1	Ag nanoparticles and wheat roots: a complex interplay
2	Supporting Information (SI)
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12	1. MATERIALS AND METHODS
13	1.1 Nanoparticle characterization
14	Metallic silver nanoparticles (Ag-NPs) were provided by NanoAmor, Nanostructured and Amorphous
15	Materials Inc. (USA). These Ag-NPs have a coating of polyvinylpyrrolidone (PVP). Ag ₂ S nanoparticles
16	(Ag_2S-NPs) were produced by the sulfidation of the PVP-Ag-NPs as stated in Levard et al. $(2011)^1$.
17	Briefly, a suspension of 1mM PVP-Ag-NPs (pH=7) was mixed with a 1mM Na2S in a 0.01M NaNO3
18	electrolyte. After 3 days of reaction, solutions were centrifuged, washed three times with MQ water
19	and dried. The total sulfidation of PVP-Ag-NPs to Ag_2S -NPs was tested by X-ray Difraction (XRD)
20	Pristine metalllic (Ag-NPs) and sulfided silver nanoparticles (Ag $_2$ S-NPs) were thoroughly
21	characterized in powder and in suspension. Nanoparticles stock suspensions were prepared in
22	double distilled water at 10 mM of Ag concentration and sonicated for 45 min in a ultrasonication
23	bath (Bioblock Scientific 88169). Shape and nominal diameter were determined by Transmission
24	Electron Microscopy (TEM-TECNAI OSIRIS). Specific Surface Area (BET) was determined by the

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25 Volumetric Gas (N₂) Adsorption Method at 77K (BELSORP-max, BelJapan.Inc). Coherent domain size 26 was determined by X-Ray diffraction (XRD). XRD patterns were recorded with a Bruker D5000 powder 27 diffractometer equipped with a SolX Si(Li) solid state detector from Baltic Scientific Instruments using CuK α 1+2 radiation. Intensities were recorded at 0.04°, 2-theta step intervals from 15 to 70° (5 s 28 29 counting time per step). Rietveld refinement of the powder diffraction patterns were carried out using the FullProf package². In suspensions of 100 mgPVP-Ag-NPs·L⁻¹ in ultrapure water, the 30 31 hydrodynamic diameter was measured by Dynamic Light Scattering (DLS), Z potential was also 32 measured over a range of pH from 2 to 10 (Zetasizer Nano ZS Malvern Instruments). In the same 33 suspensions, ionic Ag⁺ concentration was measured with an ion specific silver/sulfide Ion selective 34 electrode (Thermo Scientific) calibrated with a series of AgNO₃ standard solutions.

35 Results of the characterization are presented in figure S1.

a)	nominal diameter (nm)	Especific surface area (m ² ·g ⁻¹)	Coherent domain size (nm)	Hydrodynamic diameter (nm)	Zeta potential (mV)	ionic Ag⁺ (mg·L⁻¹)
PVP-Ag-NPs	52±1	6.3	23±3	61±10	-0.7	<0.01
nano Ag₂S	42±5	4.2	57±5	*	-17	<0.01
b)	Intensity (a.u)) Exp Calc Diffe	eremental ulated erence 70 90 90 (C) (C) (C) (C) (C) (C) (C) (C) (C) (C)			Experimental Calculated Difference
		ig+ were measured in	Nutrient solutio	n	s were determine	u ili tile noagialiu
Figure S1 : Ag-NPS	and Ag ₂ S-NPs	characterization: a)) Summary of th	ne main characteristics	(shape and nom	ninal diameter
were investigated ir	120 particles)	, b) TEM image of t	he PVP-Ag-NPs	, c) Experimental and R	ietveld refined 2	K-ray diffraction
patterns of PVP-AgN	IPs, d) TEM ima	age of the Ag ₂ S-NP	s, e) Experimen	tal and Rietveld refined	I X-ray diffractic	n patterns of

42 Ag₂S-NPs

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44 **1.2. Plant culture**

45 Seeds of wheat (Triticum aestivum L.) were germinated in moisture vermiculite after sterilization on 46 NaClO 15% for 10 min. Once the cotyledon was developed, plants were transferred to a hydroponic 47 system. The hydroponic system consisted on glass pots containing 2L of the corresponding solution. 48 12 plants were placed in each pot by floating in a polystyrene cover. Plants were acclimated 3 days in 49 ¼ strength Hoagland Nutrient solution containing the following constituents: 0.75 mM KNO₃, 0.5 mM 50 Ca(NO₃)₂·4H₂O , 0.25 mM NH₄H₂PO₄, 0.13 mM MgSO₄·7H₂O, 12.5 μM NaCl, 6.25 μM H₃BO₃, 0.5 μM 51 ZnSO₄·7H₂O, 0.5 μM MnSO₄·H₂O, 0.03 μM CuSO₄·5H₂O, 0.13 μM (NH₄)6Mo₇O₂₄·4H₂O, 5 μM Fe(III)-52 EDTA. Then, plants were randomly selected to be treated as follow: i) control, no Ag addition; ii) 30 53 μ M Ag-NP; iii) 30 μ M Ag₂S-NPs and 4) 30 μ M AgNO₃. The system was continuously aerated and 54 agitated in order to keep the nanoparticles in suspension. Solutions were replenish with MQ water 55 daily and totally renewed twice a week.

Plants were harvested after 3 weeks of treatment. Roots and shoots were separated and thoroughly rinsed with MQ water and EDTA 20mM and fresh weights were recorded. Samples for phtytotoxicity studies were immediately frozen in liquid nitrogen and stored at -80°C until analysis. Samples for synchrotron analysis and for X-ray computed-tomography were keep fresh until specific sample preparation.

61 **1.3** Silver distribution and speciation in roots cross sections (synchrotron μ-XRF and μ-XANES)

μ-XANES and XANES spectra were treated by using ATHENA software³, backgrounds were subtracted
and spectra were normalized. Principal components Analysis (PCA) of all the spectra was performed
as stated in Ressler et al. (2000)⁴. The number of components needed to describe the system was set
in 4 based in the statistical IND indicator. Then, the Ag model compounds that constitute the most
probable components of the original XANES spectra were identified by Target Transformations (TT).
Ag⁰, Ag-NPs, Ag₂S, Ag-GSH, Ag- DEDTC and AgNO₃ were the reference compounds better
reconstructed by the TT.

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Once main Ag species were identified, their proportion in the samples was determined by Least-Squares Combination Fitting (LCF) of the experimental spectra. The XANES spectra for Ag⁰ and Ag-NPs were indistinguishable so Ag⁰ was selected to perform the LCF. Similarly, the XANES spectra of Ag-GSH and Ag-DEDTC were also very similar giving the same results in the LCF so they were considered as proxy for Ag-Thiol.

The quality of the fits was quantified by the normalized sum-squares residuals NSS = $\Sigma(\mu \mu \pi)^2 = 2 (\mu \mu \pi)^2 + 2 (\mu \pi)^2 + 2$

78 1.4 Silver concentrations

Immediately after harvest, shoots and roots were separated, fresh weights were recorded and they were frozen in liquid N2 and lyophilized. Then, samples were mineralized in a HNO3 and HF mixture at 120°C for 4h. After acid evaporation, samples were diluted in HNO3 5% until analysis. Ag concentrations were determined by Inductively Coupled Plasma Mass Spectrometry ICP-MS (Elan DRC II Perkin Elmer). NIST 2782 reference material (Industrial Sludge) was used to test the efficiency of the process. The recovery percentages were: 105 ± 2% for Ag.

85 **1.5. Phytotoxicity**

86 **1.5.1 Estimation of lipid peroxidation: malondialdehyde (MDA).**

Lipid peroxidation was evaluated on fresh tissues by the quantification of thiobarbituric acid reactive species (TBARS) by a modified method of Reilly and Aust as stated in Pradas del Real. $(2014)^5$. Briefly, fresh samples (0.1 g) of roots were homogenized on ice with 1 mL of deionised water and centrifuged at 16,000 g for 10 min. The supernatants were removed, and the pellets were re-suspended in 500 µL of solution containing 0.01% butylated hydroxy-toluene (BHT) in 80% ethanol. Then, 900µL of TBA $(2.57 \times 10^{-2}$ M), TCA (9.18 × 10⁻¹M) and HCl (3.20 M) were added to each sample. Samples were 93 vortexed, incubated in a water bath at 70°C for 30 min, cooled on ice and then centrifuged at 16,000
94 g for 10 min. The absorbance of supernatants was measured at 532 nm. Absorbance at 600 nm was
95 subtracted from this measure to eliminate the interference of soluble sugars in the samples.
96 Absorbances were determined by UV-vis light spectrophotometer (Perkin Elmer, Lambda 35).

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98 2. RESULTS



Figure S2: Fitted μ-XRF spectrum extracted from the cell wall and the cytoplasm of the cortex of root

101 exposed to AgNO₃ (root AgNO₃), Ag-NPs (root Ag-NP-1.f), and Ag₂S-NPs (Ag₂S-2)





103 Figure S3: Phytotoxicity: a) silver concentrations in shoots and roots of wheat plants exposed to the

104 different Ag sources (no replicates); b) biomass(g FW·plant⁻¹) (n=3, biological replicates) and c)

105 oxidative stress as Thiobarbituric acid Reactive species (µmol TBARS·g⁻¹ F.W, n=3, biological

106 replicates)



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Figure S4. Roots of wheat plants exposed to different sources of Ag. Ag-NPs-2: a) μ -XRF tricolor map of a frozen hydrated root cross section, b) μ -XRF map of the area indicated in map a, white narrows indicate points where μ XANES were collected. Ag-NPs-3: a) μ -XRF tricolor map of a frozen hydrated 111 root cross section, b) μ -XRF map of the area indicated in map a, c) μ -XRF map of the area indicated in 112 map b, red squares indicate areas where μ -XANES were collected by the μ -mapping mode. g) Ag₂S-113 NP: μ -XRF tricolor map of a cross section a fresh lateral root, narrows indicate points where XANES 114 were collected. Silver in red, Sulfur in green and phosphorous in blue.

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117 Figure S5. Roots of wheat plants exposed to Ag-NPs a) 3D reconstructed image by μ -CT (voxel size 0.9

118 μ m), b) virtual nano-CT slice extracted from a), c) nano-CT reconstructed image of the root surface

119 (voxel size of 63.5 nm). Red arrows indicate zones of Ag accumulation.

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122 Figure S6: 3D reconstructed image of dehydrated roots by μ -CT. a) control, b) Ag-NPs, c) AgNO₃, d)

123 Ag₂S-NPs.

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127 Figure S7: 3D reconstructed image of dehydrated roots by μ-CT (voxel size 0.9 μm). a) control, b) Ag-NPs, c)

- 128 Ag₂S-NPs, c) AgNO₃ . Red arrows show root hairs and/or Ag accumulation zones
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Figure S8: virtual μCT slice (1 μm thickness) from dehydrated roots by μ-CT. a) control, b) Ag-NPs (red narrows

- 133 indicate Ag accumulation in root hairs), c) Ag₂S-NPs (red narrows indicate Nps aggregates), c) AgNO₃.
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136 **Table S1**: List of genes which expression have been studied by qPCR. Encoded genes names, accession number

137 in NCBI database (National Center for Biotechnology: https://www.ncbi.nlm.nih.gov/) and the list of primers

138 used.

Encoded				
	gene	Gene ID	Forward primer	Reverse primer
Gene name	name			
Catalase	CAT	D86327.1	GGACTATGAGGAGCGGTTCG	TTGTTGTGGTGGGAGCACTT
Iron superoxide dismutase	FeSOD	JX398977.1	TGGTTGGGTTTGGCTTGTCT	TCCCCAAAAGCAAGTGGGTT
Glutathione peroxidase 1	GPX	FJ797431.1	CACGACTTCACCGTCAAGGA	TTGGAGTTGGTCAAGCCACA
Metalbthionein-like protein (wali1)	Wali	L11879.1	TGCAACCCCTGCAACTGTTA	ACACACAAGGACACCAAGGG
Phosphoethanolamine N-methyltransferase (PEAMT2)	PEAMT	FJ803924.1	TTGCTGAAGACCG CACTGAT	ATAGTCCTCCTGGCCGAAGT
Pathogen-inducible ethylene-responsive element-binding protein (PIEP1) mRNA	PIEP	EF583940.1	CGAATCTACCG GGATG GTG G	ATGATCACCCGTCATCGTCG
ETTIN-like auxin response factor (ETT1-alpha)	ETT1	AY376128.1	GCGATCGACGTCCAACAATG	AACCAGCTAAATGGCCTCCC
Actin (ACT1)*	Actin	AF326781	AAATCTGGCATCACACTTTCTAC	GTCTCAAACATAATCTGGGTCATC
18S rRNA*	185R	GI223036846	CAAGCCATCG CTCTG GATACATT	CCTGTTATTGCCTCAAACTTCC
Glyceraldehyde-3-phosphate dehydrogenase (GAPDH) *	GAPDH	EF592180	TTTTCACCGACAAGGACA	AAGAGGAGCAAGGCAGTT

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145 **Table S2**: Ag species distribution (%) plant roots obtained by Linear Least squares Combination Fitting of Ag L_{III}

146 edge XANES spectra

		% of Ag				
SAMPLE	Ag-Np	Ag₂S-NP	Ag-Thiol	AgNO ₃	SUM	NSS(%)
AgNO3-Cortex 1			84	16	100	0.12
AgNO3-Cortex 2	14		76	12	102	0.21
Ag2S-NP-Epiderm-main root 1	40	62			102	0.02
Ag2S-NP-Epiderm-main root 2	17	84			101	0.13
Ag2S-NP-Epiderm-leteral root 1		75	26		101	0.04
Ag2S-NP-Epiderm-leteral root 2		89	13		102	0.21
Ag-NP-Epidermis (3-2)	80		24		104	0.52
Ag-NP-Epidermis (3-1)	97				97	0.18
Ag-NP-Endodermis (2-1)			92	1	93	0.29
Ag-NP-Endodermis (2-2)			85	16	101	0.45
Ag-NP-Cortex (1-2)			86	14	100	0.18
Ag-NP-Endodermis (1-1)			77	26	103	0.38

148 Proportions are expressed as mean percentage (%) calculated for these fits. Residual between fit and

149 experimental data: NSS =
$$\sum [k^3 \chi_{exp} - k^3 \chi_{fit}]^2 / \sum [k^3 \chi_{exp}]^2 \times 100$$
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