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Environmental Impact

Among the materials to immobilize heavy metals, nano-hydroxyapatite (NHAp) was found to be effective in immobilizing heavy metals due to its moderate solubility and their high surface area and reactivity in soil. There are some new and significant results found in the manuscript. The results showed that NHAp could effectively reduce the CaCl2-extractable Pb, Cu, Cd, Zn and significantly reduce the metal content in ryegrass over time. Treatment with NHAp increased the *Stenotrophomonas sp.* and *Bacteroides* and enzyme activities including urease, phosphatase and dehydrogenase. The results from this study can be very useful for assess the role of NHAp on heavy metals remediation in soil.

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Nano-hydroxyapatite alleviates the detrimental effects of heavy metals on plant growth

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Abstract

The crude recycling activities of e-waste have led to the severe and complex contamination of soil in e-waste workshop topsoil (0-10 cm) by heavy metals. After nano-hydroxyapatite (NHAp) application in June 2013, plant and soil samples were obtained in November 2013, December 2013, March 2014 and June 2014, respectively. 43 The results showed that NHAp effectively reduced the CaCl₂-extractable Pb, Cu, Cd, and Zn in the topsoil, significantly reduced the metal content in ryegrass and also increased the plant biomass compared with the control. Moreover, the concentrations of CaCl2-extractable metals in the soil decreased with the increasing NHAp. NHAp application also increased the activities of soil urease, phosphatase and dehydrogenase. Moreover, soil bacterial diversity and community structure were also altered after NHAp application. Particularly, *Stenotrophomonas sp.* and *Bacteroides* percentages were increased. Our work proves that NHAp application can alleviate the detrimental effects of heavy metals on plants grown in e-waste-contaminated soil and soil enzyme activities, as well as soil microbial diversity.

Keywords

e-waste, heavy metals, plant growth and biomass, soil microbial communities, enzyme activity

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1 Introduction

The rapid development of electrical technology has markedly increased the production of

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electronic waste (e-waste). The majority of e-waste is exported to developing countries, such as China, India, and Pakistan, for recycling and burning, where they are mostly treated by land filling, cyanide leaching and open burning[1]. These crude recycling activities have led to the severe and complex contamination of the soil by heavy metals (Cd, Pb, Cu, and Hg)[2].Milojkovic & Litovski have reported that 70% of heavy metals (including Hg and Cd) found in the soil are of electronic origin [3]. In an e-waste recycling slum in Bangalore, India, the soil concentration of Cd, In, Sn, Sb, Hg, Pb and 68 Bi were up to 39, 4.6, 957, 180, 49, 2850, and 2.7 mg, kg^{-1} respectively, which were approx. 100-fold higher than those at a nearby control site[4]. In 2005, Tang et al. have investigated the soil heavy metal content in soil samples from farmlands nearest an e-waste recycling area in Taizhou and found that the soil heavy metal contents exceeded the standard levels by 100% for Cd, 87.5% for Cd, 37.5% for Hg, and 25% for Zn[5]. It is difficult and costly to remove heavy metals from soil and sediment[6]. As an

alternative, researchers have attempted to stabilize heavy metals in soil or sediment using materials that make these contaminants less mobile and bioavailable, thereby reducing the ecological risk of these metals. Among the materials used to immobilize heavy metals, nano-hydroxyapatite (NHAp) is an efficient heavy metal-immobilizing agent because of its high sorption capacity for heavy metal, low water solubility, high stability under reducing and oxidizing conditions, availability and cost-effect[7]. Many studies have confirmed the efficiency of NHAp in immobilizing Pb and Cd in the contaminated 81 sediment or soil [7, 8]. However, there is limited information on the effects of NHAp on plant growth and soil microbes, especially the long-term effects of NHAp application.

This study was designed to investigate the long-term effects of NHAp on immobilizing

heavy metals, plants growth and biomass, as well as on soil microbes in the e-waste recycling area where the soil was contaminated by e-waste.

2 Materials and methods

2.1 Measurement of soil properties

88 The study area is located in north China (N $39^{\circ}15'$, E117°15'). Many simple household e-waste recycling and burning workshops are distributed across farmlands and riversides in this area, and most of them are currently operational. The bulk density, water content, organic carbon content, cation exchange capacity (CEC) and pH of surface soil were measured before the experiments were started. The bulk density was measured using an soil density instrument (SDG200, TransTech, USA); pH was measured using a glass 94 electrode after the soil was suspended in H₂O (1:2.5 w/v). Water content was measured as follows: 0.1 g of soil was collected and then dried in 105℃ for 6-8 h until a constant weight was obtained, of the dried soil was used as the water content. Organic carbon content was calculated by subtracting the inorganic carbon content from the total carbon content, each being measured using a carbon measurement instrument. CEC was measured as follows: First, 1.00 g of dried soil was weighed and then mixed in EDTA-ammonium acetate solution repeatedly, followed by centrifugation at 3000g/min. The precipitate was retained and was transferred into a 150 ml volumetric flash with deionized water, with a final volume of 80-100ml. Then, 2 ml liquid paraffin and 1 g MgO were added, followed by distilling using an azotometer. Finally, CEC was 104 calculated according to the equation: CEC (cmol/kg \pm)=M \times (V-V0)/soil sample weight. The total concentrations of Cd, Cu, Pb, and Zn in soils were measured as follows: Briefly, 106 6 mL HNO₃ and 3 mL HF were added to each soil sample (1 g) , and the mixture was

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subjected to microwave digestion (120°C for 3 min and 180°C for 15 min). Subsequently, the acids were removed by using an acids-driving instrument (PH60-460, CIF, USA), and the total heavy metal concentration was detected using ICP-MS (inductively coupled plasma mass spectrometry).

2.2NHAp application and sowing of ryegrass seeds

Nano-hydroxyapatite (NHAp) (purity > 98%) was purchased from Nanjing Emperor Nano Materials Co., Ltd (Nanjing, China).The average unit cell size of NHAp used in the present study was 3 nm. Transmission electron microscopy(TEM, Tecnai G2 20 S-TWIN, FEI, USA)revealed that the NHAp material 116 had a nano rod structure, with dimensions of 20 nm (i.d.) \times 200 nm (length). The specific 117 surface area of NHAp was calculated as 130 m^2 g according to its structural geometry.

A random block design was generated for three treatments with five replicates each. The 119 total field area was 560 m², including 9 plots and each plot had an area of 50 m²(10) $m \times 5m$). NHAp was manually spread onto the topsoil in June 2013at 3t ha⁻¹ and $5t \cdot ha^{-1}$ respectively; no NHAp was spread in the control. After spreading, the soil was superficially tilled into interrows using a tiller at a 7-10 cm depth. The ryegrass seeds 123 were directly sown in soil at 1.5 g·m⁻² in June 2013. Afterwards, the soil was never plowed anymore.

Measurement of plant biomass and heavy metal concentration in plants

Ryegrass was harvested four times in November 2013, December 2013, March 2014, and June 2014 respectively, and corresponding biomass was measured separately. Briefly, the aboveground part of all ryegrass plants was cut directly in each plot and placed into paper bags. Then, all the plants were washed first under tap water and then with deionized

water, followed by drying in a 60℃ oven for at least 48 h. The dried plants were then finely ground and subjected to microwave digestion in nitric acid as described by Mackieet al. (2015). Metal content in the plants was analyzed by in ductively coupled plasma mass spectrometry (ICP-MS).

Soil sampling and measurement of CaCl2-extractable heavy metal

Soil was sampled in the same day when plants were harvested. Briefly, 500 g of soil was collected randomly from five locations in each plot at the depth of 10cm beneath, 100 g each. Each sample was collected among ryegrasses not grown on the edge the plot to reduce the edge effect. Then, the five samples were mixed to generate a mixed sample. Among them,20 g was stored in a 4°C refrigerator for later soil enzyme activities analyses; 10 g soil was stored in a -20°C freezer for subsequent microbial community analyses. The rest soil samples treated with NHAp were sieved using a nylon mesh (2 mm in diameter) and homogenized, followed by air-drying and measurement of the following parameters: pH, CEC and heavy metal content.

Heavy metal content was measured as previously described [9, 10]. Briefly, 2.5 g of each 145 soil sample was put in a polypropylene centrifugation tube containing $25 \text{ ml } CaCl₂ (0.01)$ M) and shaken for 2 h at 20°C. Subsequently, the mixture was centrifuged at 3000 g for 15 min and the supernatant was retained. Finally, the concentration of each metal in the supernatant was measured using Agilent 7500a ICP-MS instrument (Agilent, USA). The 149 detection limits for Cd, Zn, Pb and Cu were lower than $1.0 \text{ ng } \text{mL}^{-1}$.

2.4 Measurement of soil enzyme activity

Soil urease, alkaline phosphatase and dehydrogenase activities were determined as described by Guan et al. (1986).

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For the assessment of soil urease activity, 5.0 g of soil was incubated with 1 mL toluene, 10 mL of 10% urea, and 20 mL citrate buffer (pH6.7) at 37°C for 24 h. Afterwards, 40 mL of deionized water, 4 mL of sodium phenolate and 3 mL of sodium hypochlorite were 156 added. The blue products were measured using a spectrophotometer at λ = 578 nm within 1 h after a 30-min color reaction. Assays without soil and urea were examined as controls. 158 The urease activity was expressed as milligrams $NH₃-N$ generated from 1 g of soil at 37°C per 24 h.

Alkaline phosphatase activity was assayed as follows: 5.0 g of soil was incubated with1 mL of toluene and 20 mL of 0.5% disodium phenylphosphate in acetate buffer (pH6.7) at 162 37°C for 24 h. The phenol produced was extracted and oxidized using 0.5 mL potassium hexacyanoferrate in alkaline buffer. The products were determined using 0.5 mL 164 of 4-aminoantipyrine through colorimetry at λ = 510 nm. An assay without soil was examined as a control. The phosphatase activity was expressed as milligrams of hydrolyzedphenol generated from 1 g soil at 37°C per 24 h.

Dehydrogenase activity was determined after incubating 5.0 g of soil with 5 mL of 2,3,5-triphenyltetrazolium chloride (TTC) solution at 30°C for 6 h in the dark. Following incubation, the soil was extracted with 40 mL of methyl alcohol for 1 h to produce 170 tetrazole red formazan (TRF). The filtrate was colorimetrically determined at λ = 485 nm. The dehydrogenase activity was expressed as microliters of hydriongenerated from 5 g of

172 soil at 30°C per 6 h.

2.5 PCR-DGGE analysis of soil bacteria community

After incubation for 360 days, the total microbial DNA was extracted from 0.5g of soil sample by the bead beating method following the manufacturer's instructions using FastDNA SPIN Kit for Soil(Bio101 Inc., USA).The purified DNA extract was fluorometrically quantified using Quantity One 4.0.1,(Fluor-S MultiImager, Bio-Rad, USA). A fluorescent DNA quantification kit (Bio-Rad, CA, USA) was used to generate the standard curve. The quantified DNA extracts were stored at -20°Cfor subsequent analysis.

The V3 region of 16S rDNA was amplified by PCR on Hybrid PCR Express thermal cycler (Bio-Rad, USA) using the primers 357F-GC clamp (5'-CGCCCGCCGCGCCCCGCGCCCGGCCCGCCGCCCCCGCCCCCCTACGGGAG GCAGCAG-3') and 518R (5'-ATTACCGCGGCTGCTGG-3'). PCR was performed in a 50-Μl volume containing 10-15 ng of DNA template, 25 pM of each primer, 2.5 mM deoxynucleotide triphosphates (dNTPs, Promega, USA), PCR buffer (Applied 187 Biosystems, USA), 0.1 mM MgCl₂ solution (Sigma) and 1 U of Taq polymerase (Applied Biosystems, USA). The negative controls contained no template DNA. Parameters for amplification were initial denaturation at 94°C for 3 min, followed by 34 cycles of denaturation at 94°C for 50 s, annealing at 56°C for 1 min and DNA extension at 72°C for 30 s, with a final extension step at 72°C for 10 min. The amplified DNA was verified 192 on 1% agarose gels stained with $SYBR^{TM}$ Green I (Sigma, USA) and visualized using a Fluor-S MultiImager (Bio-Rad, USA).

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194 DGGE (denaturing gradient gel electrophoresis) was performed using a DcodeTM Universal Detection System instrument following the manufacturer's instructions (Bio-Rad, USA). Briefly, a polyacrylamide gel (8% acrylamide/bisacrylamide (37.5:1) in 1X TAE buffer (pH 8.0) was prepared with a denaturing gradient of 35~60% (100% 198 denaturant contained 7 M urea and 40% (v/v) formamide). Thirty microliters of each

PCR product was mixed with loading dye (0.08% bromophenol blue (w/v), 0.08% xylene 200 cyanol (w/v) and 30% glycerol (v/v)), loaded onto the gels and electrophoresed in $1X$ TAE buffer at 60°C for 5 h at a constant voltage of 160 V (DcodeTM Universal Detection System, Bio-Rad, USA). After electrophoresis, the gels were stained with 203 SYBRTM Green I (Sigma, USA) for 30 min and photographed under UV light using a Fluor-S MultiImager (Bio-Rad, USA).

Cluster analysis of the 9 samples based on all DGGE fingerprints was performed using the SAS program (SAS Institute, Cary, NC).The Shannon diversity index of each replicate was calculated following the equation, $H' = \sum_{i=1}^{S} p_i$ lnp_i, where *S* is richness or the total number of band, pi is the proportion of total intensity accounted for by the ith band, and ln is the natural logarithm. The mean value in the three replicates was used as final Shannon diversity index.

2.6Identification of featured bands by sequencing

After DGGE, the bands that varied notably between NHAp-treated soil samples and the control soil samples were excised from the gel. DNA from each band was extracted using the FastDNA SPIN Kit (Bio101 Inc., USA). Then, the extracted DNA was re-amplified with the primer set without a GC clamp. The qualified PCR products were sent to the Beijing Huada Gene Company (Beijing, China) for sequencing. The sequences were aligned using the software MAGE4.0 (Tokyo, Japan).

2.7 Quantitative real-time PCR assays

Quantitative real-time PCR (qPCR) was performed on an iCycler IQ(BioRad, Hercules, 220 CA) using the SYBR Green Jump StartTM Taq Ready MixTM (Sigma, Milan, Italy) following the manufacturer's instructions. Amplification of 16S rRNA genes was

performed the universal primers 341F and 534R [27], respectively. Amplification was performed in a 25 mL total volume containing 12.5 mL of 2X SYBR Green Jump-Start Taq mix, 2.5 mL of each primer (0.05 and 0.9 mM for the primers 341F and 534R respectively), and 7.5 mL of template DNA. To avoid PCR amplification problems due to the presence of inhibitors, the environmental DNA samples were diluted 10 to 100 times.The amplification cycle included an initial denaturation step at 95°C (5 min), followed by 50 cycles of denaturation at 95°C for 30 s, annealing at 55°C for 30 s and 229 elongation at 72° C for 30 s, with a final extension step at 72° C for 7 min. At the end of the qPCR, the melting curve analysis was conducted, measuring the SYBR Green I signal intensities for a 0.5°C temperature increment every 10 s from 50°C to 95°C. The target gene abundance in the microcosms was investigated, and the results were expressed as changes (fold) with respect to the relative zero-time point, according to the expression:

Fold $=2 \frac{(Ct)}{x}$ $2 \frac{(Ct)}{0}$

235 where C_t ⁰ and C_t ^x are the threshold cycles for the zero and successive time-points, respectively. The threshold cycle (Ct) is the cycle number at which the fluorescence generated within are action crosses the threshold. The specificity of the qPCR assays was confirmed based on the occurrence of both single melting peaks and the unique bands of expected sizes on agarose gels.

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2.6 Statistical analysis

All data was expressed as Mean± SD. Differences between different treatments or groups were statistically calculated using ANOVA and Tukey's *t*-testusing SPSS 11.5 (SPSS for Windows, Version 11.5, USA). P<0.05 was considered to indicate a significant difference. Pearson correlation analyses between heavy metal concentrations and

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enzyme activities were also performed using SPSS 11.5.

3 Results

Soil properties prior to the experiments

The bulk density, water content, organic carbon content, cation exchange capacity (CEC) 249 and pH of the soil before any treatment were 1.03 g·cm³, 47.3%, 2.54%, 17.3 cmol·kg⁻¹ 250 and 5.16, respectively. The soil contained 2.65 $g \cdot kg^{-1}$ total N and 0.47 $g \cdot kg^{-1}$ total P. The total concentrations of Cd, Cu, Pb, and Zn in the soil were 3.76, 472.7, 2016.3 and 3076.5 252 mg·kg $^{-1}$, respectively.

Effect of NHAp on CaCl2-extractablemetal concentration in the soil

Before the experiments started, pH was 5.16±0.1, 5.77±0.1 and 6.96±0.2 in the control

255 soil, in the 3 t·ha⁻¹ NHAp-treated soil and 5 t·ha⁻¹ NHAp-treated soil respectively (Table 1). Generally, the pH in each soil was increasing, and that of the soil treated with NHAp was larger than that of the control soil at each time point,suggesting that after NHAp application, the soil pH was affected and the metal immobilization in the soil was enhanced.

With reference to the thresholds in table 2, the concentration of each heavy metal before NHAp treatment exceeded the values of Grade II soil quality standards of the State Environmental Protection Administration (SEPA) of China.

And there was no significant difference in the concentration of each metal among different groups before NHAp treatment (Fig.1). The concentration of each metal 265 decreased markedly over time in the soil treated with t·ha⁻¹ or 5 t·ha⁻¹ NHAp, each being significant lower as compared to that in the control soil in June 2014.

Effect of NHAp on plant growth and metal accumulation in ryegrass

Symptoms of toxicity were observed in the control ryegrass, such as filemotnecrotic spots on the young leaves; by contrast, no symptoms of toxicity were visually observed in the ryegrass grown in the soil treated with NHAp. Meanwhile, the biomass of the control ryegrass harvested at each time point was lower than that of ryegrass grown in soil treated with NHAp, although without significant difference (Table 1).

Overall, ryegrass grown in the control soil had lower metal contents as compared to those grown in the NHAp-treated soil, with significant difference in Cu, Pb and Zn content at each time point (Table 1).

Effects of NHAp on soil enzyme activity

Changes in soil dehydrogenase, urease, and acid phosphatase after the addition of NHAp were determined in the present study (Table 3).

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280 Either dehydrogenase, urease or phosphatase activity in the soil to be treated by $5 \text{ t} \cdot \text{ha}^{-1}$ was the highest prior to NHAp treatment. However, the enzyme activity in the control soil was lower than its initial level, indicating a detrimental effect of heavy metals on soil enzyme activities, whereas the enzyme activity in the soil treated by either NHAp was higher than their initial level, indicating a beneficial role of NHAp treatment. Thus, the 285 dehydrogenase activity in soil treated by $3t \cdot ha^{-1}$ NHAp, urease activity in the soil treated 286 with 5 t·ha⁻¹ NHAp, and phosphatase activity in the soil treated with either 3 or 5 t·ha⁻¹ NHAp was significantly higher than their counterpart in the control soil in June 2014.

Pearson correlation analyses revealed a significant positive correlation between 289 dehydrogenase and phosphatase ($R^2 = 0.758$), indicating a similar sensitivity of the two enzymes to heavy metal contamination in the study area. It was also found that there was

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291 a significant negative correlation between areas activity with either soil Cu (
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R^2 = -0.897
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)
292 or Cd ($R^2 = -0.911$) content (Table 4).

Effect of NHAponsoil microbial diversity

DGGE revealed that the DGGE fingerprints were similar in the three replicates in the soil subject to the same treatment, indicating a relatively higher reproducibility(Fig. 2). 296 Cluster analysis showed that the bacterial communities treated with NHAp $(3t.ha^{-1})$ and 5 $t.ha^{-1}$ were separated from the control group (Fig. 3), indicating similar microbial composition in the soil treated with NHAp.

The Shannon diversity index of microbial commnunities in the control soil was 3.41, 300 while that in the soil treated with t·ha⁻¹ and $5t$ ·ha⁻¹ NHAp was 3.69 and 3.88 , respectively, indicating that soil microbial diversity is increasing with NHAp concentration.

Identification of featured bands in the NHAp-treated soil microbial samples

Sequence analysis showed that most clones in the soil treated by NHAp belonged to *Stenotrophomonas sp* and *Bacteroides*accounting for 40% and 28% respectively (Fig. 3); and the rest bands mostly belonged to *Enterobacter sp.* and *Acidobacteria*.

4 Discussion

Effects of NHAp on CaCl2-extractablemetal concentration in the soil and metal concentration in plants

pH affects the chemical forms of the metals in the soil [11]. CEC is a commonly used indicator of soil fertility, nutrient retention capacity[12]. In the present study, it was found that NHAp didn't alter the change in pH over time, while higher concentration NHAp altered the change in CEC over time. *in situ* immobilization of heavy metals using NHAp is a cost-effective and environmentally sustainable remediation approach by reducing

their mobility and availability. Here, we examined the effect of NHAp on immobilizing 316 heavy metals by using CaCl₂ to extract exchangeable metal heavy metals that were not immobilized in the soil. For eachCaCl₂-extractable metal, its concentration in the soil was significantly lower than that in the NHAp-treated soil, thus it seemed that NHAp 319 especially 5 t·ha⁻¹ NHAp can significantly decreased the content of exchangeable heavy metals, which were available by plants. This was consistent with the finding that lower heavy metal content was detected in plants grown in the NHAp-treated soil, as well as better performance of ryegrass in growth and biomass in the soil treated by NHAp. Previously, Boisson et al. have also reported that hydroxyapatite decreased the concentrations of `toxic' metals in the leaves of the test plants; however, they also found that too higher hydroxyapatite application rate reduced the uptake of some essential trace elements, thus leading to deficiency problems[13].

Effects of NHAp on soil enzyme activity

Dehydrogenase is an intercellular enzyme in the soil, which catalyzes the removal of hydrogen atom from different metabolites[14], urease hydrolyzes urea intracellularly, leading to a shift in soil pH Phosphatase activity. Alkaline phosphatase is involved in soil phosphorus metabolism[15]. Numerous studies have confirmed that the activities of soil enzymes were decreased with the increasing heavy metal pollution [16-18].Dehydrogenase activity was particularly sensitive to heavy metals [19].Kandeler et al. have reported that phosphatase activities were dramatically decreased by heavy metal pollutants [17].Tyler has declared that urease and acid phosphatase activity was closely negatively correlated with log Cu+Zn concentration[20]. Decrease in the activities of three soil enzymes was also observed in the present study. Heavy metals in

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the soils react with active the protein groups of enzymes, such as sulfhydryl groups, to form metal-sulfide equivalents, or sequester the enzyme binding sites through the formation of complexes with the substrates, thereby inactivating or inhibiting enzyme activity[21]. By contrast, the beneficial role of NHAp was also observed here. Previously, Bert et al. (2012) have reported that addition of hydroxyapatite can reduce sediment ecotoxicity and improved the growth of the total bacterial population.

Effect of NHAp on soil microbial diversity

Here, comparison of Shannon diversity indices between the soil subject to different treatments indicates that microbial diversity was decreased in the soil contaminated by heavy metals, while NHAp has a beneficial effect on soil microbial diversity. Previously, Oliveira et al. have also reported that quantitative analysis of soil microbial populations shows a marked decrease in total culturable numbers of the different microbial groups of the soil samples contaminated by Hg and As[19].Du et al. further reported that the microbial diversity index of microbial community in the treatments amended with NHAp was significantly higher than that of control[22]. Thus, it can be concluded that NHAp can improve the microbial diversity in the metal-contaminated soil. Furthermore, sequencing discovered that microbes belonging to *Stenotrophomonas sp* and *Bacteroides* showed an elevated abundance in the NHAp-treated soil as compared to the control soil. Previously, Pages et al. have presented that another *Stenotrophomonas* species *S. maltophilia* can develop tolerance to overcome metal toxicity in the presence of heavy metals[23]. Thus, it is presumably that NHAp may increase the microbial populations that are tolerant to metal toxicity.

In the present study, NHAp significantly decreased the exchangeable heavy metals contents in the e-waste-contaminated soil, also reduced the metal concentration in plants and increased plant biomass, suggesting NHAp has a good performance on immobilizing the heavy metals in produced by e-waste, accordingly alleviating the detrimental effects of heavy metals on plant growth. NHAp has also alleviated the detrimental effects of heavy metals on soil enzyme activities. In addition, NHAp application also has a positive role on soil microbial diversity and microbial composition possibly via increasing the percentage of metal- tolerant populations, such as *Stenotrophomonas sp* and *Bacteroides*.

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501 Table 1

503 according to the results of ANOVA and Tukey test (P<0.05);* indicates significant differences 504 between different treatments at the same sampling time point according to the results of ANOVA and 505 Tukey test(P<0.05). 506

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581 Note: Mean values followed by the same letter are not significantly different between different sampling time in the same treatment plot according to 582 ANOVA and multiple comparisons with Tukey test (p≥0.05). * means the significantly different between different treatment at the same sampling time 583 according to ANOVA and multiple comparisons with Tukey test ($p \ge 0.05$).

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