., Landersdorfer, C.B., and Li, J. (2013). PK/PD  
 637 models in antibacterial development. *Curr. Opin. Microbiol.* *16*, 573-579.  
 638 10.1016/j.mib.2013.06.010.
- 639 22. Aguilar-Ayala, D.A., Sanz-Garcia, F., Rabodoarivelo, M.S., Susanto, B.O., Bailo, R.,  
 640 Eveque-Mourroux, M.R., Willand, N., Simonsson, U.S.H., Ramon-Garcia, S., Lucia, A.,  
 641 and consortium, E.T. (2024). Evaluation of critical parameters in the hollow-fibre system  
 642 for tuberculosis: A case study of moxifloxacin. *Br. J. Clin. Pharmacol.* *90*, 1711-1727.  
 643 10.1111/bcp.16068.
- 644 23. Hollow Fiber System of Tuberculosis (HFS-TB): A Laboratory Manual to Guide System  
 645 Engineering, Study Design and Execution. (2018). 1-30.
- 646 24. Silakari, O., and Singh, P.K. (2021). ADMET tools: Prediction and assessment of  
 647 chemical ADMET properties of NCEs. *Concepts and Experimental Protocols of*  
 648 *Modelling and Informatics in Drug Design*, 299-320. [https://doi.org/10.1016/B978-0-12-  
 649 \*820546-4.00014-3\*.](https://doi.org/10.1016/B978-0-12-820546-4.00014-3)
- 650 25. Pajouhesh, H., and Lenz, G.R. (2005). Medicinal chemical properties of successful  
 651 central nervous system drugs. *NeuroRx : the journal of the American Society for*  
 652 *Experimental NeuroTherapeutics* *2*, 541-553. 10.1602/neurorx.2.4.541.
- 653 26. Perveen, S., Kumari, D., Singh, K., and Sharma, R. (2022). Tuberculosis drug discovery:  
 654 Progression and future interventions in the wake of emerging resistance. *Eur. J. Med.*  
 655 *Chem.* *229*, 114066. 10.1016/j.ejmech.2021.114066.
- 656 27. Van der Paardt, A.L., Akkerman, O.W., Gualano, G., Palmieri, F., Davies Forsman, L.,  
 657 Aleksa, A., Tiberi, S., de Lange, W.C., Bolhuis, M.S., Skrahina, A., et al. (2017). Safety  
 658 and tolerability of clarithromycin in the treatment of multidrug-resistant tuberculosis.  
 659 *Eur. Respir. J.* *49*. 10.1183/13993003.01612-2016.
- 660 28. Conradie, F., Bagdasaryan, T.R., Borisov, S., Howell, P., Mikiashvili, L., Ngubane, N.,  
 661 Samoilova, A., Skornikova, S., Tudor, E., Variava, E., et al. (2022). Bedaquiline-  
 662 Pretomanid-Linezolid Regimens for Drug-Resistant Tuberculosis. *N. Engl. J. Med.* *387*,  
 663 810-823. 10.1056/NEJMoa2119430.

- 664 29. Alfarisi, O., Alghamdi, W.A., Al-Shaer, M.H., Dooley, K.E., and Peloquin, C.A. (2017).  
665 Rifampin vs. rifapentine: what is the preferred rifamycin for tuberculosis? Expert review  
666 of clinical pharmacology 10, 1027-1036. 10.1080/17512433.2017.1366311.
- 667 30. Deshkar, A.T., and Shirure, P.A. (2022). Bedaquiline: A Novel Diarylquinoline for  
668 Multidrug-Resistant Pulmonary Tuberculosis. Cureus 14, e28519.  
669 10.7759/cureus.28519.
- 670 31. Raaijmakers, J., Salillas, S., Aarnoutse, R., Svensson, E., Te Brake, L., Stemkens, R.,  
671 Wertheim, H., Hoefsloot, W., and van Ingen, J. (2025). Bedaquiline does not enhance a  
672 clofazimine-azithromycin-ethambutol regimen against Mycobacterium avium in the  
673 hollow-fiber system. Antimicrob. Agents Chemother. 69, e0146424. 10.1128/aac.01464-  
674 24.
- 675 32. Lounis, N., Vranckx, L., Gevers, T., Kaniga, K., and Andries, K. (2016). In vitro culture  
676 conditions affecting minimal inhibitory concentration of bedaquiline against M.  
677 tuberculosis. Med. Mal. Infect. 46, 220-225. 10.1016/j.medmal.2016.04.007.
- 678 33. TB-Platform for the Aggregation of Preclinical Experiments Data (TB-APEX). (2024)  
679 <https://c-path.org/tools-platforms/tb-apex/TB-Platform>
- 680 34. Gumbo, T., Srivastava, S., Deshpande, D., Pasipanodya, J.G., Berg, A., Romero, K.,  
681 Hermann, D., and Hanna, D. (2023). Hollow-fibre system model of tuberculosis  
682 reproducibility and performance specifications for best practice in drug and combination  
683 therapy development. J Antimicrob Chemother 78, 953-964. 10.1093/jac/dkad029.
- 684 35. Al-Sallami, H.S., Cheah, S.L., Han, S.Y., Liew, J., Lim, J., Ng, M.A., Solanki, H., Soo,  
685 R.J., Tan, V., and Duffull, S.B. (2014). Between-subject variability: should high be the  
686 new normal? Eur. J. Clin. Pharmacol. 70, 1403-1404. 10.1007/s00228-014-1740-8.
- 687 36. Drusano, G.L., Sgambati, N., Eichas, A., Brown, D.L., Kulawy, R., and Louie, A. (2010).  
688 The combination of rifampin plus moxifloxacin is synergistic for suppression of  
689 resistance but antagonistic for cell kill of Mycobacterium tuberculosis as determined in a  
690 hollow-fiber infection model. mBio 1. 10.1128/mBio.00139-10.
- 691 37. Deshpande, D., Srivastava, S., Nuermberger, E., Pasipanodya, J.G., Swaminathan, S., and  
692 Gumbo, T. (2016). Concentration-Dependent Synergy and Antagonism of Linezolid and  
693 Moxifloxacin in the Treatment of Childhood Tuberculosis: The Dynamic Duo. Clin Infect  
694 Dis 63, S88-S94. 10.1093/cid/ciw473.
- 695 38. Hou, Y., Mi, K., Sun, L., Zhou, K., Wang, L., Zhang, L., Liu, Z., and Huang, L. (2022).  
696 The Application of Hollow Fiber Cartridge in Biomedicine. Pharmaceutics 14.  
697 10.3390/pharmaceutics14071485.
- 698 39. C, H., AJ, L., and D, H. (1995). Exploring QSAR: Fundamentals and Applications in  
699 Chemistry and Biology. American Chemical Society. 10.1021/jm950902o.
- 700 40. Boorgula, G.D., Singh, S., Shankar, P., Gumbo, T., Heysell, S.K., and Srivastava, S.  
701 (2023). Isoniazid pharmacokinetics/pharmacodynamics as monotherapy and in  
702 combination regimen in the hollow fiber system model of Mycobacterium kansasii.  
703 Tuberculosis 138, 102289. 10.1016/j.tube.2022.102289.
- 704 41. Park, Y., Tung, P.M., Anh, N.K., Cho, Y.S., and Shin, J.G. (2024). Application of the  
705 Hollow-Fiber Infection Model to Personalized Precision Dosing of Isoniazid in a Clinical  
706 Setting. J. Korean Med. Sci. 39, e104. 10.3346/jkms.2024.39.e104.
- 707 42. Tao, M., Liu, F., and Xue, L. (2012). Hydrophilic poly(vinylidene fluoride) (PVDF)  
708 membrane by in situ polymerisation of 2-hydroxyethyl methacrylate (HEMA) and micro-  
709 phase separation. J. Mater. Chem. 22. 10.1039/c2jm30695f.
- 710 43. Etale, A., Onyianta, A.J., Turner, S.R., and Eichhorn, S.J. (2023). Cellulose: A Review  
711 of Water Interactions, Applications in Composites, and Water Treatment. Chem. Rev.  
712 123, 2016-2048. 10.1021/acs.chemrev.2c00477.
- 713 44. Long, H.P., Lai, C.C., and Chung, C.K. (2016). Polyethylene glycol coating for  
714 hydrophilicity enhancement of polydimethylsiloxane self-driven microfluidic chip. Surf.  
715 Coat. Technol. 320, 315-319. 10.1016/j.surfcoat.2016.12.059.
- 716 45. Wang, L., Wei, J., and Wu, B. (2016). Enhancing hydrophilicity performance of  
717 polysulfone hollow fiber membrane by surface modification via UV-induced graft  
718 polymerization of HEA. Desalination Water Treat. 57, 16269-16276.

- 719 46. Said, N., Hasbullah, H., Ismail, A.F., Othman, M.H.D., Goh, P.S., Zainol Abidin, M.N.,  
720 ..., and & Ng, B.C. (2017). Enhanced hydrophilic polysulfone hollow fiber membranes  
721 with addition of iron oxide nanoparticles. *Polym. Int.* *66*, 1424-1429.
- 722 47. Peechmani, P., Othman, M.H.D., Kamaludin, R., Puteh, M.H., Jaafar, J., Rahman, M.A.,  
723 ..., and & Djuli, S.M. (2021). High flux polysulfone braided hollow fiber membrane for  
724 wastewater treatment role of zinc oxide as hydrophilic enhancer. *J. Environ. Chem. Eng.*  
725 *9*, 105873.
- 726 48. Yoo, S.M., and Ghosh, R. (2018). A method for coating of hollow fiber membranes with  
727 calcium alginate. *J. Membr. Sci.* *558*, 45-51. 10.1016/j.memsci.2018.04.044.
- 728 49. Stadler, J.A.M., Maartens, G., Meintjes, G., and Wasserman, S. (2023). Clofazimine for  
729 the treatment of tuberculosis. *Front. Pharmacol.* *14*, 1100488.  
730 10.3389/fphar.2023.1100488.
- 731 50. Jayaram, R., Shandil, R.K., Gaonkar, S., Kaur, P., Suresh, B.L., Mahesh, B.N., Jayashree,  
732 R., Nandi, V., Bharath, S., Kantharaj, E., and Balasubramanian, V. (2004). Isoniazid  
733 pharmacokinetics-pharmacodynamics in an aerosol infection model of tuberculosis.  
734 *Antimicrob. Agents Chemother.* *48*, 2951-2957. 10.1128/AAC.48.8.2951-2957.2004.
- 735 51. Gumbo, T., Louie, A., Deziel, M.R., Liu, W., Parsons, L.M., Salfinger, M., and Drusano,  
736 G.L. (2007). Concentration-dependent Mycobacterium tuberculosis killing and  
737 prevention of resistance by rifampin. *Antimicrob. Agents Chemother.* *51*, 3781-3788.  
738 10.1128/AAC.01533-06.
- 739 52. Mallikaarjun, S., Chapagain, M.L., Sasaki, T., Hariguchi, N., Deshpande, D., Srivastava,  
740 S., Berg, A., Hirota, K., Inoue, Y., Matsumoto, M., et al. (2020). Cumulative Fraction of  
741 Response for Once- and Twice-Daily Delamanid in Patients with Pulmonary Multidrug-  
742 Resistant Tuberculosis. *Antimicrob. Agents Chemother.* *65*. 10.1128/AAC.01207-20.
- 743 53. Beraldi-Magalhaes, F., Parker, S.L., Sanches, C., Sousa Garcia, L., Souza Carvalho, B.K.,  
744 Fachi, M.M., de Liz, M.V., Pontarolo, R., Lipman, J., Cordeiro-Santos, M., and Roberts,  
745 J.A. (2021). Is Dosing of Ethambutol as Part of a Fixed-Dose Combination Product  
746 Optimal for Mechanically Ventilated ICU Patients with Tuberculosis? A Population  
747 Pharmacokinetic Study. *Antibiotics* *10*. 10.3390/antibiotics10121559.
- 748 54. Ahmad, Z., Peloquin, C.A., Singh, R.P., Derendorf, H., Tyagi, S., Ginsberg, A., Grosset,  
749 J.H., and Nuermberger, E.L. (2011). PA-824 exhibits time-dependent activity in a murine  
750 model of tuberculosis. *Antimicrob. Agents. Chemother.* *55*, 239-245.  
751 10.1128/AAC.00849-10.
- 752 55. Bateson, A., Ortiz Canseco, J., McHugh, T.D., Witney, A.A., Feuerriegel, S., Merker,  
753 M., Kohl, T.A., Utpatel, C., Niemann, S., Andres, S., et al. (2022). Ancient and recent  
754 differences in the intrinsic susceptibility of Mycobacterium tuberculosis complex to  
755 pretomanid. *J. Antimicrob. Chemother.* *77*, 1685-1693. 10.1093/jac/dkac070.
- 756 56. Cadwell, J. (2015). The Hollow Fiber Infection Model: Principles and Practice. *Adv*  
757 *Antibiotics Antibodies 1:1*. 10.4172/aaa.1000101.
- 758 57. Omar, S.V., Ismail, F., Ndjeka, N., Kaniga, K., and Ismail, N.A. (2022). Bedaquiline-  
759 Resistant Tuberculosis Associated with Rv0678 Mutations. *N. Engl. J. Med.* *386*, 93-94.  
760 10.1056/NEJMc2103049.
- 761 58. Rodriguez-Ochoa, J.L., Saucó-Carballo, C., Perez-Palacios, P., Merino-Bohorquez, V.,  
762 Velazquez-Escudero, A., Lopez-Cerero, L., Rodriguez-Bano, J., Rodriguez-Martinez,  
763 J.M., Pascual, A., and Docobo-Perez, F. (2025). Pharmacokinetic/pharmacodynamic  
764 optimization of temocillin treatment against CTX-M-15-producing *Klebsiella*  
765 *pneumoniae* isolates in a hollow-fiber infection model. *Antimicrob. Agents Chemother.*,  
766 e0094625. 10.1128/aac.00946-25.
- 767 59. van Heeswijk, R.P., Dannemann, B., and Hoetelmans, R.M. (2014). Bedaquiline: a  
768 review of human pharmacokinetics and drug-drug interactions. *J. Antimicrob.*  
769 *Chemother.* *69*, 2310-2318. 10.1093/jac/dku171.
- 770 60. Alghamdi, W.A., Al-Shaer, M.H., Kipiani, M., Barbakadze, K., Mikiashvili, L.,  
771 Kempker, R.R., and Peloquin, C.A. (2021). Pharmacokinetics of bedaquiline, delamanid  
772 and clofazimine in patients with multidrug-resistant tuberculosis. *J. Antimicrob.*  
773 *Chemother.* *76*, 1019-1024. 10.1093/jac/dkaa550.

- 774 61. Moodliar, R., Aksenova, V., Frias, M.V.G., van de Logt, J., Rossenu, S., Birmingham,  
775 E., Zhuo, S., Mao, G., Lounis, N., Kambili, C., and Bakare, N. (2021). Bedaquiline for  
776 multidrug-resistant TB in paediatric patients. *Int. J. Tuberc. Lung Dis.* *25*, 716-724.  
777 10.5588/ijtld.21.0022.
- 778 62. Svensson, E.M., Dosne, A.G., and Karlsson, M.O. (2016). Population Pharmacokinetics  
779 of Bedaquiline and Metabolite M2 in Patients With Drug-Resistant Tuberculosis: The  
780 Effect of Time-Varying Weight and Albumin. *CPT Pharmacometrics Syst. Pharmacol.* *5*,  
781 682-691. 10.1002/psp4.12147.
- 782 63. Stephens, R.H., Benjamin, A.R., and Walters, D.V. (1996). Volume and protein  
783 concentration of epithelial lining liquid in perfused in situ postnatal sheep lungs. *J. Appl.*  
784 *Physiol.* *80*, 1911-1920. 10.1152/jappl.1996.80.6.1911.
- 785 64. Ngwalero, P., Brust, J.C.M., van Beek, S.W., Wasserman, S., Maartens, G., Meintjes, G.,  
786 Joubert, A., Norman, J., Castel, S., Gandhi, N.R., et al. (2021). Relationship between  
787 Plasma and Intracellular Concentrations of Bedaquiline and Its M2 Metabolite in South  
788 African Patients with Rifampin-Resistant Tuberculosis. *Antimicrob. Agents Chemother.*  
789 *65*, e0239920. 10.1128/AAC.02399-20.
- 790 65. Guay, D.R., Gustavson, L.E., Devcich, K.J., Zhang, J., Cao, G., and Olson, C.A. (2001).  
791 Pharmacokinetics and tolerability of extended-release clarithromycin. *Clin. Ther.* *23*,  
792 566-577. 10.1016/s0149-2918(01)80060-6.
- 793 66. Ramirez, G., Pham, A.C., Clulow, A.J., Salim, M., Hawley, A., and Boyd, B.J. (2021).  
794 Sustained absorption of delamanid from lipid-based formulations as a path to reduced  
795 frequency of administration. *Drug. Deliv. Transl. Res.* *11*, 1236-1244. 10.1007/s13346-  
796 020-00851-z.
- 797 67. Bekker, A., Schaaf, H.S., Draper, H.R., van der Laan, L., Murray, S., Wiesner, L.,  
798 Donald, P.R., McIlleron, H.M., and Hesselning, A.C. (2016). Pharmacokinetics of  
799 Rifampin, Isoniazid, Pyrazinamide, and Ethambutol in Infants Dosed According to  
800 Revised WHO-Recommended Treatment Guidelines. *Antimicrob. Agents. Chemother.*  
801 *60*, 2171-2179. 10.1128/AAC.02600-15.
- 802 68. Tikiso, T., McIlleron, H., Abdelwahab, M.T., Bekker, A., Hesselning, A., Chabala, C.,  
803 Davies, G., Zar, H.J., Rabie, H., Andrieux-Meyer, I., et al. (2022). Population  
804 pharmacokinetics of ethambutol in African children: a pooled analysis. *J. Antimicrob.*  
805 *Chemother.* *77*, 1949-1959. 10.1093/jac/dkac127.
- 806 69. Lee, C.S., Brater, D.C., Gambertoglio, J.G., and Benet, L.Z. (1980). Disposition kinetics  
807 of ethambutol in man. *J. Pharmacokinet. Biopharm.* *8*, 335-346. 10.1007/BF01059382.
- 808 70. Srivastava, S., Musuka, S., Sherman, C., Meek, C., Leff, R., and Gumbo, T. (2010).  
809 Efflux-pump-derived multiple drug resistance to ethambutol monotherapy in  
810 *Mycobacterium tuberculosis* and the pharmacokinetics and pharmacodynamics of  
811 ethambutol. *J. Infect. Dis.* *201*, 1225-1231. 10.1086/651377.
- 812 71. Ginsburg, A.S., Lee, J., Woolwine, S.C., Grosset, J.H., Hamzeh, F.M., and Bishai, W.R.  
813 (2005). Modeling in vivo pharmacokinetics and pharmacodynamics of moxifloxacin  
814 therapy for *Mycobacterium tuberculosis* infection by using a novel cartridge system.  
815 *Antimicrob. Agents Chemother.* *49*, 853-856. 10.1128/AAC.49.2.853-856.2005.
- 816 72. Kim, M.K., and Nightingale, C.H. (2000). Pharmacodynamics and pharmacokinetics of  
817 the fluoroquinolones. In V. T. Andriole (ed.), *The quinolones*, 3rd ed., 169-202.
- 818 73. Lyons, M.A. (2018). Modeling and Simulation of Pretomanid Pharmacokinetics in  
819 Pulmonary Tuberculosis Patients. *Antimicrob. Agents Chemother.* *62*.  
820 10.1128/AAC.02359-17.
- 821 74. Liu, Y., Tan, Y., Wei, G., Lu, Z., Liu, Y., Yang, B., Hui, A.M., and Li, K. (2022). Safety  
822 and pharmacokinetic profile of pretomanid in healthy Chinese adults: Results of a phase  
823 I single dose escalation study. *Pulm. Pharmacol. Ther.* *73-74*, 102132.  
824 10.1016/j.pupt.2022.102132.
- 825 75. Mehta, K., Guo, T., van der Graaf, P.H., and van Hasselt, J.G.C. (2023). Predictions of  
826 Bedaquiline and Pretomanid Target Attainment in Lung Lesions of Tuberculosis Patients  
827 using Translational Minimal Physiologically Based Pharmacokinetic Modeling. *Clin.*  
828 *Pharmacokinet.* *62*, 519-532. 10.1007/s40262-023-01217-7.

- 829 76. Karakitsios, E., Della Pasqua, O., and Dokoumetzidis, A. (2025). Extrapolation of lung  
830 pharmacokinetics of bedaquiline across species using physiologically-based  
831 pharmacokinetic modelling. *Br. J. Clin. Pharmacol.* *91*, 3167-3178. 10.1002/bcp.70163.
- 832 77. Mehta, K., Guo, T., van der Graaf, P.H., and van Hasselt, J.G.C. (2024). Model-based  
833 dose optimization framework for bedaquiline, pretomanid and linezolid for the treatment  
834 of drug-resistant tuberculosis. *Br. J. Clin. Pharmacol.* *90*, 463-474. 10.1111/bcp.15925.
- 835 78. Stott, K.E., Pertinez, H., Sturkenboom, M.G.G., Boeree, M.J., Aarnoutse, R.,  
836 Ramachandran, G., Requena-Mendez, A., Peloquin, C., Koegelenberg, C.F.N., Alffenaar,  
837 J.W.C., et al. (2018). Pharmacokinetics of rifampicin in adult TB patients and healthy  
838 volunteers: a systematic review and meta-analysis. *J. Antimicrob. Chemother.* *73*, 2305-  
839 2313. 10.1093/jac/dky152.
- 840 79. Clewe, O., Goutelle, S., Conte, J.E., Jr., and Simonsson, U.S. (2015). A pharmacometric  
841 pulmonary model predicting the extent and rate of distribution from plasma to epithelial  
842 lining fluid and alveolar cells--using rifampicin as an example. *Eur. J. Clin. Pharmacol.*  
843 *71*, 313-319. 10.1007/s00228-014-1798-3.
- 844 80. European Medicines Agency. Investigation of bioequivalence - Scientific guideline.  
845 Available online. (2010).

846

847

848 **STAR★METHODS**849 **METHOD DETAILS**

850 **Culture media and drugs.** Unless otherwise stated, Middlebrook 7H9 broth (Becton  
851 Dickinson, New Jersey, USA) supplemented with 10% (vol/vol) Middlebrook oleic acid-  
852 albumin-dextrose-catalase (OADC) (Becton Dickinson, New Jersey, USA), 0.5%  
853 (vol/vol) glycerol (Scharlau, Hamburg, Germany) and 0.05% (vol/vol) tyloxapol (Sigma-  
854 Aldrich, St. Louis, USA) (from now on abbreviated as 7H9/OADC/Gly/Tx) or  
855 Middlebrook 7H9 broth supplemented with 0.0004% (vol/vol) cholesterol (Sigma-  
856 Aldrich, St. Louis, USA), dissolved in ethanol:tyloxapol mixture 1:1, 0.085% (vol/vol)  
857 NaCl (PanReac Química S.L.U., Barcelona, Spain) and 0.0004% (vol/vol) catalase  
858 (Sigma-Aldrich, St. Louis, USA) (from now on abbreviated as 7H9/Chol/NaCl/Cat) were  
859 used at 37°C without CO<sub>2</sub>.

860 EMB, INH, MXF (European Pharmacopeia Reference Standard I0500000, E1850000 and  
861 Y0000703, respectively) and LZD (Sigma-Aldrich, St. Louis, USA, USP Reference  
862 Standard 1367561) were dissolved in sterile water on the day of starting the experiment.  
863 RMP (European Pharmacopeia Reference Standard, R0700000), RPT, CLA (Sigma-  
864 Aldrich, St. Louis, USA, R0533-25MG and A3487-100MG, respectively), BDQ, PTD  
865 (RTI International/TB Alliance, FF07153068 and 11356-156, respectively) and DLM  
866 (MedChemExpress, New Jersey, USA, HY-10846/CS-5866) were dissolved in dimethyl  
867 sulfoxide on the day of starting the experiment. When needed, freshly prepared stock  
868 solutions were frozen and thawed once after a maximum of four days at -20°C for single  
869 use. Drug aliquots were diluted to the required concentration in 7H9/OADC/Gly/Tx or  
870 7H9/Chol/NaCl/Cat (in the case of INH and the optimization approaches with BDQ and  
871 BPaL) and used immediately. Bioanalyses included a drug stock control and the syringe's

872 solution, which contained the compound dissolved in broth media. Except for PTD-PS  
873 and PTD-cellulose compatibility tests' syringes, and the one containing LZD for the  
874 BPaL approach, these controls barely diverged from their theoretical concentrations.  
875 When this occurred, expected concentrations and PK parameters were recalculated,  
876 thereby not affecting conclusions from compatibility assessments. They were prepared  
877 before the freeze-thaw cycle of the samples and analyzed after that, which validates the  
878 stability of the drugs in these conditions.

879 **HFS experimental set up.** A schematic depiction of the HFS-TB can be found in **Figure**  
880 **S1.** Materials, technical specifications and HFS-TB assembly is described by Aguilar-  
881 Ayala DA, *et al.*<sup>22</sup>. Medium-size PS, PVDF and cellulose (FiberCell Systems, New  
882 Market, USA. Refs C2011, C7011 and C8008, respectively) cartridges were used in this  
883 study. Bottles, caps, tubing, fittings, filters and media were autoclaved before each assay.  
884 As a sterility test, culture media was incubated first at 37°C for 2 days and, then, at room  
885 temperature for 2 additional days. On the day the HFS was assembled, the peristaltic  
886 pump was switched on at a high rate to ensure there were no leakages throughout the  
887 circuit. In compliance with manufacturer's recommendations (FiberCell Systems, New  
888 Market, USA), Duet pumps were set at a rate of 25 U (~100-110 mL/min) to promote fast  
889 distribution and equilibration of broth media and drug between the ICS and the ECS. HFS  
890 compatibility tests included a pre-conditioning step of 3-4 days recirculating broth, so as  
891 to accurately hydrate the fibers. Total volume of the HFS was 200 mL. Before each  
892 sampling, the ECS content was thoroughly mixed through the cartridge ports using 3-  
893 bodies Luer-Lock syringes of 20 mL (BBraun, Melsungen, Germany). The same 20 mL  
894 syringes, or 30 mL ones (Rays, Osimo, Italy), were used for drug administration.  
895 Compound concentrations within syringes and pump settings are enclosed in **Table S5.**  
896 Polypropylene 3-bodies Luer-Lock 1 mL syringes (Thermofisher Scientific, Waltham,

897 USA) were used for sampling. Regarding the optimization approach for BDQ and BPaL,  
898 previous data on binding to plastic -especially polypropylene- and silicone were taken  
899 into account<sup>31,32</sup>. Therefore, 1/16" and 3/16" ID polycarbonate fittings (Masterflex,  
900 Gelsenkirchen, Germany), 3-bodies Luer-Lock 20 mL (SAI Infusion Technologies, Lake  
901 Villa, USA) and 1 mL (BD, Becton Dickinson, New Jersey, USA) polycarbonate  
902 syringes, glass tubes and polystyrene plates (TPP®, Trasadingen, Switzerland) were used  
903 in these experiments. PK samples were stored at -80 °C until quantification.

904 Syringe and peristaltic pumps rates were obtained using standard PK equations. The  
905 elimination rate constant ( $k$ ) was calculated based on **equation 1**:  $k = \ln(2)/T_{1/2}$ , where  
906  $T_{1/2}$  is the half-life of the drug under study. The clearance (CL) was derived according to  
907 **equation 2**:  $CL = k \cdot V$ , where  $V$  is the total volume of the system (200 mL). The compound  
908 concentration at time ( $t$ ) after infusion ( $C_t$ ) was obtained using **equation 3**:  $C_t = C_s \cdot e^{-kt}$ ,  
909 where  $C_s$  is the concentration in the infusion syringe divided by the clearance.

#### 910 **Mimicking a pharmacokinetic profile in the HFS for drug-fiber compatibility tests.**

911 Drug-fiber compatibility tests consist in infusing a compound into the system -in absence  
912 of bacteria- to mimic a specific PK profile and gauging the drug concentration in the  
913 system over time. There are two alternatives to address this assay: on the one hand, a  
914 known concentration of the antibiotic is infused into the HFS-TB so as to reach  $C_{max}$  at  
915 the targeted  $T_{max}$ , and the system is kept static for 1.5-2 hours before drug dilution. On  
916 the other hand, a clinically plausible PK profile of more than one dose is simulated. In  
917 both approaches, drug binding is spotted if target concentrations are not achieved and/or  
918 they vary between compartments.

919 In this study, we mainly centered on the second approach; thereby mimicking PK profiles  
920 resembling the *in vivo* context for *M. tuberculosis* therapy: i.e., drug free concentrations  
921 (unbound to proteins) in the site of infection (lungs), as long as these data were available.

922 A summary of target PK parameters is encompassed in **Table 4**. A 3 days QD (*quaque*  
923 *die*; once a day) dosing was designed for all drugs except for CLA (3 days QAD; *quaque*  
924 *altera die*: every other day), DLM (4 days BID; *bis in die*: twice a day) and BDQ (3 days;  
925 single dose). In each case, the drug was infused into the system at a rate and concentration  
926 that led to the steady state  $C_{\max}$  ( $C_{\max, ss}$ ) at the predetermined  $T_{\max}$ . In the case of BDQ,  
927  $C_{\max, ss}$  was maintained for 1.5 hours before drug dilution. In all cases, drug clearance was  
928 set at the corresponding rate (see afore-mentioned equation), and ca. 15 time points were  
929 sampled from both the ICS and the ECS.

930 A drug was deemed within the compatibility range with a specific cartridge when the  
931 following criteria were met in the ECS:  $AUC_{0-24h, real}/AUC_{0-24h, expected}$  ( $AUC_{0-24h, ECS, ratio}$ )  
932  $= 0.8-1.25$ ;  $C_{\max, ss, real}/C_{\max, ss, expected}$  ( $C_{\max, ss, ECS, ratio}$ )  $= 0.7-1.3$ , and  $T_{1/2, real}/T_{1/2, expected}$   
933 ( $T_{1/2, ECS, ratio}$ )  $= 0.6-1.4$ , based on EMA bioequivalence guidelines<sup>80</sup>, as well as the  
934 convention on acceptable coefficients of variation in between-subject variance of PK  
935 parameters, extracted from Al-Sallami HS, *et al*<sup>35</sup>. If these criteria were not met, a drug  
936 was considered outside the compatibility range with a specific cartridge.

937 **Optimization approach to mimic a pharmacokinetic profile of bedaquiline in the**  
938 **HFS-TB.** As a proof of concept to optimize the use of highly lipophilic molecules in the  
939 HFS-TB, a median concentration at protein-unbound clinical steady state ( $C_{med, ss}$ ) in the  
940 lung was targeted for BDQ (200 mg TIW; three times a week) (**Table 4**). To obtain that  
941 value, we followed the same procedure described by Raaijmakers J, *et al.*<sup>31</sup>, taking into  
942 consideration BDQ's average plasma concentrations<sup>60,61,64</sup>, protein binding<sup>59</sup> and  
943 plasma/lung ratio of albumin<sup>62,63</sup>. To keep  $C_{med, ss}$  along 3 days in the ECS, six infusions  
944 a day of BDQ (1.65 mL/min of a 100  $\mu$ g/mL syringe stock on the first day, and 1.65  
945 mL/min of a 50  $\mu$ g/mL syringe stock on the second and third days) were administered  
946 directly into the ECS ( $T_{\max} = 1$  min). These values were calculated based on data from

947 BDQ's compatibility tests and the TB-APEX<sup>33</sup>. Drug clearance was set at an arbitrary  
948 rate (corresponding to  $T_{1/2} = 14.6$  h; see afore-mentioned equation) and 19 time points  
949 were sampled from the ECS.

950 **Optimization approach to mimic a pharmacokinetic profile of bedaquiline-**  
951 **pretomanid-linezolid (BPaL) combination in the HFS-TB.** A 3-days unbound PK  
952 profile of BDQ, PTD and LZD in lung was targeted in the HFS-TB (**Table 4**). An  
953 unbound  $C_{med, ss}$  in the lung was aimed for BDQ in a cellulose cartridge, following a  
954 slightly modified version of the infusions' schedule detailed above. Data from BDQ alone  
955 in cellulose fibers was used to inform the design of the combination experiment (1.65  
956 mL/min of a 20  $\mu\text{g/mL}$  syringe stock on the first day, and 1.65 mL/min of a 10  $\mu\text{g/mL}$   
957 syringe stock on the second and third days;  $T_{max} = 1$  min). PTD's PK profile was aimed  
958 by adding thrice the amount of drug into the ICS in each infusion (**Table S5**). This was  
959 done to counteract the unspecific binding registered from experiments with PTD alone in  
960 cellulose fibers. Plus, five top-up infusions a day of PTD (1.20 mL/min of a 400  $\mu\text{g/mL}$   
961 syringe stock;  $T_{max} = 1$  h) were administered to compensate its profile under LZD's  
962 clearance, which held a shorter half-life (**Table 4**). 22 time points were sampled only  
963 from the ECS, because it is the compartment where bacteria will be housed in PKPD  
964 experiments and where BDQ was added, thereby assuming that BDQ's levels in the ICS  
965 would be negligible.

966 **Preparation of drug samples and drug quantification.** A seven-point calibration curve  
967 covering the range of expected drug concentrations in the experiment was established in  
968 duplicate under the same matrix composition as the samples to be quantified.

969 The LC analysis was performed using a UPLC Acquity I-Class system (Waters®,  
970 Milford, USA) equipped with an Acquity Ethylene Bridge Hybrid (BEH) C18 analytical  
971 column (1.7  $\mu\text{m}$ , 50 mm·2.1 mm). The mobile phase consisted of 5 mM ammonium

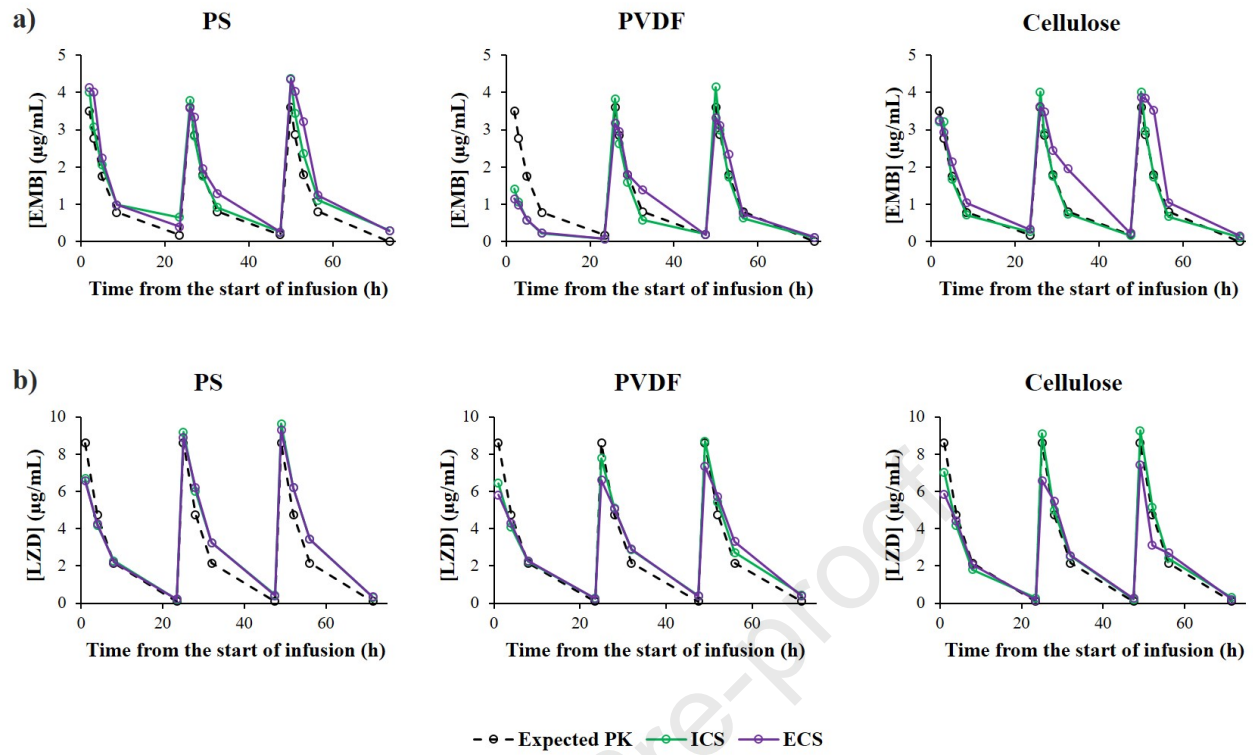
972 formate in water (phase A) or in acetonitrile (phase B) and the elution was performed  
973 with a 4 min gradient from 2% to 98% phase B, with an injection volume of 1  $\mu$ L and a  
974 flow rate set at 600  $\mu$ L/min. The UPLC system was coupled online to a Xevo TQD system  
975 (Waters®, Milford, USA) equipped with an ESI source. The mass spectrometer was  
976 operated in positive ion mode with a specific Multiple Reaction Monitoring (MRM)  
977 method for each compound (**Table S6**). The raw data were processed within the  
978 MassLynx software version 4.2 (Waters®, Milford, USA). In addition, bioanalytical LC-  
979 MS/MS methods for the compounds were validated for: i) selectivity and specificity -  
980 optimized LC-MSMS method with specific MRM transitions for each drug- (**Table S6**),  
981 ii) accuracy and precision -based on the  $R^2$  values obtained for each calibration curves  
982 models-, iii) recovery -barely a difference between the expected and the calculated  
983 concentrations- and, iv) matrix effect -calibration curves built using the broth to mimic  
984 the samples to quantify-.

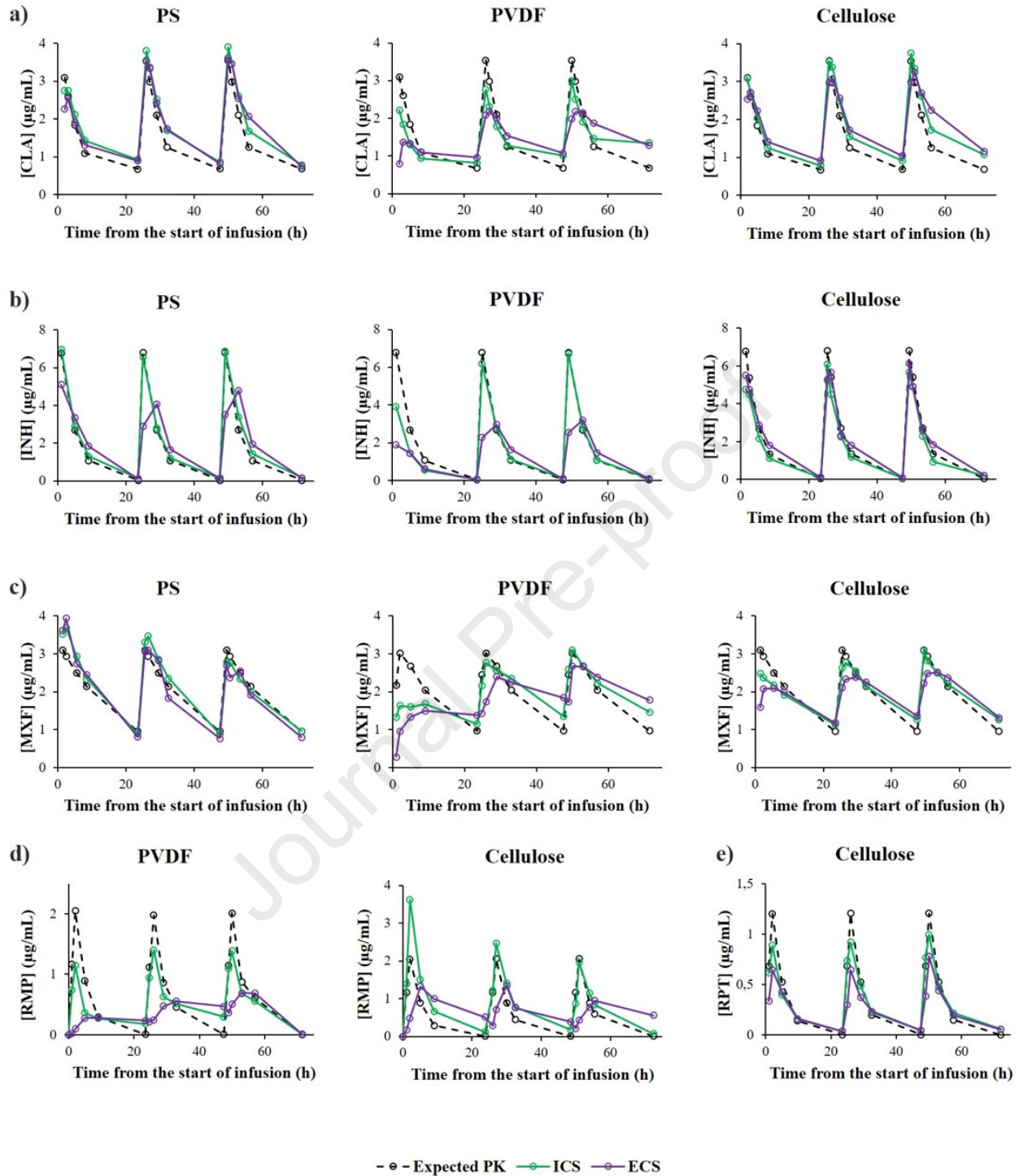
#### 985 **QUANTIFICATION AND STATISTICAL ANALYSIS**

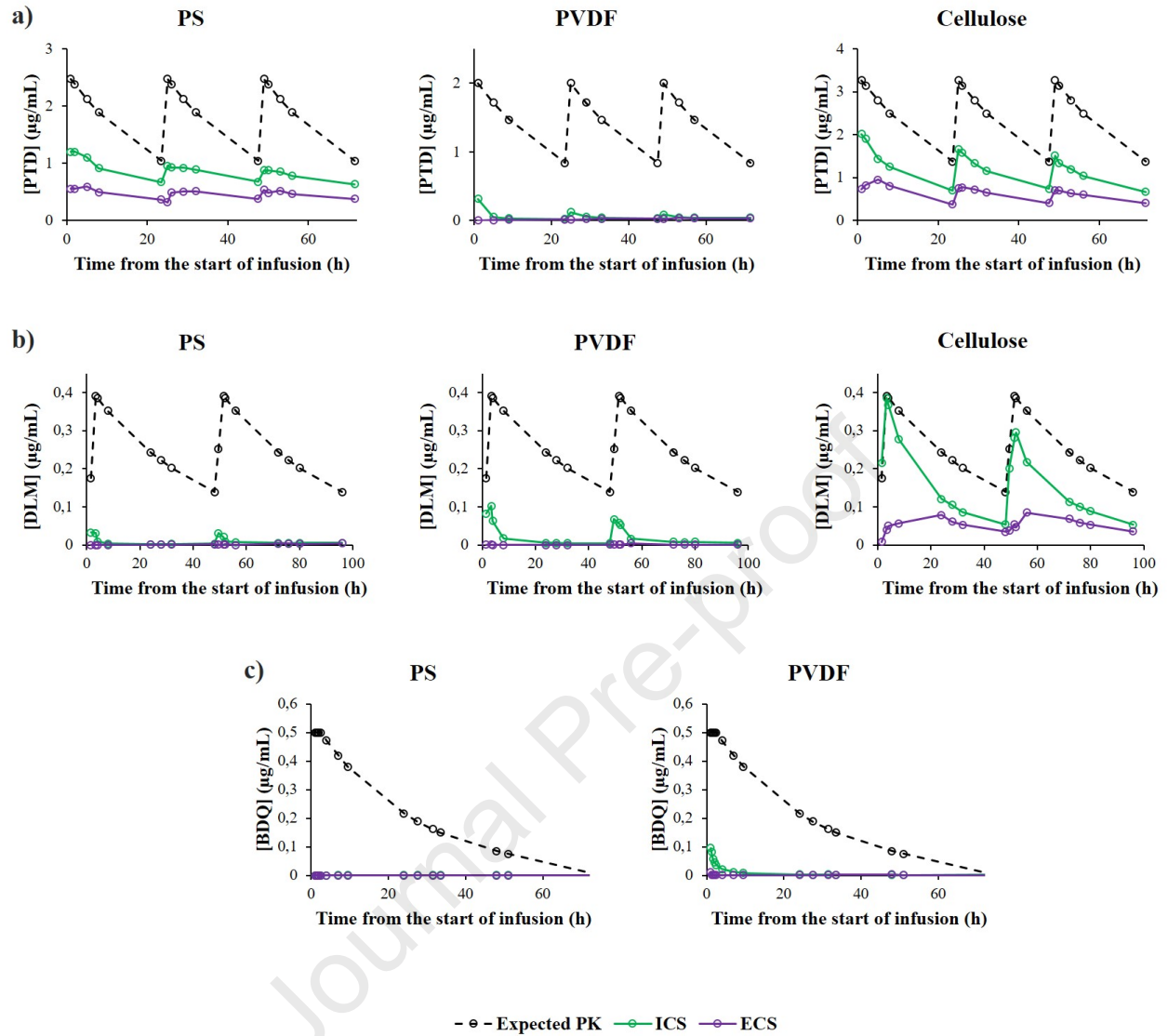
986 Accuracy and precision of the calibration curves for drug quantification -0.3 nM to 500  
987 or 1000 nM, depending on the range of expected concentrations- were evaluated based  
988 on  $R^2$  values of linear regression ( $\geq 0.95$ ). Each point of said curves was performed in  
989 duplicate. Accuracy of the  $T_{1/2}$  calculated for drug clearance after each dose (2-3,  
990 depending on the targeted PK) was evaluated based on  $R^2$  values of exponential  
991 regression ( $\geq 0.95$ ).

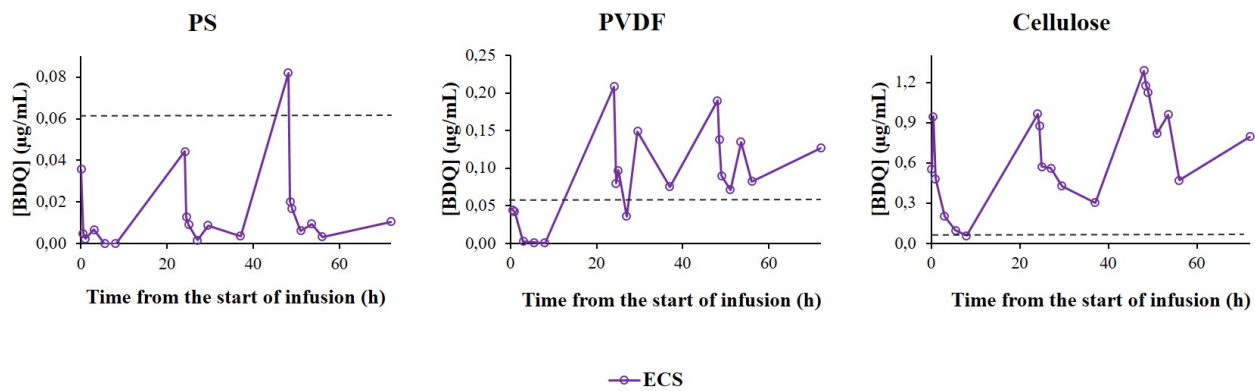
#### 992 **SUPPLEMENTAL INFORMATION**

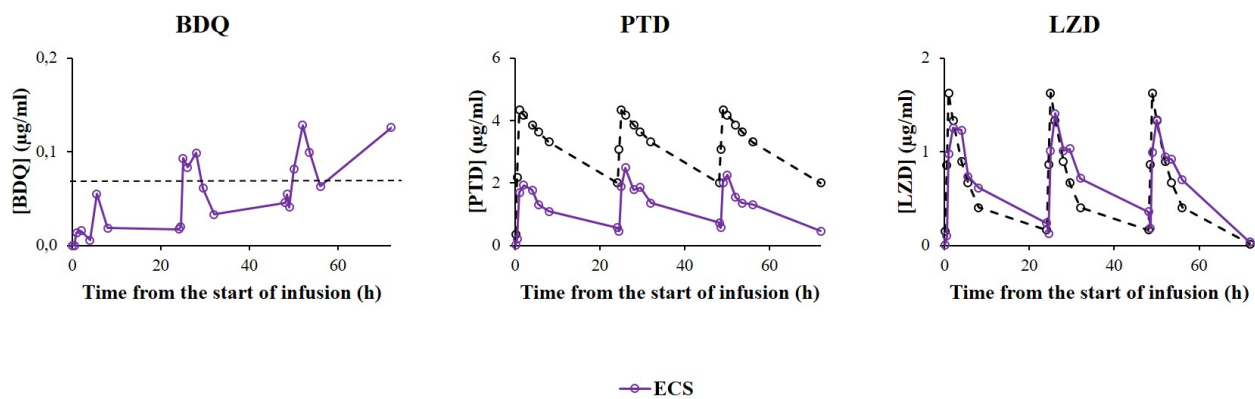
993 **Table S1.** Expected and measured concentrations from drug-fiber compatibility tests in  
994 the HFS-TB.











Drug	Polysulfone	PVDF	Cellulose
INH			
EMB			
LZD			
RMP			
MXF			
CLA			
RPT	ND	ND	
PTD			
DLM			
BDQ			*


 LIPOPHILICITY

Within the compatibility range	Outside the compatibility range (optimizable)	Outside the compatibility range
--------------------------------	-----------------------------------------------	---------------------------------

- A comprehensive assay toolkit for the HFS-TB is provided.
- Polysulfone and cellulose fibers are the most suitable for compounds in the toolkit.
- A strategy to mimic BPaL exposure in the HFS-TB has been successfully implemented.
- Assessing drug concentrations in the extra-capillary space is a key verification step.

Journal Pre-proof

## KEY RESOURCES TABLE

REAGENT OR RESOURCE	SOURCE	IDENTIFIER
<b>Chemicals, peptides, and recombinant proteins</b>		
Middlebrook 7H9 broth	Becton Dickinson	Cat# 271310
Middlebrook oleic acid-albumin-dextrose-catalase	Becton Dickinson	Cat# 212351
Glycerol	Scharlau	Cat# 15523-1L-M
Tyloxapol	Sigma-Aldrich	Cat# T8761-50G
Cholesterol	Sigma-Aldrich	Cat# C3045-25 G
NaCl	Panreac Química S.L.U.	Cat# S1679
Catalase	Sigma-Aldrich	Cat# C40-500MG
Ethambutol	European Pharmacopeia Reference Standard	Cat# I0500000
Isoniazid	European Pharmacopeia Reference Standard	Cat# E1850000
Moxifloxacin	European Pharmacopeia Reference Standard	Cat# Y0000703
Linezolid	Sigma-Aldrich	Cat# 1367561
Rifampicin	European Pharmacopeia Reference Standard	Cat# R0700000
Rifapentine	Sigma-Aldrich	Cat# R0533-25MG
Clarithromycin	Sigma-Aldrich	Cat# A3487-100MG
Bedaquiline	RTI International/TB Alliance	Cat# FF07153068
Pretomanid	RTI International/TB Alliance	Cat# 11356-156
Delamanid	MedChemExpress	Cat# HY-10846/CS-5866
Dimethyl sulfoxide	Sigma-Aldrich	Cat# 276855
<b>Software and algorithms</b>		
MassLynx software version 4.2	Waters	NA
<b>Other</b>		
Medium-size polysulfone cartridges	FiberCell Systems	Cat# C2011
Medium-size polyvinylidene fluoride cartridges	FiberCell Systems	Cat# C7011
Medium-size cellulose cartridges	FiberCell Systems	Cat# C8008
250 mL polycarbonate bottles	VWR	Cat# VWRI215-2204
4 L polycarbonate bottles	VWR	Cat# DS2205 -0010
PKPD reservoir cap	FiberCell Systems	Cat# A1007
Reservoir cap	FiberCell Systems	Cat# A1006
Silicone pump tubing	Masterflex	Cat# 96410-16
Polytetrafluoroethylene-based, hydrophobic filter unit	Millex-FG	Cat# SLFG05010
C-flex tubing 0.8 mm ID	Masterflex	Cat# 06427-13
C-flex tubing 1.6 mm ID	Masterflex	Cat# 06427-14
BPT tubing 1.52" ID	Ismatec Saint-Gobain Pharmed BTP, Masterflex	Cat# S0732
Polypropylene female Luer-lock fittings 1/16" ID	Masterflex	Cat# 14845571
Polypropylene male Luer-lock fittings 1/16" ID	Masterflex	Cat# 13546720
Polypropylene female Luer-lock fittings 3/16" ID	Masterflex	Cat# 15296618

Polypropylene male Luer-lock fittings 3/16" ID	Masterflex	Cat# 14860962
Polycarbonate female Luer-lock fittings 1/16" ID	Masterffex	Cat# 12197750
Polycarbonate male Luer-lock fittings 1/16" ID	Masterflex	Cat# 15819911
Polycarbonate female Luer-lock fittings 3/16" ID	Masterffex	Cat# 15216088
Polycarbonate male Luer-lock fittings 3/16" ID	Masterflex	Cat# 13538810
Sterile polyamide microcrystalline T-joints	BBraun	Cat# 16494C
Clave needle-free connectors	Icumedical	Cat# 011-C1000
Duet pumps	FiberCell Systems	Cat# P3202
Peristaltic Ismatec Reglo pumps	Masterflex	Cat# ISM4408
Masterflex LS pumps	Masterflex	Cat# 07522-30
One Channel Programmable Syringe Pumps	New Era Pump Systems, Inc.	Cat# NE-1000
Polypropylene 3-bodies Luer-Lock syringes of 20 mL	BBraun	Cat# 4617207V
Polypropylene 3-bodies Luer-Lock syringes of 30 mL	Rays	Cat# 30LL
Polypropylene 3-bodies Luer-Lock syringes of 1 mL	Thermo Scientific	Cat# 309628
Polycarbonate 3-bodies Luer-Lock syringes of 20 mL	SAI Infusion Technologies	Cat# 847.356.0321
Polycarbonate 3-bodies Luer-Lock syringes of 1 mL	Becton Dickinson	Cat# 309628
Polystyrene p96-well plates	TPP	Cat# 92096
TB-Platform for the Aggregation of Preclinical Experiments Data (TB-APEX)	NA	NA