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Silica Nanoparticles Enhance Disease Resistance in *Arabidopsis* Plants

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Abstract

In plants, pathogen attack can induce an immune response known as systemic acquired resistance (SAR) that protects against a broad spectrum of pathogens. In the search for safer agrochemicals, silica nanoparticles (SiO₂-NPs, food additive E551) have recently been proposed as a new tool. However, initial results are controversial, and the molecular mechanisms of SiO₂-NP-induced disease resistance are unknown. Here, we show that SiO₂-NPs, as well as soluble orthosilicic acid (Si(OH)₄), can induce SAR in a dose-dependent manner, that involves the defence hormone salicylic acid. Nanoparticle uptake and action occurred exclusively through stomata (leaf pores facilitating gas exchange) and involved extracellular adsorption in leaf air spaces of the spongy mesophyll. In contrast to treatment with SiO₂-NPs, induction of SAR by Si(OH)₄ was problematic, since high concentrations caused stress. We conclude that SiO₂-NPs have the potential to serve as an inexpensive, highly efficient, safe, and sustainable alternative for plant disease protection.

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Author Contributions

M.E.-S., F.S., and F.M. conceived and designed the study. F.S. led the team, rationally designed the SiO₂-NPs to induce optimal plant defence, performed initial germination tests to establish the dosing regimen, contributed the mechanistic understanding of silica and with plant TEM, and has drawn the artwork. M.M. synthesized and characterized the SiO₂-NPs. A.M. cultured the *Arabidopsis* plants and conducted the *C. elegans* experiments. D.R. provided access to his microtome and a technician that trained F.S. in microtoming. M.E.-S. performed all the *Arabidopsis* experiments and their statistical evaluation and wrote the manuscript draft with contributions by F.S. (figures and text) and F.M. (text). F. M. contributed the mechanistic understanding of the gene expression results and molecular mechanisms of SAR. The manuscript was critically reviewed by A.P.-F., B.R.-R., and D.R. All co-authors read and approved the manuscript prior to submission.

Competing Interests

Any opinions, findings, conclusions, or recommendations expressed in this material are those of the authors and do not necessarily reflect the views of the Swiss NSF or the government. This work has not been subjected to SNSF review, and no official endorsement should be inferred. F.S. and M.M. have a patent pending on a SiO₂-NP plant growth enhancer. F.S. was supported by Innosuisse (project No. 38515.1 IP-EE). Other than that, the authors have declared no conflict of interest and are responsible for the content and writing of the article.

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Nanoagrochemicals are a promising tool to improve crop yield and thus, global food security¹. Silica nanoparticles (SiO₂-NPs) have been proposed for the controlled nanodelivery of silicon and other active ingredients to plants but have never been systematically tested for this purpose. Silicon from orthosilicic acid (Si(OH)₄, monosilicic acid), the hydrolytic degradation product of SiO₂-NPs, is the only known form of silicon bioavailable for plants, and is ubiquitous in soil pore water²⁻⁴. Orthosilicic acid can promote plant growth and plant resistance against biotic and abiotic stresses^{3,5}, thereby protecting plants against pathogen attacks or agricultural damages related to severe climate conditions^{3,6,7}. The uptake and movement of SiO₂-NPs as well as other engineered nanomaterials in plants have been studied intensively in the last decade⁷⁻¹⁰. However, it is uncertain how the nanoparticles interact with leaves at the subcellular level. Direct evidence by nm-resolution imaging for the entrance of intact nanoparticles into leaves, or intercellular movement of SiO₂-NPs within leaves, is mostly missing¹⁰. It is also not known whether SiO₂-NPs can induce resistance in plants, whether their performance differs from dissolved silicon species, and which molecular pathways they may induce.

To fend off potential pathogens, plants have evolved disease resistance mechanisms that share mechanistic principles with the innate immunity of animals¹¹. An especially interesting form of plant disease resistance is the so-called induced resistance in which disease resistance of the plant can be enhanced by previous exposure to beneficial rhizosphere microorganisms, avirulent and virulent pathogens, or specific resistance-inducing chemical compounds¹²⁻¹⁴. A hallmark of induced resistance is its activity against a broad spectrum of pathogens. While the induction of plant disease resistance using chemical compounds is relatively well understood¹², the benefit of using slow nano-enabled delivery systems for the same purpose has not been investigated in systematic experiments^{1,7}.

A special form of induced resistance is systemic acquired resistance (SAR) that is characterized by the spread of locally induced disease resistance to the whole plant^{15,16}. Systemic acquired resistance is induced in all plant parts after challenging the plant locally with a pathogen or by local application of so-called resistance-inducing compounds. Both treatments induce signal transduction pathways that lead to the production of signals that move to distant tissues¹⁴. A key signalling compound that contributes to SAR is the plant hormone salicylic acid (SA) that is responsible for the activation of pathogenesis-related (PR) genes^{16,17}. Other factors include, *e.g.*, nitric oxide and reactive oxygen species^{18,19}. The fact that SAR can be activated by the application of resistance-inducing compounds^{12,13} makes SAR an attractive alternative strategy for controlling crop pests without the use of irreversible genetic modifications or environmentally problematic pesticides.

Systemic acquired resistance-inducing compounds such as benzothiadiazole successfully enhanced disease resistance, but also reduce crop yields^{20,21}. Interestingly, silicon-based compounds also seem to have the capacity to induce disease resistance *via* a broad range of different and partially still unknown mechanisms, including the mechanical reinforcement of defensive structures of the plant architecture, most notably the cell wall^{3,22}, but also by the activation of biochemical defenses^{3,23}. Biochemically, root-applied silicon led, for example, to a broad-spectrum resistance against powdery mildew pathogen by increasing the activity

of defence-related enzymes in leaves²⁴. It is important to note that the protective effect of silicon seems to have, in contrast to other biostimulants such as benzothiadiazole, no negative effects on the growth and yield of plants^{3,25}. All of this makes Si an attractive candidate to strengthen plant stress tolerance. Initial studies found that SiO₂-NPs may induce stress tolerance similarly to conventional Si products, but a clear mechanistic understanding of the underlying processes is still lacking^{7,8,26,27}.

In this work, we demonstrate the potential of SiO₂-NPs in inducing local and systemic disease resistance in the widely used model plant *Arabidopsis thaliana* against the bacterial pathogen *Pseudomonas syringae*. Silicic acid was assessed in parallel to disentangle the potential differences in the mode of action of dissolved Si species compared to SiO₂-NPs. We assessed the role of salicylic acid and reactive oxygen species (ROS) defence-related genes, established the therapeutic concentration range of SiO₂-NPs to induce the desired beneficial effects in plants, compared the laboratory setup (infiltration of selected leaves) with the more realistic spray application, and visualized the nanoparticle-leaf interactions using transmission electron microscopy (TEM), with important implications for future strategies to apply nanoscale active ingredients for slow release on leaves.

SiO₂-NPs and Subcellular Distribution within the Leaf

The SiO₂-NP suspensions used for the dosing of the plants (Fig. 1) were well dispersed with a hydrodynamic particle size of 76.7±0.8 nm and a polydispersity index of 0.07. The primary particle size, as determined by TEM, was 54±7 nm. The interaction of the nanoparticles with the plant was assessed by TEM (Fig. 2) 2 d after application of SiO₂-NPs. Preliminary experiments had shown that at this time point the SiO₂-NP-exposed plants already had developed resistance. The size range of ~50-70 nm of the nanoparticles allowed them to enter the leaf exclusively through the stomata and distribute within the large extracellular air spaces of the spongy mesophyll without penetrating any cell walls (Fig. 2, Supplementary Fig. S1). The SiO₂-NPs remained in the leaf air spaces during the 2 d between their application and the time point of TEM observation. At the same time, the size of the nanoparticles prevented (undesirable) nanoparticle uptake into the cytoplasm as well as cell-to-cell translocation through the plasmodesmata (Fig. 2b). This is in line with previous literature based on nm-resolution imaging of nanoparticles in plants suggesting that the cut-off for root-shoot nanoparticle translocation is at approx. <36 nm, and for cell-to-cell plasmodesmata transport <15-40 nm (basal size exclusion limits of ~3–4 nm)¹⁰. Compared to the fully closed stomata in the control plants (samples were kept in the dark for fixation), the nanoparticle-treated plants showed incompletely closed stomata due to nanoparticles stuck in between the guard cells (Fig. 2b).

Exogenous Application of SiO₂-NPs confers SAR

The local defence responses of *Arabidopsis* sprayed with SiO₂-NPs or a control treatment to virulent *P. syringae* were quantified *via* the bacterial growth on leaves (Fig. 3a). Due to the lack of the *avrRpt2* gene in the virulent *P. syringae* that is needed by the *RPS2* resistance gene in *Arabidopsis* to induce a strong plant defence against *P. syringae*^{28,29}, a severe infection would be expected. However, a pronounced infection only occurred in the control

treatment. Plants sprayed with SiO₂-NPs showed an 8-fold improved basal resistance compared with 4 (2 hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES) buffer-treated control plants (Fig. 3a), demonstrating that the SiO₂-NPs induced a local defence in the plant within 24 h (the nanoparticles were applied 24 h before inoculation with virulent *P. syringae*). The number of bacteria was reduced 8-fold in SiO₂-NP-treated plants compared to the control.

The systemic responses of wild type *Arabidopsis* plants to SiO₂-NPs and dissolved Si species are reflected in the inhibited bacterial growth in Fig. 3b-c. The positive control showed that plants previously infiltrated with the avirulent *P. syringae* that is known to induce systemic acquired resistance, contained as expected 10-fold less virulent *P. syringae* compared with MgCl₂ or HEPES-preinfiltrated plants (Fig. 3b-c). Remarkably, treating local leaves with SiO₂-NPs led to comparable systemic protection against virulent *P. syringae* as observed in the positive avirulent *P. syringae* control (Fig. 3a), which is equal to >90% bacterial inhibition. It is highly unlikely that a local response in distal tissue to SiO₂-NPs or Si(OH)₄ has caused this resistance because of the observed distribution and very slow dissolution of SiO₂-NPs (Fig. 2, Supplementary Fig. S1), and the passive transport³⁰ and high reactivity of Si(OH)₄. This shows that treating *Arabidopsis* with SiO₂-NPs induced local and systemic resistance to *P. syringae*.

It is well known that Si(OH)₄ improves plant defences against different plant pathogens such as fungi, bacteria, and viruses^{5,7}. We therefore also tested SAR in response to Si(OH)₄ (Fig. 3c) and found that treatment with Si(OH)₄ was also able to induce SAR. These results suggest that Si(OH)₄ released from SiO₂-NPs is at least partially responsible for the SAR-inducing ability of SiO₂-NPs, and that the SiO₂-NPs can act as a slow release source for Si(OH)₄.

Measuring the exact amount of free Si(OH)₄ directly *in planta* is challenging due to the low concentrations and fragile equilibrium of dissolved Si(OH)₄ and Si oligomers, and solid SiO₂ species^{2,4,31} (refer to the Supplemental Information, section 'Details on Si(OH)₄ Analytics'). We therefore resorted to direct TEM imaging of the nanoparticles in the plants that revealed at high resolution abundant intact SiO₂-NPs in stomata 2 d after the SiO₂-NP treatments (Fig. 2). This demonstrates that the plants could not degrade the nanoparticles at the time point of inoculation with virulent *P. syringae* (in all assays nanoparticles were applied at least 24 h before inoculation). The slow nanoparticle dissolution is in line with the slow dissolution kinetics of the SiO₂-NPs we measured previously in water (half-life ~66 d at pH 7)³².

To test whether SiO₂-NPs and Si(OH)₄ have a direct toxic effect on bacterial growth, virulent *P. syringae* was cultivated *in vitro* in the presence or absence of SiO₂-NPs, or Si(OH)₄ at the lowest fully effective dose of SiO₂-NPs at 100 mg L⁻¹. At these concentrations that induced strong defence in plants, neither SiO₂-NPs nor Si(OH)₄ harmed the growth of the virulent *P. syringae* bacteria alone (Fig. 3d), demonstrating that SiO₂-NPs induce resistance in plants by activating defence responses in plants and not by directly inhibiting bacterial growth.

Dose Dependence of SAR Response

The SAR was further tested in response to different concentrations of SiO₂-NPs or Si(OH)₄, using for additional validation a second bacterial growth quantification method³³ based on bacterial DNA (Fig. 4a-b, Supplementary Tab. S1). Treatment with SiO₂-NPs at a concentration of 25 mg SiO₂ L⁻¹ already resulted in a partial reduction of 29% of bacterial growth in systemic leaves, and treatment with 100 mg L⁻¹ resulted in maximal protection (>90%) compared to positive control plants preinfiltrated with avirulent *P. syringae* (Fig. 4a). As the series of concentrations in Fig. 4a shows, higher concentrations of SiO₂-NPs >1600 mg SiO₂ L⁻¹ led to increased bacterial infection and were thus less effective in activating SAR. Pre-treatment with a concentration of 5 mg SiO₂ L⁻¹ of Si(OH)₄ (concentration normalized to SiO₂ L⁻¹ for the sake of comparability) led to a reduction in bacterial numbers of 81% compared to the positive control. Maximal protection with a reduction similar to control plants preinfiltrated with avirulent *P. syringae* was achieved at concentrations between 20 to 320 mg SiO₂ L⁻¹. A higher concentration of 640 mg SiO₂ L⁻¹ was less effective, and a concentration of 2560 mg SiO₂ L⁻¹ was ineffective in inducing SAR, demonstrating a detrimental effect of high concentrations of Si(OH)₄ on SAR induction.

The data of Fig. 4 served to establish a dose-response relationship between SAR and SiO₂-NP concentration (Fig. 5a). Using a standard log-logistic dose-response model, the dynamic range and the effective concentration at 50% bacterial inhibition (EC₅₀) was determined at 0.4±0.04 mM Si for SiO₂-NPs (*i.e.*, 24 mg SiO₂ L⁻¹, refer to Supplementary Fig. S2 for the residual analysis and Supplementary Tab. S2 for the fitting parameters) in a range of 25 to 100 mg SiO₂ L⁻¹. For spraying, the EC₅₀ may be similar to injected SiO₂-NPs, as local (sprayed) and systemic (injected) assays at 100 mg SiO₂ L⁻¹ both resulted in disease resistance (Fig. 3, Fig. 6a-b).

The results based on counting bacterial colonies were confirmed by estimating the bacterial biomass based on qPCR analysis of the bacterial outer membrane protein gene *oprF* (Fig. 4b). The bacterial DNA levels were in good agreement with the bacterial colony counting results in Fig. 4a, in line with previous research that compared the two techniques³³.

In contrast to SiO₂-NPs, high concentrations of Si(OH)₄ adversely affected the phenotype of treated plants (Fig. 3e). At a concentration of 1000 mg SiO₂ L⁻¹, leaves of plants treated with Si(OH)₄ showed signs of chlorosis (yellowing), whereas leaves of plants treated with SiO₂-NPs looked healthy (Fig. 3e). This different behaviour at higher concentrations prompted us to further investigate the negative effect of high concentrations of SiO₂-NPs and Si(OH)₄. The expression level of the heat shock protein *AtHSP17.4C1*, a molecular marker for oxidative stress³⁴, was analysed by reverse transcription-quantitative real-time polymerase chain reaction (RT-qPCR). The *HSP17.4C1* transcript levels were determined in response to avirulent *P. syringae*, SiO₂-NPs, or Si(OH)₄ (100 and 1000 mg SiO₂ L⁻¹; Fig. 4c) 2 d after the treatments. Treatment with avirulent *P. syringae* caused a minor increase (2.7-fold) in *AtHSP17.4C1* expression compared to the control. Similarly, treatment with SiO₂-NPs led to a 1.6-fold (100 mg SiO₂ L⁻¹) and 2-fold (1000 mg SiO₂ L⁻¹) increase in transcript abundance relative to control treatment that was not statistically significant.

However, treatment with high concentrations of Si(OH)_4 caused stress, as transcript levels of the oxidative stress marker gene *HSP17.4CI* were induced 9-fold at a concentration of 100 mg $\text{SiO}_2 \text{ L}^{-1}$ and 18-fold at 1000 mg $\text{SiO}_2 \text{ L}^{-1}$.

SiO_2 -NP mediated SAR depends on Salicylic Acid

The plant hormone salicylic acid (SA) plays a core regulatory role in plant immunity³⁵. We therefore tested the possibility that SiO_2 -NP-mediated SAR might function *via* the salicylic acid-dependent defence pathway by testing the ability of SiO_2 -NPs to induce local disease resistance and SAR in an *Arabidopsis* mutant defective in salicylic acid biosynthesis (*salicylic acid deficient 2 (sid2)*³⁶). Notably, neither Si(OH)_4 nor SiO_2 -NPs induced local disease resistance or SAR in *sid2* mutant plants, while they induced basal disease resistance and SAR in wild type plants (Fig. 6a,b), demonstrating that salicylic acid-dependent defence signalling is essential for Si(OH)_4 - and SiO_2 -NP-induced disease resistance. To further support this result, we next quantified the expression of the salicylic acid-responsive marker genes Pathogenesis-related protein 1 (PR-1, gene *AtPR-1*) and PR-5 (gene *AtPR-5*) in wild type plants (Fig. 6c-d). Similar to treatment with avirulent *P. syringae*, treatment with Si(OH)_4 and SiO_2 -NPs resulted in an up to 30-fold and 6-fold increase in transcript abundance of *AtPR-1* (Fig. 6c) and *AtPR-5* (Fig. 6d), respectively, in comparison to control treatments. Hence, both Si(OH)_4 and SiO_2 -NPs activated salicylic acid-dependent defence reactions. Although SiO_2 -NPs triggered lower *AtPR-1* and *AtPR-5* expression levels in comparison with avirulent *P. syringae* infiltrated plants and Si(OH)_4 treated plants, the inducing effect was sufficient to confer SAR.

Implications on the Mode of Action of Leaf-Applied SiO_2 -NPs

The pathosystem involving *Arabidopsis* and the hemibiotrophic bacterial pathogen *P. syringae* offers an ideal model to investigate the effect of SiO_2 -NPs and Si(OH)_4 on plant defence. Our results summarized in the model in Fig. 5b-c show that the protective effect of SiO_2 -NPs and Si(OH)_4 is based on the ability to induce basal resistance and SAR (Fig. 3a-c) and not on direct toxic effects as neither SiO_2 -NPs nor Si(OH)_4 inhibited bacterial growth (Fig. 3d). Our data is in line with initial results that suggested that Si(OH)_4 and sometimes SiO_2 -NPs can protect plants from different plant pathogens^{7,26,27,37}, nevertheless here we show that the mechanism involved no toxic effect on the pathogen, but rather induced the defences of the plant.

Both SiO_2 -NPs and Si(OH)_4 induce SAR in a dose-dependent manner, leading to bacterial inhibition of >90% compared to control plants treated with HEPES buffer or MgCl_2 only. These results are consistent with previous results suggesting that SiO_2 -NPs and Si(OH)_4 function in a dose-dependent manner in plants and animals^{26,27,38}. However, instead of the previously proposed pesticidal action of SiO_2 -NPs, we show here that the nanoparticles caused an increase of the plant defence. Our data suggest that the SiO_2 -NPs used in the present study can be successfully used to slowly release Si(OH)_4 to the plant from within the spongy mesophyll (Fig. 2) in close direct interaction with the diffusion layer on the plant cell walls, which is at least partially responsible for the SAR-inducing ability of SiO_2 -NPs. Water (vapour) secreted from the plant cell wall, or plant-induced dissolution of SiO_2 -NPs

linked to increased secretory activity¹⁰ (exudates) may have promoted further dissolution of Si(OH)_4 . Based on the release rates of SiO_2 -NPs that were determined earlier under conditions optimized for dissolution in a continuously depleted ultrapure water system (half-life of ~66d at pH 7)³², max. ~13% of the particles could have dissolved within the 48 h of SiO_2 -NP exposure. Si-containing reaction byproducts of the nanoparticle synthesis were ruled out to play a significant role in the induction of defence (refer to section ‘Si Reaction Byproducts’ in the Supplemental Information). The max. released Si(OH)_4 from SiO_2 -NPs could thus explain the bacterial inhibition; however, it cannot fully explain the lack of oxidative stress responses and higher bacterial DNA levels for SiO_2 -NPs in the plants (Fig. 4b). Probably, the absence of peak Si(OH)_4 concentrations resulted in lower Si(OH)_4 toxicity for both bacteria and plants. Other effects, such as modulated evapotranspiration due to the blockage and incomplete closure of the stomata by the nanoparticles (Fig. 2) which can cause salicylic-acid related responses similar to drought stress³⁹, and the close interaction of the nanoparticles with cells in the spongy mesophyll may play an important role, in line with earlier research about stomata as ports of entry for pollutants and nanoparticles^{40,41}. The exact relative contribution of each effect remains to be elucidated in follow-up studies. It is important to note that the cell walls in the mesophyll air spaces have very thin, or lack, cuticular waxes¹⁰, and therefore, in contrast to the external leaf surface, a direct interaction of the nanoparticles can take place with the cell wall and thus apoplast transport system including the xylem. Irrespective of the detailed mechanism of the nanoparticles, this is of importance for any nanoagrochemical application aiming at slow release of active ingredients, because nanoparticles in the extracellular spongy mesophyll air spaces (Fig. 2, Supplementary Fig. S1) can interact with the leaf for extended periods without being washed away by rain.

High concentrations of Si(OH)_4 caused chlorotic yellowing of leaves indicative of stress (Fig. 3e). Increased expression of the oxidative stress marker gene *AtHSP17.4C1*³⁴ confirmed stress in the Si(OH)_4 treatment at 100 and 1000 mg $\text{SiO}_2 \text{ L}^{-1}$, as transcript levels of *AtHSP17.4C1* were more strongly induced in comparison with avirulent *P. syringae*, or SiO_2 -NP treatments (Fig. 4c). Together, these data show that the Si(OH)_4 was more toxic to plants than SiO_2 -NPs. Hence, impaired SAR in plants treated with high concentrations of Si(OH)_4 (Fig. 4a) might be linked to enhanced oxidative stress, consistent with the fact that high levels of NO and reactive oxygen species were shown to impair the induction of SAR^{19,23}. For SiO_2 -NPs, no significant increase of the oxidative stress marker gene was found (Fig. 4c). Impaired SAR for SiO_2 -NPs occurred only at very high concentrations in the g L^{-1} range (Fig. 4a), likely due to excess release of Si(OH)_4 causing oxidative stress, or too intense clogging of the stomata (Fig. 2) disrupting evapotranspiration. While the low polydispersity index measured by DLS (Fig. 1c) indicates well-dispersed SiO_2 -NP suspensions even at high concentrations, upon contact with the leaf, heteroaggregation with mucilage in stomata, and at higher nanoparticle concentrations, probably also homoaggregation, appeared to promote the clogging of the stomata (Fig. 2a, red arrows). These results are in line with Slomberg *et al.*⁸, who found that SiO_2 -NP concentrations up to 1000 mg $\text{SiO}_2 \text{ L}^{-1}$ were not phytotoxic despite the uptake of the SiO_2 -NPs into the root system of *A. thaliana*. Our results are also consistent with initial studies^{42,43} that found

better effects of SiO₂-NPs on plant growth than conventional silica fertilizers. In conclusion, the application of SiO₂-NPs can reduce the risk of overdosage.

Our data demonstrate that SiO₂-NPs and Si(OH)₄-mediated SAR acts *via* the activation of the salicylic acid-dependent defence pathway, which is a key component of basal disease resistance and SAR^{44,45}. Neither SiO₂-NPs nor Si(OH)₄ induced resistance in the salicylic acid-deficient mutant *sid2* that has a defect in salicylic acid biosynthesis (Fig. 6a-b). The induction of resistance by SiO₂-NPs was comparable to the effect of Si(OH)₄ at intermediate concentrations, albeit the soluble fraction of Si(OH)₄ in this treatment was far lower, as the particles dissolved only partially in the plant, if at all (Fig. 2), suggesting that SiO₂-NPs can induce salicylic acid-dependent defence pathways as intact particles. Furthermore, the expression levels of two salicylic acid-responsive marker genes *AtPR-1* and *AtPR-5* encoding the Pathogenesis-related protein 1 (PR-1) and Pathogenesis-related protein 5 (PR-5) were induced in response to SiO₂-NPs and Si(OH)₄ (Fig. 6c-d). These results are in line with Fauteux *et al.*⁴⁶, who reported that exogenous application of Si(OH)₄ induced salicylic acid biosynthesis in leaves exposed to the fungal pathogen *Erysiphe cichoracearum*. In addition, silicon-primed tomato plants were protected against *Ralstonia solanacearum* *via* the upregulation of salicylic acid-controlled defence gene expression⁴⁷. Although SiO₂-NPs triggered lower *AtPR-1* and *AtPR-5* expression levels than plants infiltrated with avirulent *P. syringae* and Si(OH)₄-treated plants, the achieved level of expression was sufficient to confer a full SAR response.

Conclusions

The present results show that low concentrations of SiO₂-NPs efficiently protect the widely used model plant *Arabidopsis* from infection by the bacterial pathogen *Pseudomonas*, and revealed the mode of action of SiO₂-NPs compared to its dissolved counterpart Si(OH)₄. The protective effect of SiO₂-NPs is mediated by the activation of salicylic acid-dependent plant immunity responses and is partially based on the slow release of Si(OH)₄ from nanoparticles entering through the stomata and distribution within the spongy mesophyll, and likely partially by intact nanoparticle-induced salicylic acid-dependent responses.

In comparison to direct Si(OH)₄ application, SiO₂-NPs proved to be safer for the plant. They did not cause phytotoxicity even at concentrations 10-fold higher than the minimal dose needed for plant protection, and therefore have a broader therapeutic range than Si(OH)₄. The lowest fully effective dose (100 mg SiO₂ L⁻¹) is promising because it corresponds to an extrapolated field dose of only 3 kg SiO₂ kg ha⁻¹, corresponding to >1000-fold material savings compared to solid bulk SiO₂ treatments. This calculation assumes a typical 300 L ha⁻¹ application (conventional aqueous spray volumes for pesticide application equipment⁴⁸), and an uncertainty factor of 100 for the concentration. Contrary to previous assumptions about the ability of nanoparticles to penetrate the cuticle, SiO₂-NP intake was clearly restricted to the stomata and extracellular spongy mesophyll, confirming our hypothesis that the leaf cuticle represents an impermeable barrier to nanoparticles¹⁰, in line with earlier fundamental research⁴⁹. The spongy mesophyll is an attractive target for long-term deposition of slow-release nanoagrochemicals. Future research should extend the investigations to a broader spectrum of defence-related genes with other plant pathogens,

and to biomechanic quantification of the physical effects of nanoparticles that affect leaf permeability and may trigger the salicylic acid-related responses. To further advance SiO₂-NPs as nanobiostimulants and fertilizers, as this should be the case with every material or organism used in agriculture, the long-term effects of SiO₂-NPs to occupationally exposed agricultural workers and non-target organisms, such as beneficial soil microorganisms or bees, must be thoroughly analysed before broad commercial application. The potential risks of nanoagrochemicals, and possible strategies for risk mitigation have been thoroughly reviewed previously^{1,50,51}. As for amorphous SiO₂-NPs, they have already been approved by the FDA as they are generally regarded as safe, and they are in use as dietary additives (E551)⁵² in a broad range of foodstuffs such as table salt. The daily intake of nano-scale silica from food is estimated to be 1.8 mg kg⁻¹⁵³. Our own initial experiments with *C. elegans* nematodes used as model non-target microorganisms (Supplementary Fig. S3) have shown a ~36-fold lower ecotoxicity of SiO₂-NPs compared to liquid Si(OH)₄ preparations that are in use for plant nutrition since decades. Thus, compared to currently used treatments, the present SiO₂-NPs alone, or in combination with other active ingredients, promise to offer a cost-effective, consumer-safe because tracelessly degradable, and sustainable alternative to protect plants against pathogens *via* the controlled induction of SAR, without negative effects on yield or non-target organisms that were associated with the action of previously described plant biostimulants or pesticides.

Methods

Plant Growth Conditions

Arabidopsis thaliana seeds were grown on Jiffy soil substrates (powered by Tref, Jiffy Products International B.V.). Two *A. thaliana* strains were grown: wild type Columbia (Col-0) plants that carry an *RPS2* locus responsible for the recognition of *P. syringae* strains expressing the avirulence gene *avrRpt2*^{28,29}; and an *A. thaliana* mutant defective in salicylic acid biosynthesis (*salicylic acid deficient 2*, *sid2*³⁶). The seeds sown on the soil were kept at 4 °C for two days and then transferred to the growth chamber (RMC Tableaux SA). The plants were grown in a 12 h photoperiod with 60% of relative humidity, with a day temperature of 22 °C and a night temperature of 18 °C (photon flux density 100 μmol m⁻² s⁻¹). The transplanted seedlings were covered with transparent plastic domes for 2-3 days to allow the seedlings to adapt to the new soil. Four-to five-week-old plants were used in experiments because previous experiments had shown that under the abovementioned growth conditions, this is the optimal age of the plant to induce SAR⁵⁴.

Culture of *Pseudomonas syringae* pv. *tomato*

Pseudomonas syringae pv. *tomato* bacteria were prepared by inoculating a single colony in 10 mL of King's B medium (1.5 g K₂HPO₄, 1.5 g MgSO₄·7H₂O, 20 g tryptone, 10 mL glycerol per 1 L of water, all from Sigma Aldrich, Switzerland, purity 99%) containing the appropriate antibiotics. A virulent and an avirulent strain of *P. syringae* were grown: *P. syringae* DC3000 (here termed virulent *P. syringae*); and *P. syringae* DC3000 expressing the avirulence gene *avrRpt2* recognized by the *A. thaliana* *RPS2* locus and inducing systemic acquired resistance (here termed avirulent *P. syringae*). The virulent *P. syringae* bacteria strain served to induce a strong infection with *P. syringae* in the plants. The avirulent *P.*

syringae strain served as a positive control to induce systemic acquired resistance and thus an actively suppressed bacterial growth in the *A. thaliana* plants *via* recognition of the bacterial *avrRpt2* gene by the plant's *RPS2* gene (refer to Chen *et al.* 2000 for a detailed description of the pathosystem²⁹). The virulent *P. syringae* was grown with rifampicin (25 $\mu\text{g mL}^{-1}$), and the avirulent *P. syringae* was grown with kanamycin (50 $\mu\text{g mL}^{-1}$) and rifampicin (25 $\mu\text{g mL}^{-1}$). After overnight incubation on a shaker at 28 °C in the dark (Kühner LT-W Lab Therm Table Top Incubator Shaker, Adolf Kühner AG, Switzerland), the cells were centrifuged at 3000 rpm for 10 min, and the pellet was suspended in 10 mM MgCl_2 . The cell density was calculated *via* measurement of the light absorption of the liquid culture at the absorption wavelength 600 nm using a spectrophotometer (BioPhotometer, Eppendorf – Netheler - Hinz GmbH, Germany) and by counting the colonies plated on King's B agar (raw data publicly available⁵⁵).

Inoculation Procedures for Local Disease Resistance

For a local disease resistance assay, three leaves per *A. thaliana* plant were inoculated with virulent *P. syringae* bacteria, and the plants were incubated at the standard *A. thaliana* growth conditions described above. The inoculation with the virulent *P. syringae* bacteria was operationally defined as 0 days *post* inoculation. After inoculation, leaf discs (4 mm) were harvested from the inoculated leaves at 0 and 3 days *post* inoculation using a cork borer (3 leaf discs from different plant leaves/sample). The leaf discs were ground and homogenized with pestles in 10 mM MgCl_2 and the undiluted (0 days *post* inoculation) or the 10^3 fold diluted (3 days *post* inoculation) homogenates were plated on King's B agar plates (King's B medium as above with 15 g L^{-1} agar). The plates were incubated at 28°C in the dark for 48 h. Then, the bacterial colonies were counted (raw data publicly available⁵⁵).

Inoculation Procedures for Systemic Acquired Resistance Assays

For a systemic acquired resistance (SAR) assay, three leaves of 4-5 week old wild type Col-0 plants were infiltrated with 10 mM MgCl_2 (negative control) or the avirulent *P. syringae* bacteria at 10^6 colony-forming unit (CFU) per millilitre in 10 mM MgCl_2 (positive control). After 48 h, the distal leaves were inoculated with the virulent *P. syringae* bacteria (10^5 CFU mL^{-1}). The inoculation with the virulent *P. syringae* bacteria was operationally defined as 0 days *post* inoculation. Leaf discs (4 mm) were harvested from the distal leaves at 0 and 3 days *post* inoculation using a cork borer (three times three leaf discs from different plant leaves were analysed for each treatment). The leaf discs were ground in 10 mM MgCl_2 , and the undiluted (0 days *post* inoculation) or the 10^3 fold diluted (3 days *post* inoculation) homogenates were plated on King's B agar and incubated at 28°C for 48 h in the dark (Salvis incubator, Switzerland). Then, the bacterial colonies were counted (raw data publicly available⁵⁵). For details about this procedure, refer to Wang *et al.* 2014.

Plant Treatments

The SiO_2 -NPs (25, 100, 400 and 1600 mg $\text{SiO}_2 \text{ L}^{-1}$, pH 7.0) and Si(OH)_4 (5, 20, 80, 100, 320, 640 and 2560 mg $\text{SiO}_2 \text{ L}^{-1}$, pH 7.0, from an aqueous potassium silicate stock solution, $\text{K}_2\text{O}:\text{SiO}_2$ 1:2.60, SiO_2 content 20.8 wt%, MonDroguiste, France) were prepared in sterile distilled water in 4-(2 hydroxyethyl)-1-piperazineethanesulfonic acid (HEPES) buffer (1-mM, pH 7.0, Sigma-Aldrich, 99.5%). The Si(OH)_4 concentrations were expressed in mg

$\text{SiO}_2 \text{ L}^{-1}$ to allow for direct comparison of the effects of dissolved $\text{Si}(\text{OH})_4$ and solid SiO_2 -NPs without having to take into account the different molecular weights.

For the local disease resistance assay, the plants were sprayed with these chemicals 24 h before inoculation with virulent *P. syringae*. For the SAR assays, all these chemicals were injected abaxially (from the bottom of the leaf) into *Arabidopsis* plant leaves 2 d before inoculation using 1 mL needleless sterile disposable syringes.

SiO_2 -NPs and Subcellular Distribution within the Leaf

The SiO_2 -NPs were synthesized and characterized according to a previously established procedure^{31,32} adapted from earlier work⁵⁶. Briefly, one equivalent of tetraethyl orthosilicate (TEOS, 10 mL, Sigma-Aldrich, >99%) was added to an equilibrated reaction mixture at 70°C containing two equivalents of ultrapure water (Milli-Q, 18.2 M Ω arium 611 DI, Sartorius Stedim Biotech, Germany), and absolute ethanol (81 mL) as a solvent under basic conditions (2.93 mL 25% NH_3). The particles resulting after 3 h of hydrolysis and polycondensation of TEOS were washed by three steps of centrifugation (15 000 $\times g$, 15 min) in ultrapure water, and by 5 or more steps of dialysis through a 14 kDa molecular weight cut-off membrane (regenerated cellulose, Carl Roth, Germany). Several batches of particles 64.8-76.7 nm in hydrodynamic diameter were prepared using the identical procedure to prevent artefacts due to suspension aging (size variability between batches 5.2 nm). Dynamic light scattering was used to quantify the hydrodynamic particle size and surface charge of diluted samples (1% v/v, Brookhaven Particle Size Analyzer Plus90, USA, scattering angle 90°, 1 min acquisition, raw data publicly available⁵⁵). Inductively coupled plasma - optical emission spectroscopy (ICP-OES) and gravimetry served to quantify the SiO_2 concentration (methods described in Bossert *et al.*³¹).

For the particle characterization, and to analyse the effects of SiO_2 -NPs, $\text{Si}(\text{OH})_4$, and control treatments in the leaves, we used transmission electron microscopy (TEM). The particle size distribution was established using ImageJ (version 1.52n) analysis of TEM micrographs (raw data publicly available⁵⁵). The plants were pre-fixed in 4% glutaraldehyde solution, gently stained in the dark with 1% OsO_4 solution that was centrifuged beforehand to remove potential precipitates, dehydrated using an ethanol series, and embedded in polymer resin (AGAR Low Viscosity Kit, Plano, Germany) without further staining according to a procedure described in detail in Stegemeier *et al.* 2015⁵⁷. The correct position of the stomata to cut cross-sections were identified by light microscopy examination of semi-thin resin sections before the ultramicrotoming. The TEM images were taken on an FEI Tecnai Spirit at an acceleration voltage of 120 kV (2048 \times 2048 px. Resolution, Veleta CCD camera, Olympus). Besides cropping and adjustment of brightness and contrast, the micrographs were not further processed; the unprocessed raw data is publicly available⁵⁵.

DNA Extraction

Plant leaf samples (five leaf discs from different inoculated plant leaves/sample) were frozen in liquid nitrogen and were homogenized using a ceramic mortar and pestle. The total DNA was extracted with a Plant DNA mini Kit (peqlab, a VWR brand, Germany). More

information about the sample preparation is available in section ‘Details on DNA Extraction’ in the Supplemental Information.

RNA Extraction and complementary DNA Synthesis

Plant leaf samples (10 leaf discs taken from different infiltrated plant leaves/sample) were flash-frozen in liquid nitrogen, and the total RNA was extracted with the Spectrum Plant Total RNA Kit (Sigma Life Science, USA). One microgram of the total RNA was used for complementary DNA synthesis using the Omniscript Reverse Transcription Kit (Qiagen, Germany). More information about the sample preparation is available in section ‘Details on RNA Extraction and complementary DNA Synthesis’ in the Supplemental Information.

Quantitative real-time polymerase chain reaction (qPCR)

To validate the SAR response based on bacterial colony counts, the bacteria were also quantified *via* the outer membrane protein *oprF* gene of *P. syringae* in inoculated leaves (raw data publicly available⁵⁵) based on a previously established method^{18,33}. For this bacterial DNA quantification, a reaction mixture for qPCR was prepared with 7.5 μ L of 2 \times SensiMix™ SYBR Hi-ROX Mastermix (No. QT605-05, Bioline, Meridian Bioscience, UK), 5 μ L of plant DNA, and 0.5 μ L of each primer (Supplementary Tab. S1) at a concentration of 10 μ M in a final volume replenished with water to 15 μ L in MIC tubes (Bio Molecular Systems, Australia). Runs were performed on a MIC qPCR machine (Bio Molecular Systems, Châtel-St-Denis, Switzerland). The conditions for the qPCR were 10 min. at 95 °C initial denaturation, and then 40 cycles (95 °C for 15 s, 62 °C for 1-min, and 72 °C for 30 s). The final PCR products were analyzed by melting point analysis. The qPCR analysis software for the melting curve analysis and amplification efficiency calculation was micPCR v2.8.13 from Bio Molecular Systems. This software is designed to meet MIQE⁵⁸ specifications and performs qPCR analysis automatically based on the real time runs. Five leaf discs from different plant leaves were sampled per each replicate, frozen in liquid nitrogen, and processed immediately for DNA extraction. The bacterial DNA levels of the bacterial *oprF* gene in *Arabidopsis* plants were calculated using At4g26410 (*expG*) as a reference gene³³ and the comparative cycle threshold method ($2^{-(C_t - C_{t0})}$)⁵⁹.

For oxidative stress and salicylic acid-responsive plant transcript levels, leaf discs were flash-frozen in liquid nitrogen and stored at -80°C for <24 h before being processed for RNA extraction and complementary DNA synthesis. Three independent technical replicates (ten leaf discs taken from different plant leaves) were used per treatment. The reaction mixture for RT-qPCR contained 7.5 μ L of 2 \times SensiMix™ SYBR Hi-ROX Mastermix (No. QT605-05, Bioline, Meridian Bioscience, UK), 5 μ L of complementary DNA (corresponding to 25 ng RNA), and 0.5 μ L of each primer (Supplementary Tab. S1) at a concentration of 10 μ M in a final volume replenished with water to 15 μ L in MIC tubes (Bio Molecular Systems, Australia). Runs were performed on a MIC qPCR machine (Bio Molecular Systems, Châtel-St-Denis, Switzerland). The conditions for the qPCR and the analysis of the final PCR products by melting point analysis were analogous to the above bacterial DNA quantification. The final PCR products were analyzed by melting point analysis. The transcript levels of the oxidative stress marker (At3g46230; *HSP17.4C1*)³⁴, and the salicylic acid-responsive genes *AtPR-1* and *AtPR-5* in *Arabidopsis* plants were

calculated with At4g26410 (*expG*) as reference gene⁶⁰ and the comparative cycle threshold method ($2^{-(C_t)}$) as mentioned above.

The *expG* gene was selected because Czechowski et al.⁶⁰ specifically recommends *expG* as one of the top five reference genes to be used in biotic stress studies due to its high stability under such conditions. This high stability was confirmed in previous work of our laboratory⁶¹ and others' work³³. In the present study, the stable expression of *expG* is reflected in the very small variation of the C_q for *expG*. In the PR1 expression experiments (Fig. 6c) for example, the average C_q ranged from 23.19-23.93 for all different tested conditions with an average relative error of only 0.63%⁵⁵. The amplification efficiencies were all very close to two and with good comparability of the reference gene to the target gene. For example, in Fig. 6c, the average amplification efficiency of *expG* and *AtPR-1* across all the different treatment conditions (1.949 ± 0.011 vs. 1.962 ± 0.027) differed by only 0.7%⁵⁵. All statistical tests hereinafter were performed using IBM SPSS Statistics (version 22).

Ecotoxicity of SiO₂-NPs and Si(OH)₄ to *C. elegans* larvae

The ecotoxicity assays were conducted on larval stage one (L1) nematodes of the *C. elegans* wild type (ancestral; N2) genotype. Synchronized *C. elegans* larvae were grown according to a previously established protocol⁶² (raw data publicly available⁵⁵). A known number of larvae (~70) per replicate were then exposed to 0, 25, 125, 250, 500, 750, 1000, 1500, or 2000 mg SiO₂ L⁻¹ of SiO₂-NPs or Si(OH)₄ in 96-well plates (Corning Costar No. 3596). A 0.1% NaN₃ solution served as a positive control. As a food source for the nematodes, the wells contained 10 μ L of living *Escherichia coli* (strain OP50; final optical density at 600 nm 1 a. u., $\sim 5 \times 10^8$ cells mL⁻¹). The total volume per well was 100 μ L, and the final pH installed in the phosphate-buffered saline test solutions was 7.4. After incubating the nematodes at 20 °C for 48 h in the dark, the surviving larvae were counted under a stereo microscope at 20 \times magnification. The resulting number of mobile nematode larvae was subtracted from the initially incubated number of larvae to calculate the percentage of immobile nematodes. The effective concentrations at 50% (EC50) were calculated using a numerically fitted standard log-logistic dose-response model (Levenberg-Marquardt iteration algorithm, Origin 2016, build 9.3.2.903, OriginLab Corporation, MA USA, Supplementary Fig. S3). The experiment comprised twelve biological replicates for each treatment and was repeated twice with comparable results.

Supplementary Material

Refer to Web version on PubMed Central for supplementary material.

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Data Availability

The datasets that support the findings of the current study are available in the Zenodo repository with the identifier doi:10.5281/zenodo.4131137 at <https://doi.org/10.5281/zenodo.4131137>. Additional data related to this study are available from M. El-Shetehy and F. Schwab upon reasonable request.

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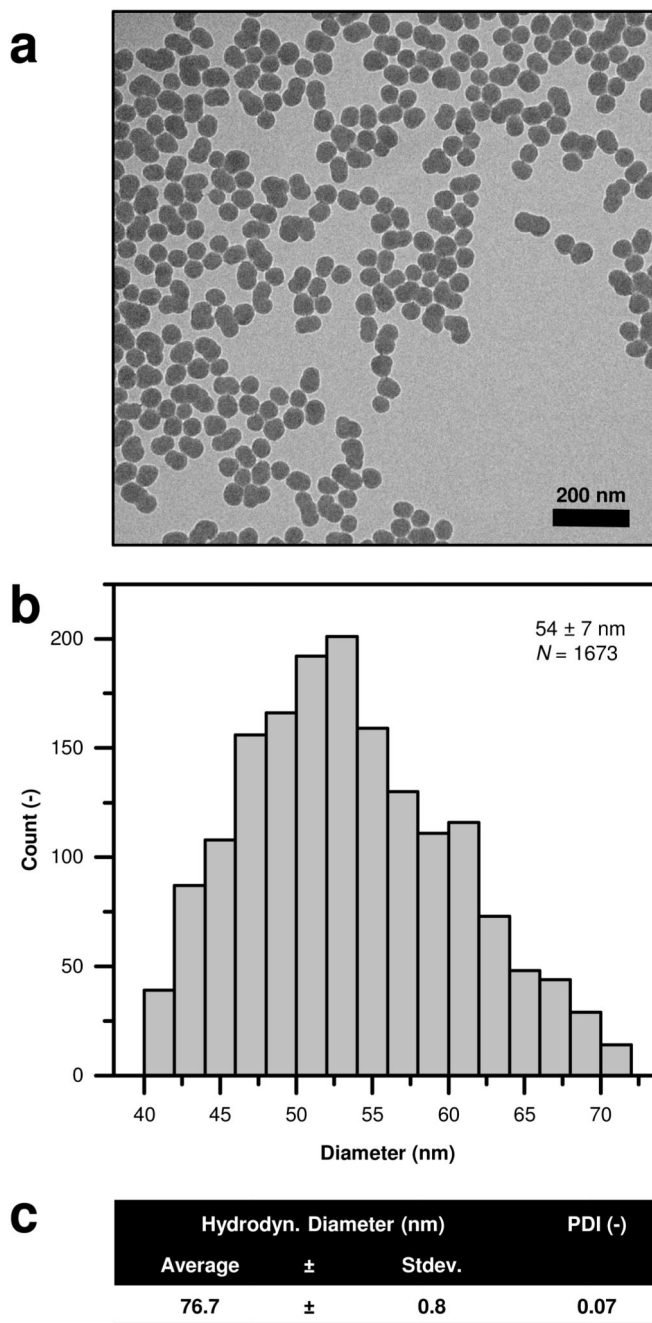


Figure 1. Silica nanoparticles (SiO₂-NPs) under investigation.

a Transmission electron microscopy (TEM) imaging showing the particles. **b** Particle size distribution based on TEM image analysis. **c** Dynamic light scattering (DLS) measurements of SiO₂-NPs. Hydrodynamic radius consistent with the primary particle size shown in **a** and **b**. PDI: Polydispersity index. Averages ± standard deviations. For DLS, $N=10$.

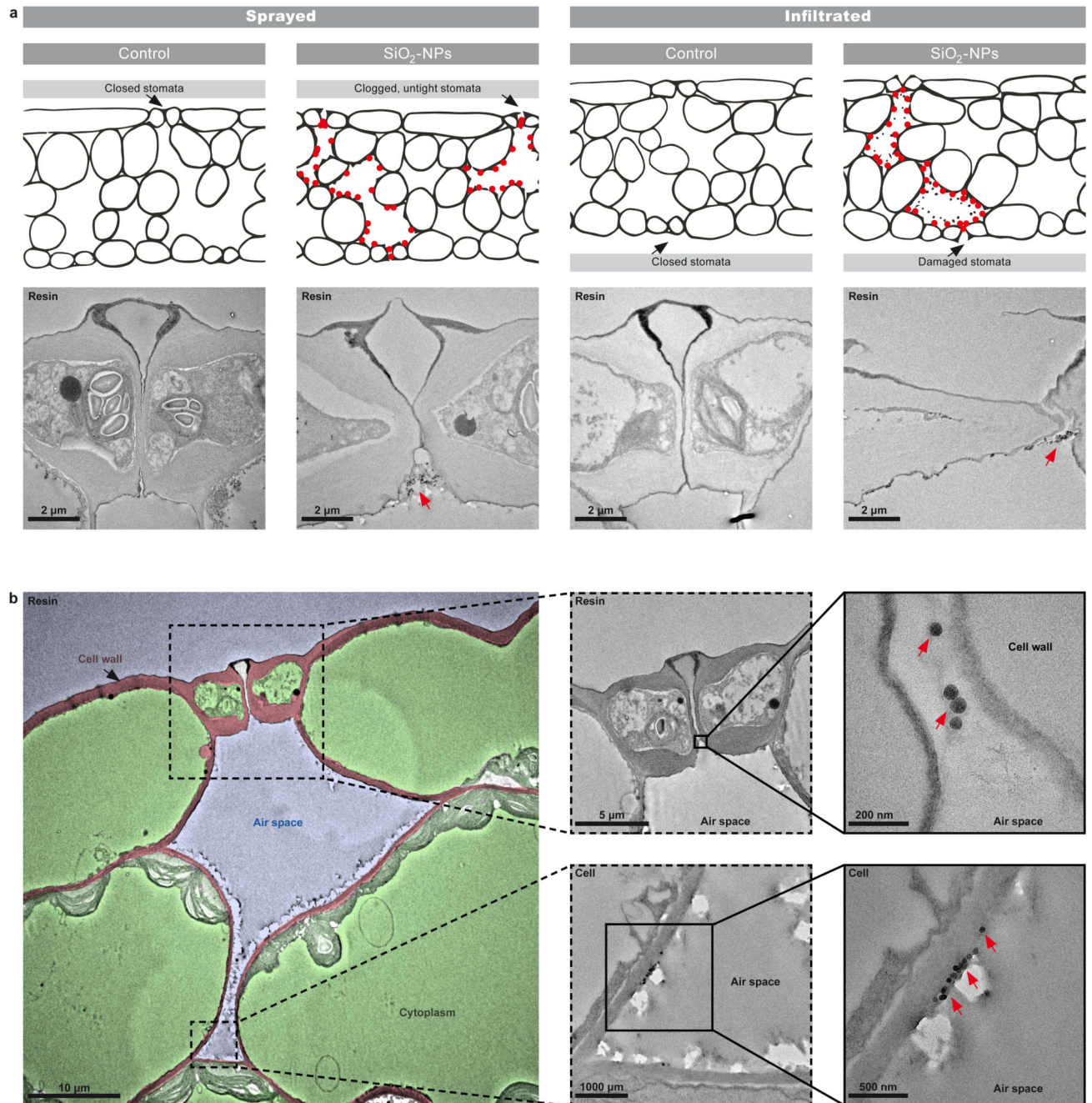


Figure 2. Transmission electron microscopy (TEM) of silica nanoparticle (SiO₂-NP) distribution and physiological effects in *Arabidopsis* leaves. Red arrows and dots: Nanoparticles. Comparison between spray application used in the field and for local defence assays, and infiltration application used in laboratory studies. Images obtained when plants already had developed resistance 2 d after exposure to SiO₂-NPs. **a** Control leaves treated with buffer solution only. **b** TEM overview image and zooms into the stoma and cell-air space interface. False colours mark, in red, the cell wall (apoplast); in green, the cytoplasm (symplast); and in blue, the spaces filled with air. SiO₂-NP-sprayed leaf

in higher resolution shows that the stomata are not tightly closed anymore due to nanoparticle uptake and clogging. Nanoparticles entered through the stomata into the leaf air spaces, were also found adsorbed extracellularly to the outer edge of the cell walls in the air gaps of the spongy mesophyll, and were absent in the cytoplasm (intracellular space). Higher-resolution TEM in Supplementary Fig. S1.

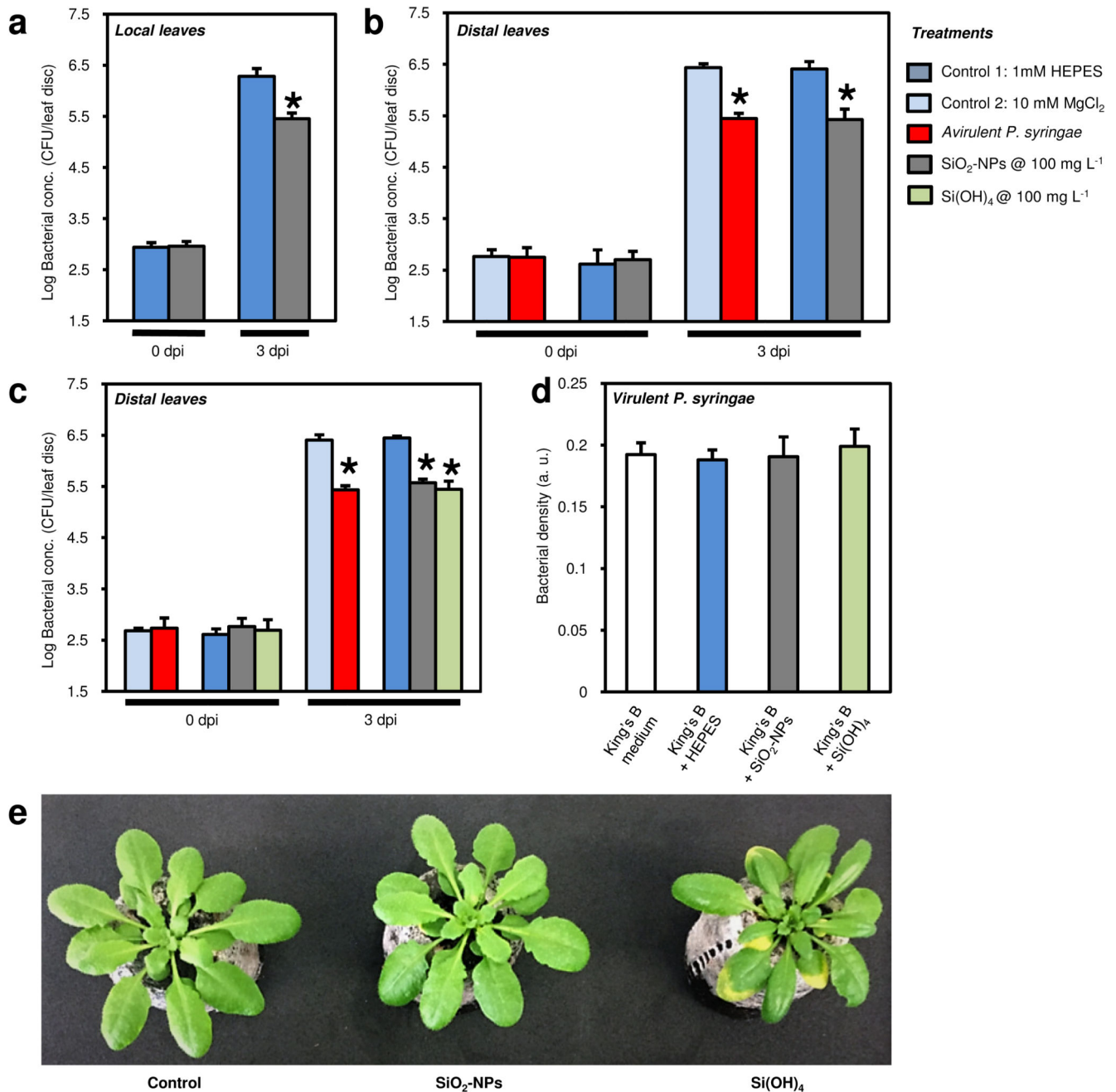


Figure 3. Enhanced local and systemic disease resistance in wild type Col-0 *Arabidopsis* to *Pseudomonas syringae* induced by silica nanoparticles (SiO₂-NPs) or orthosilicic acid Si(OH)₄. CFU: Colony-forming units. dpi: days *post* inoculation with virulent *P. syringae*. The bacteria in leaves were quantified 0 and 3 dpi. **a** Growth of virulent *P. syringae* in leaves. Plants were sprayed with different treatments, and virulent *P. syringae* was inoculated 24 h later. **b** Systemic acquired resistance (SAR) in distal leaves. Plants infiltrated locally with different treatments. 48 h later, virulent *P. syringae* was inoculated on untreated systemic leaves. **c** SAR in distal leaves, repetition of experimental setup in b with an additional Si(OH)₄ treatment. **d** No effect of SiO₂-NPs and Si(OH)₄ on *in vitro* growth of virulent *P.*

syringae bacteria in absence of plant. **e** Phenotype of *Arabidopsis* plants. Plants pre-treated with HEPES buffer (control), SiO₂-NPs, or Si(OH)₄ (1000 mg SiO₂ L⁻¹ each). Note the yellow leaves in the plant exposed to Si(OH)₄ coinciding with the upregulated expression of the oxidative stress marker gene in Fig. 4c. In a–d, all experiments were performed twice with comparable results. Bars and whiskers are averages and standard deviations, *N*=3, 1-way ANOVA; *post-hoc* LSD, *p*<0.01.

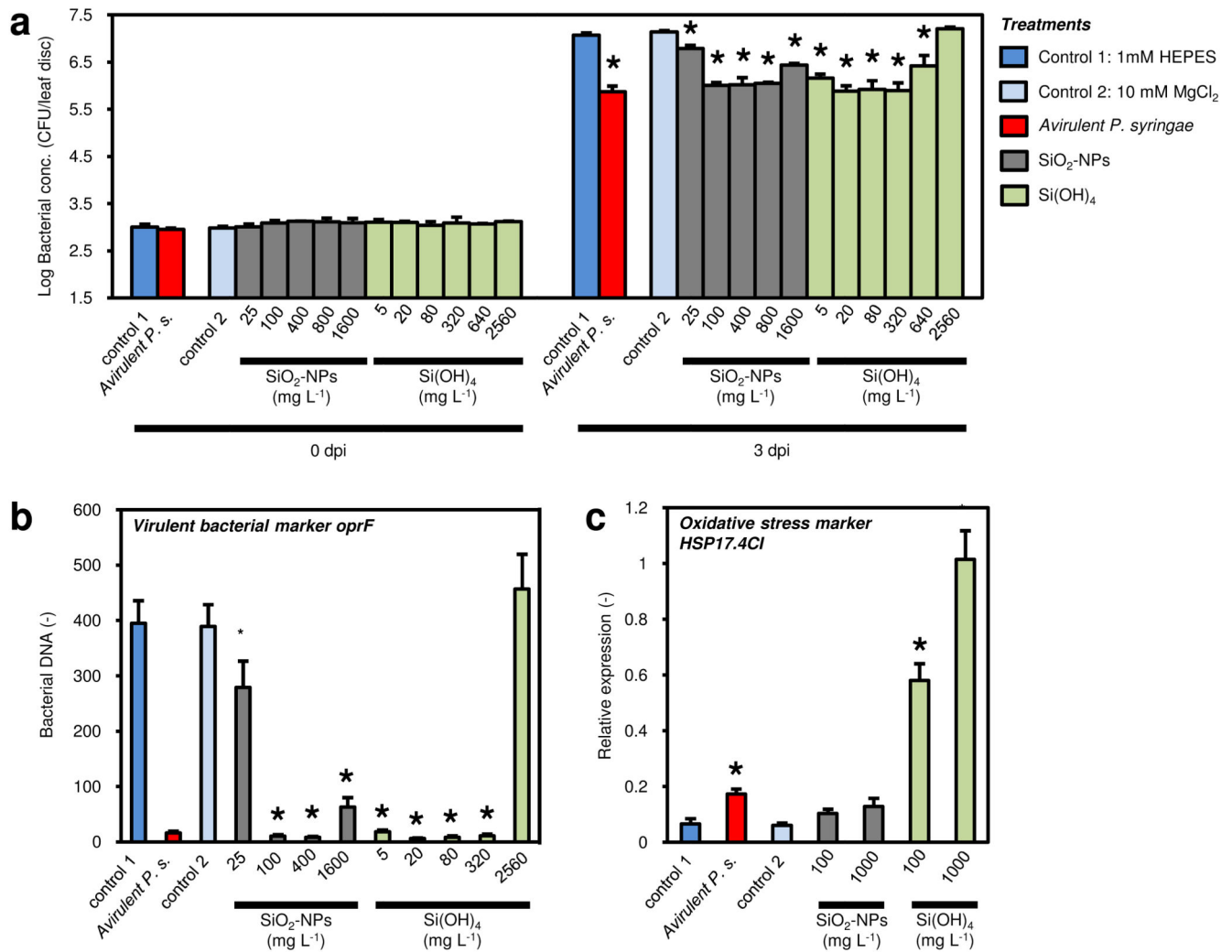


Figure 4. Silica nanoparticles confer systemic acquired resistance (SAR) in a dose-dependent manner.

Distal leaves of wild type Col-0 *Arabidopsis* treated with controls or silica nanoparticles (SiO₂-NPs) or orthosilicic acid Si(OH)₄. CFU: Colony-forming units. dpi: days *post* inoculation with virulent *Pseudomonas syringae*. **a** SAR in plants infiltrated locally with different treatments. 48 h later, virulent *P. syringae* was inoculated on untreated systemic leaves. The bacteria in leaves were quantified 0 and 3 dpi. **b** qPCR transcript levels of *oprF* gene from virulent *P. syringae* using DNA templates extracted from inoculated leaves. **c** RT-qPCR transcript levels of oxidative stress marker gene *AtHSP17.4CI* in response to different treatments. Plants were infiltrated locally with different treatments. Leaves sampled 48 h after treatments. Reference gene: At4g26410 (*expG*). Bars and whiskers are averages and standard deviations, $N=3$, 1-way ANOVA; *post-hoc* LSD, $p<0.05$. All experiments in a–c were performed twice with comparable results.

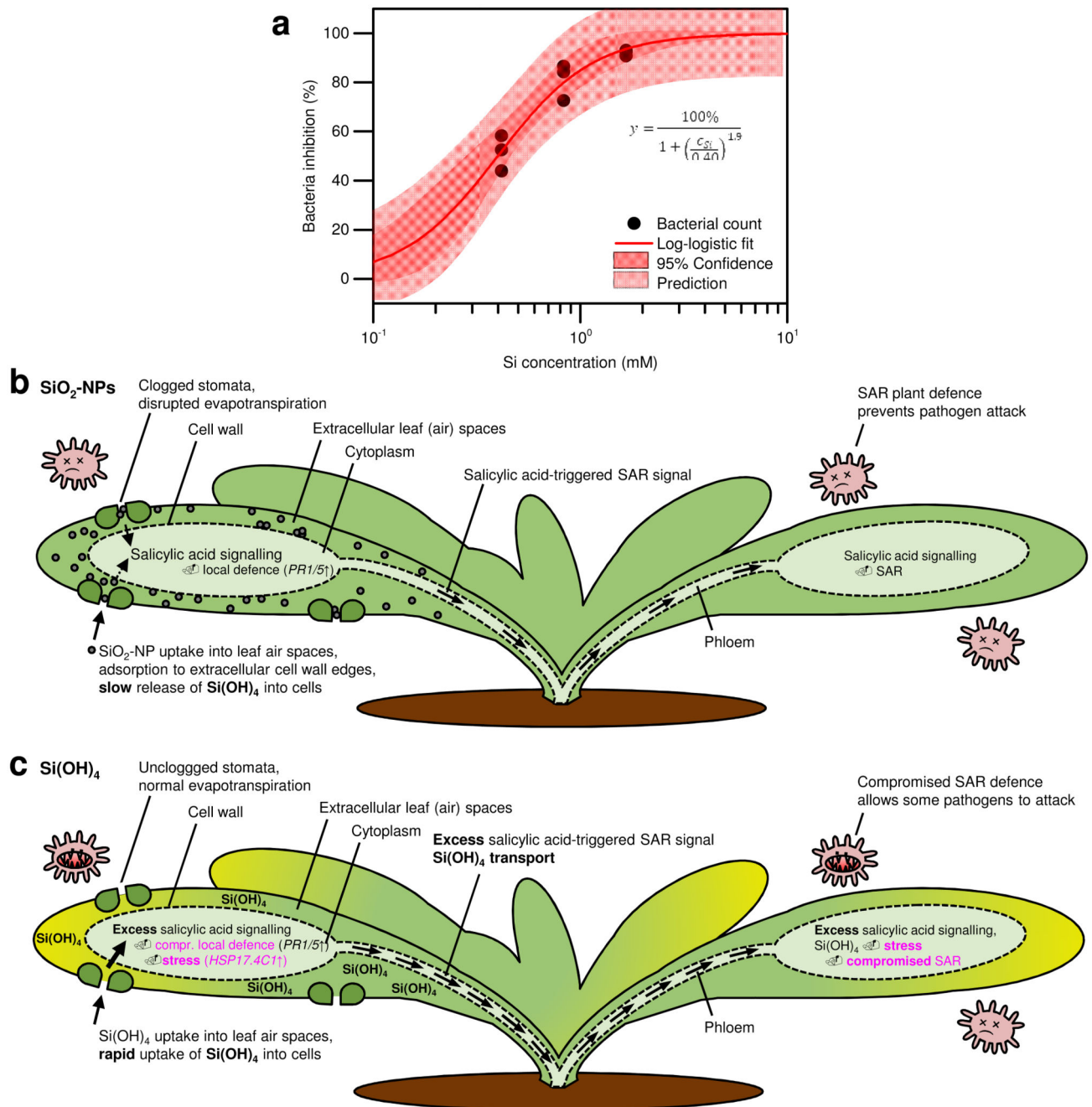


Figure 5. Dynamic range for systemic acquired resistance (SAR) induced in distal leaves by silica nanoparticles (SiO₂-NPs) in *Arabidopsis thaliana*, and model summarizing the observed plant defence-enhancing actions of SiO₂-NPs and orthosilicic acid Si(OH)₄.

a Data from Fig. 4a. SiO₂-NP-triggered dose-dependent bacterial inhibition 3 d after infection of wild type *A. thaliana* with virulent *Pseudomonas syringae*. The effective concentration at 50% bacterial inhibition (EC₅₀) was 0.40±0.04 mM Si for SiO₂-NPs (*i.e.* 24 mg SiO₂ L⁻¹). Above the dynamic range, bacterial infection can increase again (Fig. 4a). Six data points at 0 mM Si are not shown due to the nature of the log axis but apparent in the detailed residual analysis in Supplementary Fig. S2 and Supplementary Tab. S2. **b** SiO₂-NPs

act by i) slowly releasing Si(OH)_4 into cells, triggering SA, and thus local defence and SAR; and ii) clogging stomata, triggering SA and subsequent defences. Absence of intracellular nanoparticles confirmed by electron microscopy (Fig. 2; Supplementary Fig. S1). **c** Si(OH)_4 instantly diffuses into cells, triggering SA and subsequent local defence and SAR. However, the instant uptake causes overdose, stress, and compromised defences. Both mechanisms are shown after treatment with the same amount of SiO_2 equivalents ($1000 \text{ mg SiO}_2 \text{ L}^{-1}$). Salicylic acid (SA): plant hormone regulating SAR and *PR1/5* gene expression. *PR1/5*: Genes encoding Pathogenesis-related protein 1 and 5. *HSP17.4C1*: Heat shock protein and oxidative stress marker gene.

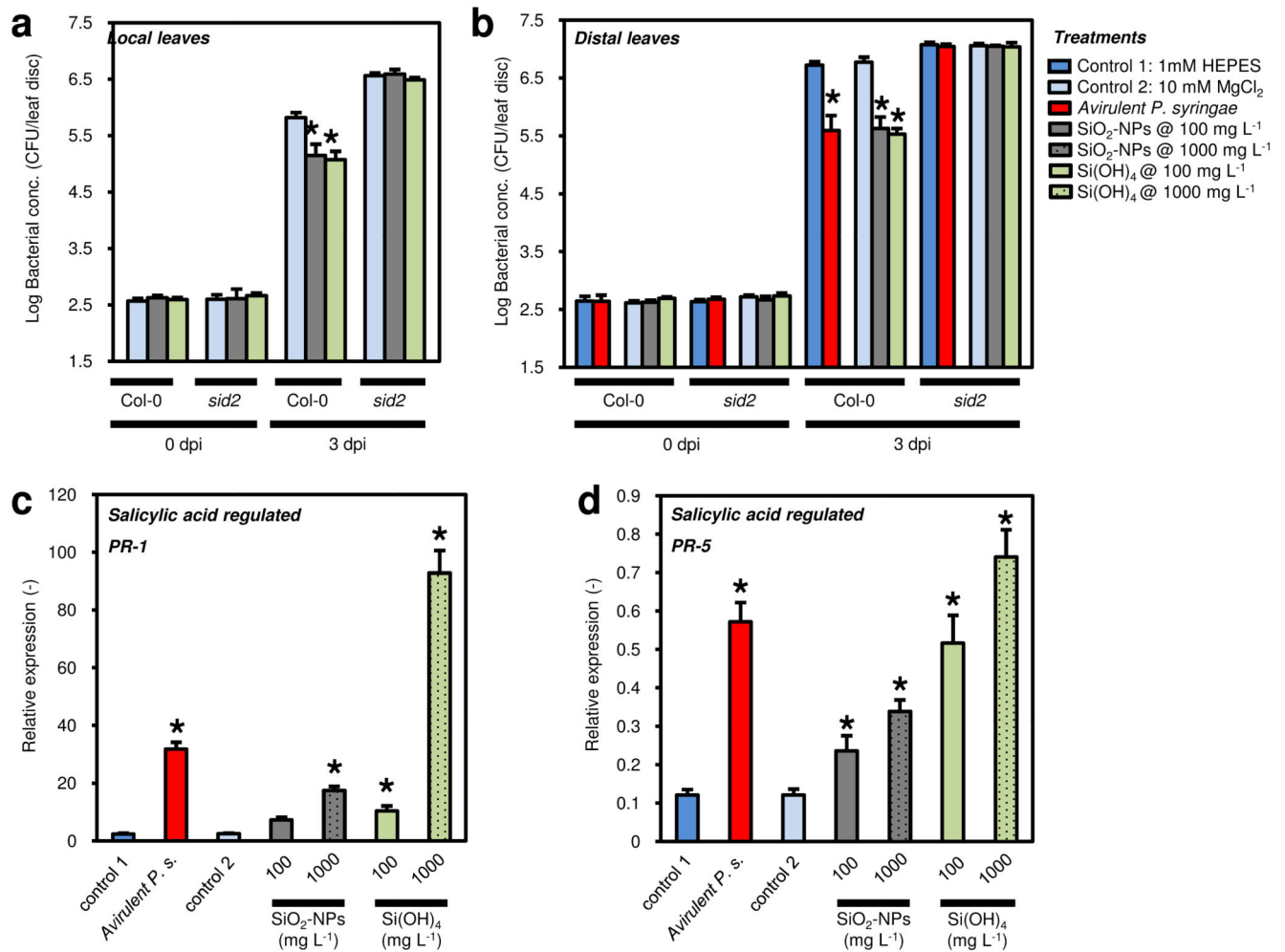


Figure 6. Silica nanoparticles (SiO₂-NPs) induce disease resistance based on salicylic acid dependent pathway.

Experiments in *Arabidopsis* wild type Col-0 and salicylic acid-deficient mutant *sid2*. CFU: Colony-forming units. dpi: days *post* inoculation with virulent *Pseudomonas syringae*. The bacteria in leaves were quantified 0 and 3 dpi. **a** *A. thaliana* wild type Col-0 and salicylic acid-deficient *A. thaliana* mutant *sid2* were infiltrated locally with different treatments. 24 h after these treatments, virulent *P. syringae* was inoculated. **b** Systemic acquired resistance (SAR) in distal leaves of Col-0 wild type and mutant *sid2*. Plants were infiltrated locally with different treatments. 48 h after these treatments, virulent *P. syringae* was inoculated. **c**, **d** RT-qPCR analysis of gene expression of the salicylic acid-regulated genes *AtPR-1* **c** and *AtPR-5* **d** in response to different local treatments of wild type *Arabidopsis*. Leaves sampled 48 h after treatments. Reference gene: At4g26410 (*expG*). Bars and whiskers are averages and standard deviations, $N=3$, 1-way ANOVA; *post-hoc* LSD, $p<0.02$. All experiments in **a-d** were performed twice with comparable results.