



OPEN Pterostilbene-loaded PLGA nanoparticles alter phenylpropanoid and oxylipin metabolism in *Solanum lycopersicum* L. leaves

Camilla Badiali^{1,5}, Marzia Beccaccioli^{1,5}, Fabio Sciubba^{1,2}, Laura Chronopoulou^{3,4}, Valerio Petrucci¹, Cleofe Palocci^{3,4}, Massimo Reverberi¹, Alfredo Miccheli^{1,2}, Gabriella Pasqua^{1,2} & Elisa Brasili^{1,2}✉

Due to the fast-changing global climate, conventional agricultural systems have to deal with more unpredictable and harsh environmental conditions leading to compromise food production. The application of phytonanotechnology can ensure safer and more sustainable crop production, allowing the target-specific delivery of bioactive molecules with great and partially explored positive effects for agriculture, such as an increase in crop production and plant pathogen reduction. In this study, the effect of free pterostilbene (PTB) and poly(lactic-co-glycolic) acid (PLGA) nanoparticles (NPs) loaded with pterostilbene was investigated on *Solanum lycopersicum* L. metabolism. An untargeted NMR-based metabolomics approach was used to examine primary and secondary metabolism whereas a targeted HPLC–MS/MS-based approach was used to explore the impact on defense response subjected to anti-oxidant effect of PTB, such as free fatty acids, oxylipins and their impact on hormone biosynthesis, in particular salicylic and jasmonic acid. In tomato leaves after treatment with PTB and PLGA NPs loaded with PTB (NPs + PTB), both NPs + PTB and free PTB treatments increased GABA levels in tomato leaves. In addition, a decrease of quercetin-3-glucoside associated with the increase in caffeic acid was observed suggesting a shift in secondary metabolism towards the biosynthesis of phenylpropanoids and other phenolic compounds. An increase of behenic acid (C22:0) and a remodulation of oxylipin metabolism deriving from the linoleic acid (*i.e.* 9-HpODE, 13-HpODE and 9-oxo-ODE) and linolenic acid (9-HOTrE and 9-oxoOTrE) after treatment with PLGA NPs and PLGA NPs + PTB were also found as a part of mechanisms of plant redox modulation. To the best of our knowledge, this is the first study showing the role of PLGA nanoparticles loaded with pterostilbene in modulating leaf metabolome and physiology in terms of secondary metabolites, fatty acids, oxylipins and hormones. In perspective, PLGA NPs loaded with PTB could be used to reshape the metabolic profile to allow plant to react more quickly to stresses.

In the current scenario of global climate change and rapid population increase, agriculture has been dealing with a wide range of challenges for food security and it demands efficient crop improvement methods to ensure food quality and quantity¹. Current strategies in modern agriculture should consider efficacy, cost affordability, environmental safety, toxicity towards non-target organisms, and sustainability of the production system especially since some pathogen species are naturally resistant to certain types of drugs, and resistance can also develop over time because of the exposure to drugs used in a large amount. A solution could be provided by lowering the amount of the drug, thus delaying the appearance of resistance, as well as seeking novel natural products with antimicrobial activity, and using innovative approaches for delivering^{2,3}. In this regard,

¹Department of Environmental Biology, Sapienza University of Rome, Rome, Italy. ²NMR-Based Metabolomics Laboratory (NMLab), Sapienza University of Rome, Rome, Italy. ³Department of Chemistry, Sapienza University of Rome, Rome, Italy. ⁴Research Center for Applied Sciences to the Safeguard of Environment and Cultural Heritage (CIABC), Sapienza University of Rome, Rome, Italy. ⁵These authors contributed equally: Camilla Badiali and Marzia Beccaccioli. ✉email: elisa.brasili@uniroma1.it

nanotechnologies are recently emerging as revolutionary approaches to renew the resilient agricultural system and transform modern agriculture into precision agriculture².

Nanoparticles (NPs) combined with bioactive compounds could offer a smart solution to improve crop growth and development as well as to control crop diseases limiting the use of conventional pesticides. In recent years, nanoparticles have also emerged as novel triggers for inducing the biosynthesis of bioactive compounds in plants⁴. Ag NP treatment increased artemisinin content by 3.9-fold in 20-day-old hairy root cultures of *Artemisia annua* L.⁵ Hydroponically grown *Bacopa monnieri* L. treated with copper-based NPs (Cu) improved antioxidant capacity and showed a hormetic increase in the content of saponins, alkaloids, flavonoids and phenols⁶. In any case, nanoparticles facilitate site-targeted delivery and controlled release of bioactive compounds, ensuring efficient utilization^{7,8}.

Among nanomaterials, polymeric NPs are composed of natural or synthetic polymeric materials, some of which have desirable features such as biodegradability⁹. Among these, poly(lactic-co-glycolic acid) (PLGA) NPs have been extensively studied, their biocompatibility has been proved⁹, and they have shown great potential for use in the development of nano-based delivery systems for plants^{10–13}. PLGA has been approved by the FDA, WHO, and other regulating agencies. However, further studies be performed to guide the rational application of PLGA nanoparticles in agriculture and ensure sustainability. Resveratrol (3,5,4'-trihydroxystilbene) and its methoxylated derivative, pterostilbene (3,5-dimethoxy-4'-hydroxystilbene) (PTB), are synthesized via the phenylpropanoid pathway as antimicrobial phytoalexin in response to pathogen attacks. Whereas resveratrol has been identified in a wide number of food plants as peanut (*Arachis hypogaea*), grape (*Vitis vinifera*), mulberry (*Morus alba*), pistachio (*Pistachia vera*) and tomato (*S. lycopersicum*), PTB has been found in a few plants and crops such as *Vitis vinifera*¹⁴, *Pterocarpus marsupium*^{15,16}, *Arachis hypogaea*¹⁷ and *Vaccinium* species (berries)¹⁸ but not in *S. lycopersicum*. In this regard, PTB could represent a possible strategy to enhance the metabolic machinery of the plant and prepare the plant to better react to subsequent environmental stress. It has been demonstrated that PTB exhibits a strong antifungal activity against common crop fungal pathogens, such as *Botrytis cinerea*, *Fusarium oxysporum*, *Sclerotinia sclerotiorum*, *Plasmopara viticola*, *Septoria nodorum*, and *Leptosphaeria maculans*^{19,20} but its effect as natural compound on tomato plant metabolism is unknown. Further studies should be carried out to evaluate the impact and toxicity of PLGA NPs loaded with natural compounds on the metabolism of plants.

In the present work, tomato leaves were analyzed after treatment with PLGA NPs or PTB or PLGA NPs loaded with PTB (NPs + PTB) by an untargeted NMR-based metabolomics approach and a targeted HPLC–MS/MS approach to evaluate the free fatty acids, oxylipins, and plant hormones. In fact, free fatty acids have a role against abiotic (metals²¹ and temperature²²) and biotic stress e.g. as signaling molecules^{23–25}. Here, the oxylipins derived from the linoleic and linolenic acid were investigated, playing an important role as signaling molecules during plant defense processes in response to biotic or abiotic stress²⁶. Since free fatty acids and the oxylipins are closely related to response towards stress an additional evaluation was carried out to define the alteration of the main hormones implicated in plant defense, that are the jasmonic acid and salicylic acid²⁷.

The valuable insights that emerge showed metabolites linked both to primary and secondary metabolism suggesting an alteration of redox signaling promoting the plant defense capacity. With this study, we demonstrate the role of PLGA nanoparticles loaded with pterostilbene in modulating leaf metabolome and physiology in terms of secondary metabolites, free fatty acids, oxylipins and hormones. Future studies considering the exposition of plant to abiotic and biotic stresses including pathogens will be useful to determine the efficacy of PLGA NPs loaded with pterostilbene in improving the plant responses to stresses.

Results

Effect of PLGA NPs, PTB, and PLGA NPs loaded with PTB on the metabolome of tomato leaves evaluated by ¹H-NMR

The analysis of the 600 MHz ¹H-NMR spectra obtained from hydroalcoholic and chloroformic extracts of tomato leaves allowed to unequivocally identify 21 molecules. A total of 21 metabolites including organic acids, amino acids, organic compounds, sugars, fatty acids, phenylpropanoids, and other compounds were integrated. Comparing the spectra obtained from tomato leaves treated with water (Ctrl), PLGA NPs (NPs), PTB and PLGA NPs loaded with PTB (NPs + PTB), it was possible to observe qualitative differences. Examples of ¹H-NMR spectra are reported in Supplementary Figures S1–S4, and the table of resonance assignment is shown in Supplementary Table S3. Quantitative analysis of the chemical composition of tomato leaves is reported in Supplementary Table S4. PLS-DA discriminant analyses were applied to highlight the most important metabolites discriminating the treatments with NPs, PTB, and NPs + PTB.

To evaluate the effect of PLGA NPs on the metabolism of tomato plants, the PLS-DA analysis was carried out by comparing plants treated with empty NPs and control plants (Ctrl) (Fig. 1). The PLS-DA score plot highlights the clear separation of the tomato plants treated with NPs (Fig. 1A) that showed higher levels of caffeic acid and lower levels of trigonelline, GABA and quercetin-3-glucoside compared to control plants ($R^2 = 0.999$, $Q^2 = 0.77$), as revealed by regression coefficients (Fig. 1B).

To investigate the effect of PTB treatment on the metabolism of tomato leaves, PLS-DA analysis was applied, and results are reported in Fig. 2. The score plot revealed a clear separation of tomato plants treated with PTB from untreated plants (Ctrl) ($R^2 = 0.998$, $Q^2 = 0.93$) (Fig. 2A). The levels of caffeic acid, valine, GABA, glucose and sucrose increased while alanine, arginine and choline levels decreased in plants treated with PTB compared to Ctrl plants (Fig. 2B).

In order to disentangle the metabolic changes due to the treatment with NPs loaded with PTB from the treatment with empty NPs, a PLS-DA was applied and a clear separation between the treatment groups was observed ($R^2 = 0.94$, $Q^2 = 0.55$) (Fig. 3A). The levels of GABA significantly increased in leaves treated with

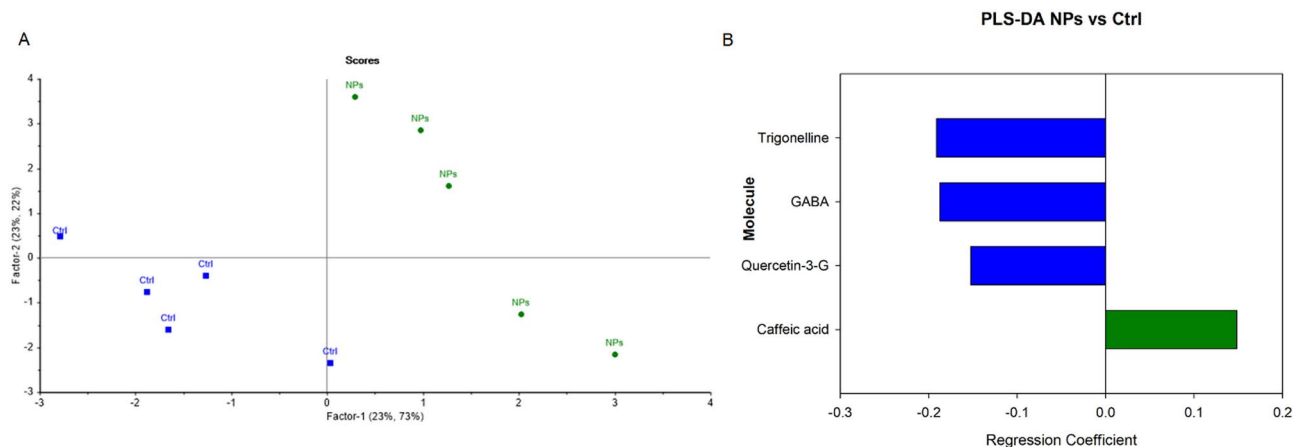


Fig. 1. **A** PLS-DA score plot of ^1H -NMR metabolomics data of hydroalcoholic extracts obtained from tomato leaves treated with NPs or water (Ctrl). **B** PLS-DA regression coefficients of significantly different metabolites in tomato leaves treated with NPs or water (Ctrl).

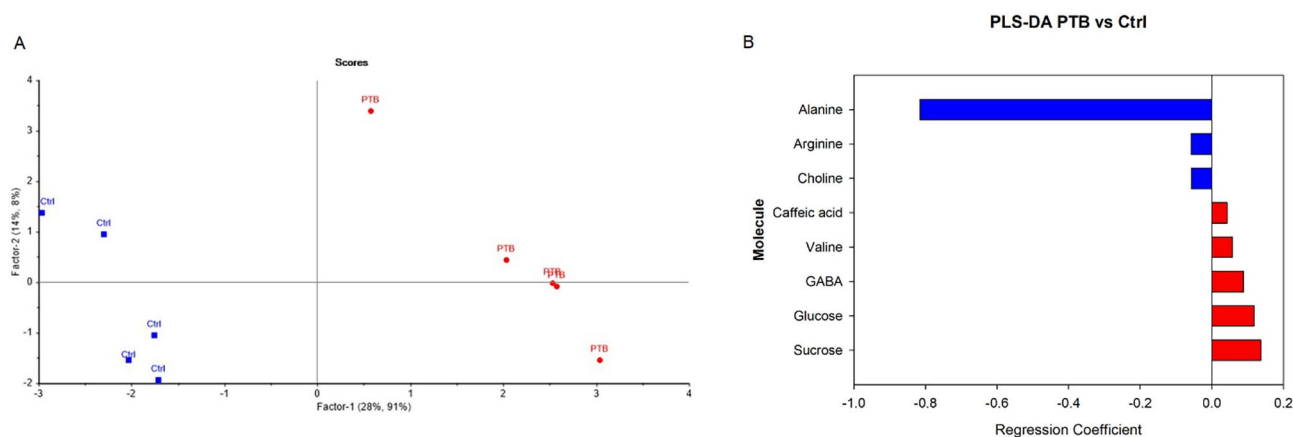


Fig. 2. **A** PLS-DA score plot of ^1H -NMR metabolomics data of hydroalcoholic extracts obtained from tomato leaves treated with PTB or water (Ctrl). **B** PLS-DA regression coefficients of significantly different metabolites in tomato leaves treated with PTB or water (Ctrl).

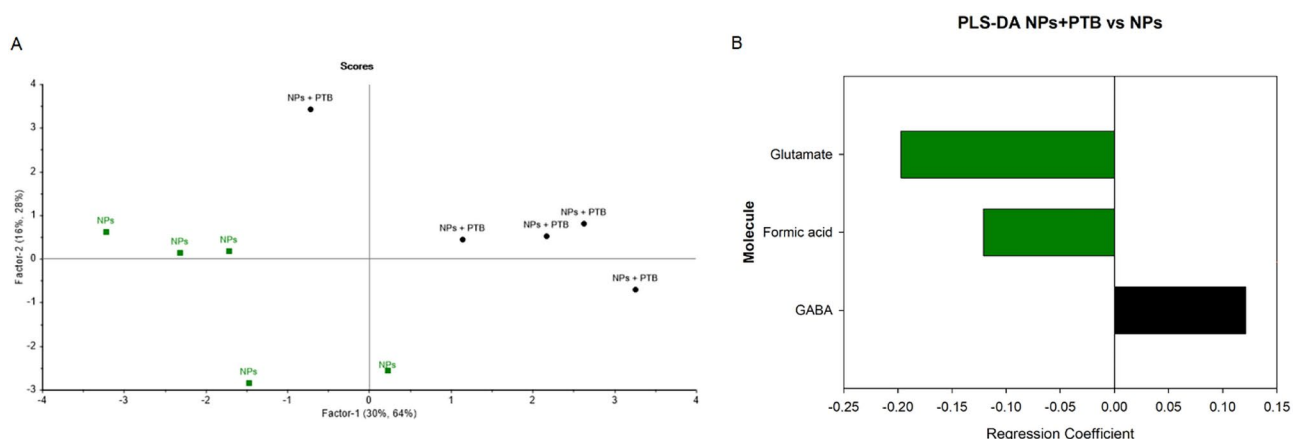


Fig. 3. **A** PLS-DA score plot of ^1H -NMR metabolomics data of hydroalcoholic extracts obtained from tomato leaves treated with NPs or NPs + PTB. **B** PLS-DA regression coefficients of significantly different metabolites in tomato leaves treated with NPs or NPs + PTB.

NPs + PTB compared to the NPs group. Conversely, the levels of glutamate and formate significantly decreased in NPs + PTB group compared to the NPs group (Fig. 3B).

Finally, a PLS-DA was carried out to discriminate the effect of PTB from NPs loaded with PTB and results are reported in Fig. 4. The PLS-DA model represented by three factors and the following indices $R^2=0.99$, $Q^2=0.80$ (Fig. 4A), showed metabolic differences between the two treatments consisting in higher levels of choline, threonine, arginine and valine, and lower levels of sucrose and glucose in NPs + PTB group compared to the PTB group (Fig. 4B).

Effect of PLGA NPs, PTB, and NPs loaded with PTB on tomato leaf: phyto-oxylipin, fatty acid and hormone levels by HPLC-MS/MS

Exists a deeply link between the redox state, membrane fluidity and defense responses of plant cells. Free fatty acids assume a double role, they are constituents of cell wall and cell membranes, but during the pathogen infection they assume a role of signaling molecules. To better understand the effect of the treatments we evaluated palmitic (C16:0), palmitoleic (C16:1), stearic (C18:0), oleic (C18:1), linoleic (C18:2), linolenic (C18:3), behenic (C22:0), lignoceric (C24:0) and nervonic (C:24:1) acids, and we noted that the treatments proposed not alter the composition of free fatty acid except for C22:0, that increase in tomato leaves treated with NPs, PTB and NPs + PTB compared to the Ctrl group (Fig. 5).

Oxylipins and hormones (salicylic acid and jasmonic acid) were also evaluated as markers linked to plant defense. Mass spectrometry analysis reveals a significant change in 9-HpODE, 13-HpODE and 9-oxo-ODE (18:2-derived), 9-HOTrE, 9-oxoOTrE (18:3-derived) and salicylic acid levels (Fig. 6). In particular, 9-HpODE and 13-HpODE significantly decreased in tomato leaves treated with PTB or NPs + PTB compared to the Ctrl group and NP treatment; 9-oxo-ODE, 9-HOTrE and 9-oxoOTrE significantly increased in plants treated with NPs compared to the Ctrl group and significantly decreased in NPs + PTB compared to NPs group. Salicylic acid levels significantly increased in plants treated with NPs compared to the Ctrl group, while in PTB and NPs + PTB decrease probably for the antioxidant effects.

Determination of total carotenoids and chlorophyll (a and b) content in tomato leaves treated with NPs, PTB, and NPs + PTB

The content of chlorophylls and carotenoids was analyzed by spectrophotometric analysis and no significant differences were observed among the treatment groups with the respect of untreated controls (Figure S5 and S6).

Discussion

In this study, an omics approach was applied to investigate the metabolic changes in tomato leaves after treatment with PLGA NPs or PTB, a natural compound, or PLGA NPs loaded with PTB. The ^1H -NMR-based metabolomic analysis allowed to highlight the metabolic effects due to PLGA NPs or PTB treatment used in its free form or loaded in NPs. A common biochemical pathway involving GABA metabolism was affected after treatment with PLGA NPs loaded with PTB or free PTB. In particular, an increase in GABA levels was observed after treatment with PLGA NPs loaded with PTB or free PTB. In addition, a decrease in glutamate levels, GABA precursor, was observed after treatment with PLGA NPs loaded with PTB. GABA is a ubiquitous four-carbon non proteinogenic amino acid that is greatly metabolized via the GABA shunt, a short pathway linked with several pathways such as the TCA cycle^{28,29} and glutamic acid decarboxylation. In plants, GABA plays important roles in carbon/nitrogen balance maintenance, regulation of cytosolic pH, protection against oxidative stress, energy production, cell elongation and leaf development regulation as well as germination, fruit ripening and senescence^{30,31}. Similarly, to other plant molecules, such as calcium, jasmonic acid, salicylic or abscisic acid, GABA levels readily increase in response to environmental biotic and abiotic stresses and its concentrations range from micromolar (low) to millimolar (high) in different tissues, organs and cell compartments^{30,32,33}.

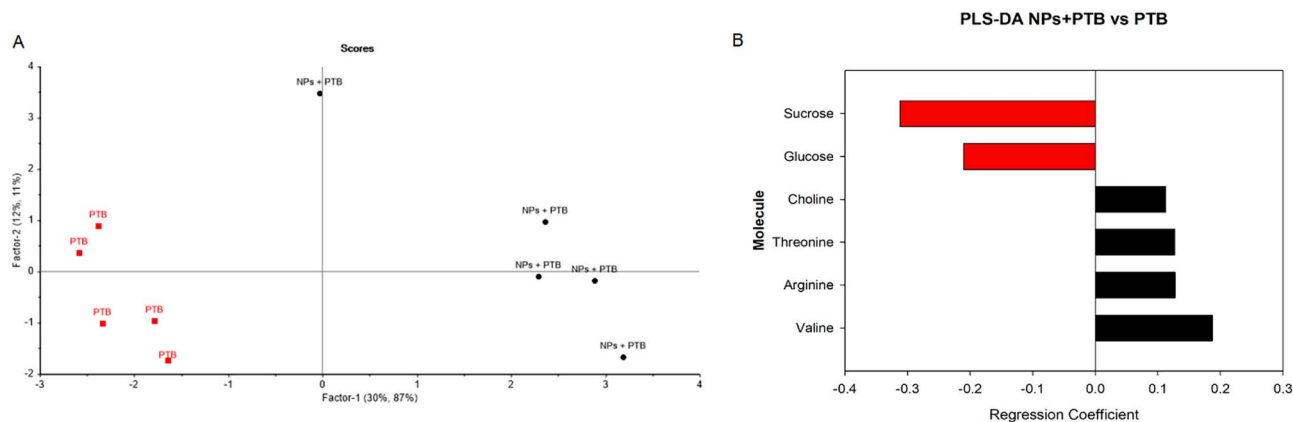


Fig. 4. **A** PLS-DA score plot of ^1H -NMR metabolomics data of hydroalcoholic extracts obtained from tomato leaves treated with PTB and NPs + PTB. **B** PLS-DA regression coefficients of significantly different metabolites in tomato leaves treated with PTB and NPs + PTB.

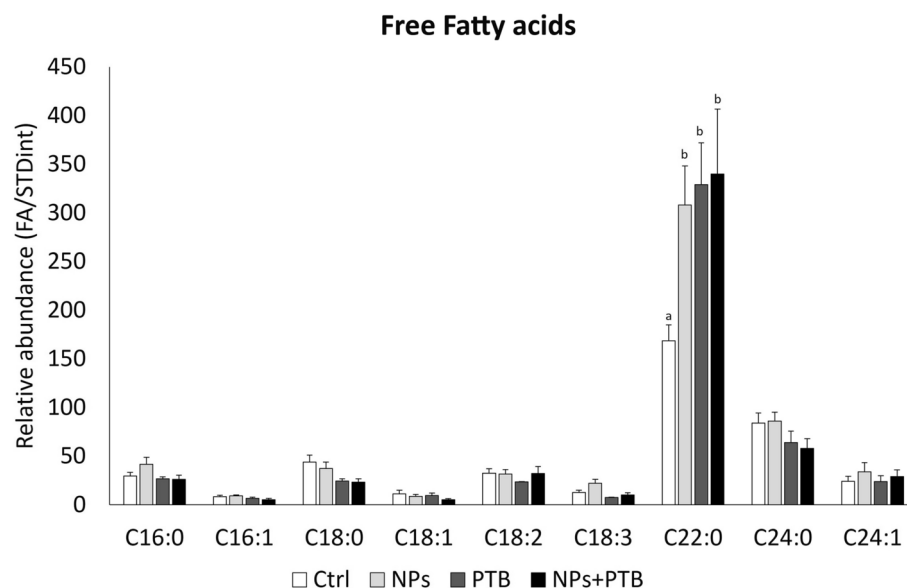


Fig. 5. Free Fatty acid relative abundances in tomato plants treated with NPs, PTB, and NPs + PTB. Profile of all fatty acids analyzed. Ctrl: control, plants treated with distilled water; NPs: PLGA NPs; PTB: pterostilbene; NPs + PTB: PLGA NPs loaded with PTB. The data were expressed as the mean of three replicates and bars represent the standard error. Different letters represent significant differences Mann–Whitney *U* test was also run to determine significant differences for considered metabolites. A *p*-value < 0.05 was considered statistically significant.

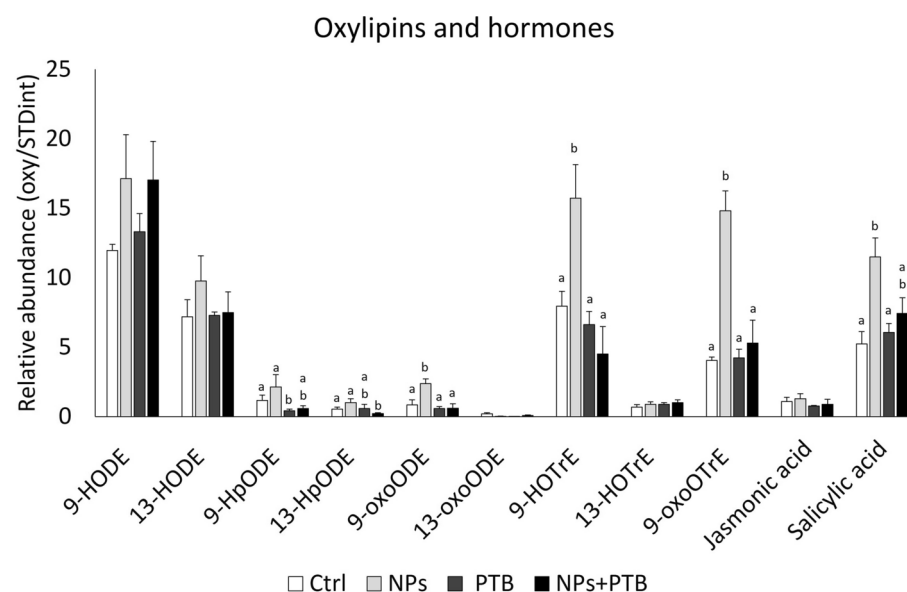


Fig. 6. Oxylipin and hormone abundances in tomato plants treated with NPs, PTB, and PTB NPs. Ctrl: control, plants treated with distilled water; NPs: PLGA NPs; PTB: pterostilbene; PTB NPs: PLGA NPs loaded with PTB; FW: fresh weight. The data were expressed as the mean of three replicates and bars represent the standard error. Different letters represent significant differences.

Natural compounds such as resveratrol have been reported as effective priming agents³⁴. Plant priming is a promising strategy consisting in an alteration of the plant's physiological state intentionally promoted by natural products or synthetic chemicals to enhance the defense response in the case of subsequent harsher stresses³⁵. When the plant reaches the “primed” state, the activated defense genes induce broad-spectrum defenses which change the metabolic profile and the accumulated defense compounds, among other processes, minimizing the negative effects on plant productivity³⁶. Unlike the results reported by Páramo et al.³⁷ in which changes in the increasing or decreasing content of chlorophylls were observed under different treatments with metal

nanoparticles, in the present study no significant differences were observed among the treatment groups with the respect of controls.

De Bona et al.³⁸ demonstrated that stilbene-rich extracts obtained from grape canes induce defense responses by activating mitogen-activated protein kinase (MAPK) and defense-related gene expression such as *PR* and *Glutathione-S-transferase 1 (GST1)* genes³⁸. Kang et al.³⁴ showed that resveratrol derivatives, as kobophenol A and hopeaphenol oligomers, can enhance priming events by promoting stronger local defense responses in *Arabidopsis thaliana* and *Nicotiana benthamiana*. As reported by Duke³⁹, severe plasma membrane damage in cucumber tissue caused by ROS and resulting in the accumulation of the protoporphyrin IX was restored through the exogenous application of resveratrol, pterostilbene, and α -tocopherol. Our results suggest that free PTB and PTB loaded in PLGA NPs could act as priming agents increasing GABA levels in tomato leaves, through the stimulation of GAD. Interestingly, tomato plants treated with free PLGA NPs showed lower levels of GABA compared to control plants confirming that the increase of GABA was due to PTB. As far as we know, there are no studies highlighting an increase in GABA levels in response to treatment with PTB in plants.

Amino acid metabolism is an important bridge between primary and secondary metabolites. Many amino acids are important precursors in the biosynthesis of alkaloids (e.g., arginine, lysine, ornithine, phenylalanine, proline, tryptophan, tyrosine), glucosinolates (e.g. methionine, leucine, isoleucine, phenylalanine, tryptophan), and phenylpropanoids (e.g., phenylalanine and tyrosine)^{40,41}. Among secondary metabolites, phenylpropanoids and flavonoids are crucial for their role in the preservation of the plant's metabolic machinery and stress tolerance. In particular, phenylpropanoids are involved in the regulation of oxidative stress, free ion chelation, cell wall lignification, and plant defense⁴². In addition, secondary metabolites are also known to be involved in pest defense⁴³ signal transduction in plant–microbe symbiosis⁴⁴ and plant innate immunity⁴⁵.

Most studies with NPs indicate their capability to modulate plant secondary metabolism, although the mechanism through which this process could occur is not understood. Accumulation of shikimate and phenylpropanoid pathway products was observed in cucumber and maize after foliar application of $\text{Cu}(\text{OH})_2$ ⁴⁶ in wheat exposed to Ag⁴⁷ in pepper exposed to SiO_2 or Fe_2O_3 ⁴⁸ and in *A.thaliana* exposed to CuO NPs⁴⁹. The concentration of benzoic acid and gallic acid was increased, while the content of hydroxycinnamic acid derivatives was reduced in *C. sativus* when exposed to CuO NPs⁵⁰. In our study, the decrease of quercetin-3-glucoside associated with the increase in caffeic acid (3,4-dihydroxycinnamic acid) observed after treatment with PLGA NPs and PTB suggests a shift in secondary metabolism towards the biosynthesis of phenylpropanoids and other phenolic compounds. In this context, both PLGA NPs and PTB acted increasing the levels of a cinnamic acid primarily involved in the synthesis of lignin, in the regulation of cell expansion, turgor pressure, phototropism, water flux, and growth processes. Caffeic acid and derivatives are known to be involved in both plant biotic and abiotic stress tolerance^{51,52}. In our study, the increase of caffeic acid was not associated with an increase in chlorogenic acid levels, being this latter an intermediate of its synthesis⁵³. The PLS-DA model comparing PLGA NPs + PTB vs PTB (Fig. 4) highlighted a decrease of carbohydrates such as sucrose and glucose in aid of synthesis of amino acids as threonine, arginine, and valine. This effect was determined by PLGA NPs. Conversely, the treatment with PTB increased sucrose and glucose levels and decreased arginine, choline and alanine levels compared to the Ctrl group (Fig. 2). In other words, PLGA NPs promoted the carbohydrate catabolism and protein synthesis as opposed to PTB.

The modulation in plant hormone levels is directly correlated with the physiological performance of the plants. The intricate cross-talk between NP exposure and phytohormone signaling results in synergistic or antagonistic interactions that play crucial roles in the response of plants to stress and are just beginning to be understood. Fe-NPs have been reported to affect the endogenous concentration of salicylic acid in plants⁵⁴.

In our study, an increase in salicylic acid levels was observed after treatment with PLGA NPs compared to control plants. Salicylic acid is a plant hormone synthesized from the shikimate pathway that plays a central regulatory role in plant immunity modulating disease resistance, stress tolerance, DNA damage/repair, fruit yield and seed germination as well as contributing to the systemic acquired resistance (SAR) and the activation of pathogenesis-related (PR) genes^{55–59}.

A strong increase of salicylic acid levels was observed after ZnONPs application in *A. thaliana* leaves⁶⁰. The remarkable role of salicylic acid in stress amelioration after NPs exposure has been demonstrated in several studies^{61,62}. Cai et al.⁶¹ demonstrated that the application of Fe_3O_4 NPs on *N. benthamiana* leaves increased salicylic acid biosynthesis and expression of salicylic acid-responsive PR genes (PR1 and PR2) leading to an improvement in plant resistance against TMV infection. On the other hand, the exposure of ZnONPs and co-treatment with exogenous salicylic acid in tobacco plants induced an increase of endogenous associated to the overexpression of salicylic acid-binding protein 2 (SABP2), which promoted the action of photosynthetic and antioxidative defense system⁶³.

Interestingly, besides the salicylic acid levels, NPs increased also the levels of caffeic acid as compared to control plants, suggesting a defense response mediated by the nanoparticles.

Free fatty acids act as signaling molecules in several biological processes, including plant stress response⁶⁴. Among the fatty acids analyzed, behenic acid (C22:0) significantly increased in the leaves of tomato plants treated with NPs, PTB and NPs + PTB compared to control plants. The C22:0 is produced via elongation system in the cytosol. Thus, it seems that NPs and PTB increased its synthesis. As C22:0 is present normally in a very low level in membrane complex lipids, it could be expected that its elevated level could be connected with increased synthesis of cuticle waxes⁶⁵, the hydrophobic layers of leaf surfaces with an important role in SAR^{66,67}. However, a detailed study of changes in lipid classes 'structures of analyzed leaves had not been done during presenting research.

Marked alteration may be observed in oxylipin category, they are biologically active compounds acting as signaling molecules for defense and development which can be generated because of the action of free radicals and reactive oxygen species. Oxylipins are lipid mediators consisting in polyunsaturated fatty acids (PUFA), such

as arachidonic acid (ARA), docosahexaenoic acid (DHA), eicosapentaenoic acid (EPA), α -linolenic acid (ALA) and linoleic acid (LA) and involved in reactions catalyzed by cyclooxygenase (COX), cytochrome P450 (CYP 450), lipoxygenase (LOX), as well as non-enzymatic oxidation pathways. Based on the position of the initial oxygenation, several enzymatic pathways can be described. Tomato has two 13-LOX genes and three 9-LOX genes. *TomloxC*, a 13-LOX gene, provides the substrate for jasmonate synthesis. *TomloxD* produces jasmonate in defense against herbivores⁶⁸. In rice, the cross-talk between the 13-LOX and 9-LOX products was observed to lead to elevated JA levels that improved the responses against herbivores. In this study, products of 13-LOX and 9-LOX such as 9-HpODE and 13-HpODE, derived from linoleic acid (C18:2), significantly decreased in tomato leaves after treatment with PTB and NPs + PTB. Conversely, 9-oxo-ODE, 9-HOTrE and 9-oxoOTrE significantly increased in plants treated with NPs compared to the control group. 9-oxylinins were usually defined as “death oxylinins” since in *A. thaliana* they are upstream the event causing HR-related PCD and therefore closely related to salicylic acid elevation into challenged tissues⁶⁹. NPs, even considering the result with behenic acid which clearly indicates a membrane-stress could trigger HR-related events. In this sense, the use of NPs charged with known antioxidants and inhibitors of LOXs (oxylinins significantly decreased in NPs + PTB compared to NPs group), suggesting the capacity of NPs and especially PTB⁷⁰ loaded in NPs to differently modulate the oxylinin metabolism and consequently HR-related events. Considering the antioxidant properties of PTB, it could compensate for the stimulating effect of NPs, leading to a reduction in the redox potential of the plant and therefore lowering the oxidation of fatty acids to oxylinins. Several evidence support the use of exogenously applied antioxidants to improve both plant growth and their resilience to stress⁷¹. Concerning this, the proper dose of PTB charged in NP could modulate in a tailored way the response of plant to stresses.

Final remarks

In this study, the impact of PLGA NPs and PTB on the metabolism of *S. lycopersicum* L. was investigated by performing untargeted metabolomics by ¹H-NMR and targeted analysis of lipidic compounds by HPLC-MS/MS. The obtained results demonstrated that both PLGA NPs and PTB affect the GABA, phenylpropanoid and lipidic metabolism of tomato plants. In this study we demonstrate the role of PLGA nanoparticles loaded with pterostilbene in redirecting leaf metabolome to boost the production of secondary metabolites, fatty acids, oxylinins and hormones. This leaf metabolic structure could prove useful to improve the plant defense response to future biotic or abiotic stress. Further studies will be needed to address the role of PLGA NPs loaded with PTB in environmental stress conditions. These data could be useful for interested researchers on developing alternative strategies to control crop plant diseases and constitute an important step forward in understanding the mechanisms of interaction between plant metabolism and NPs. In this context, we propose a possible use that should be further explored of PTB as a priming agent.

It should be emphasized that the impacts of whole NPs as well as of PTB on the yield and nutritional properties of crops are not exhaustively clarified. For this purpose, a coordinated research plan considering standardized experimental procedures, such as PTB concentrations, NPs application path and plant development stage, is necessary not only to guarantee the effect and to improve the crop production, but also to enhance plant tolerance towards environmental stresses and optimize the utilization of nutrients. Last, but not least, the future advances in phytonanotechnology will allow to make up for the vital gaps in regulatory and marketing policies driving a faster and safer development of nanotechnology for sustainable agriculture in the near future.

The ¹H-NMR-based untargeted metabolomics and LC-MS-based lipidomics approach allowed to disentangle the contribution of metabolic variations due to PLGA NPs and PTB. These aspects are relevant to better understand the metabolic impact of the interaction between nanoparticles and bioactive compounds. The fact that the observed metabolic changes were not severe indicates that the treatment of tomato plants with PLGA NPs and PTB is safe but could also improve the content of phenols in *S. lycopersicum* L. in terms of antioxidant and anti-inflammatory compounds.

Materials and methods

Chemicals

All solvents and standards used in this study were bought from VWR International (Milan, Italy) and Merck—Sigma-Aldrich (Darmstadt, Germany), unless specified otherwise. Pterostilbene was purchased from Chemodex (St. Gallen, Switzerland).

Synthesis, characterization, and in vitro release studies on PLGA NPs

Poly(D,L)-lactic-co-glycolic acid (PLGA), with a proportion 50:50 of lactide:glycolide and molecular weight of 50 kDa, was used for NPs preparation. 50 nm PLGA NPs, empty or loaded with PTB, were obtained by using the microfluidic reactor, as previously reported by Palocci et al.¹¹ The reactor has a flow-focusing configuration described previously by Chronopoulou et al.¹² PLGA NPs characterization and in vitro release studies of PTB from PLGA NPs have been conducted as described by De Angelis et al.⁸

Plant material

Tomato plants (*S. lycopersicum* cv ciliegino) with 3–4 true leaves were purchased from a commercial producer (Ortoflora Valter Finocchietti, Rome, Italy), and were cultivated in all-purpose soil (Compo Sana, Compo Italia Srl) in round pots (diameter 10 cm) under controlled conditions (photoperiod 16:8 light:dark at 24 °C during light hours and 20 °C during dark hours) in a phytotron before the experiments. The plants have not been subjected to any treatment by the producer. Each group (5 replicates) was treated with distilled water (control) or PLGA NPs (0.1 mg mL⁻¹), or pterostilbene (PTB) (0.04 mg mL⁻¹) or NPs loaded with PTB (0.04 mg mL⁻¹, respectively), at the stage of 9 true leaves. The treatments were administered by spraying water or NPs suspension

in water on the plants. The leaves were harvested 72 h after the treatment, immediately frozen with liquid nitrogen, and stored at -80°C until HPLC–MS/MS or ^1H -NMR analysis. Otherwise, leaves were extracted immediately after harvesting for chlorophyll and carotenoid analysis by spectrophotometric methods.

Sample preparation for ^1H -NMR analysis

Tomato leaves were sampled from nodes of the same age for each treatment. A total of 1.5 g of ground tomato leaves for each plant was extracted using a modified Bligh-Dyer protocol.⁷² In brief, five replicates for each treatment group were ground in a mortar with liquid nitrogen and added to a cold mixture composed of chloroform (3 mL), methanol (3 mL), and water (1.2 mL). The samples were stirred, stored at 4°C overnight, and then centrifuged for 30 min at 4°C with a rotation speed of 11,000 rpm. The upper hydrophilic and the lower organic phases were carefully separated and dried under a gentle nitrogen flow. A mixture of $\text{CD}_3\text{OD}/\text{D}_2\text{O}$ in a ratio of 1:2 containing 3-(trimethylsilyl)-propionic-2,2,3,3-d4 acid sodium salt (TSP, 2 mM) as an internal chemical shift and concentration standard was added to the hydrophilic phase. The hydrophilic phase was then analyzed by ^1H -NMR.

NMR experiments

NMR experiments were carried out at 298 K on a JNM-ECZ 600R spectrometer operating at the proton frequency of 600 MHz and equipped with a multinuclear z-gradient inverse probe head, and the monodimensional ^1H and bidimensional ^1H - ^1H TOCSY experiments were acquired according to Spinelli et al.⁷³.

Free fatty acid, oxylipin and hormone extraction and HPLC–MS/MS analysis

All solvents used for the extraction and HPLC–MS/MS analysis were of HPLC/MS grade. Mass spectrometry analyses were carried out by a HPLC 1200 series rapid resolution coupled to a triple quadrupole MS (G6420 series triple quadrupole, QqQ; Agilent Technologies, Santa Clara, CA, USA), with an electrospray ionization source (ESI). Chromatographic column and analysis software were purchased by Agilent Technologies (Santa Clara, CA, USA). The nitrogen flow was at 10 L/min, the nebulization pressure at 20 psi, the temperature was set at 350°C , and the voltage at 4000 V.

Tomato leaves sampled from nodes at the same age were lyophilized and ground with liquid nitrogen. Simultaneous extraction method was used for free fatty acids, oxylipins and hormones following the protocol described in Beccaccioli et al.⁷⁴. Briefly, 2 mL of extraction solution constituted by Isopropanol:Water:Ethyl Acetate (1:1:3 v/v) were added to 30 mg of powder leaf in presence of 1 μL of butylated hydroxytoluene (0.0025% w/v) to avoid oxidation and 2 μL of 9-HODEd₄ (Cayman Chemical) as internal reference standard (final concentration 2 μM calculated on the volume of final resuspension, 100 μL). The samples were vortexed (5 min), centrifuged (10 min, 4°C , 10,000 rpm), and the supernatant taken. The extraction was repeated on the matrices with 1.2 mL of ethyl acetate. The collected supernatants were dried under nitrogen flux and the dried samples were resuspended with 100 μL of methanol.

The chromatographic separation of free fatty acids, oxylipins and hormones was performed with a Zorbax ECLIPSE XDB-C18 rapid resolution HT 4.6×50 mm 1.8 μm column (Agilent technologies, Santa Clara, CA, USA). The extract was separated at a flow rate of 0.6 mL/min. The mobile phase consisted of A: water/acetonitrile (97:3 v/v containing 0.1% formic acid), and B: acetonitrile/isopropyl alcohol (90:10 v/v). The elution program was: 0–2 min 20% B, 2–4 min 35% B, 4–6 min 40% B, 6–7 min 42% B, 7–9 min 48% B, 9–15 min 65% B, 15–17 min 75% B, 17–18.50 min 85% B, 18.50–19.50 min 95% B, 19.50–24 min 95% B, 24–26 min 99% B, 26–30 min 99% B, 30–34 min 20% B. Sample was separated at a flow rate of 0.6 mL/min (0–24 min), 1 mL/min (24–30 min), 0.6 mL/min (30–34 min). The column temperature was set at 60°C . The injection volume was 10 μL . Free fatty acids were identified by single ion monitoring (SIM) approach, while oxylipins and hormones were identified by Multiple Reaction Monitoring (MRM) approach. Detailed data on all the identified metabolites are shown in S1 and S2 tables.

Extraction and quantification of chlorophylls and carotenoids of tomato leaves

The content of chlorophylls and carotenoids was analyzed by spectrophotometric analysis⁷⁵. A total of 50 mg of tomato leaves were sampled from nodes of the same age for each treatment. The samples were added to a 96% hydroalcoholic methanol solution (v/v), in a 1:50 ratio (w/v) and maintained at 4°C under dark conditions for 72 h. The extract was then separated and analyzed using the Shimadzu UV-1280 spectrophotometer (Japan). The analysis for quantifying chlorophyll a (Chl a) and b (Chl b) was carried out at 666 nm and 653 nm, respectively. The total content of carotenoids (Cars) in leaves was determined at 470 nm. After obtaining the absorbances (abs) of the samples, chlorophylls and carotenoids were quantified with the following formulas and the quantity was expressed in $\mu\text{g gFW}^{-1}$ (fresh weight):

$$\text{Chla} = 15.65 \times (\text{abs } 666 \text{ nm}) + 7.34 \times (\text{abs } 653 \text{ nm})$$

$$\text{Chlb} = 27.05 \times (\text{abs } 666 \text{ nm}) - 11.21 \times (\text{abs } 653 \text{ nm})$$

$$\text{Cars} = \frac{1000 \times (\text{abs } 470 \text{ nm}) - (2.86 \times \text{Ca}) - (129.2 \times \text{Cb})}{245}$$

Statistical analysis

Multivariate partial least square analysis (PLS-DA) was applied on the metabolomic data matrix with using the Unscrambler ver. 10.5 software (Camo Software AS, Oslo, Norway).

Univariate t-test and one-way ANOVA were carried out on the metabolomics and lipidomics data matrix, chlorophyll and carotenoids with SigmaPlot 14.0 software (Systat Software Inc., San Jose, CA, USA). The Shapiro–

Wilk test was performed on each variable to assess data normality prior to one-way ANOVA. A Kruskal–Wallis test was applied to identify significant differences between categories ($p < 0.05$). Mann–Whitney U test was also run to determine significant differences for considered metabolites. A p -value < 0.05 was considered statistically significant.

Data availability

Data supporting the results are included in the article. The datasets used and/or analyzed during the current study are available from the corresponding author on reasonable request.

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Author contributions

CB, MB, VP and LC performed the experiments; EB, CB and MB wrote the manuscript; MB, FS, EB, CB, VP analyzed the data, MR, AM, CP, GP critically evaluated the manuscript. All the authors approved the final version of the manuscript.

Declarations

Competing interests

The authors declare no competing interests.

Additional information

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Correspondence and requests for materials should be addressed to E.B.

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