

# Biocomposite Materials Derived from *Andropogon halepensis* – Eco-design and Biophysical Evaluation

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**Table S1.** FTIR ATR band assignment for vegetal extract and phyto-derived metallic nanoparticles

Sample	FT-IR Bands (cm <sup>-1</sup> )	Assignment	Ref.
EAh	3277 (sharp, narrow, strong)	O–H bending and stretching vibrations of hydroxyl groups belonging to phenolic compounds and polysaccharides; Stretching vibrations of the primary and secondary amines	[1] [2]
	2920 (weak)	C–H anti-symmetric stretching vibration	[2]
	2845 (weak)	C–H symmetrical stretch vibration of alkyl chains	[2]
	1645 (shoulder), 1585 (strong)	Amide I, due to carbonyl stretch in proteins; Carboxylate groups (–COO <sup>−</sup> ); Stretching C=C (aromatic ring); Secondary amine, N–H bend	[1] [2]
	1517 (weak, sharp)	Amide II due to N–H bending and C–N stretching in proteins	[3] [4]
	1369 (medium, sharp)	Carboxylates; Phenol or tertiary alcohol, O–H bend; Primary or secondary, O–H in-plane bend; C–H bend	[2]
	1261 (weak)	Primary or secondary O–H in-plan bend; Aromatic ethers, aryl–O stretch; Aromatic primary amine, C–N stretch	[2]

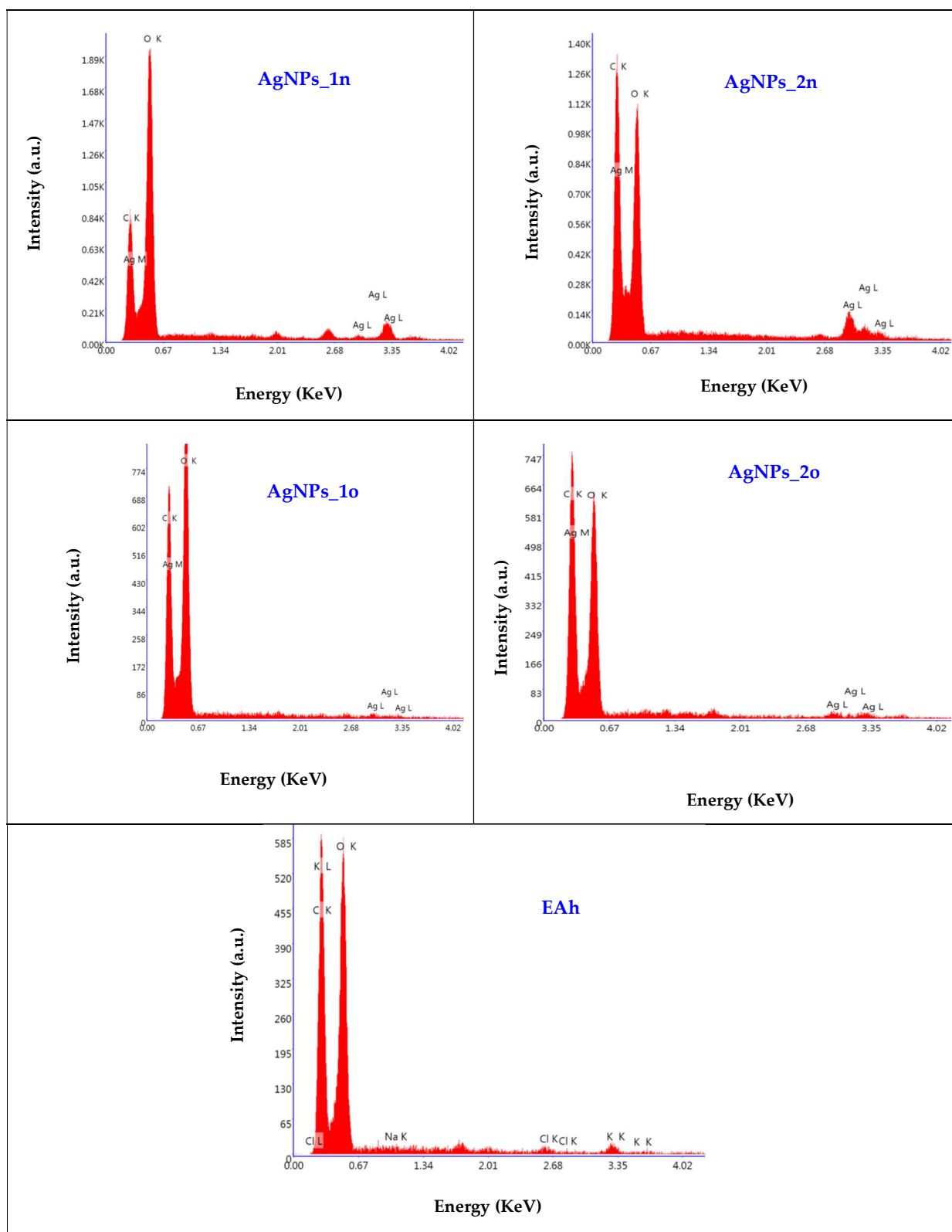
	1030 (strong, sharp)	Cyclic ethers, large rings, C–O stretch; –C–O–C– ether Antisymmetric stretching of –C–O group of polysaccharides and/or chlorophyll; C–O bending in esters; Secondary alcohol, C–O stretch; Primary amine, C–N stretch	[1] [2]
AgNPs_1n/AgNPs_1o	3286/3270 (sharp, strong)	O–H stretching alcohols, polysaccharides and phenolic compounds/ polyphenols; Stretching vibrations of the primary and secondary amines	[1] [2]
	2925/2930 (weak)	C–H anti-symmetric stretching vibration	[2]
	2851/2880 (shoulder)	C–H symmetrical stretch vibration of alkyl chains	[2]
	1591/1585 (strong, sharp)	Amide I, due to carbonyl stretch in proteins; Carboxylate groups (–COO <sup>−</sup> ); Stretching C=C (aromatic ring); Primary amine, N–H bend	[1] [2]
	1512/1517 (weak, sharp)	Amide II, due to N–H bending and C–N stretching in proteins	[3] [4]
	1398/1394 (medium, sharp)	Carboxylates; Phenol or tertiary alcohol, O–H bend; Primary or secondary, O–H in-plane bend; C–H bend	[2]
	1261/1265 (shoulder)	Aromatic primary amine, C–N stretch; Primary or secondary O–H in-plane bend; Aromatic ethers, aryl–O stretch	[2]
	1030/1035 (strong, sharp)	Cyclic ethers, large rings, C–O stretch; –C–O–C– ether Antisymmetric stretching of –C–O group of polysaccharides and/or chlorophyll; C–O bending in esters; Secondary alcohol, C–O stretch; Primary amine, C–N stretch	[1] [2]
AgNPs_2n/AgNPs_2o	3319/ 3348 (very broad)	Very broad intense band overlapping the frequencies of the following groups: Hy- drogen-bonded O–H (the bending and stretching vibrations of hydroxyl groups in alcohols, polysaccharides and phenolic compounds)	[1] [2]
	2930/2920 (very weak, sharp)	C–H anti-symmetric stretching vibration; Transition metal carbonyls	[2]
	2861/2851 (very weak, sharp)	C–H symmetrical stretch vibration of alkyl chains	[2]
	1605/1605 (broad, very weak)	Amide I, arising due to carbonyl stretch in proteins; Carboxylate groups (–COO <sup>−</sup> ); Stretching C=C (aromatic ring); Primary amine, N–H bend	[1] [2]
	1354/1354 (medium, broad)	Carboxylates; Phenol or tertiary alcohol, O–H bend; Primary or secondary, O–H in-plan bend; C–H bend	[2]
	1074, 1040/1079, 1044 (very weak)	Primary amine, C–N stretch; Alkyl-substituted ether; –C–O–C stretching vibration; Primary alcohol, C–O stretch	[2] [5]

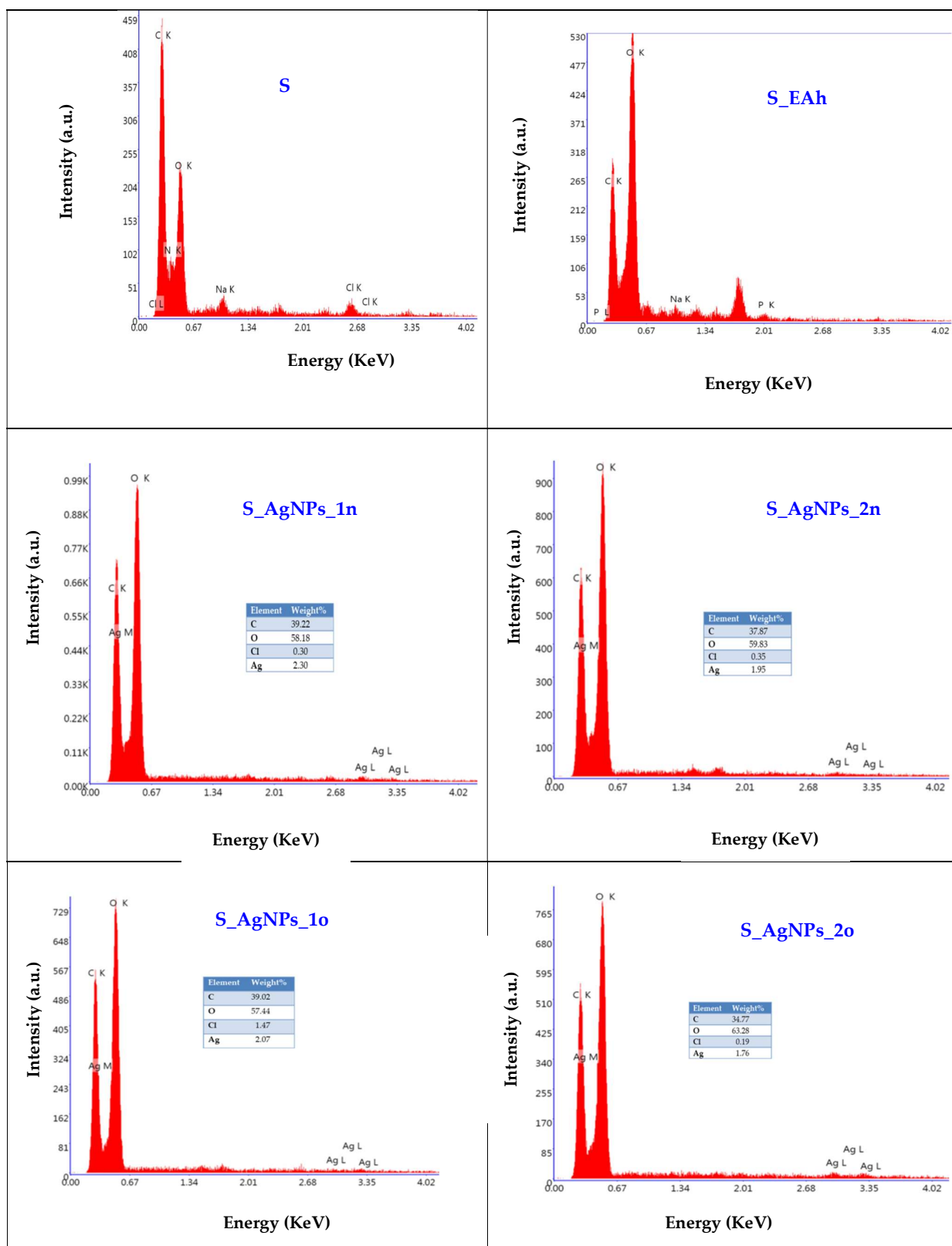
**Table S2.** FTIR ATR band assignment for untreated and treated spider silk

Sample	FT-IR Bands (cm <sup>-1</sup> )	Assignment	Ref.
S/ S_EAh	3332, 3286 (strong narrow band, splitted in 2 bands)/3281(broad)	Normal “polymeric” OH stretch; bending and stretching vibrations of hydroxyl groups intermolecularly hydrogen bonded, in alcohols, polysaccharides and phenolic compounds/ polyphenols; Stretching vibrations of the primary and secondary amines	[1] [2]
	2910 (broad, medium)/2918 (medium, sharp)	C–H anti-symmetric stretching vibration	[2]
	2854 (shoulder)/ 2852 (weak, sharp)	C–H symmetrical stretch vibration of alkyl chains	[2]
	1711 (very weak)/ 1714 (strong, sharp)	Carboxylates, C=O stretching	[6]
	1643 (medium, broad)/ 1655,1601 (medium)	Amide I, arising from carbonyl (–C=O) stretch in spider silk proteins	[3]
	1515, 1540/1517, 1503 (very weak)	N–H bending belonged to the Amide II regions of spider silk proteins	[3]
	1428 (weak, sharp)/ 1450 (weak, broad)	C–H bending	[2]
	1409, 1372, 1338, 1315 (weak, broad)/ 1409, 1373, 1341 (weak, sharp)	Phenol or tertiary alcohol, O–H bend ; C–H bend	[2]
	1278-1239 (very broad, weak)/1260, 1242 (medium)	Amide III band in proteins	[7]
	1161 (medium, sharp)/ 1155 (shoulder)	C–N stretch	[2]
	1106-1031 (sharp)/ 1077 (shoulder), 1019 (strong)	C–H and O–H bond vibrations because of the prevalence of hydroxyl amino acid side chains.	[3]
	898 (weak)/ 897-795 (weak bands), 722 (medium, sharp)	Hydrogen bonded O–H out-of-plan bending; Aromatic C–H out-of-plan bending	[2]
	665 (weak)/ 663 (weak, sharp)	C–H bend; Alcohol, O–H out-of-plan bend	[2]
S_AgNPs_1n/ S_AgNPs_1o	3332, 3281/3330, 3285 (strong narrow band, splitted in 2 bands)/	Hydrogen-bonded O–H (the bending and stretching vibrations of hydroxyl groups in alcohols, polysaccharides and phenolic compounds); Stretching vibrations of the primary and secondary amines	[1] [2] [6]
	2918/ 2915 (medium, sharp)	C–H anti-symmetric stretching vibration	[2]
	2852 (shoulder)/ 2850 (weak,	C–H symmetrical stretch vibration of alkyl chains	[2]

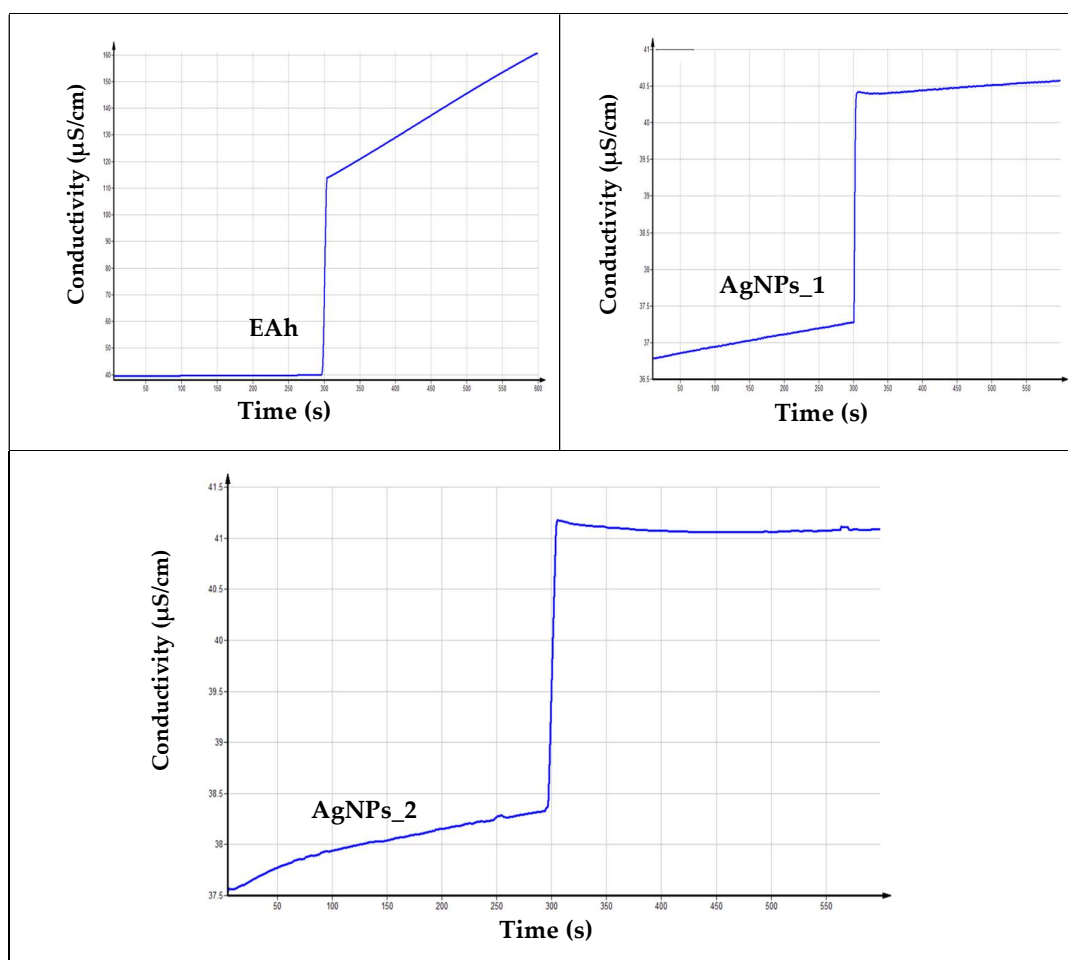
	sharp)		
	1712 (medium, sharp)/ 1709 (shoulder)	Carboxylates, C=O stretching (in silk proteins)	[6]
	1640 (medium)/ 1634 (strong)	Amide I, due to carbonyl (C=O) stretch in silk proteins and in proteins which cover AgNPs	[3] [8]
	1519 (weak, broad)/ 1517 (medium)	Amide II arising from N-H bending and C-N stretching in proteins	[1] [3] [4]
	1450, 1429/1452, 1431 (weak bands)	C-H bending	[2]
	1407-1315/ 1409-1317 (weak)	Phenol or tertiary alcohol, O-H bend ; C-H bend	[2]
	1242/ 1242 (medium, broad)	Amide III band in proteins	[7]
	1161/1158 (medium, sharp)	C-N stretch	[2]
	1102-1018/ 1158-1028 (medium bands)	C-H and O-H bond vibrations because of the prevalence of hydroxyl amino acid side chains.	[3]
	895-793/ 897-845 (weak bands)	Hydrogen bonded O-H out-of-plan bending; Aromatic C-H out-of-plane bending	[2]
	722, 664/ 702, 662 (weak)	C-H bend; Alcohol, O-H out-of-plan bend; vibration of Ag...O bond between AgNPs and O-H groups of serine of spider silk proteins	[2] [9]
S_AgNPs_2n/ S_AgNPs_2o	3334, 3292/ 3337, 3291 (a splitted band, strong)	Very broad intense band overlapping the frequencies of the following groups: Phenols, O-H stretch; Hydrogen-bonded O-H (the bending and stretching vibrations of hydroxyl groups in alcohols, polysaccharides and phenolic compounds)	[1] [2]
	2918/2917 (medium, sharp)	C-H anti-symmetric stretching vibration	[2]
	2852 (medium, sharp)/ 2851 (very weak)	C-H symmetrical stretch vibration of alkyl chains	[2]
	1714 (strong, sharp)/1715 (very weak)	Carboxylates, C=O stretching (in silk proteins)	[6]
	1637 (broad, medium)/ 1635 (strong)	Amide I, arising from carbonyl (C=O) stretch in silk proteins and in proteins which cover AgNPs	[3] [8]
	1517 (medium)/ 1518 (weak)	Amide II band due to N-H bending and C-N stretching in proteins	[1] [3] [4]

	1472 (shoulder), 1454 and 1427 (very weak)/ 1450 and 1432 (very weak)	C–H bending	[2]
	1409 (weak, sharp), 1370 (weak), 1338 (medium, sharp) and 1315 (weak, sharp)/ 1409, 1338 and 1316 (weak, sharp)	Phenol or tertiary alcohol, O–H bend ; C–H bend	[2]
	1245 (strong, sharp)/ 1245 (weak, broad)	Amide III band in proteins	[7]
	1160 / 1162 (medium, sharp)	C–N stretch	[2]
	1104-1018/ 1105-1029 (strong bands)	C–H and O–H bond vibrations because of the prevalence of hydroxyl amino acid side chains.	[3]
	898-793 (weak bands)/ 899 (weak), 873 (shoulder)	Hydrogen bonded O–H out-of-plan bending; Aromatic C–H out-of-plane bending	[2]
	722 (medium, sharp), 663 (weak, sharp)/ 704 (very weak, broad), 665 (weak, sharp)	C–H bend; Alcohol, O–H out-of-plan bend; vibration of Ag...O bond between AgNPs and O–H groups of serine of spider silk proteins	[2] [9]





**Figure S1.** The EDX spectra of all samples: *Andropogon halepensis* extract (EAh) and *Andropogon*-derived AgNPs (AgNPs\_1n, AgNPs\_1o, AgNPs\_2n & AgNPs\_2o), spider silk (S) and spider silk composites with plant extract (S\_EAh) or with silver nanoparticles (S\_AgNPs\_1n, S\_AgNPs\_1o, S\_AgNPs\_2n & S\_AgNPs\_2o), at 10 kV.



**Figure S2.** Investigation of urease inhibitory action of *the Andropogon halepensis* extract and of the developed phytometallic nanoparticles, estimated by the conductometric method.

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