Supporting Information for

## Rice Exposure to Silver Nanoparticles in a Life Cycle Study: Biphasic Dose-Responses on Grain Metabolomic Profile, Yield, and Soil Bacteria

Xin Yan<sup>1</sup>, Zhengyan Pan<sup>2</sup>, Si Chen<sup>1</sup>, Nibin Shi<sup>1</sup>, Tonghao Bai<sup>1</sup>, Liqiang Dong<sup>2</sup>,

Dongmei Zhou<sup>1</sup>, Jason C. White<sup>3</sup>, Lijuan Zhao<sup>1</sup>\*

<sup>1</sup>State Key Laboratory of Pollution Control and Resource Reuse, School of
Environment, Nanjing University, Nanjing 210023, China

<sup>2</sup>Liaoning Rice Research Institute, Shenyang, 110101, China

<sup>3</sup>The Connecticut Agricultural Experiment Station (CAES), New Haven, Connecticut

06504, United States

‡Xin Yan and Zhengyan Pan contribute equally to this work

\*Corresponding author. Email address: <u>ljzhao@nju.edu.cn</u>

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Table S1. Ag content in rice tissues (mg/kg dry weight)

	Control	1 mg/kg AgNPs	10 mg/kg AgNPs
Polished grain	$1.38 \pm 0.32$	$0.98\pm0.10$	$1.10 \pm 0.14$
Bran	$1.16\pm0.22$	$1.16 \pm 0.23$	$3.86\pm1.83$
Rice hull	$1.50\pm0.46$	$3.12 \pm 1.13*$	$4.70 \pm 1.75^*$
Leaf	$1.55\pm0.50$	$1.78 \pm 0.61$	$1.86\pm0.31$
Stem	$0.67\pm0.19$	$1.26 \pm 0.19*$	$2.98 \pm 1.16$ *
Root	$5.68\pm0.78$	$14.40 \pm 4.31^*$	$29.24 \pm 14.77*$

All data are averages of 4 replicates. \*represent statistical difference at p < 0.05

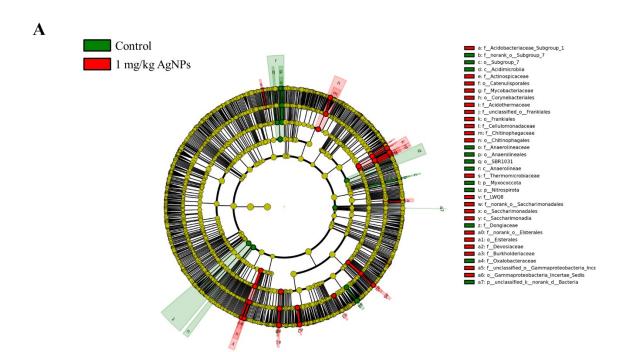
Table S2. Mineral nutrient content in polished rice grain (mg/kg dry weight)

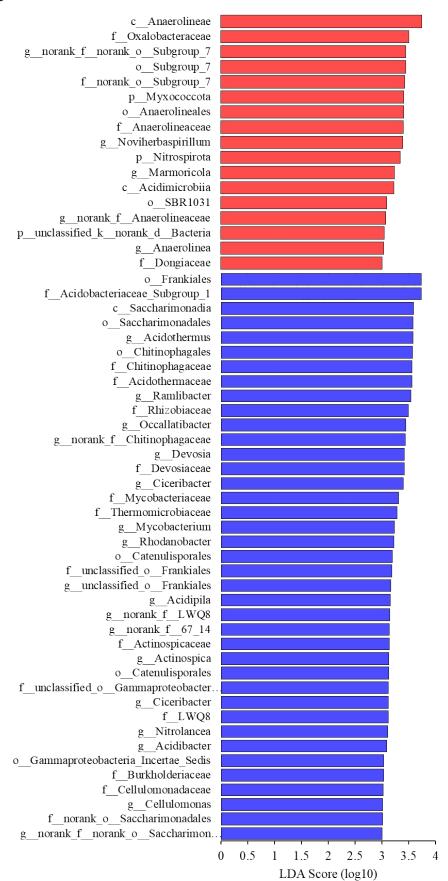
	Control	1 mg/kg AgNPs	10 mg/kg AgNPs
K	$208.5 \pm 4.18$	$169.0 \pm 33.17$	168.1 ± 8.64*
Ca	$219.6 \pm 32.89$	$193.9\pm68.82$	$199.9\pm41.90$
Mg	$262.6 \pm 57.43$	$242.1 \pm 50.76$	$279.0 \pm 9.10$
Fe	$77.04 \pm 11.84$	$58.48 \pm 8.72*$	$64.40 \pm 12.27$
Zn	$15.69 \pm 3.11$	$11.50 \pm 3.11$	$11.86 \pm 2.84$
Mn	$9.49\pm0.94$	$10.72 \pm 3.35$	$9.62 \pm 1.50$
Cu	$7.90 \pm 1.69$	$6.28 \pm 0.75$	$7.23\pm0.14$
Mo	$1.07\pm0.20$	$0.73 \pm 0.11^*$	$0.91\pm0.22$

All data are averages of 4 replicates. \*represent statistical difference at p<0.05

VIP Overlap		T-test
Ethylene glycol Glutaric acid Hydrocinnamic acid L-arabitol Linolenic acid Sorbitol 1-heptanol 1-kestose 3-hydroxymethylglutaric acid D-arabinose Gluconic acid Glutathione Glycocyamine L-asparagine L-tryptophan L-tyrosine Salicylaldehyde Udp-glucuronic acid	1,2,4-benzenetriol 2,4-hexadienedioic acid 3-deoxyhexitol 4-(4-hydroxyphenyl)-4-methyl-2-pentanone Adipic acid Behenic acid Benzylalcohol Cellobiose Citric acid L-cysteine-glycine Metharbital Naproxen Ribonic acid Thymine Beta-hydroxymyristic acid Citraconic acid Citrazinic D-myo-inositol 4-phosphate Itaconic acid L-proline Maleic acid Palatinitol	2-monoolein 3-fluoro-2-hydroxyprop-2-enoate Arachidic acid Dodecanol Glycolic acid Guanosine Isocitric acid Isomaltose Lauric acid Octanol Quinic acid Terephthalic acid 3,4-dihydroxycinnamic acid Chlorogenic acid D-mannose Fumaric acid Phenylalanine Sophorose Uracil

**Figure S1.** Up- and Down- regulated metabolites upon exposure to 1 mg/kg AgNPs. Red and green represent up- and down- regulation of metabolites respectively. The results are the combination of non-variate t-test analysis (p<0.05) and top 40 discriminant metabolites screened by VIP score from PLS-DA model.





**Figure S2.** (**A**) LEfSe analysisat multiple taxonomic levels indicates the taxonomic groups with significant differences in abundance between control and 1 mg/kg AgNPs. Each ring of cladogram represents a taxonomic level, with phylum, class, order, family and genus emanating from inside to the outside. (**B**) LDA score shows taxa enriched in control (red) and 1mg/kg AgNPs (blue) with LDA score>3 and significance of p<0.05 determined by Wilcoxon signed-rank test.