

Supplementary information

Extracellular probiotic lipase capped silver nanoparticles as highly efficient broad spectrum antimicrobial agents

Imran Khan^{1†}, Nivetha Sivasankaran^{1†}, Ravikiran Nagarjuna², Ramakrishnan Ganesan^{2*} and Jayati Ray Dutta^{1*}

¹Department of Biological Sciences, BITS Pilani, Hyderabad Campus, Jawahar Nagar, Kapra Mandal, Hyderabad – 500078, Telangana, India. Tel.: +91 40 6630 3542;

E-mail: jayati@hyderabad.bits-pilani.ac.in

²Department of Chemistry, BITS Pilani, Hyderabad Campus, Jawahar Nagar, Kapra Mandal, Hyderabad – 500078, Telangana, India. Tel: +91 40 6630 3602;

E-mail: ram.ganesan@hyderabad.bits-pilani.ac.in

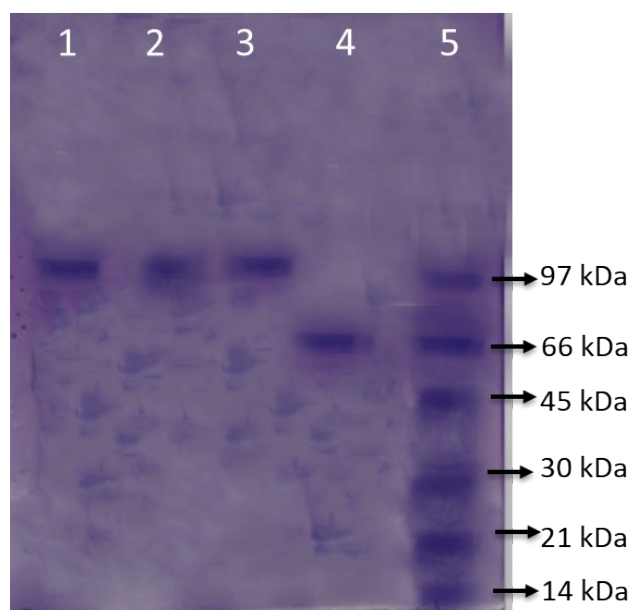


Figure S1. SDS-PAGE analyses of (1) 100, (2) 50 and (3) 25 $\mu\text{g/mL}$ lipase-capped Ag NPs synthesized from 1 mM AgNO_3 solution. The SDS-PAGE analyses of (4) pristine lipase and (5) 100 kDa protein ladder (Sigma Aldrich) are given for comparison.

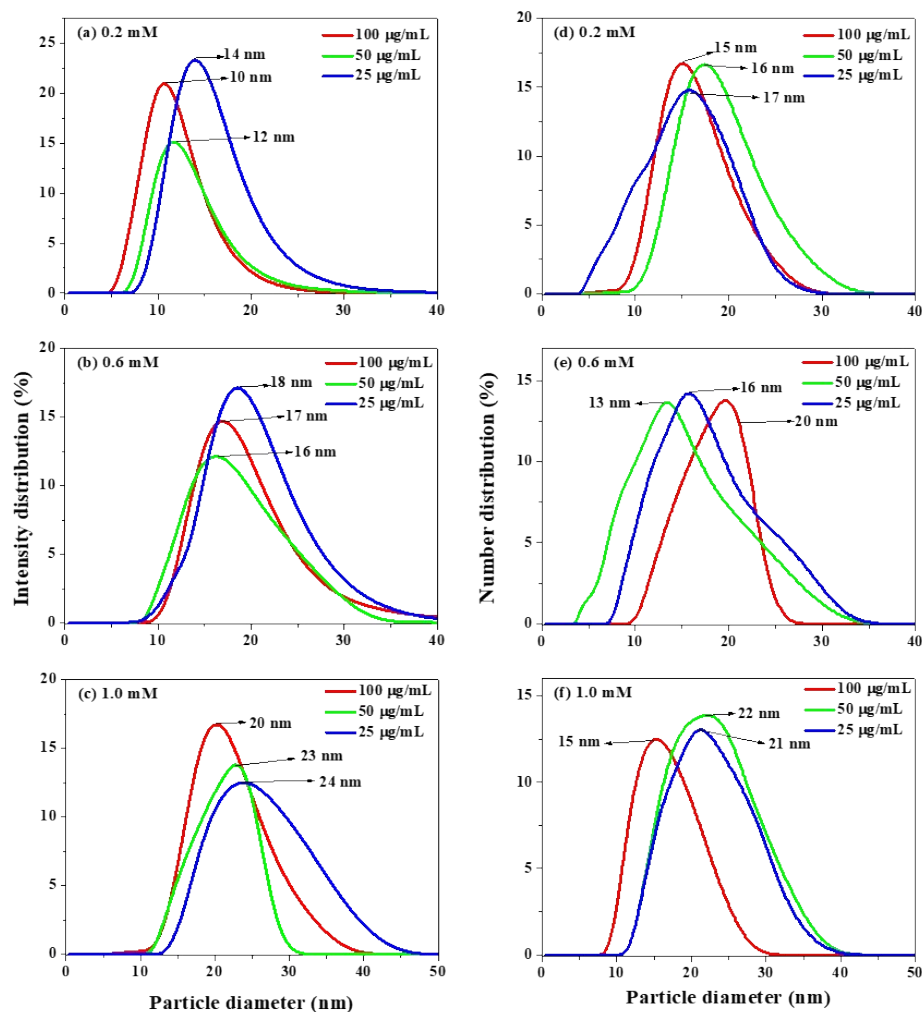


Figure S2. DLS measurements showing intensity (a-c) and number (d-f) distribution of Ag NPs synthesized from 0.2, 0.6 and 1.0 mM AgNO_3 solutions with varying amount of lipase capping agent.

Table S1. Comparative MIC values of Ag NPs against various microbial strains.

Microbes	Initial Concentration (CFU/mL)	MIC of Ag NPs	Reference
<i>Escherichia coli</i> <i>Staphylococcus aureus</i> Yeast	10^7	0.56 ng/mL 5.6 ng/mL 2.24 ng/mL	31
<i>Bacillus subtilis</i> <i>Staphylococcus aureus</i> <i>Pseudomonas aeruginosa</i> <i>Escherichia coli</i>	N/A	6.8 $\mu\text{g/mL}$ 5.1 $\mu\text{g/mL}$ 1.70 $\mu\text{g/mL}$ 3.4 $\mu\text{g/mL}$	35
<i>Escherichia coli</i> J53 <i>Escherichia</i> J53pMG101 <i>Salmonella</i> sp. <i>Staphylococcus aureus</i>	2.5 to 4×10^5	3.7 $\mu\text{g/mL}$ 4.0 $\mu\text{g/mL}$ 3.0 $\mu\text{g/mL}$ 4.0 $\mu\text{g/mL}$	36
<i>Pseudomonas aeruginosa</i>	10^6	2 $\mu\text{g/mL}$	37

<i>Escherichia coli</i> <i>Staphylococcus aureus</i>	10 ⁵	6.25 µg/mL 7.5 µg/mL	38
<i>Escherichia faecalis</i>	10 ⁸	5 mg/mL	39
<i>Escherichia coli</i> MTCC 443 <i>Escherichia coli</i> MTCC 739 <i>Bacillus subtilis</i> MTCC 441 <i>Staphylococcus aureus</i> NCIM 5021	10 ⁵	20 µg/mL 60 µg/mL 30 µg/mL 70 µg/mL	40
<i>Escherichia coli</i> 116S, J53 <i>Escherichia coli</i> J53pMG101	10 ⁸	0.216 ng/mL 8.64 ng/mL	41
<i>Escherichia coli</i>	5.0 × 10 ⁶	5 µg/mL	42
BCG <i>Escherichia coli</i>	10 ⁵	5 µg/mL 10 µg/mL	43
<i>Escherichia coli</i> <i>Acinetobacter baumannii</i> <i>Pseudomonas aeruginosa</i> <i>Bacillus subtilis</i> <i>Mycobacterium smegmatis</i> <i>Mycobacterium bovis</i> <i>Staphylococcus aureus</i>	10 ⁵	0.5 µg/mL 0.4 µg/mL 0.4 µg/mL 1.7 µg/mL 0.5 µg/mL 1.1 µg/mL 0.7 µg/mL	44
<i>Escherichia coli</i>	10 ⁸	1.35 ng/mL	45
<i>Salmonella enterica</i>	10 ⁵	9-14 µg/mL	46
<i>Escherichia coli</i> <i>Staphylococcus aureus</i> <i>Bacillus anthracis</i> <i>Candida albicans</i>	10 ⁵	1.7 µg/mL 1.9 µg/mL 3.0 µg/mL 19 µg/mL	47
<i>Staphylococcus aureus</i> <i>Pseudomonas aeruginosa</i> <i>Salmonella enterica</i> <i>Escherichia coli</i>	10 ⁵	4.7 µg/mL 2.3 µg/mL 2.3 µg/mL 1.2 µg/mL	48
<i>Escherichia Coli</i> <i>Salmonella typhimurium</i>	10 ⁵	4.0 µg/mL 6.0 µg/mL	49
<i>Bacillus cereus</i> <i>Bacillus subtilis</i> <i>Staphylococcus aureus</i> <i>Micrococcus luteus</i> <i>Enterobacter cloacae</i> <i>Serratia marcescens</i> <i>Shigella dysentery</i>	10 ⁸	60 µg/mL 58 µg/mL 59 µg/mL 60 µg/mL 49 µg/mL 50 µg/mL 48 µg/mL	50
<i>Escherichia coli</i> MTCC 443 <i>Bacillus subtilis</i> MTCC 441 <i>Escherichia coli</i> MTCC 739 <i>Staphylococcus aureus</i> NCIM 5021	10 ⁵	20 µg/mL 60 µg/mL 30 µg/mL 70 µg/mL	51
<i>Escherichia coli</i> <i>Staphylococcus aureus</i>	10 ⁵	16.9 µg/mL 8.45 µg/mL	52
<i>Pseudomonas putida</i> <i>Staphylococcus aureus</i> <i>Aspergillus niger</i>	10 ⁵	1.08 ng/mL 1.08 ng/mL 1.62 ng/mL	This work

The literature reporting the MIC values in the ng/mL are highlighted in red color.