

**Table S1:** Preliminary identification of microbial isolates and their biofilm forming ability

Isolate number	Categorization of Isolates Based on Culturing and Gram Staining <sup>1</sup>	Biofilm Formation (Average Optical Density at 630 nm) <sup>2,3</sup>	Classification of Biofilm Production <sup>4</sup>
1	G-ve LF	0.346	++
2	G-ve LF	0.170	+
3	G-ve NLF	0.155	0
4	G-ve LF	0.291	+
5	G-ve NLF	0.203	+
<b>6*</b>	<b>yeast</b>	<b>0.870</b>	<b>+++</b>
7	yeast	0.490	++
<b>8*</b>	<b>G-ve LF</b>	<b>0.953</b>	<b>+++</b>
9	yeast	0.432	++
10	yeast	0.501	++
11	G-ve LF	0.568	++
12	G-ve LF	0.209	+
13	G-ve LF	0.153	0
14	G-ve NLF	0.175	+
15	yeast	0.214	+
16	yeast	0.370	++
17	G-ve LF	0.180	+
18	G-ve LF	0.350	++
19	G+ve cocci	0.439	++
20	yeast	0.321	++
21	G-ve LF	0.388	++
22	G-ve LF	0.135	0
23	G-ve NLF	0.470	++
24	G-ve NLF	0.599	++
25	G-ve LF	0.165	+
<b>26*</b>	<b>G-ve NLF</b>	<b>0.947</b>	<b>+++</b>
27	yeast	0.468	++
28	yeast	0.193	+
29	G-ve LF	0.229	+
30	yeast	0.395	++
31	G-ve LF	0.131	0
32	G-ve LF	0.270	+
33	G-ve LF	0.098	0
34	G+ve cocci	0.114	0
<b>35*</b>	<b>G+ve cocci</b>	<b>0.751</b>	<b>+++</b>
<b>36*</b>	<b>G-ve LF</b>	<b>0.702</b>	<b>+++</b>
37	yeast	0.207	+

<sup>1</sup> culturing on nutrient agar, MacConkey no. 3 and Sabaroud agar followed by Gram staining, G+ve: Gram-positive, G-ve: Gram-negative, LF: lactose fermenters, NLF: non-lactose fermenters. <sup>2</sup>Average Blank = 0.125; <sup>3</sup>Optical density cut-off value (ODc) = 0.158; <sup>4</sup> 0: non-biofilm forming, +: weak, ++: moderate, and +++: strong-biofilm producing. \*selected strong-biofilm producing isolated.

Table S1: continued

Isolate number	Categorization of Isolates Based on Culturing and Gram Staining <sup>1</sup>	Biofilm Formation (Average Optical Density at 630 nm) <sup>2,3</sup>	Classification of Biofilm Production <sup>4</sup>
38	G-ve LF	0.406	++
39	G-ve LF	0.356	++
<b>40*</b>	<b>G-ve NLF</b>	<b>0.884</b>	<b>+++</b>
41	yeast	0.191	+
42	G+ve cocci	0.350	++
43	G-ve LF	0.145	0
44	G-ve LF	0.197	+
45	yeast	0.489	++
46	G-ve LF	0.149	0
47	G-ve LF	0.244	+
48	G+ve cocci	0.319	++
<b>49*</b>	<b>G-ve LF</b>	<b>0.869</b>	<b>+++</b>
50	G+ve cocci	0.380	++
51	G-ve LF	0.310	+
52	G-ve NLF	0.159	+
53	yeast	0.174	+
54	G-ve LF	0.138	0
<b>55*</b>	<b>G-ve LF</b>	<b>0.859</b>	<b>+++</b>
<b>56*</b>	<b>G-ve NLF</b>	<b>0.801</b>	<b>+++</b>
57	yeast	0.405	++
58	yeast	0.233	+
59	G+ve cocci	0.490	++
60	G+ve cocci	0.260	+
<b>61*</b>	<b>G-ve NLF</b>	<b>1.069</b>	<b>+++</b>
62	G-ve LF	0.350	++
63	G-ve LF	0.449	++
64	G-ve NLF	0.576	++
65	yeast	0.346	++
66	G-ve NLF	0.190	+
67	G+ve cocci	0.400	++
68	G-ve LF	0.317	++
69	G-ve NLF	0.163	+
70	G-ve NLF	0.360	++
71	yeast	0.178	+
72	G-ve LF	0.174	+
73	G-ve LF	0.12	0
74	G+ve cocci	0.37	++

<sup>1</sup> culturing on nutrient agar, MacConkey no. 3 and Sabaroud agar followed by Gram staining, G+ve: Gram-positive, G-ve: Gram-negative, LF: lactose fermenters, NLF: non-lactose fermenters. <sup>2</sup>Average Blank = 0.125; <sup>3</sup>Optical density cut-off value (ODc) = 0.158; <sup>4</sup> 0: non-biofilm forming, +: weak, ++: moderate, and +++: strong-biofilm producing. \*selected strong-biofilm producing isolated.

**Table S2.** Summary of the classification of microbial isolates recovered from urinary silicone catheters according to their growth on culture media, Gram-staining reaction, and degree of biofilm formation

Microbe	Biofilm production				Total
	Strong biofilm	Moderate biofilm	Weak biofilm	Non-biofilm	
Gram Negative Bacilli	8	13	16	9	46
Gram Positive Cocci	1	7	1	1	10
Yeast	1	10	7	0	18
Total	10	30	24	10	74

**Table S3:** Biochemical Identification of Selected Strong-biofilm forming Clinical Isolates using VITEK 2 Compact system

Type	Microbial Type	Isolate Number	Strain Identification by Vitek2 compact	Bionumber
Gram-negative bacilli	Lactose Fermenters	8	<i>Klebsiella pneumoniae</i>	6607730773565010
		49		6607734753564010
		55		6667735753765011
	Non-Lactose Fermenters	36	<i>Escherichia coli</i>	0405610554566211
		26	<i>Pseudomonas aeruginosa</i>	0003043003500252
		40		0003453143500210
		61		0003453143500210
56	<i>Alcaligenes faecalis</i>	0000001103500042		
Gram-positive cocci	cocci	35	<i>Staphylococcus aureus</i>	3063153303520050
Fungal isolates	yeast	6	<i>Candida tropicalis</i>	6627734553566010

**Table S4:** Antimicrobial Susceptibility Testing of Gram negative Strong-biofilm forming Clinical Isolates and their Interpretation<sup>1</sup>

Clinical Isolate Antibiotic tested	<i>E. coli</i>	<i>A. faecalis</i>	<i>K. pneumonia</i>			<i>P. aeruginosa</i>		
	Isolate 36	Isolate 56	Isolate 55	Isolate 49	Isolate 8	Isolate 26	Isolate 40	Isolate 61
<b>Beta-lactams</b>								
Ampicillin	≥ 32 / R <sup>2</sup>	-	≥ 32 / R	≥ 32 / R	≥ 32 / R	-	-	-
Ampicillin/Sulbactam	≥ 32 / R	-	≥ 32 / R	≥ 32 / R	≥ 32 / R	-	-	-
Piperacillin/Tazobactam	≥128 / R	≥ 128 / R	≥ 128 / R	16 / S <sup>3</sup>	16 / S	≥ 128 / R	≥ 128 / R	≥ 128 / R
Cefazolin	≥ 64 / R	16 / I <sup>4</sup>	≥ 64 / R	≥ 64 / R	≥ 64 / R	≥ 64 / R	≥ 64 / R	≥ 64 / R
Cefoxitin	≥ 64 / R	-	≥ 64 / R	≥ 64 / R	≥ 64 / R		-	-
Ceftazidime	32 / R	4 / S	≥ 64 / R	8 / I	8 / I	≥ 64 / R	32 / R	32 / R
Ceftriaxone	≥ 64 / R	≤ 1 / S	≥64 / R	≤ 1 / S	≤ 1 / S	-	-	-
Cefepime	≥ 64 / R	8 / S	≥ 64 / R	≤ 1 / S	≤ 1 / S	≥ 64 / R	≥ 64 / R	≥ 64 / R
Meropenem	≤ 0.25 / S	≤ 0.25 / S	≤ 0.25 / S	≤ 0.25 / S	≤ 0.25 / S	≤ 0.25 / S	≤ 0.25 / S	≤ 0.25 / S
ESBL	NEG	-	NEG	NEG	NEG	-	-	-
<b>Aminoglycoside</b>								
Amikacin	≤ 2 / S	4 / S	4 / S	≤ 2 / S	≤ 2 / S	4 / S	≤ 2 / S	≤ 2 / S
Gentamicin	≤ 1 / S	2 / S	≥ 16 / R	≤ 1 / S	≤ 1 / S	≥ 16 / R	≤ 1 / S	≤ 1 / S
Tobramycin	≤ 1 / S	2 / S	≥ 16 / R	≤ 1 / S	≤ 1 / S	≥ 16 / R	≤ 1 / S	≤ 1 / S
<b>Quinolones</b>								
Ciprofloxacin	≤ 0.25 / S	1 / S	≥ 4 / R	0.5 / S	0.5 / S	≥ 4 / R	≤ 0.25 / S	≤ 0.25 / S
Levofloxacin	1 / S	1 / S	≥ 8 / R	1 / S	1 / S	≥ 8 / R	1 / S	1 / S
Nitrofurantoin	≥ 512 / R	-	≤ 16 / S	64 / I	64 / I	-	-	-
<b>Folate inhibitors</b>								
Trimethoprim/Sulfamethoxazole	≥ 320 / R	≤ 20 / S	≥ 320 / R	≤ 20 / S	≤ 20 / S	-	-	-

<sup>1</sup> interpretation according to Performance Standards for Antimicrobial Susceptibility Testing, 32nd ed. CLSI supplement M100. Clinical and Laboratory Standards Institute; 2022.<sup>2</sup> R: resistant, <sup>3</sup>S: sensitive, <sup>4</sup>I: intermediate.

**Table S5:** Antimicrobial Susceptibility Testing of Gram positive Strong-biofilm forming Clinical Isolate and its Interpretation<sup>1</sup>

Antibiotics tested	<i>Staphylococcus aureus</i> (Isolate 35)	
	MIC µg/mL	Interpretation
<b><i>Beta-lactams</i></b>		
Benzylpenicillin	≤ 0.03	S
Oxacillin	1	S
Cefoxitin Screen	Positive	
<b><i>Aminoglycoside</i></b>		
Gentamicin	≤ 0.5	S
<b><i>Quinolones</i></b>		
Ciprofloxacin	≤ 0.5	S
Levofloxacin	0.25	S
Moxifloxacin	≤ 0.25	S
<b><i>Macrolides</i></b>		
Erythromycin	0.5	S
Clindamycin	≤ 0.25	S
Inducible Clindamycin Resistance	Negative	
<b><i>Streptogramin</i></b>		
Quinupristin/Dalfopristin	0.5	S
<b><i>Oxazolidinones</i></b>		
Linezolid	4	S
<b><i>Glycopeptide</i></b>		
Vancomycin	1	S
<b><i>Tetracycline</i></b>		
Tetracycline	≤ 1	S
<b><i>Glycylcycline</i></b>		
Tigecycline	≤ 0.12	S
<b><i>Others</i></b>		
Nitrofurantoin	≤ 16	S
<b><i>Antimycobacterials</i></b>		
Rifampicin	≤ 0.5	S
<b><i>Folate inhibitors</i></b>		
Trimethoprim/Sulfamethoxazole	≤ 10	S

<sup>1</sup> interpretation according to Performance Standards for Antimicrobial Susceptibility Testing. 32nd ed. CLSI supplement M100. Clinical and Laboratory Standards Institute; 2022.

**Table S6.** Summary of the classification of the strong biofilm forming isolates according to the interim standard definitions for acquired resistance

<b>Strong Biofilm-producing clinical isolates Classification of Antibiotics Susceptibility</b>		
<i>Klebsiella pneumoniae</i>	#08	S <sup>1</sup>
	#49	S
	#55*	MDR <sup>2</sup>
<i>Pseudomonas aeruginosa</i>	#61	S
	#40	S
	#26*	XDR <sup>3</sup>
<i>Alcaligenes faecalis</i>	#56*	S
<i>Escherichia coli</i>	#36*	MDR
<i>Staphylococcus aureus</i>	#35*	S

\* Molecularly identified isolates, <sup>1</sup> S: susceptible, <sup>2</sup> Multidrug-resistant, <sup>3</sup> extensively drug-resistant.

**Table S7.** Effect of ZnO NPs at subMICs concentrations on growth and biofilm formation by selected isolates

Microbe	ZnO NPs (mg/mL)						
	Control	0.062	0.125	0.25	0.5	1	
<i>P. aeruginosa</i>	+	+	+	+	+	+	Growth <sup>1</sup>
	0.947 ± 0.1	0.306 ± 0.091	0.257 ± 0.095	0.189 ± 0.007	0.171 ± 0.041	0.165 ± 0.067	OD630
<i>K. pneumoniae</i>	+	+	+	+	+	+	Growth
	0.859 ± 0.1	0.225 ± 0.009	0.352 ± 0.026	0.448 ± 0.007	0.488 ± 0.117	0.478 ± 0.120	OD630
<i>E. coli</i>	+	+	+	+	-	-	Growth
	0.702 ± 0.1	0.104 ± 0.004	0.112 ± 0.004	0.109 ± 0.006	0.109 ± 0.011	0.111 ± 0.003	OD630
<i>A. faecalis</i>	+	+	+	+	+	-	Growth
	0.801 ± 0.1	0.306 ± 0.043	0.350 ± 0.084	0.218 ± 0.089	0.262 ± 0.091	0.198 ± 0.068	OD630
<i>S. aureus</i>	+	+	+	-	-	-	Growth
	0.751 ± 0.1	0.760 ± 0.115	0.406 ± 0.194	0.259 ± 0.131	0.163 ± 0.004	0.291 ± 0.079	OD630
<i>C. tropicalis</i>	+	+	+	+	+	+	Growth
	0.870 ± 0.1	0.259 ± 0.016	0.251 ± 0.016	0.177 ± 0.053	0.149 ± 0.031	0.142 ± 0.026	OD630

1: growth either as + = growth or - = no growth observed

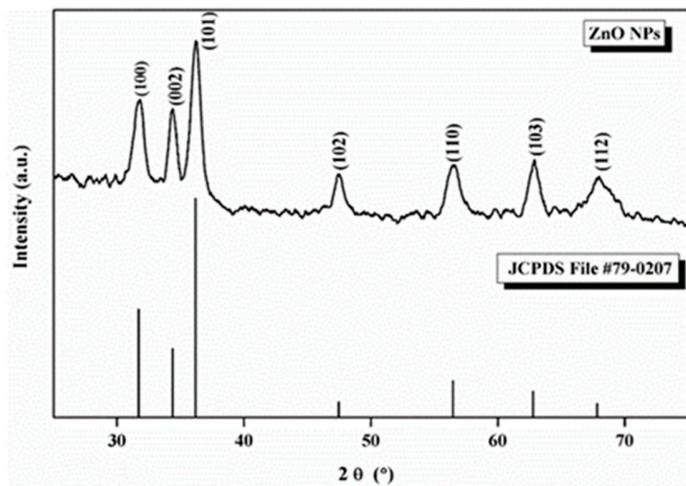


Figure S1. X-ray Diffraction Analysis patterns of the synthesized ZnO NPs relative to the standard ZnO NPs

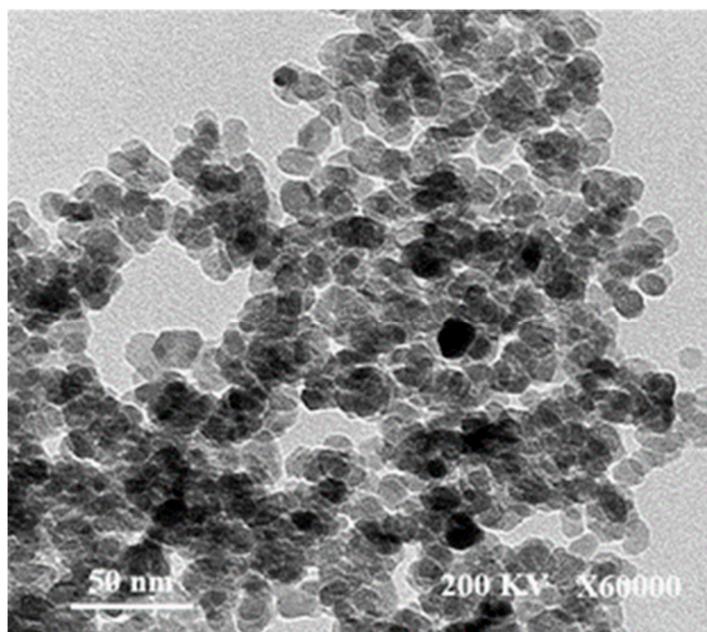
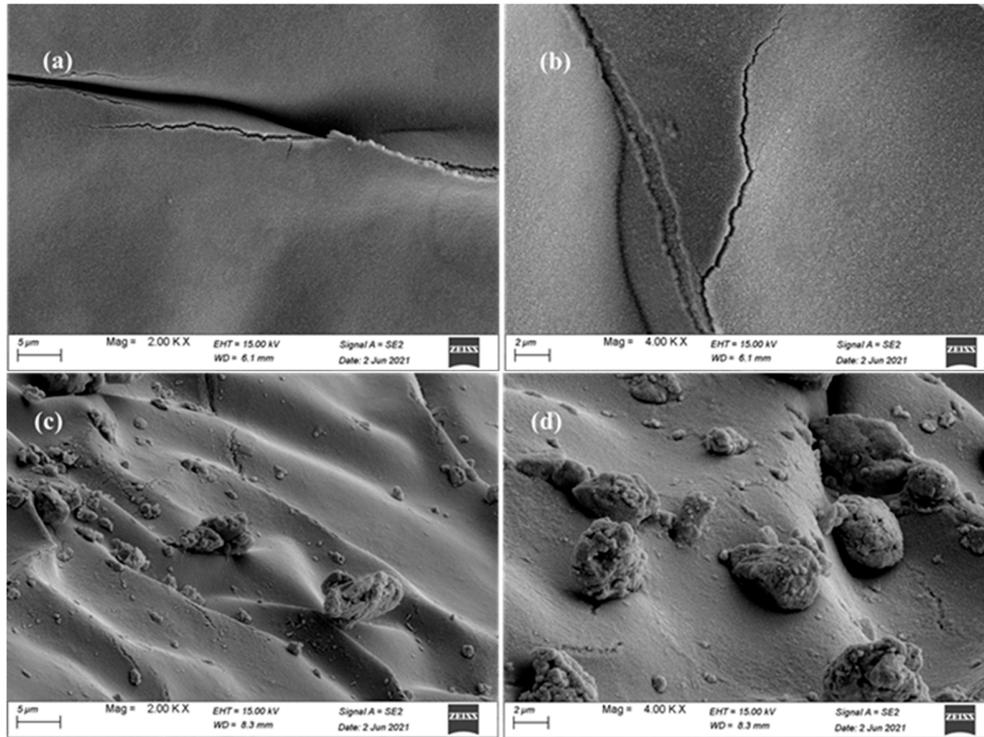


Figure S2. Transmission Electron Microscopy Image of the synthesized ZnO NPs



**Figure S3.** Scanning electron microscopy (SEM) microphotograph of SR and SR-g-AAc. (a) SR at at 2.00 KX and (v) at 4.00 KX, SR-g-AAc at (c) 2.00 KX and (d) 4.00 KX.