

## Supplementary Material

# Bactericidal Activity of Non-Cytotoxic Cationic Nanoparticles Against Clinically and Environmentally Relevant *Pseudomonas* spp. Isolates

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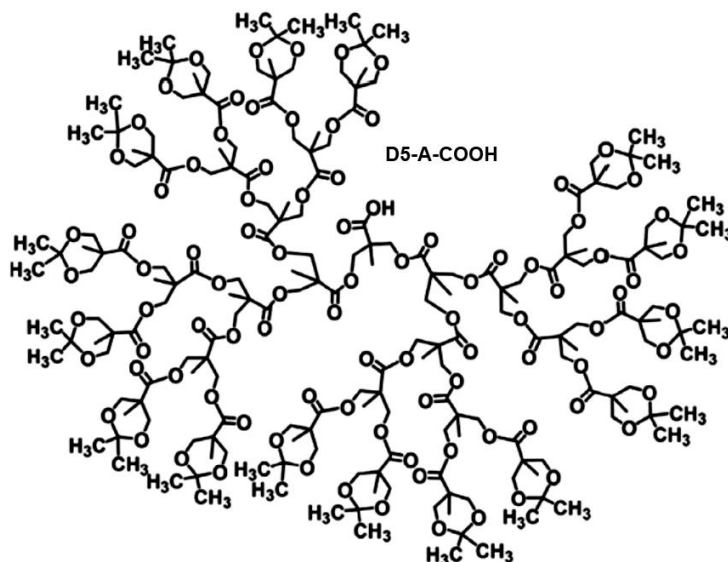
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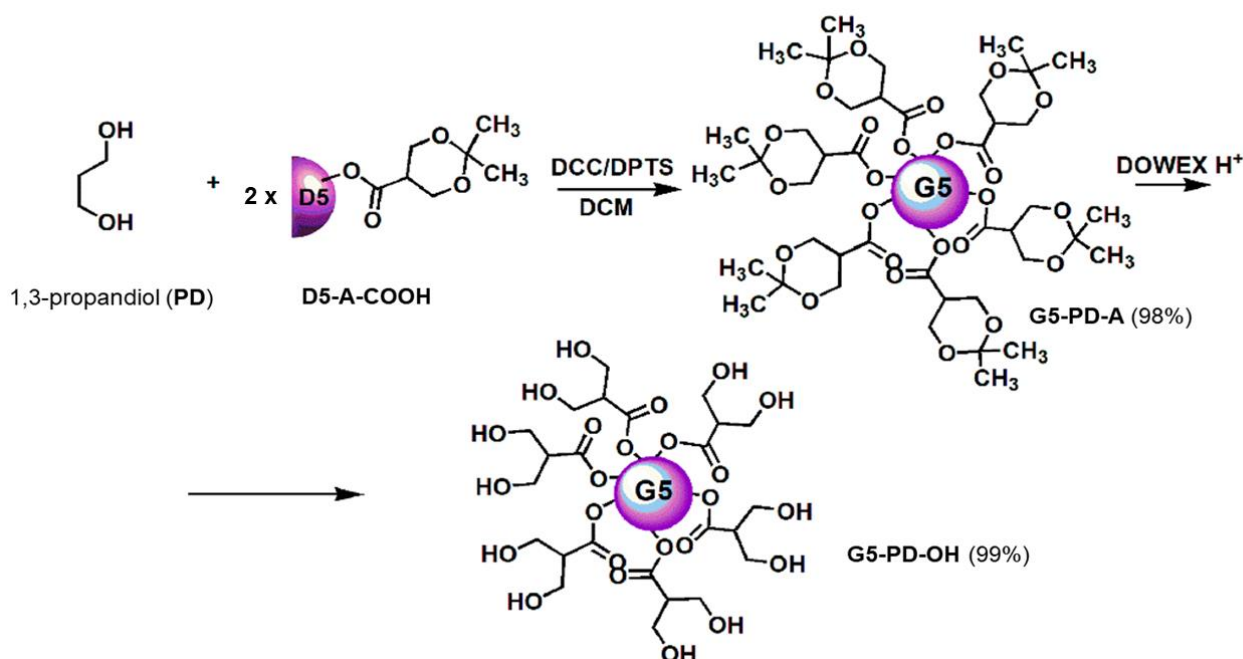
### Section S1. Synthesis of dendrimer G5-PDK [1-6].

#### S1.1. Synthesis of the Uncharged Fifth Generation Inner Scaffold of G5-PDK (G5-PD-OH)

Performing previously reported procedures [2-6], starting from the AB<sub>2</sub> monomer known as *bis*-hydroxymethyl propanoic acid *bis*-HMPA, firstly, we prepared the fifth generation dendron D5-A-COOH (Figure S1), and then, according to Scheme S1, we synthesized the uncharged dendrimer G5-PD-OH [4].



**Figure S1.** Structure of dendron intermediate (D5-A-COOH), prepared to synthesize dendrimer G5-PD-A.



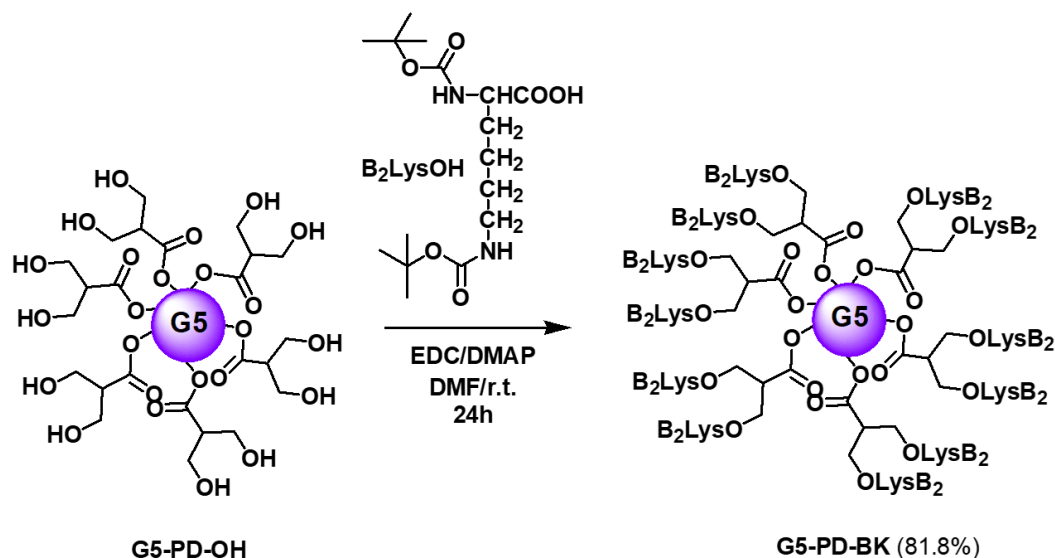
**Scheme S1.** Synthetic procedure for achieving dendrimer G5-PD-OH. D = Dendron; A = acetone protected; COOH = free carboxylic group; D5 = generations number of dendron; G5 = generations number of dendrimers; PD = propanediol; DCC = *N,N'*-dicyclohexylcarbodiimide; DPTS = 4-(dimethyl-amino)pyridinium 4-toluene-sulfonate; DCM = dichloromethane; DOWEX H<sup>+</sup> = acid resins.

#### S1.1.1. FTIR, NMR spectral data and Elemental analysis results of G5-PD-OH [4]

FTIR (KBr, cm<sup>-1</sup>): 3436 (OH), 2936, 1737 (C=O). <sup>1</sup>H NMR (400 MHz, DMSO-*d*<sub>6</sub>), δ (ppm): 1.01, 1.16, 1.18, 1.23, 1.34 (five s signals, 186H, CH<sub>3</sub> of generations), 1.70 (m, 2H, CH<sub>2</sub> propanediol), 3.52 (dd, 128H, CH<sub>2</sub>OH), 3.56 (partially overlapped signal, 2H, CH<sub>2</sub>O propanediol), 3.98 (partially overlapped signal, 2H, CH<sub>2</sub>O propanediol), 4.08-4.18 (m, 120H, CH<sub>2</sub>O of four generations), 4.37 (br s, 64H, OH). <sup>13</sup>C NMR (100 MHz, DMSO-*d*<sub>6</sub>), δ (ppm): 173.94, 171.73 (C=O), 64.27, 63.55 (CH<sub>2</sub>O), 50.13 (quaternary C of fifth generation), 46.12 (other generations detectable quaternary C), 17.05, 16.61 (CH<sub>3</sub> of generations). Found: C, 51.71; H, 7.01. C<sub>313</sub>H<sub>504</sub>O<sub>188</sub> requires C, 51.67; H, 6.98%.

S1.2. Synthesis of Lysine-Modified Cationic Dendrimer G5-PDK (128 HCl) [1]

S1.2.1. Synthesis of Lysine-Modified Boc-Protected Dendrimer G5-PD-BK

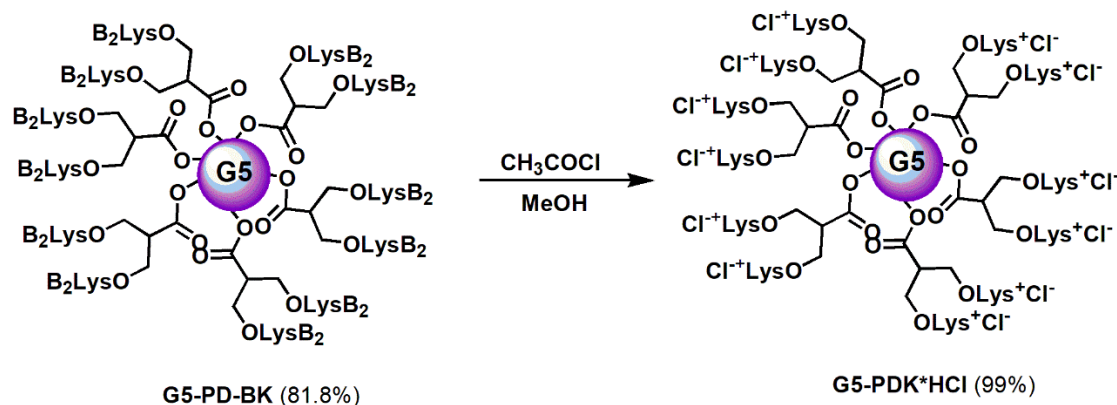


**Scheme S2.** Synthesis of Boc-protected dendrimer G5-PD-BK. PD = propanediol; B = Boc-protecting group; Lys or K = Lysine; G5 = generations number [1].

A solution of G5-PD-OH (68.3 mg; 0.0094 mmol) in dry DMF (1.5 mL) was added with Boc<sub>2</sub>-Lys-OH (76.8 equiv; 250.0 mg; 0.7209 mmol), 4-dimethylaminopyridine (DMAP) (38.4 equiv.; 44.1 mg; 0.3610) and *N*-ethyl-*N*-(3-dimethylamino)propyl carbodiimide hydrochloride (EDC) (76.8 equiv.; 111.9 mg; 0.7209 mmol). The solution was kept under magnetic stirring at r.t. for 24 h then added with 15 mL of ethyl acetate (EtOAc) to produce a suspension which was washed with 10% aq. KHSO<sub>4</sub> (3x15 mL). The aqueous washings were extracted with EtOAc and the combined organic phases were washed with aq. 15% NaOH followed by water then dried on MgSO<sub>4</sub> overnight. The removal of the solvent at reduced pressure afforded the Boc-protected lysine-modified dendrimer G5-PD-BK as off white glassy solid (217.5 mg; 0.0077 mmol; 81.8% yield).

FTIR (KBr, cm<sup>-1</sup>): 3380 (NH), 1747 (C=O ester), 1710 (C=O urethane), 1527 (NH). <sup>1</sup>H NMR (400 MHz, CDCl<sub>3</sub>), δ (ppm): 0.95-1.90 (m, 572H, CH<sub>3</sub> of dendrimer + CH<sub>2</sub>O propanediol + CH<sub>2</sub>CH<sub>2</sub>CH<sub>2</sub> of Lys), 1.43 (s, 576H, CH<sub>3</sub> of Boc), 1.44 (s, 576H, CH<sub>3</sub> of Boc), 3.10 (m, 128H, CH<sub>2</sub>NH of Lys), 3.56 (partially overlapped signal, 2H, CH<sub>2</sub>O propanediol), 4.25 (m, 314H, CH<sub>2</sub>O of dendrimer and of propanediol + CHNH of Lys), 4.70-5.50 (m, 128H, αNHBoc + εNHBoc of Lys). <sup>13</sup>C NMR (CDCl<sub>3</sub>, 100 MHz), δ (ppm): 14.20-17.90 (CH<sub>3</sub> of G1, G2, G3, G4, G5), 22.57 (CH<sub>2</sub>), 28.36 (CH<sub>3</sub> of Boc), 28.47 (CH<sub>3</sub> of Boc), 29.57 (CH<sub>2</sub>), 31.84 (CH<sub>2</sub>), 40.04 (CH<sub>2</sub>NH), 46.42 (quaternary C), 53.37 (CHNH), 65.41-65.60 (CH<sub>2</sub>O of G1, G2, G3, G4, G5), 79.02 (quaternary C of Boc), 79.80 (quaternary C of Boc), 155.63 (C=O urethane), 156.17 (C=O urethane), 172.32 (C=O amino acid + C=O ester of G1, G2, G3, G4, G5), CH<sub>2</sub> of propanediol not detectable. Found: C, 56.78; H, 8.30; N, 6.00. C<sub>1337</sub>H<sub>2296</sub>N<sub>128</sub>O<sub>508</sub> requires C, 56.76; H, 8.18; N, 6.34%.

### S1.2.2. Acidic Deprotection of G5-PD-BK to obtain G5-PDK \* 128 HCl



**Scheme S3.** Synthesis of cationic dendrimer G5-PDK having 128 protonated nitrogen atoms. PD = propanediol; B = Boc-protecting group; Lys or K = Lysine; G5 = generations number [1].

A solution of G5-PD-BK (208.8 mg; 0.0074 mmol) in 1 mL methanol (MeOH) was cooled to 0 °C and treated with acetyl chloride (2 equiv./Boc-groups to be removed; 148.3 mg; 1.8893 mmol; 134.8  $\mu\text{L}$ ). The solution was kept at r.t. under magnetic stirring for 24 h, then it was concentrated at reduced pressure, taken with MeOH, and precipitated into acetone. The dendrimer in the form of hydrochloride was recovered as oil after centrifugation, washed repeatedly with fresh acetone, recovered all times by centrifugation, and finally dried at reduced pressure. G5-PDK \* 128 HCl was obtained as white highly hygroscopic solid, which was stored under vacuum over  $\text{P}_2\text{O}_5$ . (147.2 mg, 0.0073 mmol, 99 % yield).

FTIR (KBr,  $\text{cm}^{-1}$ ): 3431 ( $\text{NH}_3^+$ ), 1744 (C=O ester), 1635 (NH).  $^1\text{H}$  NMR (400 MHz,  $\text{DMSO}-d_6$ ),  $\delta$  (ppm): 1.03-1.99 (m, 570H,  $\text{CH}_3$  of dendrimer +  $\text{CH}_2\text{CH}_2\text{CH}_2$  of Lys), 1.70 (m, 2H,  $\text{CH}_2$  propanediol), 2.76 (m, 128H,  $\text{CH}_2\text{NH}_3^+$  of Lys), 3.56 (partially overlapped signal, 2H,  $\text{CH}_2\text{O}$  propanediol), 3.99 (m, 64H,  $\text{CHNH}_3^+$  of Lys), 4.10-4.50 (m, 250H,  $\text{CH}_2\text{O}$  of propanediol and of dendrimer +  $\text{CHNH}_3^+$  of Lys), 8.20 (br s, 192H,  $^{\alpha}\text{NH}_3^+$ ), 8.82 (br s, 192H,  $^{\epsilon}\text{NH}_3^+$  of Lys).  $^{13}\text{C}$  NMR ( $\text{DMSO}-d_6$ , 100 MHz),  $\delta$  (ppm): 19.33 ( $\text{CH}_3$ ), 23.14 ( $\text{CH}_2$ ), 28.01 ( $\text{CH}_2$ ), 31.01 ( $\text{CH}_2$ ), 40.02 ( $\text{CH}_2\text{NH}_3^+$ ), 47.70 (quaternary C), 53.55 ( $\text{CHNH}_3^+$ ), 67.65-67.82 ( $\text{CH}_2\text{O}$  and of G1, G2, G3, G4), 170.68-173.33 (C=O of amino acid + ester of G1, G2, G3, G4),  $\text{CH}_2$  of propanediol not detectable. Found: C, 56.78; H, 8.30; N, 6.00.  $\text{C}_{697}\text{H}_{1400}\text{N}_{128}\text{O}_{252}\text{Cl}_{128}$  requires C, 41.56; H, 7.00; N, 8.90; Cl, 22.53%.

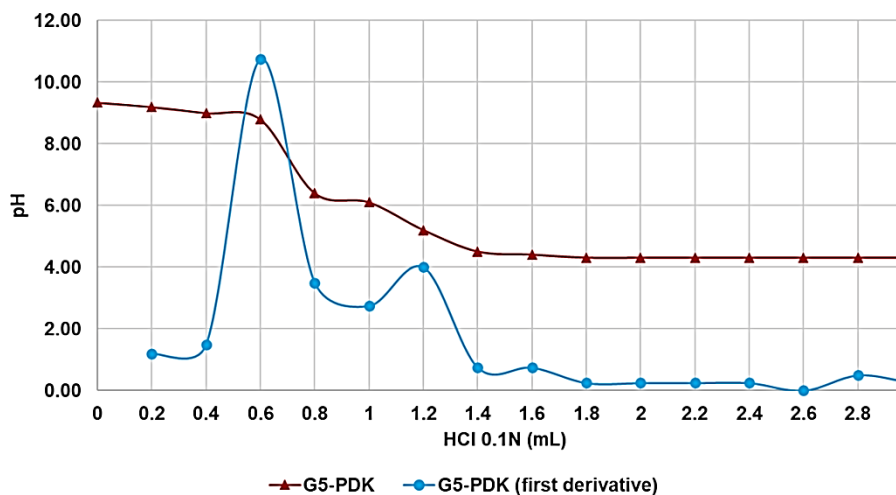
Section S1.2.3. Potentiometric titration of G5-PDK [1].

**Table S1.** Data of potentiometric titration used for constructing the titration curve and those of dpH/dV used for constructing the first derivative curve.

mL HCl 0.1N	pH	dpH/dV
0.0	9.54 ± 0.02	---
0.2	9.30 ± 0.03	1.2
0.4	9.00 ± 0.02	1.5
0.6	6.85 ± 0.05	10.75
0.8	6.15 ± 0.01	3.5
1.0	5.60 ± 0.04	2.75
1.2	4.80 ± 0.02	4
1.4	4.65 ± 0.02	0.75
1.6	4.50 ± 0.03	0.75
1.8	4.45 ± 0.02	0.25
2.0	4.40 ± 0.04	0.25
2.2	4.35 ± 0.02	0.25
2.4	4.30 ± 0.01	0.25
2.6	4.30 ± 0.009	0
2.8	4.20 ± 0.009	0.5
3.0	4.15 ± 0.01	0.25

G5-PDK		
Max dpH/dV	10.75	4.0
HCl (mL)	0.6	1.2
pH	6.85	4.80



**Figure S2.** Titration curve of G5-PDK (red line); first derivative line of the titration curve (light blue line), which shows separate peaks in correspondence of each end point of the titration curve [1].

**Table S2.** Main physical features of G5-PDK [1].

Physical characteristics	G5-PDK	
N <sup>1</sup>	128	
MW (calc.) <sup>2</sup>	20145.3	
MW (obs.) <sup>3</sup>	19961.2±480.2	
Error (%)	-0.9%	
Z-Ave (nm) <sup>4</sup>	203.0±2.6 <sup>5</sup>	203.0±2.6 <sup>6</sup>
PDI <sup>7</sup>	0.282±0.028 <sup>5</sup>	0.282±0.028 <sup>6</sup>
ζ-p (mV) <sup>8</sup>	+19.2±7.3	

<sup>1</sup> Number of protonated nitrogen atoms; <sup>2</sup> molecular weight (MW) computed according to the structure of G5-PDK confirmed by NMR analysis and elemental analysis; <sup>3</sup> MW obtained by volumetric titration; <sup>4</sup> particle size by dynamic light scattering (DLS) analysis; <sup>5</sup> intensity-weighted mean hydrodynamic diameters; <sup>6</sup> number-weighted mean hydrodynamic diameters; <sup>7</sup> polydispersity index (DLS); <sup>8</sup> Z-potential (DLS).

**Section S2.** Antibacterial properties of G5-PDK evaluated against the most relevant MDR representative of both Gram-negative and Gram-positive species [1].

**Table S3.** MIC values of G5-PDK on relevant representatives of Gram-positive and Gram-negative bacteria obtained from experiments carried out in triplicate<sup>1</sup>, expressed as μM and as μg/mL.

Strains	G5-PDK (20145) <sup>2</sup>	Ciprofloxacin
	MIC μM (μg/mL)	MIC μM (μg/mL)
<i>E. faecalis</i> *	>25.4 (>512)	193.2 (64)
<i>E. faecium</i> *	>25.4 (>512)	772.7 (256)
<i>S. aureus</i> **	>25.4 (>512)	386.4 (128)
<i>S. epidermidis</i> **	>25.4 (>512)	193.2 (64)
<i>E. coli</i> #	>25.4 (>512)	96.6 (32)
<i>K. pneumoniae</i> #	>25.4 (>512)	96.6 (32)
<i>A. baumannii</i>	6.3 (128)	193.2 (64)

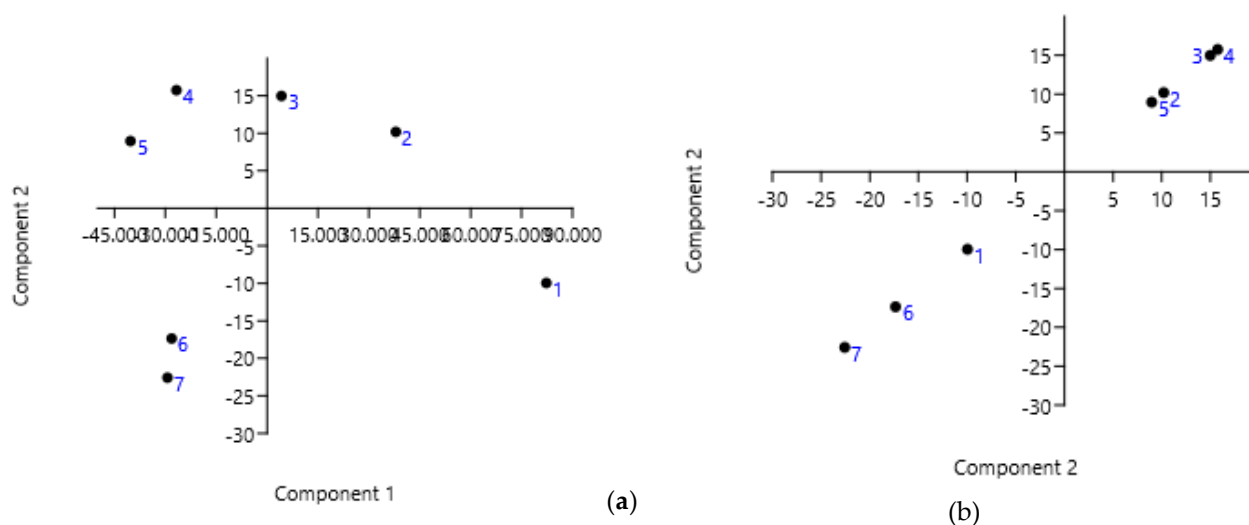
<sup>1</sup> The degree of concordance was in all the experiments 3/3, and standard deviation (±S.D.) was zero; <sup>2</sup> MW of G5-PDK; \* denotes vancomycin resistant isolates (VRE); \*\* denotes methicillin resistant isolates; # denotes a carbapenemase (KPC)-producing bacterium; *A. baumannii* was a MDR strain.

**Table S4.** MICs of G5-PDK obtained on isolates of the genus *Acinetobacter* from experiments conducted in triplicate<sup>1</sup> (expressed as μM and as μg/mL) compared to the MICs of ciprofloxacin.

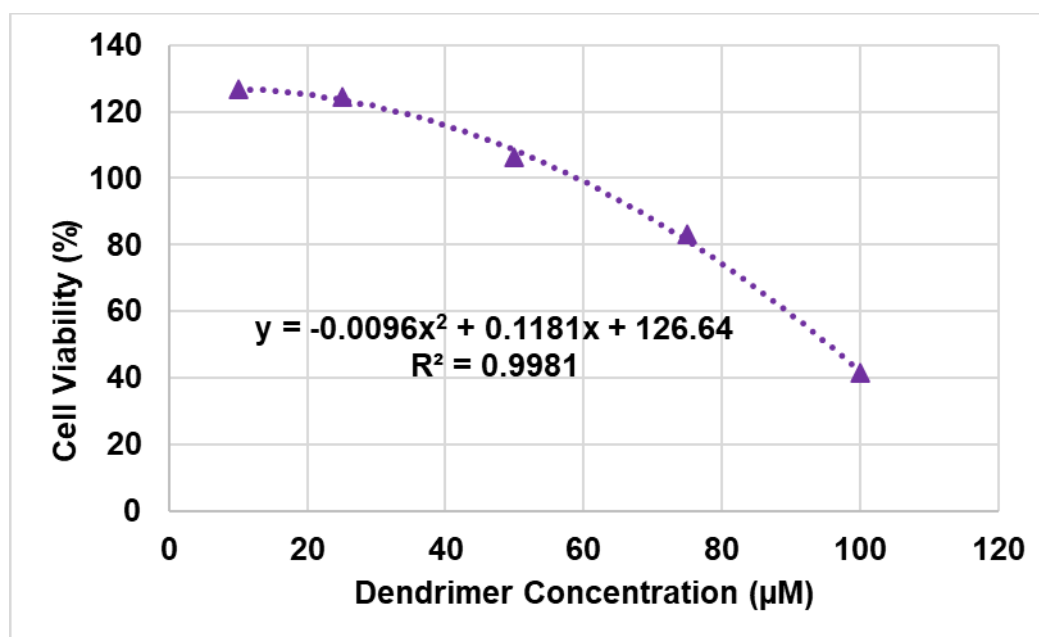
Strains	G5-PDK (20145) <sup>2</sup>	Ciprofloxacin
	MIC μM (μg/mL)	MIC μM (μg/mL)
<i>A. baumannii</i> 236	6.3 (128)	193.2 (64)
<i>A. baumannii</i> 245	6.3 (128)	1.6 (0,5)
<i>A. baumannii</i> 257	6.3 (128)	96.6 (32)
<i>A. baumannii</i> 279	12.7 (256)	48.3 (16)
<i>A. baumannii</i> 383	6.3 (128)	193.2 (64)
<i>A. baumannii</i> 406	6.3 (128)	96.6 (32)
<i>A. johnsonii</i> 387	6.3 (128)	0.9 (0.3)
<i>A. junii</i> 389	12.7 (256)	0.4 (0.125)
<i>A. pittii</i> 263	6.3 (128)	3.2 (1)
<i>A. pittii</i> 272	6.3 (128)	1.6 (0.5)
<i>A. ursingii</i> 388	3.2 (64)	0.4 (0.125)
<i>A. ursingii</i> 408	6.3 (128)	0.8 (0.25)

<sup>1</sup> The degree of concordance was in all the experiments 3/3, and S.D. was zero; <sup>2</sup> MW of G5-PDK; *Acinetobacters* are all MDR bacteria.

### Section S3. Cytotoxicity Experiments



**Figure S3.** PCA results on data concerning 12 h of cells exposure: score plot showing data locations on PC1 vs PC2 (a); score plot showing data locations on PC2 vs PC2 (b).



**Figure S4.** Second-degree polynomial regression model obtained from data concerning 12 h of cells exposure.

### References

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