

Supplementary Material

Absence of light exposure increases pathogenicity of *Pseudomonas aeruginosa* pneumonia-associated clinical isolates

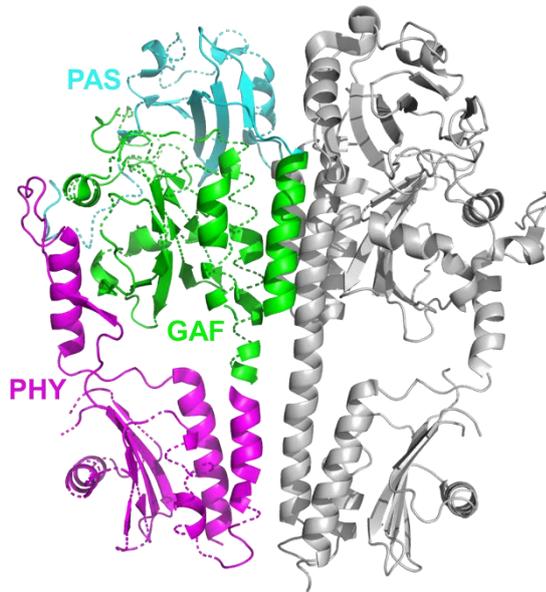
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(a)



(b)

PA4117	MTSITPVTLAN	CEDEPIHVPGAIQPHGAL	MLRADGMVLAASENIQALLGFVASPSSYLA	60	
PA13_1006205	MTSITPVTLAN	CEDEPIHVPGAIQPHGAL	MLRADGMVLAASENIQALLGFVASPSSYLA	120	
PA15_0309150	MTSITPVTLAN	CEDEPIHVPGAIQPHGAL	MLRADGMVLAASENIQALLGFVASPSSYLA	60	
consensus	*****	*****	*****	*****	
PA4117	QEQVGPVLRMLEEGL	TCNGPWSNSVETRIGEHLFDVIGHSYKEVFYLEFEIR/TADTLSI	120		
PA13_1006205	QEQVGPVLRMLEEGL	TCNGPWSNSVETRIGEHLFDVIGHSYKEVFYLEFEIR/TADTLSI	120		
PA15_0309150	QEQVGPVLRMLEEGL	TCNGPWSNSVETRIGEHLFDVIGHSYKEVFYLEFEIR/TADTLSI	120		
consensus	*****	*****	*****	*****	
PA4117	TSFTLNAQR	IIAQVQLHNDTASLLSNVTDELRR	TGYDRVMAYRFRHDDSGEVVAESRRE	180	
PA13_1006205	TSFTLNAQR	IIAQVQLHNDTASLLSNVTDELRR	TGYDRVMAYRFRHDDSGEVVAESRRE	180	
PA15_0309150	TSFTLNAQR	IIAQVQLHNDTASLLSNVTDELRR	TGYDRVMAYRFRHDDSGEVVAESRRE	180	
consensus	*****	*****	*****	*****	
PA4117	DLESYLQQRYPASDI	IPAQARRLYIQNP	IRLIAADVAYTPMRVFPALNPETNESFDLSYSVLI	240	
PA13_1006205	DLESYLQQRYPASDI	IPAQARRLYIQNP	IRLIAADVAYTPMRVFPALNPETNESFDLSYSVLI	240	
PA15_0309150	DLESYLQQRYPASDI	IPAQARRLYIQNP	IRLIAADVAYTPMRVFPALNPETNESFDLSYSVLI	240	
consensus	*****	*****	*****	*****	
PA4117	RSVSPHICEYLT	NMGVRAASMSISIVVGGKLGWLF	SCHHMSPKLIPYVFRMSQIFFSQVCS	300	
PA13_1006205	RSVSPHICEYLT	NMGVRAASMSISIVVGGKLGWLF	SCHHMSPKLIPYVFRMSQIFFSQVCS	300	
PA15_0309150	RSVSPHICEYLT	NMGVRAASMSISIVVGGKLGWLF	SCHHMSPKLIPYVFRMSQIFFSQVCS	300	
consensus	*****	*****	*****	*****	
PA4117	AIVERLEQQR	IAELLRWSTERR	ILARRBARADDLFGALAHDDGTAALI	PCDGAIVMLG	360
PA13_1006205	AIVERLEQQR	IAELLRWSTERR	ILARRBARADDLFGALAHDDGTAALI	PCDGAIVMLG	360
PA15_0309150	AIVERLEQQR	IAELLRWSTERR	ILARRBARADDLFGALAHDDGTAALI	PCDGAIVMLG	360
consensus	*****	*****	*****	*****	
PA4117	GRTLSIRGDF	FERQAGNWLQRLQRD	PERDIYHTLNNWQPESESDPDGGDCCOQVLA	IRFHRQK	420
PA13_1006205	GRTLSIRGDF	FERQAGNWLQRLQRD	PERDIYHTLNNWQPESESDPDGGDCCOQVLA	IRFHRQK	420
PA15_0309150	GRTLSIRGDF	FERQAGNWLQRLQRD	PERDIYHTLNNWQPESESDPDGGDCCOQVLA	IRFHRQK	420
consensus	*****	*****	*****	*****	
PA4117	SGWIFWRH	EEVHRIRWGGKPEKLLT	IGPSPGRLTPRGSFAWEVVRGHSTPWS	ETDLA	480
PA13_1006205	SGWIFWRH	EEVHRIRWGGKPEKLLT	IGPSPGRLTPRGSFAWEVVRGHSTPWS	ETDLA	480
PA15_0309150	SGWIFWRH	EEVHRIRWGGKPEKLLT	IGPSPGRLTPRGSFAWEVVRGHSTPWS	ETDLA	480
consensus	*****	*****	*****	*****	
PA4117	IAEKLR	LDMEICLNFAAEVD	MRQRILAVLGHDLRNPLOQISMAAALLSSSDT	FTTETLR	540
PA13_1006205	IAEKLR	LDMEICLNFAAEVD	MRQRILAVLGHDLRNPLOQISMAAALLSSSDT	FTTETLR	540
PA15_0309150	IAEKLR	LDMEICLNFAAEVD	MRQRILAVLGHDLRNPLOQISMAAALLSSSDT	FTTETLR	540
consensus	*****	*****	*****	*****	
PA4117	QHISASSSR	MERLVSQLDMSRLQSGI	GLTVNVPVTDVSQLV	QIVCETDVAYPGLVIEI	600
PA13_1006205	QHISASSSR	MERLVSQLDMSRLQSGI	GLTVNVPVTDVSQLV	QIVCETDVAYPGLVIEI	600
PA15_0309150	QHISASSSR	MERLVSQLDMSRLQSGI	GLTVNVPVTDVSQLV	QIVCETDVAYPGLVIEI	600
consensus	*****	*****	*****	*****	
PA4117	AIDPQVRAVVD	PDRYAQVAANLLSNAR	HGGLGPRPVLVTLTRQGEVCLSVLNETSGLSE	660	
PA13_1006205	AIDPQVRAVVD	PDRYAQVAANLLSNAR	HGGLGPRPVLVTLTRQGEVCLSVLNETSGLSE	660	
PA15_0309150	AIDPQVRAVVD	PDRYAQVAANLLSNAR	HGGLGPRPVLVTLTRQGEVCLSVLNETSGLSE	660	
consensus	*****	*****	*****	*****	
PA4117	AQLANLFEP	FKRESADNQRN	RNGLGIGLYISQATAQAHQGRIDVDCRDDVITFC	CLRLPVE	720
PA13_1006205	AQLANLFEP	FKRESADNQRN	RNGLGIGLYISQATAQAHQGRIDVDCRDDVITFC	CLRLPVE	720
PA15_0309150	AQLANLFEP	FKRESADNQRN	RNGLGIGLYISQATAQAHQGRIDVDCRDDVITFC	CLRLPVE	720
consensus	*****	*****	*****	*****	
PA4117	QAE	TGSSS	728		
PA13_1006205	QAE	TGSSS	728		
PA15_0309150	QAE	TGSSS	728		
consensus	*****	*****	*****		

Figure S1. Bacteriophytochrome BphP: the photoreceptor identified in *P. aeruginosa* PAO1 found to be present in the clinical isolates HB13 and HB15 by in silico analysis. (a) Determined crystal structure of the photosensory core domain of the dimeric BphP (PDB: 3C2W) [1]. The Per-ARNT-Sim (PAS), cGMP phosphodiesterase/adenyl cyclase/FhlA (GAF) and phytochrome (PHY) domains of one monomer are highlighted in cyan, green, and magenta, respectively. A structural alignment between the C-terminal of this core domain and the N-terminal of the histidine kinase (HK) domain is proposed by the authors. The bacteriophytochrome BphP (PA13_1006205, PA15_0309150, PA4117) assembles with biliverdin, its chromophore, produced by the heme oxygenase BphO (PA13_1006200, PA15_0309145, PA4116) to generate a photosensing HK that is activated by light. (b) Sequence alignment with representation of the respective locus_tag. Boxes identify the residues corresponding to the four described domains: PAS, GAF, PHY and HK [1]. The “*” indicates positions which have a single, fully conserved residue (black); The “:” indicates the conservation between groups of strongly similar properties (grey). The numbering of each sequence represents the position of each residue in the original protein sequence. According to PROVEAN, none of the variations has a deleterious effect.

1. Yang, X.; Kuk, J.; Moffat, K. Crystal structure of *Pseudomonas aeruginosa* bacteriophytochrome: Photoconversion and signal transduction. *Proc. Natl. Acad. Sci.* **2008**, *105*, 14715–14720.

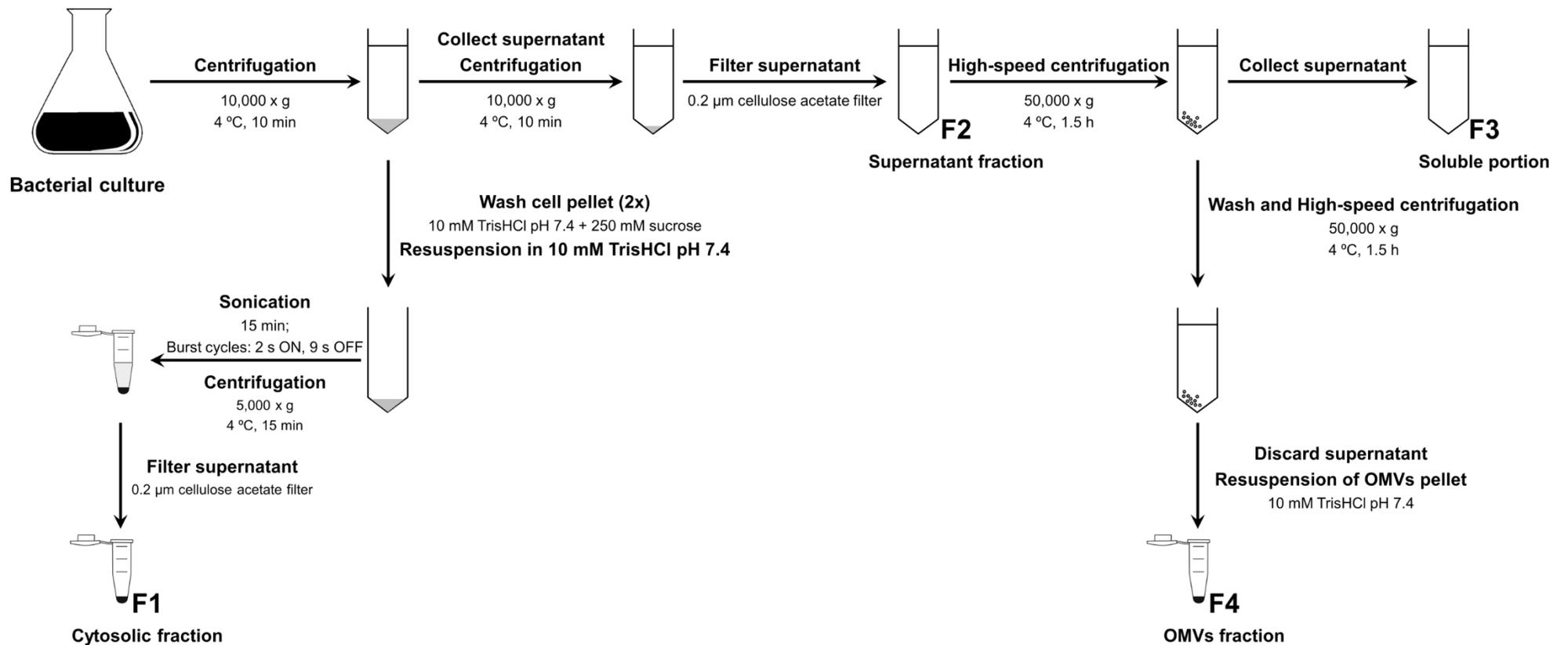


Figure S2. Procedure for collection of four different *P. aeruginosa* fractions. The reference strain PAO1 and the clinical isolates HB13 and HB15 were assayed. From each isolate, four fractions were collected, corresponding to the cytosolic fraction (F1), the whole supernatant (F2), the further fractioned soluble portion (F3) and the OMVs fraction (F4).

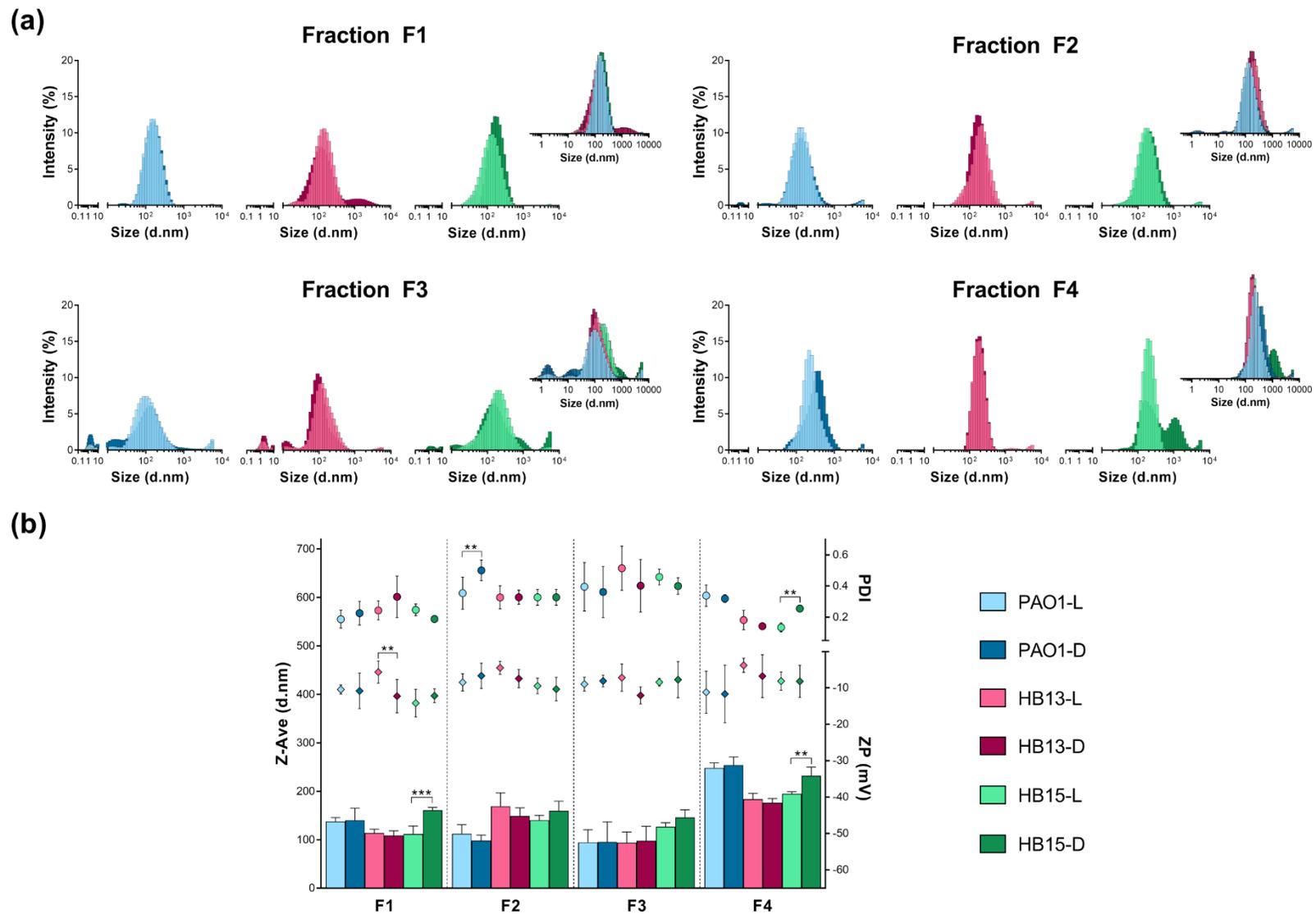


Figure S3. Characterization of the fractions of *P. aeruginosa* grown in constant exposure to full-spectrum light (L) and total absence of light (D) by DLS. **(a)** Particle size distribution by intensity organized by fraction. Combined view of the distribution is represented. **(b)** Analysis of physicochemical properties, particularly Z-ave, ZP and PDI. Standard deviation bars are represented for three independent experiments conducted in triplicate. Statistical comparisons performed by two-way ANOVA, followed by Tukey's post-hoc test for multiple comparisons. Significant differences comparing the two growth conditions are indicated as: ** $P < 0.01$, *** $P < 0.001$.

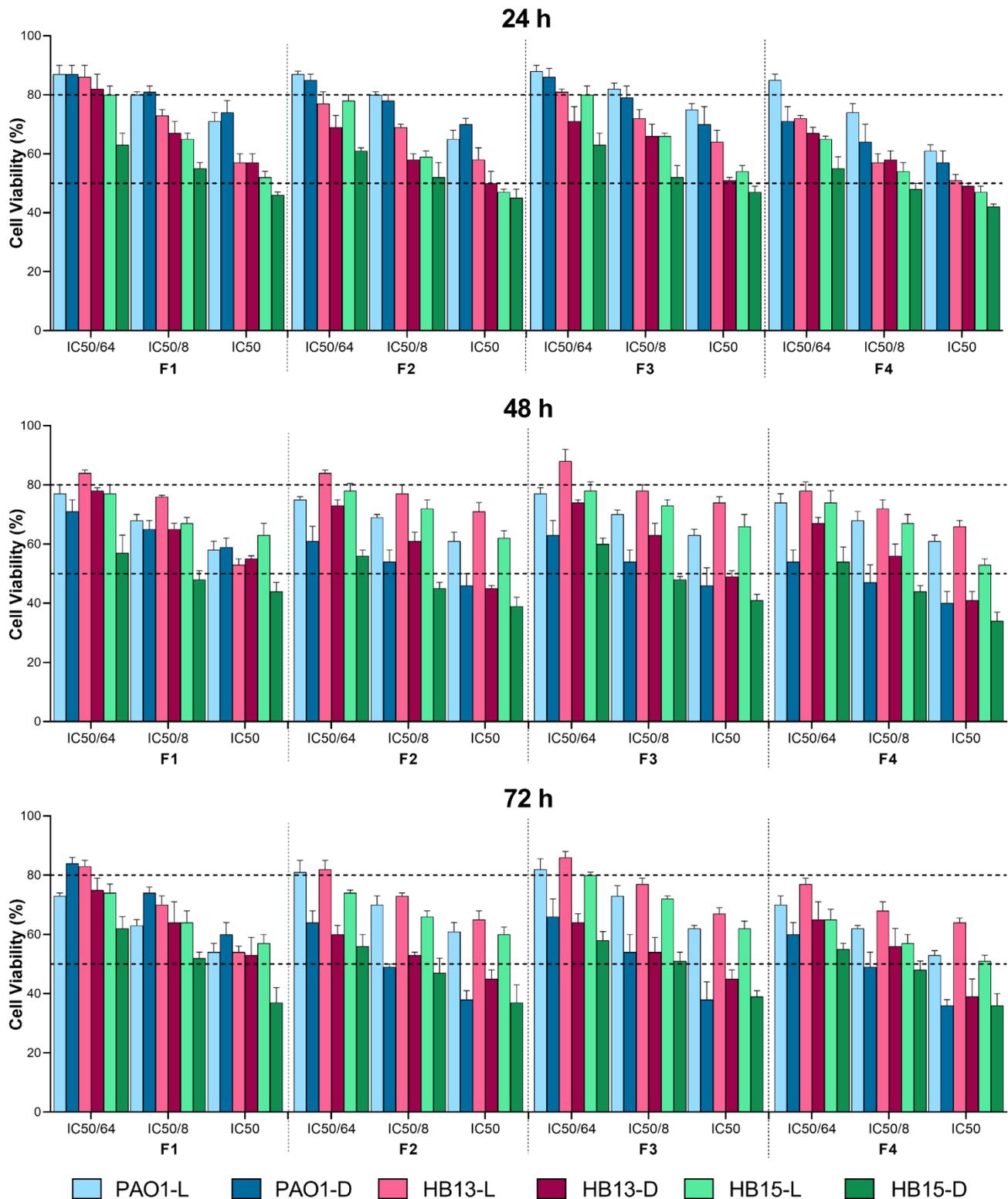


Figure S4. Effect on A549 cell viability (%) induced by fractions of *P. aeruginosa* grown in constant exposure to full-spectrum light (L) and total absence of light (D). Results of the MTT assay are depicted as the cell viability percentage of A549 cells exposed to a serial range of three concentrations (from a 64x dilution of the IC50 to the IC50) of each fraction. Horizontal lines highlight a cellular viability of 50% and 80%. Three MTT endpoints were assayed (24, 48 and 72 h of contact) and standard deviation bars are represented for three independent experiments.

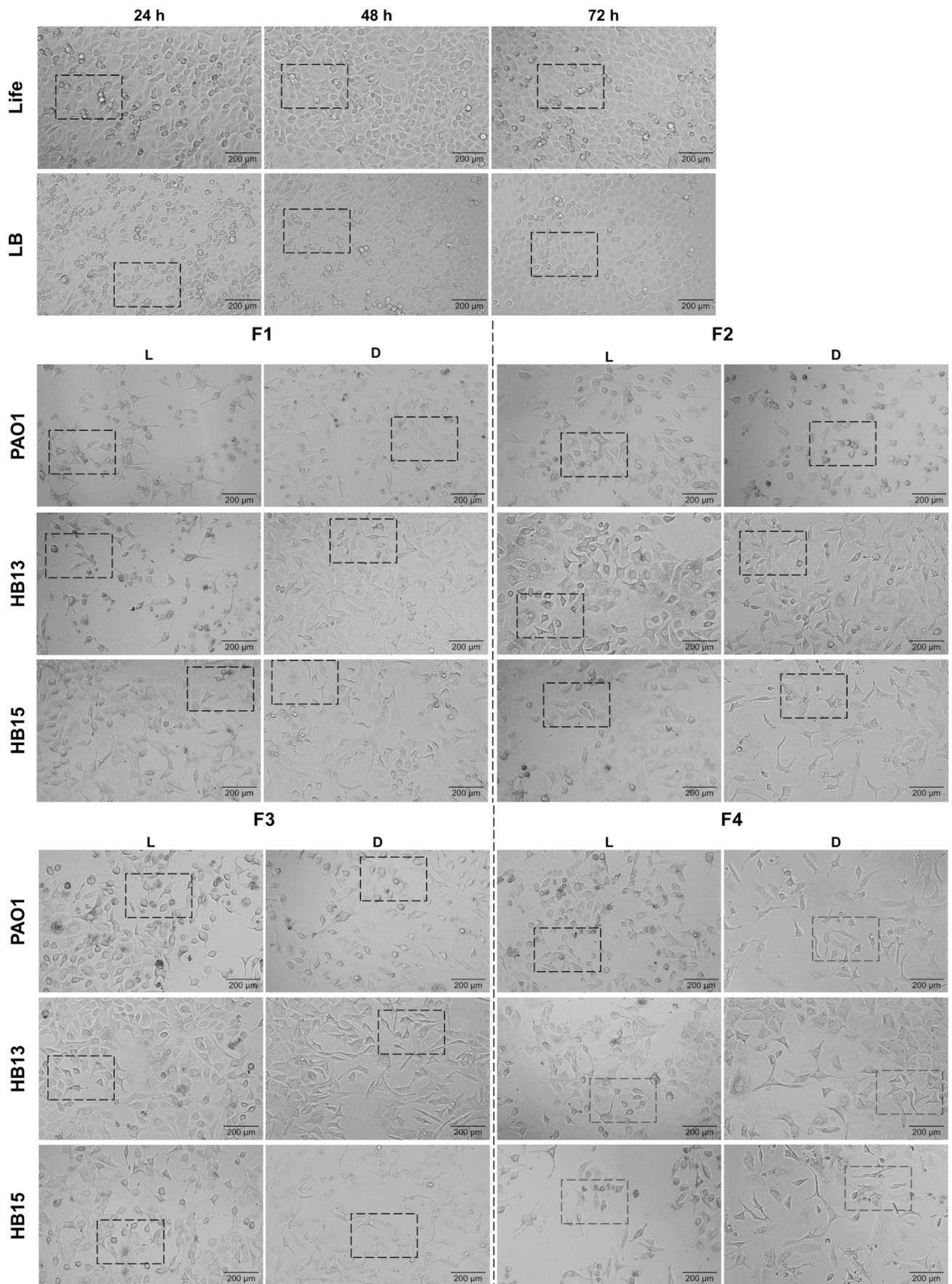


Figure S5. Contrast phase micrography of A549 cells exposed to a specific concentration (Table 1) at 48 h endpoint. Morphologic alterations induced in A549 cells by fractions of *P. aeruginosa* grown in constant exposure to full-spectrum light (L) and total absence of light (D). A representative Life control for the three MTT endpoints is presented. The scale bar is set for 200 µm. Section of the contrast phase micrography presented in Figure 2 is identified by a dashed rectangle.