*Research Article*

# **Actinobacterial-mediated synthesis of silver nanoparticles and their activity against pathogenic bacteria**

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**Abstract:** In this study, silver nanoparticles (AgNPs) were biosynthesised by using acidophilic actinobacterial SH11 strain isolated from pine forest soil. Isolate SH11 was identified based on 16S rRNA gene sequence to *Streptomyces kasugaensis* M338-M1T and *S*. *celluloflavus* NRRL B-2493T (99.8% similarity, both). Biosynthesised AgNPs were analysed by UV–visible spectroscopy, which revealed specific peak at *λ* = 420 nm. Transmission electron microscopy analyses showed polydispersed, spherical nanoparticles with a mean size of 13.2 nm, while Fourier transform infrared spectroscopy confirmed the presence of proteins as the capping agents over the surface of AgNPs. The zeta potential was found to be −16.6 mV, which indicated stability of AgNPs. The antibacterial activity of AgNPs from SH11 strain against gram-positive (*Staphylococcus aureus* and *Bacillus subtilis*) and gram-negative (*Escherichia coli*) bacteria was estimated using disc diffusion, minimum inhibitory concentration and live/dead analyses. The AgNPs showed the maximum antimicrobial activity against *E. coli*, followed by *B. subtilis* and *S. aureus*. Further, the synergistic effect of AgNPs in combination with commercial antibiotics (kanamycin, ampicillin, tetracycline) was also evaluated against bacterial isolates. The antimicrobial efficacy of antibiotics was found to be enhanced in the presence of AgNPs.

# **1** Introduction

Resistance of clinical pathogens to commercial antibiotics is increasing problem, which is intensified by a lack of new therapeutic agents [[1](#page-5-0)]. Hence, there is search for novel antimicrobial compounds by many researchers and pharmaceutical companies [\[2\]](#page-5-0). Silver nanoparticles (AgNPs) provide an alternative to commercial antibiotics by developing novel applications in the pharmaceuticals and medicine [[3](#page-5-0)].

Nanoscience and nanotechnology are emerging fields, which involve the synthesis and application of nanoscale materials and structures usually in the range of 1–100 nm [\[4](#page-5-0)]. One of the most important aspects of nanotechnology is synthesis of metal nanoparticles. The chemical procedures for synthesis of AgNPs are expensive and use toxic solvents extensively [[5](#page-5-0), [6](#page-5-0)]. Unlike chemical procedures, the biological systems for synthesis of AgNPs such as bacteria, fungi and plants are cheap, easy and ecofriendly  $[7-10]$ .

Microorganisms play an important role in remediation of toxic metals through reduction of metal ions and consequently has been considered as nanofactories [\[11\]](#page-5-0). One of the most promising organisms for synthesis of metal nanoparticles is actinobacteria, which is known as main producer of diverse secondary metabolites, such as antibiotics and many others biologically active compounds [\[6\]](#page-5-0). Since the last decade these organisms have being intensively isolated from extreme environments and screened for novel bioactive compounds [[12,](#page-5-0) [13](#page-5-0)]. However, the synthesis of AgNPs by actinobacteria from extreme habitats such as acidophilic soils ( $pH < 4.0$ ) has been not well studied.

Currently, the AgNPs are explored and extensively investigated as potential antimicrobials [[14–](#page-5-0)[16](#page-6-0)]. The small size and the high surface area to volume ratio of nanoparticles enhance their interaction with the microbial cells to carry out a broad range of potential antimicrobial activities [\[17](#page-6-0)]. AgNPs have been found to be active against gram-positive as well as gram-negative bacteria, including multidrug resistant strains as well as biofilm formation [[18–20](#page-6-0)].

Moreover, some authors have studied the synergism of AgNPs in combination with antibiotics such as amoxicillin and polymyxin

B and found higher bactericidal activity by binding of AgNPs with antibiotics [[21–24](#page-6-0)].

The aim of this study was to synthesise AgNPs using acidophilic actinobacterium and to evaluate their activity against human pathogenic bacteria.

# **2Materials and methods**

## *2.1 Isolation of actinobacterial strain*

The acidophilic actinobacterial SH11 strain was isolated from humic (H) layer of pine forest soil collected from southern slope of inland dune, near Torun, Poland ( $52^{\circ}55'37''N$ ,  $18^{\circ}42'11''E$ ) by the dilution plate procedure on the acidified starch–casein agar [[25\]](#page-6-0) as described previously by Golinska *et al.* [[26\]](#page-6-0). The isolate was maintained on acidified yeast extract–malt extract (ISP2) agar slants [\[27](#page-6-0)] at room temperature and as suspension of spores and mycelial fragments in 20% glycerol (v/v) at −80 °C.

## *2.2 Identification and phylogenetic analysis of actinobacterial SH11 strain*

Biomass for the molecular study was prepared by growing of SH11 strain in yeast extract–malt extract broth (ISP2; pH 5.5) at 28 °C for 2 weeks. Cells were harvested by centrifugation and washed twice in distilled water. The extraction of genomic DNA and amplification of 16S rRNA gene from SH11 strain were carried out according to the method described by Golinska *et al.* [\[28](#page-6-0)]. The amplified product was purified by GenEluate<sup>TM</sup>PCR Clean-Up Kit (Sigma) and sent for sequencing at the Institute of Biochemistry and Biophysics Polish Academy of Sciences, Warsaw, Poland. The closest phylogenetic neighbours of isolate SH11 based on 16S rRNA gene sequences were found using the EzTaxon server (http:// eztaxon-e.ezbiocloud.net/) [[29\]](#page-6-0). The resultant gene sequences were aligned with corresponding sequences of the type strains of the genera *Streptomyces* using ClustalW. Phylogenetic analyses were carried out using MEGA6 software [\[30](#page-6-0)]. The phylogenetic tree was inferred by using the neighbour-joining [[31\]](#page-6-0), the maximum-likelihood [\[32](#page-6-0)] and maximum-parsimony [[33\]](#page-6-0) tree-making algorithms. Further, the tree topology was evaluated by a bootstrap

analysis [\[32](#page-6-0)] of the neighbour-joining data based on 1000 resampling using MEGA6 software [[30\]](#page-6-0). The root position of the unrooted tree was estimated using the 16S rRNA gene sequence of *Streptomyces albus* subsp. *albus* DSM 40313T (GenBank accession number AJ621602).

# *2.3 Actinobacterial synthesis of AgNPs*

For synthesis of AgNPs, the 100 ml of ISP2 broth (pH 5.5) in 250  ml Erlenmeyer flask was inoculated with SH11 strain and incubated at 28 °C in the orbital shaker (150 rpm) for 14 days. The actinobacterial biomass was then centrifuged at 6000*g* for 15 min and washed three times with sterile distilled water. Subsequently, the cell pellet was suspended in 100 ml sterilised distilled water and incubated at  $28\text{ °C}$  for  $24\text{ h}$ . After that the cell biomas was separated by centrifugation at 6000*g* for 20 min, and obtained supernatant was challenged with  $AgNO<sub>3</sub>$  (1 mM final concentration) and kept at room temperature for 2–3 days. The supernatant without AgNO<sub>3</sub> was used as a control.

## *2.4 Characterisation of AgNPs*

*2.4.1 Visual detection and spectroscopy analysis:* The biosynthesis of Ag<sup>+</sup> ions into AgNPs in the reaction solution was observed by colour change from light-yellow to dark-brown. The presence of biosynthesised AgNPs was confirmed by UV–visible spectroscopy analysis in the wavelength range of 200–800 nm using Nano Drop ND 2000 (Thermo Scientific, USA).

To determine the biomolecules responsible for the reduction of silver ions and stabilisation of AgNPs in the solution, the Fourier transform infrared spectroscopy (FTIR) analysis was carried out. For FTIR analysis the powder of biosynthesised AgNPs were combined with dry KBr in the ratio of 1:100. The FTIR spectrum of sample was recorded by FTIR instrument (Spectrum 2000, Perkin-Elmer, USA). The measurements were carried out in the range of 4000–400  $\text{cm}^{-1}$  at a resolution of 4  $\text{cm}^{-1}$ .

*2.4.2 Transmission electron microscopy (TEM) analysis:* The TEM analysis was performed to determine the size and morphology of biosynthesised AgNPs by using FEI Tecnai F20 X-Twintool Electron Microscope (USA) operated at an accelerating voltage of 100 kV. For analysis the biosynthesised AgNPs solution was dropped on copper grids coated with carbon (400 µm mesh size) and kept at room temperature to dry. TEM measurements were then performed and the obtained data were evaluated by Statistica Software (StatSoft, USA).

*2.4.3 Zeta potential analysis:* To determine the stability of AgNPs, the zeta potential analysis was conducted by using the Zetasizer Nano ZS 90 (Malvern Instrument Ltd, UK). For the analysis the 25 µl of AgNPs solution was diluted ten times and sonicated for 15 min at 20 Hz. Then sample was filtered through of 0.22 µm millipore filter and used for study.

## *2.5 Determination of minimum inhibitory concentration (MIC) of antibiotics against tested bacteria*

The MIC is defined as the lowest concentration of chemicals that inhibits the growth of the organism. The MIC assay of antibiotics (ampicillin, kanamycin or tetracycline) against gram-positive (*Staphylococcus aureus* ATCC6338 and *Bacillus subtilis* PCM2021) and gram-negative (*Escherichia coli* ATCC8739) bacteria was performed using eTest strips (Biomerieux) in the range of 0.016–256 µg/ml of antibiotic. Briefly, 100 µl of bacterial inoculum  $(1 \times 10^6 \text{ CFU/ml})$  was spread on to the surface of the tripticase soy agar (Becton Dickinson) medium with sterile spreader. Subsequently, eTest stripe was placed onto the surface of inoculated medium and incubated for 24 h at 37 °C. The assay was performed in triplicate.

## *2.6 Determination of the MIC and minimum bactericidal concentration (MBC) of AgNPs*

The synthesised AgNPs were tested (in triplicate) for determination of MIC by microtiter broth dilution method. Trypticase soy broth (TSB, Becton Dickinson) was used as diluents for bacterial strains. The final concentration of bacteria in each well was  $1 \times 10^6$ CFU/ml, and various concentrations (from  $5.0$  to  $200 \mu g/ml$ ) of AgNPs were used. The positive and negative controls were also maintained. The microtiter plates were incubated at 37 °C for 24 h and then read at 450 nm using BIOLOG multimode reader to determine the MIC and MBC values. The MIC was defined as the lowest concentration of AgNPs that inhibited the growth of microorganisms, while the MBC end point was defined as the lowest concentration of AgNPs that killed ≥90% of bacterial cells.

#### *2.7 Assessment of antibacterial activity of AgNPs individually and in combination with antibiotics*

The MIC values of AgNPs from actinobacterial strain SH11 and commercial antibiotic were used to estimate accurate synergistic effect of AgNPs on antibiotic (kanamycin, ampicillin and tetracycline) activity. The assay was performed using 96-well culture plates in triplicate. The final concentration of bacteria in each well was  $1 \times 10^6$  CFU/ml. The TSB (Becton Dickinson) was used as diluent for bacterial strains, antibiotics and AgNPs. The positive and negative controls were maintained. The microtiter plates were read at 450 nm on multimode reader (BIOLOG, USA) after incubation at 37 °C for 24 h to determine the bacterial growth inhibition (%).

## *2.8 Bacterial viability assay after treatment with biosynthesised AgNPs*

The viability assays of bacterial pathogens after treatment with AgNPs were carried out by using Live/Dead baclight kit (Invitrogen). To determine viability of bacterial cells, 1 ml of bacterial suspension (*E. coli*, *B. subtilis* or *S. aureus*) in TSB (1 ×  10<sup>6</sup>  CFU/ml) was treated with biosynthesised AgNPs (at final concentration of MIC of AgNPs for each tested bacteria) and incubated for 24 h in the orbital shaker (100 rpm) at 37 °C. The bacterial cells in TSB  $(1 \times 10^6 \text{ CFU/ml})$  were used as a control. After incubation time the SYTO 9 (green-fluorescent nucleic acid stain) and propidium iodide (the red-fluorescent nucleic acid stain) dyes were added (both 1.5 µl/ml) into each experimental sample and controls. Then all samples were mixed thoroughly and incubated at room temperature for 30 min in the darkness. Subsequently, the samples were 1000 times diluted with sterile distilled water and the 100 µl of each stained sample was filtered through 0.2 μm black polycarbonate filters (Isopore; Millipore). Each filter, containing labelled bacteria, was placed on a glass slide with two drops of BacLight mounting oil and covered with a clear glass cover slip. The sample counting was performed with an E800 epifluorescence microscope (Nikon Eclipse E200, Melville, New York, USA) with a  $100\,\text{W}$  mercury light source. Images were captured with a Cool Cam camera (Cool Camera, Decatur, Georgia, USA) and associated software (MultiScanBase v.18.03, Image Pro Plus V.4.5; Media Cybernetic, Silver Spring, Maryland, USA). The numbers of dead and live cells were estimated from a count of five or more randomly chosen microscopic fields (×1000). At least 400 bacteria were counted per sample. The SYTO 9 and propidium iodide dyes differ both in their spectral characteristic and in their ability to penetrate healthy bacterial cells. Thus, with an appropriate mixture of the SYTO 9 and propidium iodide stains, bacteria with intact cell membranes stain fluorescent green, whereas bacteria with damaged membranes strain fluorescent red.

# **3Statistical analysis**

To determine whether there are any significant differences among the activities of AgNPs, antibiotics and the combination of AgNPs and antibiotics, we applied 'one way analysis of variance' for the relative variability within above parameters.



**Fig. 1** *Neighbour-joining tree based on nearly complete 16S rRNA gene sequences (1374–1485 nucleotides) showing relationships between the isolate SH11 and the type strains of Streptomyces sp. Numbers at the nodes are percentage bootstrap values based on 1000 resampled datasets, only values above 30% are given. T-type strain. Bar 0.005 substitutions per nucleotide position. The root position of the tree was determined using S. albus subsp. albus DSM 40313T as outgroup*



**Fig. 2** *Detection of AgNPs from SH11 strain by UV–visible spectrophotometer – experimental (a), control (b). Visual detection of synthesised AgNPs – (A) experimental sample after treatment with AgNO3 after 72 h of incubation, (B) control before treatment with AgNO3*

## **4Results**

#### *4.1 Identification and phylogenetic analysis of actinobacterial SH11 strain*

Almost complete 16S rRNA gene sequence (1423 nucleotides [nt]) of the isolate SH11 was determined (GenBank accession number: KU587966). Isolate SH11 formed a distinct subclade in the *Streptomyces* 16S rRNA gene tree together with the type strains of *S. kasugaensis* M338-M1T and *S. celluloflavus* NRRL B-2493T, the taxons which were supported by N-J tree-making algorithm and by a bootstrap value of 100% (Fig. 1). The isolate shared its highest 16S rRNA similarity with the type strains of *S. kasugaensis* M338- M1T and *S. celluloflavus* NRRL B-2493T, namely 99.8%, a value which corresponded to three nucleotide differences at 1421 locations.

#### *4.2 Synthesis and characterisation of AgNPs*

The AgNPs were successfully synthesised from SH11 actinobacterial strain which was observed by colour change of the cell free filtrate from light-yellow to dark-brown after challenging with 1 mM silver nitrate (Fig. 2). Furthermore, the UV–visible spectroscopy analysis of AgNPs revealed sharp narrow peak at *λ* =  420 nm and confirmed presence of biosynthesised AgNPs (Fig. 2).

The FTIR spectroscopy analysis showed intense absorption bands at 3440.7, 2927.4, 2858.6, 1631.9, 1437.2, 1387.6 and 1030.1 cm−1 which corresponded to different functional groups on nanoparticles and indicated the presence of stabilising protein molecules on the surface of AgNPs (Figs. [3](#page-3-0)*a* and *b*).

The TEM analysis indicated that the biosynthesised AgNPs from SH11 strain were spherical in shape and polydispersed. However, few aggregates of AgNPs were also found at some places. The AgNPs showed the size in the range of 4–24 nm, with a mean size of 13.2 ( $\pm$ 2.9) nm (Fig. [4\)](#page-4-0).

The zeta potential of biosynthesised AgNPs was found to be −16.6 mV, which indicated stability of Ag nanoparticles (Fig. [5\)](#page-4-0).

#### *4.3 Antibacterial activity of biosynthesised AgNPs individually and in combination with antibiotics*

The AgNPs exhibited the lowest MIC against *S. aureus* and *B. subtilis*, both at 40 μg/ml, followed by *E. coli* at 70 μg/ml. The MBC was estimated for all tested bacteria at [1](#page-4-0)40 µg/ml (Table 1).

The results of antibiotic sensitivity test for bacterial strains were presented in Table [1.](#page-4-0) All the bacterial pathogens exhibited the highest sensitivity to ampicillin, followed by kanamycin and tetracycline.

The enhanced antibacterial activity of antibiotics in the presence of AgNPs was observed against all tested pathogens. However, antibacterial activity of tetracycline was most enhanced. The highest synergistic activity of AgNPs was found against *S. aureus*, followed by *B. subtilis* and *E. coli* (Table [1\)](#page-4-0).

#### *4.4 Bacterial viability assay after treatment with AgNPs*

Live/Dead bacLight kit was used to estimate the per cent survival rate of bacterial cells after treatment with AgNPs. The ratio of live to dead bacterial cells of analysed control samples (without AgNPs) was found to be 91.9 and 8.1% for *S. aureus*, 90.2 and 9.8% for *B. subtilis* then 90.1 and 9.9% for *E. coli*. While, in experimental samples after treatment with AgNPs the ratio of live/ dead bacterial cells was recorded as follows 85, 7 and 14.3% for *S. aureus*, 60.0 and 40.0% for *B. subtilis* and 30.8 and 69.2% for *E. coli* (Fig. [6\)](#page-5-0).

## **5Discussion**

Antibiotic resistance of the pathogenic bacteria has been a great problem since the last decade [[23\]](#page-6-0). Hence, there is a need for the development of new antibacterial agents. The researchers are moving towards nanoparticles especially AgNPs to solve the

<span id="page-3-0"></span>

**Fig. 3** *FTIR analysis of AgNPs synthesised from SH11 (a)* Experimental, *(b)* Control

problem of emerging pathogens including multi-drug resistant bacteria [\[34](#page-6-0)]. Hence, the AgNPs have appeared as a promising antibacterial candidate in the medical field [\[23](#page-6-0), [34, 35](#page-6-0)].

In this study, SH11 actinobacterial strain was studied for synthesis of AgNPs by using a simple biological synthesis process.

The colour change from pale-yellow to dark-brown indicates the formation of AgNPs. The brown colour is the result of excitation of surface plasmon vibration in the metal nanoparticles and is typical of the AgNPs presence [[36\]](#page-6-0). The visible observation of synthesis of AgNPs by colour change were also reported by Zarina and Nanda [\[2](#page-5-0)], and Golinska *et al.* [\[37](#page-6-0)] with a cell filtrate of *Streptomyces*-MS 26 and *Pilimelia columellifera* subsp. *pallida* after treatment with 1 mM AgNO<sub>3</sub>. Such observation clearly indicates that SH11 strain had the potential to reduce the silver ions to AgNPs.

The UV–visible spectrum in the form of sharp peak, at around 420 nm is specific for the synthesis of AgNPs [[2](#page-5-0), [38\]](#page-6-0). Our results are similar to findings of Otari *et al.* [\[38](#page-6-0)] and Zarina and Nanda [[2](#page-5-0)]. Authors reported the absorption peak at about 420 nm when studied *Rhodococcus* sp. and *Streptomyces* sp., respectively.

FTIR spectroscopy studies reveal the presence of specific functional groups constituting organic molecules [\[39,](#page-6-0) [40](#page-6-0)]. The band at 3440.7 cm−1 is characteristic for vibrations of functional groups such as amino groups (N–H) [[41](#page-6-0)]. The peaks in the range of 2850–2960 cm<sup>-1</sup> are associated with stretching vibration of C−H and the absorbance peak at 1387.6 cm<sup>-1</sup> can be assigned to CH<sub>3</sub> sym. bending which is characteristic for alkanes group [[4](#page-5-0)]. The absorption band at 1631.9 cm−1 resulted from the amide I vibration, which originated mainly from stretching of carbonyl groups (C=O) [\[41](#page-6-0)]. Moreover, the peak at  $1437.2 \text{ cm}^{-1}$  can be assigned to the C–H deformation vibration, which represents the presence of alcoholic bonded [[19\]](#page-6-0). The peak at around 1030.1 cm  $^{-1}$  can be assigned as C–N stretching vibrations and these results showed similarity with findings of Zarina and Nanda [\[4\]](#page-5-0) and Chauhan *et al.* [[42\]](#page-6-0). The FTIR spectroscopy study confirmed the presence of proteins on the surface of AgNP.

It is claimed that the carbonyl group of amino acid residues and peptides of proteins has the stronger ability to bind to metal ions [[7,](#page-5-0) [43,](#page-6-0) [44\]](#page-6-0). Therefore, proteins are responsible for formation of AgNPs, form a coat which covers the surface of metal nanoparticles (named capping agent of the AgNPs) and effect on its stabilisation by preventing the agglomeration of nanoparticles [[34,](#page-6-0) [43, 45](#page-6-0)].

The TEM study clearly image the shape and size of biosynthesised nanoparticles [[46\]](#page-6-0). The AgNPs synthesised from SH11 actinobacterial strain were spherical and small sized (26 nm). The obtained results are similar to findings of Sadhasivam and coworkers [\[47](#page-6-0)], who reported nanoparticles size range of 20–30  nm synthesised from *Streptomyces* sp. culture. Sukanya *et al.* [[48\]](#page-6-0) and Abdeen *et al.* [[49\]](#page-6-0) also reported spherical and small sized Ag nanoparticles from actinobacterial strains, which were found to be in size range of 5–40 and 10–20 nm, respectively. Moreover, the small amount of aggregations of biosynthesised AgNPs from SH11 strain suggests the inherent stability of nanoparticles in the reaction mixture, which might be due to presence of capping agents formed by proteins or other biomolecules, as mentioned previously [[36\]](#page-6-0).

The zeta potential is one of the main factors affecting the interaction between the particles in the liquid. Zetasizer analysis revealed negative zeta potential of AgNPs from SH11 strain (−16.6 mV). It has been reported that the particles with higher negative or positive zeta potential possess a force to repel each

<span id="page-4-0"></span>

**Fig. 4** *TEM micrograph and selected area diffraction pattern of AgNPs synthesised from SH11 strain*



**Fig. 5** *Zeta potential graph of the AgNPs synthesised from SH11 strain (−16.6 mV) at pH7*

other and do not form aggregates. It has been reported that the particles with higher negative or positive zeta potential possess a force to repel each other and do not form aggregates. The low zeta potential minimize the electrostatic repulsive force and therefore maximize aggregation [\[40](#page-6-0), [50](#page-6-0)]. The AgNPs biosynthesised in this study were found to be more stable than AgNPs synthesised by the marine isolate of *S. albidoflavus* studied by Prakasham *et al.* [[51\]](#page-6-0). The zeta potential of obtained AgNPs were found to be −8.5 mV [[51\]](#page-6-0).

Among the different nanosized antibacterial agents, AgNPs have proved to be the most effective because of its broad-spectrum activity against bacteria, viruses and eukaryotic microorganisms [[35,](#page-6-0) [52, 53](#page-6-0)]. The antibacterial efficacy of the nanoparticles depends on the shape and size of AgNPs [\[54](#page-6-0), [55\]](#page-6-0). Moreover, the nanoparticles smaller than 10 nm interact with bacteria and produce electronic effects, which enhance the reactivity of nanoparticles. Thus, it is proven that the bactericidal effect of AgNPs is size dependent [[56\]](#page-6-0).

The highest antibacterial activity of AgNPs was noticed against *E. coli* followed by *B. subtilis* and *S. aureus.* These findings were compared with studies of Selvakumar *et al.* [\[57](#page-6-0)]. Authors also

**Table 1** MIC of antibiotics and AgNPs synthesised from actinobacterial strain SH11 and synergistic effect of AgNPs combined with antibiotics against bacterial isolates by dilution plate method. Bacterial growth inhibition (%)

with antipology against bacterial isolates by allation plate method. Dacterial growth immotion (70)											
Bacteria	MIC of AgNPs, $\mu q/ml$	MIC of antibiotics, µg/ml			AaNPsa	Kanamycin <sup>a</sup>		Ampicillina		Tetracycline <sup>a</sup>	
		Kan	Amp	Tet		A	$A + AqNPs$	А	$A + AaNPs$	A	$A + AqNPs$
E. coli ATCC8739	$70 \pm 0.06$		$1.5 \pm 0.5$ 2.0 $\pm 0.0$	0.79 ±0.35	$71.9 \pm 0.01$	39.62 ±0.01	72.65 ±0.03	73.94 ±0.06	72.31 ± 0.02 23.13± 0.02 74.3 ± 0.03		
S. aureus ATCC6338	$40 \pm 0.05$	1 1 6 ±0.28			$0.1 \pm 0.03$ 0.25 $\pm 0.0$ 15.8 $\pm 0.05$	39.54 ±0.01	$86.55 \pm 0.02$	47.4 ±0.04	86.05 ±0.005	$44.2 \pm 0.03$	87.04 ±0.004
<b>B.</b> subtilis <b>PCM2021</b>	$40 \pm 0.15$	29.3 ±0.17	$0.1 + 0.01$	0.15 ±0.08	$39.4 \pm 0.01$	85.9 ±0.01	$86.13 \pm 0.01$	34.62 ±0.22			51.9 ±0.02 17.14±0.04 51.4 ±0.004

Mean value are significantly different at  $p \le 0.05$ ; values are expressed in Mean  $\pm$  SD.

MIC, minimal inhibitory concentration; Kan, kanamycin; Amp, ampicillin; Tet, tetracycline; A, antibiotic; AgNPs, silver nanoparticles.

aGrowth inhibition (%) in the presence of MICs of AgNPs or/and antibiotic.

<span id="page-5-0"></span>

**Fig. 6** *Live/dead analysis of effective activity of AgNPs on bacterial pathogens*

Control – *(a) S. aureus*, *(b) B. subtilis*, *(c) E. coli*; Experimental – *(d) S. aureus*, *(e) B. subtilis*, *(f) E. coli*

examined the antimicrobial activity of AgNPs synthesised from actinobacteria against either gram-negative (*Pseudomonas aeruginosa*, *E. coli*, *K. pneumoniae*, *Enterobacter faecalis*) or gram-positive (*S. aureus*) human pathogenic bacteria. However, they recorded the maximum antimicrobial activity of synthesised AgNPs against *S. aureus* followed by *P. aeruginosa*, *E. faecalis* and *E. coli*, and the least against *Klebsiella pneumoniae*. Similarly Prakasham *et al.* [\[51](#page-6-0)] studied the antimicrobial activity of AgNPs from *S. albidoflavus* against gram-positive (*B. subtilis* and *Micrococcus luteus*) and gram-negative (*E. coli* and *K. pneumoniae*) bacterial strains. The maximum antimicrobial activity of AgNPs was found against *K. pneumoniae*, followed by *M. luteus*, *B. subtilis* and the least *E. coli*.

Biogenic AgNPs are intensively tested for their antimicrobial activity against a broad range of microorganisms including grampositive and gram-negative bacteria, fungi, yeasts and microbial biofilms [10]. It is believed that DNA loses its replication ability and cellular proteins become inactivated upon silver ion treatment [4].

The results of minimal inhibitory and bactericidal concentrations confirmed that AgNPs biosynthesised from actinobacteria SH11 strain were efficient against tested bacteria. However, both gram-positive bacterial pathogens were found to be more sensitive than gram-negative strain of *E. coli*. Similarly, live/ dead analysis displayed effective activity of AgNPs on bacterial pathogens.

Singh *et al.* [[58\]](#page-6-0) studied MIC of silver NPs synthesised from *Acinetobacter calcoaceticus* and reported activity of AgNPs against human pathogenic bacteria in much higher concentration range of 150–600 μg/ml when compared with our findings.

However, Manivasagan *et al.* [6] assessed the MIC of AgNPs synthesised from *Nocardiopsis* sp. MBRC-1 against various pathogenic bacteria and fungi. The authors determined lowest MIC against *B. subtilis* at 7 µg/ml, *B. subtilis* at 10 μg/ml and *Candida albicans* at 10 µg/ml, suggesting the broad-spectrum nature of their MIC. It seems that, the higher sensitivity of pathogens could be affected by chemical nature of biosynthesised nanoparticles or nature of microbial cells.

The combination of antibiotics with biosynthesised AgNPs against clinical pathogens offers valuable contribution to nanomedicine [10]. Enhanced effect of bacterial AgNPs on antimicrobial activity of antibiotics has been also observed in studies of Manikprabhu and Lingappa [[59\]](#page-6-0). It is claimed that synergistic effect against bacteria depends on the interaction of AgNPs with a chemical structure of antibiotic [10, [60](#page-6-0)]. Chauhan *et*

*IET Nanobiotechnol.*, 2017, Vol. 11 Iss. 3, pp. 336-342 © The Institution of Engineering and Technology 2016 *al.* [\[61](#page-6-0)] suggested that the increase in synergistic effect may be caused by the bonding reaction between antibiotic and nanomaterials. The antibiotic molecules contain many active groups such as hydroxyl and amide groups, which react easily with nanosilver by chelation.

Combination of AgNPs with antibiotics reduces the toxicity of both toward human cells by decreasing the required dosage with enhanced microbicidal properties. Furthermore, such combination restores the ability of the drug to kill bacteria that have acquired resistance to them [10, [58, 62](#page-6-0)].

Among tested antibiotics, tetracycline showed best synergistic activity against gram-positive and gram-negative bacteria when combined with biosynthesised AgNPs. The effective enhanced activity of ampicillin was noticed against *S. aureus* and *B. subtilis*, and kanamycin against *S. aureus*. The synergetic effect of AgNPs biosynthesised from *Streptomyces*-MS26 in combination with ampicillin, azithromycin, cefotaxime, erythromycin and ofloxacin against bacterial pathogens was reported by Zarina and Nanda [4]. The maximum synergistic effect was observed for amoxicillin, cefixime, amoxicillin and erythromycin against *M. luteus*, *Pseudomonas* sp., *S. aureus* and multi-drug resistant pathogen, respectively

In this study, the actinobacteria SH11 strain isolated from acid soil has shown potential for efficient synthesis of AgNPs by ecofriendly and simple process. The spherical, small sized and stable AgNPs showed significant antibacterial activity. The efficacy of nanosilver increased when used in combination with commercially used antibiotics. Therefore, such AgNPs can be used as antimicrobial agent alone or in combination with antibiotics after more trials in experimental animals.

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