



Comprehensive Study of Si-Based Compounds in Selected Plants (*Pisum sativum* L., *Medicago sativa* L., *Triticum aestivum* L.)

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Abstract: This review describes the role of silicon (Si) in plants. Methods of silicon determination and speciation are also reported. The mechanisms of Si uptake by plants, silicon fractions in the soil, and the participation of flora and fauna in the Si cycle in terrestrial ecosystems have been overviewed. Plants of Fabaceae (especially *Pisum sativum* L. and *Medicago sativa* L.) and Poaceae (particularly *Triticum aestivum* L.) families with different Si accumulation capabilities were taken into consideration to describe the role of Si in the alleviation of the negative effects of biotic and abiotic stresses. The article focuses on sample preparation, which includes extraction methods and analytical techniques. The methods of isolation and the characterization of the Si-based biologically active compounds from plants have been overviewed. The antimicrobial properties and cytotoxic effects of known bioactive compounds obtained from pea, alfalfa, and wheat were also described.

Keywords: analytical methods; antimicrobial properties; bioactive compounds; cytotoxic effects

1. Introduction

Silicon (Si) is the second most plentiful and stable element present in the Earth's crust; it has a strong affinity for oxygen [1], and it is mainly present in the form of silicon dioxide (SiO₂), quartz, and silicates. Silicon plays an important role in soil; it increases soil exchange capacity, enhances water and air patterns, and participates in the metabolization of phosphorus-containing minerals and the formation of aluminosilicates and heavy metal silicates (decreasing soil toxicity) [2]. Additionally, silicon is a beneficial mineral for plants regarding their growth, development, and environmental conditions [3], as it enhances the bioavailability of many beneficial macro- and micronutrients for plants [4]. Silicon participates in regeneration and acts against the negative effects of stress on plants (biotic and abiotic stresses), but the mechanism of this action is still not fully understood [1].

Silicon, mainly in the form of silica, can be taken up by the plant's root system and transported along the plant's various compartments (e.g., the structure of the cell walls), providing greater rigidity and flexibility, as well as utility values for agriculture [5]. Plants uptake Si in the form of monosilicic acid (H_4SiO_4) [3,6]; the transport is mediated through passive (transpirational stream) and active (from roots to shoots, mediated by specific transporter proteins) modes [3].



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Copyright: © 2023 by the authors. Licensee MDPI, Basel, Switzerland. This article is an open access article distributed under the terms and conditions of the Creative Commons Attribution (CC BY) license (https:// creativecommons.org/licenses/by/ 4.0/). Plants produce a broad range of metabolites with diverse functions [7], such as alkaloids, steroids, tannins, glycosides, volatile oils, fixed oils, resins, phenols, and flavonoids, which are accumulated in different compartments of the cells [8]. The extraction of these bioactive compounds constitutes the first step in their characterization and utilization [7]. Si concentration in plants varies according to species, in a way that plants belonging to the Poaceae family and some species belonging to the family Fabaceae [9] show specific allocations of Si in all tissues (passive or active Si transport).

The main aim of extraction methods is to obtain the highest yield of the desired extract. Thus, the composition and concentration of the final extract will depend on the method selected. Some popular methods for extracting polar bioactive compounds from plant materials include maceration or percolation, microwave-assisted extraction (MAE) or solid-phase extraction (SPE), ultrasound-assisted extraction (UAE), and supercritical fluid extraction (SFE) [10]. Further, chromatographic analyses (such as thin layer chromatography, high-performance column liquid chromatography, gas chromatography, and so on) play a very important role in the identification of plant metabolic components, owing to several advantages, such as the specificity and possible use of such assays in qualitative and quantitative analyses [11]. Since bioactive compounds occur in plant material consisting of multicomponent mixtures, their separation, extraction, purification, and combination of chromatographic techniques are essential [12].

This paper describes the action of silicon in three plants belonging to the Fabaceae (especially *Pisum sativum* L. and *Medicago sativa* L.) and Poaceae (particularly *Triticum aestivum* L.) families regarding their physiology, strength, and utility for phytoremediation activities. Moreover, this review covers current analytical techniques, instruments, and methodologies used to understand the influence of silicon in plant development. Finally, it focuses on the isolation and characterization of the bioactive compounds in plant material, antimicrobial properties, and cytotoxic effects.

2. Silicon in Soil

Si compounds in soil are usually present as silicon dioxide (SiO₂), which comprises about 50–70% of the soil mass [13,14]. Other sources of Si include calcium and magnesium silicates, silicate slag, dolomite, rock phosphate, and diatomite [15]. Si compounds exist in the liquid phase, adsorbed phase, and solid phase of soil. In the solid phase, Si compounds can be divided into amorphous, micro-, poorly crystalline, and crystalline forms (Figure 1). The largest fraction of Si in the solid phase is crystalline forms, which occur as primary and secondary silica materials and silicates. Crystalline forms are poorly soluble and include feldspars, micas, clay minerals, and quartz [16]. The second most abundant fraction in the solid phase is amorphous forms originating from plant residues and remains of microorganisms (biogenic) or litho/pedonic materials [2].

The content of amorphous forms of Si in soils varies and usually ranges from 1 to 30 mg g^{-1} on a total soil basis. This amount depends on the plant cover, climate regions, and parental material [17]. The final forms of Si in the solid phase are poorly crystalline and microcrystalline, which include allophane, chalcedony, imogolite, and secondary quartz. The Si form available for plants in the soil (the liquid phase) depends on the solubility of Si in the solid phase. The liquid phase is comprised of dissolved Si, such as orthosilicic acid (H₄SiO₄) and polysilicic acid, and the polymerized and complexed silicic acid in the soil solution (complexes with inorganic and organic compounds or organo-silicon compounds) [18]. H_4SiO_4 is dissolved in the soil solution and can be adsorbed to soil mineral particles, especially iron and aluminum oxides/hydroxide, such as goethite and hematite. The total content of H₄SiO₄ in the adsorbed phase increases with pH and varies depending on the type of soil [19]. In general, the silicic acid content in the upper 20 cm soil layer contains only an average of 0.1 to 1.6 kg Si per ha as H_4SiO_4 [20]. The concentration of H₄SiO₄ can increase in the soil solution through the fertilization of farmlands where soils are inherently low in soluble silicon [19]. Silicic acid is also used by prokaryotes and eukaryotes in biosilification. The evolution of organisms revealed their capacity to use Si to synthesize



siliceous structures. These organisms use H_4SiO_4 to synthesize hydrated amorphous silica, known as biogenic amorphous silica (bASi), in addition to other compounds [21].

Figure 1. Schematic overview of the silicon fraction in soil and participation of biota in the Si cycle of terrestrial ecosystems, according to [16] and [18] with some modifications.

Plant growth is dependent on the presence of various nutrients in the soil, which can be divided into essential, beneficial, and toxic elements [22]. Essential elements are necessary for plant growth, while toxic elements negatively affect plant development by disrupting various metabolic processes. Si, although present in plants, is not considered an essential element. Most plant species can complete their life cycle without it [23]. There is a lack of information on its physiological and biochemical role in plant biology [24]. However, various studies demonstrate that Si is beneficial for plant growth and development and in the mitigation of both abiotic and biotic stresses [14]. In addition, Si ions influence the increase in the availability of calcium, copper, manganese, molybdenum, phosphorus, sulfur, and zinc in the soil. Greger et al. [4] also concluded that chlorine and iron tended to increase, while potassium and magnesium were not much affected by Si.

3. Silicon Uptake by Plants

All plants grown in soil contain some amount of Si in their tissues. The Si concentration in plants ranges from 0.1% to 10% of dry weight. The concentration of Si in plant tissues depends on various factors, including the characteristics of Si uptake and transport. The Si uptake by plants varies between species and cultivars [25]. The Si content in plant tissues is high in monocotyledons, for example, in grasses (0.3–1.2% of dry weight) and very high in rice (up to 10%). Si taken up by plants is usually found in plant tissues as hydrogen-bound Si in organic complexes. The Si complex role is to impregnate the walls of the epidermis and xylem vessels, thus strengthening plant tissues and reducing water transpiration. It has also been demonstrated that Si associates with components of the cell wall (such as polysaccharides, lignins, and proteins) [26].

Silicon deficiency in soils can lead to negative impacts on intensive agricultural practices. The intensive cultivation of plants with high silicon accumulation (such as rice and sugar cane) in tropical areas significantly reduced soil plant-available Si (PAS). A silicon reservoir in soils can be maintained by phytoliths present in plants (amorphous SiO₂), which are an important Si pool in the biogeochemical Si cycle. The recycling of silicon-rich crop residues, such as straw, is used to replenish the Si pool in the soil [27]. Among crop residues, rice straws (39%) and husks (24%) are the most abundant groups contributing to the total global return of Si phytoliths from crops. However, the use of plant residues only partially returns the Si taken up by plants to the soil [28]. Exogenous Si application significantly increases the Si content in the soil. Si is introduced into soils by the application of Si fertilizers (e.g., basalt powder) and biochar, which can induce the formation of phytolites and phytolith-occluded (PhytOC) organic carbon in plants [29]. Organic carbon can be occluded in 0.2% to 5.8% during the formation of phytolites. Dying plants introduce PhytOC into the soil, which becomes a reservoir of Si in the soil [30]. Appropriate application of Si fertilizers improves the bioavailability of nitrogen and phosphorus. Additionally, the soil application of Si has a positive effect on the diversity of microbial communities in the soil and their activity [31].

It is thought that the only form of Si taken up by plants is H_4SiO_4 [25], and the mechanisms of its uptake differ among plant species [14]. H₄SiO₄ is taken up as an uncharged monomeric molecule by roots from a soil solution below pH 9.0, and subsequently, it is transported through one of the possible mechanisms (passive or active) for the Si uptake to higher plants [32]. Silicic acid taken up by plants is not very mobile in plant tissues and has no electric charge [33]. The uptake mechanism of Si requires specific transmembrane proteins identified as Si transporters. The protein channels are passive transporters and do not require energy to transport Si across a cell membrane. In some plants (such as rice, barley, corn, wheat, banana, and cucumber), the transport of H₄SiO₄ through the plasma membrane is dependent on energy [34]. The first identification of Si transporters in monocot plants took place in rice. It was demonstrated that the mechanism of the Si uptake is faster than that of water. There is a specific system in rice roots facilitating the uptake of Si. This system facilitates silicic acid transport across the plasma membrane [35]. Silicic acid transport takes place using the Lsi1 (Low silicon rice 1) transporter, which belongs to a subfamily of aquaporin-like (NIP3). Lsi1 is located in the membranes of the main and lateral roots but not in the root hairs. This means that root hairs do not participate in the Si uptake from the soil solution [36]. H₄SiO₄ is taken up through the symplastic or apoplastic pathway and transported by the xylem to the shoots of the plant. It polymerizes into silica gel due to a loss of water (transpiration) and is deposited in different cellular compartments. However, Si in the xylem sap is present in the form of H_4SiO_4 [37]. Plants accumulate Si in their tissues to varying concentrations. On this basis, three categories of plants were distinguished: accumulators (rice, wheat, lentils, sugarcane, spinach, ferns, mosses, conifers) that accumulate more than 1.5% Si content in their tissues, intermediate accumulators (cucumber, soybean, pumpkins, rose, marigold, chrysanthemums) having Si in the range of 0.5–1.5%, and non-accumulators (tomato, grapes, sunflower, gerbera, petunia, begonia) which have a Si content below 0.5% [32,38].

4. Silicon Role in Plant Stress Alleviation

Si treatment mitigates many abiotic and biotic stresses, including heavy metal toxicity, salt, and drought (Table 1). Silicon can act by mechanical and biochemical or molecular mechanisms to aid plant growth. Plants deposit silicon in the form of spontaneous cell wall silicification, directed cell wall silicification, and directed paramural silicification in silica cells. Si modulates cell wall properties (thickness, rigidity, permeability, etc.) to adapt to the prevailing external conditions (silicification of the trichome and epiderma causes a higher ability to survive drought conditions; in exodermis and endodermis, it binds heavy metals), but Si also indirectly influences phytohormones, and production and activity of antioxidant enzymes [39]. However, the mechanism or mechanisms of these processes are still under investigation. In this review, the effect of Si-mediated stress alleviation in wheat, pea, and alfalfa plants is discussed.

Stress	Example	Plant Species	Role of Si
Abiotic stress			
Drought	-	T. aestivum L.	Improves the water relations of wheat leaves, photosynthetic gas exchange, and carboxylase activities in field drought conditions.
Salinity	-	T. aestivum L.	Improves the defense by antioxidant enzymes
Salinity	-	M. sativa L.	Regulates the K ⁺ /Na ⁺ homeostasis and the leaf water balance, also improving antioxidant enzymes' performance in leaves.
Heavy metals	Cd _	P. sativum L.	Accumulation of glyoxalase; it enhances the uptake of macronutrients and micronutrients in shoots and roots
		M. sativa L.	Cd binding in both roots and shoots
Biotic stress			
Fungal, bacterial, virus, and herbivores diseases	Powdery mildew (Blumeria graminis)	T. aestivum L.	Causes a higher concentration of elicitors

Table 1. The role of Si in abiotic and abiotic stress mitigation in *T. aestivum* L., *Medicago sativa* L., *P. sativum* L. [39].

5. Si-Mediated Alleviation of Stress Effects in Wheat

Wheat (*T. aestivum*) is a Si accumulator [34]. The Poaceae family, including wheat, can deposit Si as silica bodies (opal phytoliths) within particular cells of the leaf epidermis. Blackman [40] described the formation of silica cells in the sheath epidermis of wheat. During leaf growth, silica is accumulated and precipitated in the cells, appearing first as a translucent peripheral mass and later as glassy silica bodies containing small vesicles, which probably represent some remnants of the disintegrated cytoplasm [40]. The silica cells are located below the epidermis but also in the epidermal appendices, such as bristles or prickle hairs, and the shape of silica bodies is determined by the shape of the silica cell [41]. An analysis of the composition of silica bodies showed that it was near to stoichiometric SiO₂ (41 w% silicon, 56 w% oxygen), and the SiO_{4/2} tetrahedrals were arranged preferentially in three-dimensional networks; a smaller proportion was in chains and layers. Moreover, the silica bodies with an overall amorphous structure contained crystalline precipitates, which could be indexed as α -quartz [41]. It was shown that the development of silicified trichomes in durum wheat depends primarily on the availability of Si in soil and is not affected by water stress. However, the accumulation of phytoliths over the veins provides better support to the leaf and ensures a better development of the whole plant [42]. Wheat plants can not only deposit Si in the aerial organs but also in the roots. Si impregnates the endodermis cell walls of the wheat seminal roots [43] and older parts of the root [44]. Si was detected in the large central metaxylem lumina in the basal zone of the root, in the smaller peripheral metaxylem and the immediately contiguous pericycle, and in the outer parenchyma cells bridging the small metaxylem vessels to the endodermal layer [44]. Casey et al. [45] identified two Si-containing species, mono- and disilicic acids (H₄SiO₄* and $(HO)_3Si(\mu-O)Si(OH)_3^*)$ in a ratio of approximately 7:1, but no organosilicate complexes in the xylem exudate of wheat plants. Furthermore, after transferring plants to the solution containing 0.02 mM silicic acid and using the silica enriched to 98.7 atom percentage in ²⁹Si, the Si concentration of the xylem exudate rises dramatically in a matter of minutes, indicating active mechanisms of Si transport across root cell membranes [45].

It was shown that different methods of Si application (soil treatment, foliar spray, and seed soaking) alleviated the Cd stress effects and significantly increased the wheat's growth, chlorophyll fluorescence, photosynthetic gas exchange, water use efficiency, membrane stability index, relative water content, and the Si content [46]. However, among the three methods, Si applied as soil addition was the best and most effective in alleviating the Cd stress effects. Shi et al. [47] showed that when wheat seedlings were supplied with Si

(through hydroponics), the treatment improved their biomass and photosynthesis but had little impact on the root morphology of plants grown under Cd stress. The application of Si nanoparticles made from sodium silicate (Na₂SiO₃) boosted the root morphology and physiology of wheat seedlings under Cd stress [48].

Moreover, amorphous silica (ASi) [49] and Si (K₂SiO₃) [50] decreased Cd concentrations in wheat shoots. The addition of Si (Na₂SiO₃) decreased Cd contents both in shoots and roots [51], and measurements of the Cd²⁺ flux showed that Si significantly inhibited the net Cd²⁺ influx in roots of wheat [47]. However, the study of Wu et al. [52] showed that Si did not accelerate Cd accumulation in cell walls or vacuoles. They concluded that Si decreased the Cd uptake by the increase in the root oxalate exudation as an avoidance mechanism. Greger et al. [50] showed that Si reduced Cd transport into the cytoplasm when Si was added both directly during the uptake measurements and to the growth medium. Silicate downregulated the expression of genes involved in the Cd uptake across the plasma membrane (i.e., LCT1) and the efflux of Cd into apoplast or vacuole from the cytosol (i.e., HMA2). Moreover, Si upregulated phytochelatin (PCS1) gene expression and enhanced the phytochelatin (PC) formation when Cd was present. The expression of iron and metal transporter genes (IRT1 and NRAMP1) was downregulated by Cd. However, the expression of IRT1 was upregulated by Si in root and shoot and NRAMP1 in roots only facilitating Fe transport in wheat [50].

Howladar et al. [46] showed that different Si application methods (soil treatment, foliar spray, and seed soaking) significantly reduced Cd²⁺ and increased the Si content. Si alleviated Cd toxicity in wheat seedlings by improving the antioxidant capacity and by decreasing the lipid peroxidation as a result of increased superoxide dismutase (SOD) and guaiacol peroxidase (POD) activity as well as decreased malondialdialdehyde (MDA) and hydrogen peroxide (H_2O_2) content in wheat leaves [47,51] and roots [53]. Howladar et al. [46] showed that the Si soil treatment, foliar spray, and seed soaking significantly alleviated the Cd stress in wheat and reduced the MDA content and electrolyte leakage, significantly increasing the content of proline and soluble sugars, antioxidative (POD, CAT, SOD) defense system activity, and the content of polyamines and their gene expression. Similarly, the incorporation of Si nanoparticles in a Cd-contaminated acidic nutrient solution increased the content of antioxidants and decreased the content of ROS in wheat roots compared with Cd single treatments in acidic pH [48]. The application of Si improved the water status of drought-stressed wheat plants [54]. The study of Pei et al. [55] also showed that the inclusion of Si in a culture solution maintained the leaf water potential of polyethylene glycol (PEG) stressed plants at the same level as that of the control plants. The addition of Si partially improved the growth of the shoot (but not the root) and increased the leaf chlorophyll concentrations of drought-stressed plants [55].

Moreover, compared with the non-Si treated wheat plants, the application of Si increased the activity of SOD, CAT, and glutathione reductase (GR), the fatty acid unsaturation of lipids, and the contents of photosynthetic pigments and soluble proteins, as well as total thiols, whereas the content of H_2O_2 , the activity of acid phospholipase (AP), and the oxidative stress of proteins were decreased by applying Si [54]. The activity of glycolate oxidase (GO), POD, and ascorbate peroxidase (APX) showed no significant difference between under-drought and drought-Si treatment [54]. At the same time, Pei et al. [55] showed that PEG stress-induced membrane lipid peroxidation, as well as the decrease in the glutathione concentration, could be partly alleviated by adding Si. These authors [55] suggested that under short-term, PEG-induced water stress conditions (1 week), antioxidant defense rather than osmotic adjustment contributed to the improved wheat growth using Si.

The ability of Si to mitigate salinity stress (NaCl application) was also investigated in wheat [56]. Alzahrani et al. [56] tested Si roles in improving salt, drought or Cd stress tolerance in wheat. They showed that under stress conditions, the Si supplementation conferred higher growth, gas exchange, tissue water and membranes stabilities, K⁺ content, and limited MDA and Na⁺ contents and electrolyte leakage relative to those obtained without Si. Compared to those results obtained without Si, enzyme (e.g., SOD, CAT, and PO) activity was improved by Si applications, which was linked to elevated contents of antioxidants and osmoprotectants (e.g., free proline, soluble sugars, ascorbic acid and glutathione). The level of 4 mM Si was most effective in mitigating the salt and drought stresses, while 6 mM Si level was most influential to alleviate the Cd stress [56].

Additionally, Silva et al. [57] showed that supplying Si to wheat plants could increase the resistance to leaf streak caused by *Xanthomonas translucens* pv. *undulosa*, possibly through an increase in tissue lignification and the participation of chitinases and peroxidases. Moreover, Si-treated wheat plants had a clear adverse effect on the greenbug, *Schizaphis graminum* (Rond.) development due to possible Si-mediated changes in the quality of phloem sap [58].

6. Si-Mediated Alleviation of Stress Effects in Pea Plants

Grasses accumulate greater amounts of silica than other species of flowering plants. Parry and Winslow [59] investigated the accumulation of Si in seedlings of peas (*P. sativum* L.). Species that do not accumulate Si have no leaf deposits of phytoliths. However, Si was observed in vascular bundles and epidermal cell walls, often in close proximity to stomatal complexes but mostly in the tendril tips [59].

The beneficial effect of Si on the yield and nutrient uptake of peas was shown under sandy soil conditions [60]. There are also several studies that investigate whether and how Si influences the alleviation of heavy metal toxicity in pea plants [61–63]. Supplementation with Si reduced Cd accumulation and enhanced the uptake of macronutrients and micronutrients in shoots and roots, which depressed Cd toxicity in pea plants [63]. Rahman et al. [62] showed that the addition of Si in Cd-stressed plants noticeably increased growth and development along with the improved total protein and membrane stability of Cd-stressed plants. Moreover, Si supplementation modulated the total chlorophyll, carotenoid, photosynthetic efficiency, photochemical quenching, leaf relative water content, and gas exchange parameters in pea plants [63]. Rahman et al. [62] showed that GSH1 (phytochelatin precursor) and MTA (metallothionein) transcripts were predominantly expressed in roots and strongly induced due to Si supplementation in Cd-stressed plants compared with Cd-free conditions. This suggests that these chelating agents may bind to Cd, leading to vacuolar Cd sequestration in roots.

Furthermore, pea Fe transporter (RIT1) showed downregulation in shoots when plants were treated with Si along with Cd compared with Cd-treatment conditions (Figures 2 and 3). In conclusion, Rahman et al. [62] suggested that the alleviation of Cd toxicity in pea plants might be associated with Cd sequestration in roots and reduced Cd translocation in shoots through the regulation of Fe transport. Additionally, the effect of Si on Cd-stressed plants showed the involvement of the antioxidant defense system. The activity of CAT, POD, SOD, and GR increased along with elevated S-metabolites (cysteine, methionine, glutathione), implicating the alleviation of Cd toxicity in peas [62]. Furthermore, Si supplementation significantly decreased the antioxidant levels: the accumulation of H_2O_2 , MDA content, electrolyte leakage, and methylglyoxal content [63]. It was also shown that the addition of silicon nanoparticles (SiNp) along with chromium (Cr (IV)) protects pea seedlings against Cr(VI) phytotoxicity by reducing Cr accumulation and oxidative stress and up-regulating the antioxidant defense system and nutrient elements [61]. Interestingly, Feng et al. [64] show that extracellular silica nanocoat formed by layer-by-layer (LBL) self-assembly confers aluminum resistance in root border cells of peas. Moreover, seed priming with Si also showed an effect on drought-stressed pea plants [65]. Si treatment improved the morphological, physio-biochemical, and yield characteristics of pea plants, such as chlorophyll a, chlorophyll b, and relative water content, regulated the up-regulation of antioxidant enzymes, increased seed yield, and decreased lipid peroxidation and reactive oxygen species, mainly superoxide and hydrogen peroxide, in drought-stressed pea plants [65]. Additionally, it is suggested that the Si application and accumulation within the plant may activate the host defenses (the increased activity of the enzymes chitinase and β -1,3-glucanase in

leaf extracts) and the subsequent resistance to the fungal pathogen (*Mycosphaerella pinodes*) in pea plants [66].



Figure 2. Transformations of Si in plant cells and mechanisms of alleviating stress caused by Cd. Adapted from Głowacka et al. [67] according to [5,68].



Figure 3. Diagram showing Cd uptake, transport, and accumulation along with selected Cd toxic effects on plants. Adapted from Głowacka et al. [67] according to [69,70].

7. Si-Mediated Alleviation of Stress Effects in Alfalfa Plants

The effect of Si application on growth, physiology [71] and the alleviation of stresses was investigated in alfalfa plants [72–75], Liu and Guo [71] showed that Si application (K_2 SiO₃) alone increased the alfalfa (*Medicago sativa* L.) forage biomass and the number of

branches. Moreover, the Si application reduced both the transpiration rate and stomatal conductance but had no effect on the photosynthetic rate. On the other hand, this promotive effect of the Si application on biomass and water use efficiency was regulated by the soil moisture conditions [71].

Kabir et al. [72] showed that the addition of Si in Cd-stressed plants caused a significant improvement in morpho-physiological features along with total protein and membrane stability. Furthermore, Si supplementation in Cd-stressed plants showed a significant decrease in the Cd and Fe concentrations in both roots and shoots compared with Cd-stressed plants, revealing that Si-mediated tolerance to Cd stress was associated with Cd inhibition in alfalfa. However, the mechanism of this process was to be established since no significant changes in the expression of two metal chelators (phytochelatin synthase and metallothionein) and PC accumulation after the Si application under Cd stress were observed. Interestingly, Kabir et al. [72] showed the Si-mediated alleviation of Cd toxicity in alfalfa limited the Fe uptake through the down-regulation of Fe acquisition mechanisms. Moreover, the increase of the activity of antioxidant enzymes (CAT, SOD, ascorbate peroxidase) and elevated methionine and proline content after the Si application may reduce H_2O_2 and provide defense against Cd stress in alfalfa [72].

The Si-mediated effect on salt stress tolerance involved the significantly increased plant biomass, nodules number, and N content in alfalfa plants [75]. Moreover, chlorophyll content, relative water content, predawn leaf water potential, water use efficiency, and relative water content increased after the Si application under salt stress [74,75]. Additionally, the concentration of Na⁺ was reduced by Si treatment with an increase in the K⁺ content [75] and increasing K⁺/Na⁺ radio to protect the leaves from Na⁺ toxicity [74]. The Si application induced the activity of antioxidant enzymes (SOD, CAT, POD) and decreased the MDA content [74]. Si treatment lowered the amounts of MDA and H₂O₂ and also reduced electrolyte leakage in salt-stressed alfalfa plants, but at the same time, increased total polyphenol, flavonoid, and carotenoid contents [75]. Moreover, compatible osmolytes (proline, glycine betaine, and soluble sugars) were found to have increased particularly after the Si treatment in comparison to Si-untreated plants. It is important to underline that alfalfa varieties reacted differently to the Si treatment [75].

Alkaline stress is one of the abiotic stresses limiting plant growth because of higher pH. Liu et al. [73] showed that the Si priming of alfalfa seedlings significantly alleviated the damage symptoms and increased biomass, chlorophyll content, photosynthesis, and water use efficiency. Moreover, the Si alleviated oxidative damage caused by alkaline stress (25 mM Na₂CO₃, pH 11.2) by decreasing membrane injury and MDA content and increasing POX and CAT activity in alfalfa leaves. In contrast with the Si effect on saline (120 mM NaCl) stressed plants, the Si priming significantly decreased the accumulation of protein and the proline content in alfalfa under alkaline stress [73]. Furthermore, they showed that Si-treated plants under alkaline stress accumulated more Na, Mg, Fe, Mn, and Zn in the roots but accumulated more K and less Na in the leaves [73].

8. Extraction Methods to Assess the Presence of Silicon in Plants

The presence of Si has beneficial effects on the crops such as sugar cane, rice, tomato, cucumber, and strawberry; hence, several procedures for the determination of the Si available in plants (e.g., amorphous silicon or total silicon in plant material) were developed [16]. The extraction methods were designed in view of providing a stable and reproducible approach, improving the extraction efficiency and selectivity. It consisted of the application of an anion to replace adsorbed Si, an effect that has been proven between the Si in the extract and the Si in the crop yield [16].

9. Sample Preparation and Si Concentration in Plant Material

Specific analytical techniques entail specific sample preparation protocols, considering their requirements [76], and silicon could be detected and quantified directly—in solid plant material (XRF, for example) or indirectly—after plant digestion into the aqueous phase

(ICP technique, in particular) (Table 2). Plant material should be dried and fragmented to obtain homogenous material [77]. Techniques that are based on the liquid sample introduction system (most of the ICP techniques) require the dissolution of a sample to the liquid phase while reducing the occurrence and effects of the matrix on the results of the analysis [78]. Both solubilization and reduction of the matrix are achieved by pyrolysis [79], ashing [80], and digestion techniques; they involve using an oxidant reagent to obtain complete dissolution [81]. Barros et al. [82] performed two steps of a microwave-assisted digestion procedure of plant material with diluted reagents, firstly in HNO₃ and secondly in NaOH, in order to reduce the matrix effect in the ICP-OES quantification. However, these techniques often require the use of toxic reagents that cause corrosion of the analytical equipment; therefore, procedures are being developed based on less toxic solvents [83]. Silicon can be dissolved by using a high concentration of alkali (Na or K-hydroxides) [84], acids (preferably HF) [85], and sodium carbonate solution [86].

Table 2. Plant species studied with Si extraction and main techniques of analysis.

Plant Species	Sample Preparation	Techniques	Reference
apple, corn, peach, pepper, watermelon, and bluegrass ground leaves	200 mg of plant tissue with 10 mL of 1 M HCl + 20 mL of 2.3 M HF for 15 h	ICP analysis	[87]
cherry laurel, potato, alfalfa, carnation sunflower, barley grain, grass, French bean, bokashi, oil palm leaf	Plant material at pyrolysis at 420 °C was used	ETV-ICP-OES	[88]
rice (O. sativa L.)	Plant tissue added to 2 mL of 30% H ₂ O ₂ and 4 mL of 50% NaOH in the oven (95 °C)	OID MBC using a UV visible - spectrophotometer	[89–91]
wheat (<i>T. aestivum</i> L.)	for 4 h. After, was added 1 mL of 5 mM ammonium fluoride (NH ₄ F)		[90,91]
plant material (29 samples)	30 mg of pulverized plant material was incubated for 4 h in 0.1 M Na ₂ CO ₃ at 80 °C. Afterward, 10 mL of the extract was filtered	NIRS	[86]
sugar maple (<i>Acer saccharum</i> Marsh.), American beech (<i>Fagus grandifolia</i> Erhr.) yellow birch (<i>Betula alleghaniensis</i> Britt.)	100 mg of dried tissue was digested in 40 mL of 0.5 M NaOH at 85 °C for 4 h. 30 mg of dry tissue was digested in 40 mL of 0.1 M Na ₂ CO ₃ at 85 °C for 4 h	DSi colorimetrically, using the molybdate-blue method	[84,92]
C. epigejos and P. australis	First digestion: 100 mg plant material digested in a mixture of 4 mL distilled water, 5 mL nitric acid (65%) and 1 mL hydrofluoric acid (40%) at 190 °C. Second digestion: to neutralize the hydrofluoric acid, was used 10 mL of a 4% boric acid solution at 150 °C	ICP-OES	[81,92]
Dillenia suffruticosa, Dipterocarpus globosus, Macaranga trachyphylla, Shorea ochracea	Plant material ashed at 450 °C. The ash is mixed with lithium meta-tetraborate at 1000 °C. The resulting bead was transferred into 10% nitric acid.	ICP-OES	[80,92]
D. caespitosa, Lolium perenne, T. aestivum	100 mg plant material homogenized to a powder; calibration was required.	XRF	[77,92]

ETV, electrothermal vaporization; OID, Oven-Induced Digestion; MBC, molybdenum blue colorimetric; NIRS, Near Infrared Reflectance Spectroscopy; DSi, dissolved Si.

These procedures are dedicated to the elemental analysis of material; they are associated with the dissolution of silicon species to ionic form Si⁴⁺ (aqua, ligands). The crucial factor is the presence of fluoride anions in the solution due to their solubilizing (silicon creates a compound with fluoride more stable than oxygen) and stabilizing properties of silicon in the solution, especially at the trace concentration level [93]. Saito et al. [94] reported that extraction with a mixture of HF (1.5 M HF and 0.6 M HCl) could dissolve 150 mg SiO₂ within 1 h of extraction from 500 mg of a rice plant sample. The volume of HBF₄ to dissolve

100 mg of sample was 0.2 mL of 48% (v/v) [95]. Taber et al. [87] compare using the mixture of acid (HCl/HF) with autoclave-induced digestion (AID). The AID method may be useful owing to the rapid determination of Si in plants without using HF, but it is not suitable for crops with a low Si concentration. The comparison of HCl/HF extraction with AID for selected leaf reference materials indicated good agreement for corn stalks and bluegrass clippings but not for apple or peach leaves. He et al. [96] used a mixture of nitric acid and hydrogen peroxide to sample preparation for silicon concentration determination from the cell wall of a rice plant.

Nowadays, modern laboratories use methods of sample preparation based on microwave-assisted mineralization [97]. This technique is based on the synergy of three features: high pressure and temperature along with the action of concentrated acids, which ensure very good recovery, reproducible results, and rapid analyses without a greater risk of sample contamination. The mixture of nitric acid (a very good solubilizing agent for most elements, which has the oxidation ability towards organic matter), hydrofluoric acid (the reason for it being applied is described above), and the same volume of water in the digestion procedure is followed by the second digestion step: the neutralization of excess HF by using boric acid (due to highly corrosive and toxic properties of HF) [81,98]. Sometimes, adding an additional oxidant reagent is helpful [99,100]. Plant material is very rich in organic carbon, and it is necessary to enhance the oxidative power of the solvent to reach complete digestion. Cross-contamination should be avoided, especially with glass materials, etc. (containing silicon), when working with HF for a quantitative analysis of silicon. Silicon can be bound to aluminum species, which makes its extraction more complicated [101].

The ashing and melting of plant material with alkaline flux constitute another possible approach to preparing samples for silicon determination. Silicon dioxide is not a volatile compound; thus, there is no risk of silicon loss. Bowen et al. [102] carried out the ashing method on plant material at a maximal temperature of 500 °C for 10 h, and next, the ash was fused with anhydrous Na₂CO₃. Pan et al. [103] prepared samples of rice plants by ashing the material at 550 °C for 3 h and then dissolving the ash in diluted hydrogen fluoride.

Raid et al. [104] presented a method based on gravimetric determination of silicon in plants as SiO_2 . After the pretreatment, a plant sample was digested in a mixture of 5 HNO_3 :H₂SO₄:2 HClO₃ and undissolved parts were filtered away; then, the sample was ignited and ashed in a porcelain crucible. The residue is SiO_2 , determined by the mass measurement of the silicon content [104].

10. Extraction Methods to Assess the Presence of Plant Biologically Active Compounds

The initial stage is to carefully prepare samples in order to preserve the biomolecules synthetized by plants; fresh and dried samples (leaves, barks, roots, stems, fruits, and flowers) could be used, but considering that a smaller particle size increases the surface contact between samples and extraction solvents, it is important to powder the plant material; it can be previously air-dried, microwave-dried, oven-dried, or freeze-dried (lyophilization) [105]. Pre-treatments are expensive and have limited influence on the process of converting biomass into target compounds, which is why it is still necessary to improve their efficiency and reduce costs [106].

The purpose of extraction techniques is to separate soluble plant metabolites, so that the initial crude extracts contain a complex mixture of plant metabolites, such as alkaloids, glycosides, terpenoids, phenolics, and flavonoids (Table 3). Extraction is an important step in the analysis of plants, being necessary to extract components for further separation and characterization, using an adequate solvent, such as methanol, ethanol or ethyl-acetate, dichloromethane, or a mixture of dichloromethane/methanol, etc. [12]. Conventional extraction methods such as maceration and Soxhlet extraction are widely used to extract compounds from plants. However, they are usually non-selective, time-consuming, and sometimes they degrade heat-sensitive substances [107]. Other modern extraction techniques with novel extraction methods are the ultrasound-assisted extraction,

the microwave-assisted extraction, the enzyme-assisted extraction, and the supercritical liquid extraction, which possess certain advantages [103], considering eco-friendly, green technologies for plant extraction.

Plant	Extraction Technique	Solvent	Conditions	Compounds	Reference
Medicago sativa L.	SFE	supercritical carbon dioxide	10 mL of 96% ethanol solution for 3 h at 40, 60, and 80 °C in the dark, and it was incubated in water bath with stirring	phenolics and flavonoids	[10]
Medicago sativa L.	Maceration	ethanol	20% aqueous ethanol (v/v) overnight. After, anhydrous ethanol was added into the soak solution to obtain 75% ethanol solution, pretreated at 50 °C with stirring for 1 h. The extracting solution was filtered, and the filter residue was extracted twice more by following the same steps as above.	flavonoids	[108]
— T. aestivum L. —	HRE	glycerin:water	0.5 g of grounded spelt was subjected to extraction with 5 mL solvent under heat reflux extraction at 60 °C in a water bath for 4 h	phenolics	[7]
	MAE		0.5 g of grounded spelt was subjected to extraction with 5 mL solvent under microwave-assisted extraction for 1 min in a microwave oven (LG MS-197H) at output power 700 W for 30 s		
	UAE		0.5 g of grounded spelt was subjected to extraction with 5 mL solvent under ultrasound-assisted extraction for 30 min in an ultrasonic bath with frequency 50/60 Hz and power 310 W		
P. sativum L.	Maceration	acetone-water	700 mL/L aqueous acetone for 30 min attraction of (1:7 <i>w/v</i>) in a shaking incubator.	phenolics	[109]
P. sativum L. (pea flour)	SFE	supercritical carbon dioxide + ethanol extraction	22% ethanol, 86 °C, and 42.71 MPa, 40-min total extraction, including a 10-min static and a 30-min dynamic extraction at a flow rate of 2 mL/min	volatile compounds (1-hexanol, 1-octanol, 1-nonanol, nonanal, and 2-alkyl methoxypyrazines)	[110]

Table 3. Extraction techniques from Medicago sativa L., T. aestivum L., and P. sativum L.

SFE, supercritical fluid extraction; HRE, heat reflux extraction; MAE, microwave-assisted extraction; UAE, ultrasound-assisted extraction.

10.1. Maceration

Maceration is the simplest extraction technique in terms of the required equipment. The process can be carried out in an analytical beaker suited for thermolabile compounds. However, because the extraction is carried out at ambient temperature and atmospheric pressure, it usually leads to low efficiency, which is why the technique requires a higher volume of reagents and a longer extraction time [111].

10.2. Pre-Treatment

The pre-treatment consists of the defatting of plant material, which is essential for sample preparation. Material can be defatted with chloroform or a chloroform:hexane (1:1) (v/v) mixture in a Soxhlet apparatus for 48 h [112].

10.3. Extraction

The maceration technique is the most common and useful method to isolate saponins from *M. sativa* L. and from other plants [113]. The solvent selection is performed based on the solubility of the biologically active compound (solvent polarity) and the solvent evaporation temperature [114]. The defatted and dried material is immersed in 80% aqueous methanol under reflux for 1 h or three times at room temperature for 24 h [112]. Tava et al. [115] used 80% methanol to extract saponins from *M. arabica*. The highest content of sapogenins (compared with other extraction techniques SE, SFE) in the methanol extract was determined in the roots of *M. sativa*; however, the method has no specificity in relation to the extraction of biologically active compounds [116]. Khaledi et al. [117] used water and butyric acid-free bitter ethanol at a 30:70 ratio for the extraction of biologically active compounds, alkaloids, and saponins.

Chegini et al. [119] used maceration to obtain a biologically active compound from *M. sativa*. Golla et al. [120] extracted peptides from seeds and used salt precipitation with a phosphate buffer against freeze-dried seeds. Ethanol solution is also a good option for extracting proteins [121] and flavonoids [108]. Rodrigues et al. [122] performed maceration with 20 mL of 50% ethanol at 40 °C for 30 min to extract active ingredients from *Medicago* spp. leaves. Sangwan et al. [123] used extraction with ethanol for *T. aestivum* and Terminalia bellirica leaves. Hadrich et al. [124] carried out sequential maceration of biologically active compounds from the pea plant with three solvents with increased polarity: ethyl acetate, methanol, and water. Sundaresan et al. [125] carried out the extraction of wheat grass bioactive compounds by using different solvents—water, methanol, ethanol, hexane, and ethyl acetate. Kim et al. [126] used chloroform to extract quinone compounds from the wheat germ. Nair et al. [127] used the maceration approach to extract lecithin from peas (P. sativum) and lentils (Lens culinaris L.), jackfruit (Artocarpus heterophyllus Lam.) and jack bean (Canavalia ensiformis (L.) DC.), while ammonium sulfate was used to extract lectins. Narendhirakannan et al. [128] used the Soxhlet apparatus with ethanol as a solvent to extract active compounds from seven traditional medicinal plants easily accessible in India. The aqueous acetone solution was used to extract phenolic compounds [109]. Pure water is also a possible reagent for extracting phenolic species from the flowers of *M. sativa*; the average content of bioactive compound in the methanol extract was 263.5 ± 1.02 mg GAE/100 g of the dry-weight lyophilized extract [129].

10.4. Purification, Quantification

Purification is followed by quantification with RP-HPLC-UV to remove sugars and some phenolics. Bialy et al. [130] obtained extract from *Medicago sativa* L. containing glycosides and triterpene compounds from the extract of saponins by chromatographically treating the n-BuOH eluent saturated with water. The peptides were purified from the solution by precipitation of ammonium sulfate and separated on a 5 kDa membrane in the spin column [120]. Nair et al. [127] used dialysis against Tris, PBS, and NaCl gradient for the purification of isolated compounds. Troszynska et al. [109] applied column chromatography with the same mobile phase composition as an extraction reagent to the fractionation and purification of the extract obtained.

10.5. Derivatization

Extracted saponins can be derivatized to enhance their solubility [131]. Abbruscato et al. [132] obtained prosapogenins from part of the fractionated solution after alkaline and acid hydrolyses.

10.6. Accelerated Solvent Extraction (ASE)

ASE is based on using a solvent above its boiling temperature alongside higher pressure in the reagent cell, which enables higher extraction efficiency, lower reagent use, and shorter process time in comparison to maceration. However, the extraction is performed with expensive equipment. The sample pretreatment is similar to the maceration procedure, but some instrumental and physicochemical parameters (temperature, pressure) can be adjusted and optimized [133]. A smaller particle size, in most cases, leads to higher extraction efficacy [134]. However, the ASE extraction yielded similar results to maceration in the anthocyanin extraction from blue wheat, purple corn, and black rice conducted at optimal conditions [135]. Aqueous ethanol was most effective in extracting phenolic species from whole grain and bran of both soft and hard wheat [136]. What is more, aqueous methanol gives a higher extraction yield in comparison to hexane and acetone when phenolics were isolated from wheat brans [134]. The ASE extraction is more efficient in yield and retains the biological activity better in comparison to hot water maceration (determined according to oxygen radical absorbance capacity, DPPH, and hydroxyl radical scavenging ability) of polysaccharides extracted from Chimonobambusa quadrangularis [137]. The ASE extract from flowers of *M. sativa* (70% ethanol) was most efficient in the extraction of compounds regarding the total phenolic content and antioxidant activity [138]. By using ASE, the lipophilic vitamin K1 can be extracted with an n-heptane:ethyl acetate (4:1) solvent from the pea plant, in which the solid phase extraction was performed after purification from other lipophilic substances [139].

10.7. Supercritical Fluid Extraction (SFE)

Using a solvent in the supercritical state provides an opportunity to reach a high extraction yield; the fast experimental process gives the highest versatility for the extraction of biologically active compounds which differ in polarity [140]. The most common fluid is carbon dioxide which is non-polar in the supercritical state; by using a co-solvent (ethanol is the most common) with hydrophilic properties, selectivity of the extraction can be obtained [10]. For example, sequential extraction was performed in the selective isolation of nonpolar and polar compounds from triticale cultivars; in the first step, carbon dioxide allowed selective isolation of alkylresorcinols, while the second step involving ethanol as a cosolvent enriched polar lipids in the second fraction [141].

A smaller particle size leads to higher extraction efficiency [110]. According to Krakowska-Sieprawska [142], enzymatic degradation of alfalfa leaves enhances the solvent's penetration of the material and consequently ensures a higher extraction yield. Using the enzymatic degradation of *M. sativa* leaves resulted in an almost 50% higher yield (quantified by HPLC-MS polyphenols analysis) compared to a nondigested sample [143].

The highest efficiency in terms of the whole group of compounds can be achieved in higher pressure and temperature extraction conditions (300 bar and 80 °C, respectively), while higher specificity can be obtained at lower values of these parameters (200 bar and 40 °C) in the extraction of lupane-type triterpenoids from *Acacia dealbata* bark [144]. In contrast, the settings of 200 bar and 40 °C were sufficient to reach the maximal and individual (specificity) efficiency in the extraction of compounds from *Pterodon* spp. [145]. The highest extraction efficiency and the content of vouacapan diterpenes were reached at 220 bar and 40 °C, while the extract of *Ptedoron* spp. plant obtained at 160 bar and 50 °C had the highest antioxidant activity [146]. The settings composed of 270 bar and 42.6 °C resulted in the highest efficiency during the extraction of compounds from *Trigonella foenum-graecum* L., while lower temperature and pressure enabled the isolation of higher amounts of sterols, vitamin E, and vitamin D [147]. The diosgenin yield from the extraction of *T. foenum-graecum* L.

seeds depended particularly on the pressure and carbon dioxide flow [146]. Ge et al. [148] concluded that the extraction of vitamin E from wheat germ with SFE was potentially better than conventional techniques. SFE is popular for extracting biologically active compounds from wheat germ oil [149].

10.8. Microwave Assisted Extraction (MAE)

MAE is used in the extraction of biologically active compounds from plants, while the proper selection of temperature, power, and time settings is crucial to retain the biological function of the compounds. The following settings: 70 °C, 300 W, and 10 min in MAE, were the most effective in extracting anthocyanins from blue wheat and purple corn, compared with 50 °C, 1200 W, and 20 min for black rice [135]. A level of 90% of the nominal microwave power, 30 s irradiation time, and 1.5 g sample size were optimal to extract phenolic antioxidants from peanut skins with 30% aqueous ethanol in accordance with the total phenolic content parameter [150]. The MAE technique (methanol as a solvent) was more effective in comparison to maceration in an antioxidant extraction study on wheat bran, taking into consideration the total phenolic compounds, catechin equivalent, total tocopherol content, and Trolox equivalent [151].

11. Techniques Applied in the Determination of Silicon in Plants

11.1. Direct Techniques

11.1.1. XRF

Dissolution always carries a risk of using a hazardous reagent and a time-consuming utilization procedure; in particular, silicon (Si) may cause trouble by solubilizing from the complex matrix, as a result of which the analysis can be charged with low accuracy [152]. Thus, silicon can be analyzed in the plant material using direct techniques, which do not require silica solubilization. Such analytical techniques do not entail the decomposition of a solid sample (the signal from a sample in the solid phase is measured). An example is the XRF technique (X-ray fluorescence). The measurement of a sample consists of an irradiation process carried out by the primary X-rays in order to subsequently analyze the secondary X-ray spectrum emitted by the sample itself. It is known to be a fast, safe, nondestructive, and potentially more accurate analysis [77]. Reidinger et al. [77] used a portable spectrometer. They showed a detection limit of 140 mg Si/kg. The technique fulfills validation parameters, and a large number of samples (200 per day) can be tested with this technique. Guerra et al. [153] carried out experiments on the determination of the real-time concentrations of nutrient elements (K, Ca, S, Si) in plants, and the results (in g/kg sample) were close in value to the validated comparative method (ICP-OES).

However, the XRF determination revealed some inaccuracies in the air absorption phenomena and evinced a high drift of the spectrometer [154]. For light elements (lighter than 31 in mass units), the measurement has to be performed in a neutral gas environment [155]. Thorne et al. [154] overcame these drawbacks by performing an analysis of silicon in wheat in a helium atmosphere (reducing the first effect, adopted from [156]) and replicating the analysis after 1 h (reducing drift, adopted from [154,157]). Van der Ent et al. [155] used the helium environment and stated that silicon accumulated in the base of the trichomes. According to Deshmukh et al. [158], soybean plants accumulated 2.5 times more Si in leaf tissue in comparison with the control plants grown without Si supplementation. Baranowski et al. [159] performed a speciation analysis in soil and showed that the form of silicon depended on the type of soil and the origin of the samples.

11.1.2. Laser Ablation

Another technique for analysis in the solid phase is LA-ICP-MS. LA uses UV-nanosecond LA to generate an aerosol from the solid surface of a sample. The most critical step of sample preparation consists of the deposition of a sample in the cell [160]. Frick et al. [160] examined the initial digestion. Purification gives the best results in terms of reproducibility and repeatability of determinations of silicon in soil and plant extracts [160]. Fleck et al. [161]

prepared samples using a modified Steedman's wax protocol. In the root of rice, the determined silicon concentration was the highest in the outer cell layers comprising the exodermis and the sclerenchyma; it gradually decreased towards the central cylinder in the endodermis of the rice roots. It was revealed that the rice plant accumulated silicon, especially in the roots, and LA made it possible to determine the silicon concentration in the interior and exterior parts of the roots; the silicon concentration was significantly higher in the interior part of the roots than in the surface [162].

11.1.3. Laser-Induced Breakdown

A high-energy laser (Nd:YAG) beam is focused on a sample, which ablates a tiny part of the material, and plasma is generated. Tripathi et al. used laser-induced breakdown spectroscopy to determine silicon in different parts of the wheat plant. They revealed that leaves had the highest silicon content [163].

11.1.4. Electrothermal Vaporization

The ETV technique transfers elements from the solid phase of a sample to a mass or optical spectrometer (ETV-ICP-MS or ETV-ICP-OES). Masson et al. achieved the limits of detection at the content range of $30 \ \mu g \ Si/g$ in the plant material. Moreover, cellulose was found to be the ideal support for matrix matching of calibration curves [88].

11.1.5. EDX

SEM-EDX is based on the generation of characteristic X-rays in atoms of a specimen using the incident beam electrons. Soukup et al. [164] used the SEM-EDX technique to detect and measure the silicon content in plant material. In this study, silica was aggregated in the non-lignified parts of the endodermal cell walls [164].

11.2. Indirect Techniques

ICP-OES is the most popular analytical technique applied to determine the concentration of silicon in a solution of plant material. Owing to its popularity, availability of the equipment, and relatively low price, the technique is adequately sensitive and insensitive to matrix components, for example, compared to ICP-MS [165]. The silicon quantification is not burdened with spectral interferences. This is the most popular technique in plant research reported in the subject literature.

11.2.1. ICP-OES

ICP-OES is an analytical technique that allows the spontaneous emission of photons from atoms and ions that have been excited in a plasma torch device. Barros et al. [82] reported limits on detection and quantification in the range of 56 and 186 mg Si/kg for the optimized sample preparation protocol, and recoveries were in the range of 91.0–109% of the silicon determination in apple leaves, tomato leaves, white cabbage powder, bush branches, and leaves of CRM materials [82]. The silicon content in plant-certified reference materials is in the range of 0.2–30 g/kg [166]. Method detection limits in the range of 0.24 and 0.024 mg/kg were determined in the matrix solution—HF-H₃BO₃ and 2% HNO₃, respectively—and used to quantify Si in sugar cane leaves (mean Si content was $5522 \pm 1410 \text{ mg/kg}$) [167].

Ramírez-Olvera et al. [168] studied the Si concentration in rice plants under conventional and osmotic stress conditions. They showed a decrease in K and Mn in roots after the silicon supplementation and an increase in Fe and Zn in shoots. Quigley et al. [165] reported the limit of detection at r 10 μ g Si/L, and the recovery of the silicon determination was higher than 95% with respect to the certified reference material, which was hay. The validated method served to carry out Si determination in more than 800 plant samples, including forbs, legumes, and raminoids [165]. Guntzer et al. [83] tested the solvent for sample preparation and tiron for electrothermal vaporization; lithium metaborate fusion gave lower recoveries. Gu et al. [169] studied the influence of zinc and silicon on rice and stated that the ameliorative mechanism of silicate on excessive zinc toxicity to rice could be attributed to an internal plant response.

11.2.2. ICP-MS

ICP mass spectrometry (ICP-MS) is a powerful tool that makes an analysis of ultratrace level concentrations possible for most elements in the solution. In this technique, plasma is used to create single positive ions of elements in the gas phase; the ions are separated by their mass-to-charge ratio (m/z) in a mass analyzer. However, silicon determination encounters many difficulties. A single quadrupole mass analyzer, due to its resolution of about 0.8–0.7 atomic mass units, is not sufficient to distinguish the analyte from spectral interferences- ¹⁴N₂, ¹²C¹⁶O; analyzing the most common silicon isotope 28Si (92% abundance), strong in intensity, is difficult at a trace level concentration [170]. In comparison with other elements, the background of the 28 m/z line is definitely higher. Mihaylova et al. [171] recommended the cold plasma ionization condition (RF power of 600 W) for ²⁸Si quantification in the plant matrix. Additionally, sample matrix (the residue of plant material after digestion and solvents) enhances observed signals for low-mass elements in particular [172], therefore, appropriate calibration methods should be used. The popular approach based on the collision cell (kinetic energy discrimination one) applied to reduce spectral interferences in many cases is insufficient. The high background can be reduced by replacing glass equipment in the spectrometer. Aureli et al. [173] showed reducing m/z 28 lines by 20% when the glass equipment (nebulizer, spray chamber, torch) was replaced by Teflon devices. Additionally, they used the dynamic reaction cell to reduce spectral interferences and achieve the limits of detection lower than 1 μ g/g for most samples. The proper selection of instrumental parameters, such as power on the plasma coil, flows of plasma/collision gases, voltages on cells and lenses, etc., is crucial for obtaining as low a background concentration as possible without a significant loss of the analyte [174].

Using the dynamic reaction cell (to reduce polyatomic interferences generated in plasma) is a very promising solution to drawbacks present in the standard set-up of ICP-MS. Liu and Jiang [175] used ammonia as a collision gas and reported an improvement of one order of magnitude in the signal to background intensities during a silicon analysis in steel. Furthermore, the limit of detection was $0.2 \,\mu g \, \text{Si}/L$, and the accuracy and repeatability were less than 3% and 5%, respectively. He et al. [96] used this instrumental set-up in their study and showed that more silicon accumulated in the cell wall with hemicellulose and pectin than in the cell wall without pectin or without hemicellulose and pectin.

The lowest detection limit of Si and, more importantly, BEC can be obtained by applying the high-resolution HR ICP-MS technique. Klemens and Heumann [176] performed silicon determination with isotope dilution calibration and showed detection limits at 0.15 μ g/g for the HNO₃ matrix and 0.2 μ g/g when HF was also present in the matrix, while the repeatability was in the range of 2–4% for homogeneous biological and clinical samples.

11.2.3. Colorimetric Method

Historically, the standard method for colorimetric determination of silicon is based on the formation of the blue complex compound between silicon and molybdenum blue and the determination of the solution's absorbance on a UV-VIS spectrometer at 660 nm wavelength. This method can be applied to a plant matrix but will only identify rather high silicon concentrations due to its limited sensitivity. Researchers have noted the tendency towards a lower silicon content than in the control after the Cd stress of rice [98].

Pan et al. [103] showed the Si-defective mutant (Lsi1) of rice did not absorb silicon from the environment, but wild rice plants adsorb this element. Mitani and Ma [177] determined Si concentrations in rice tissue and reported its absorption of up to 10% of dry weight in the plant shoots (rice has a high ability to load Si into the xylem). Liang et al. [178] showed that both active and passive Si-uptake co-occurs in *O. sativa*, *Z. mays*, *H. annuus*, and *B. hispida*. Hodson et al. [179] collated and compared silicon concentrations in many

plants (data from 125 studies and species in relation to the Poales family) and showed that, in general, ferns, gymnosperms, and angiosperms accumulated less silicon in shoots than non-vascular plant species and horsetail.

11.2.4. Silicon Speciation

A coupled separation device (liquid chromatography equipment is the most popular) for detecting the sensitivity to silicon compounds enables research on silicon speciation. Chemical speciation deals with the identification and quantification of the distribution of elements in a sample [180]. Silicon can be present in many chemical forms, especially in plant tissues. Analysis of many silicon forms may be hampered by several analytical problems because of differences between the chemical and physical properties of these compounds, most significantly solubility, viscosity, and surface tension of dispersion; the same concentrations of Si in various forms of silicon did not induce the same detector's response by intensity (OES detection) [181]. The discrepancies among signals may have been caused by the solution transport process, thus, the knowledge of the sample matrix and matrix of standards to the silicon chemical form must be taken into consideration to guarantee reliable results [181]. Ebdon et al. [182] performed polar silicon speciation by reverse-phase HPLC with the ICP-OES axial and radial view in the 1990s and reported that BEC values were similar and were about 0.9 mg/L for the axial and radial analytical mode. The sensitivity of the axial mode was significantly higher for the water matrix of the solution but similar for organics. The limits of detection were 0.1 mg/L to 0.5 mg/L for inorganic Si and hexamethyldisiloxane, respectively [182]. Carter et al. [183] used the size exclusion and reverse phase mode of chromatography coupled with the sector field ICP-MS and showed the limits of detection for silicon compounds at 12–30 μ g/L and 0.1–4 μ g/L for the PDMS and silanol compounds, respectively.

NMR can be used to obtain information about chemical forms of silicon in plants. The identification is based on the number of peaks and their position in a scale (chemical shift) and comparison with the standards. Si was detected in the xylem of *Oryza sativa* L. in the chemical form of monosilicic acid [177]. Ma [184] expanded research in this area and demonstrated the same silicon form in the root, xylem, and leaf in low-silicon rice mutant (Lsi1) and wild-type rice. However, Park et al. [185] showed three silicon forms in four rice cultivars, namely the homologs Si(OSi)₄, Si(OH)₁(OSi)₃, and Si(OH)₂(OSi)₂, were detected in rice plants. Casey et al. [45] showed two forms: mono and disilicic acid, in a ratio of approximately 7:1, present in wheat (*T. aestivum* L.). Schaller et al. [186] detected tri- and tetra-silicic acid in common reed (*Phragmites australis*).

In addition, Cabrera et al. [187] studied the effect of lignin-silica complexing by characterizing lignin and silica coprecipitates with FTIR and solid-state NMR using soda lignins, such as protobind 1000 and 6000 (plant source: wheat straw), and organosolv lignin (plant source: mix of maple/birch/poplar), reporting interactions between lignin and silica through hydrogen bonds, and reaffirming that lignin induced the substitution pattern changes in the silica surface.

11.2.5. Organic Silicon Compounds in Plants

Direct evidence for the presence of organic derivatives of silicon compounds is difficult to find in the present literature. The adsorption of oleic acid onto silica gel taken from ashed rice hulls is an exergonic reaction with ΔG equal to -23.1 kJ/mol [188]. Furthermore, it should be noted that the derivatives of silicon compounds in plants resulting from covalent interactions (bonds) are not yet known or described. However, Kinrade et al. [189] showed that silicon could create a stable compound with polyols, wherein the compound is fiveor six-coordinated and ligands are specifically localized in space. The aqueous catechol solution incites the formation of a Si-catechol complex with a coordination number of six [190]. Kinrade et al. [189] provided evidence in the form of an NMR spectrum for the existence of gluconate derivate of a silicon compound in the dilute solution with a pH similar to the pH of the soil solution. Silicate cross-linking to the cell wall of xyloglucan has also been reported [191].

In accordance with Korndörfer et al. [192], who collected works about organo-silicon compounds in plants, Inanga et al. [193] analyzed the cell wall of rice plants subjected to the silicon solution and demonstrated using IR and UV spectrometry the possible presence of silicon bound to some organic compounds. According to Matychenkov et al. [194], organo-silicon compounds are classified as the soluble fraction of silicon, and these compounds were created by hydrogen bonding. In general, biomacromolecules are involved in the transport and polymerization process of silicon in the cell wall [195,196]. Ishii and Matsunaga [197] isolated an alcohol-insoluble fraction from rice seedlings. Subsequently, a water fraction was obtained by enzymatic degradation of the cell wall. In this water fraction, silicon was found through SEC-ICP-OES to occur as bound to high molecular weight compounds (probably polysaccharides) [197]. Moreover, a relevant role of peptides and proteins in the accumulation, transport, and metabolism of silicon compounds is revealed in the literature. Peptides and amino acids could form polysilicic species through interactions. Sahebi et al. [198] showed a key role of some proteins rich in serine and proline in the absorption and accumulation of silicon in the epidermal root cell walls. Kauss et al. [199] showed that specific proline-rich proteins could significantly enhance the silica deposition in the cell wall. Although several reasons for the presence of organic derivatives of silicon have been suggested, there is still a lack of data on the sequencing of individual compounds from plants using coupled techniques, for example, data from MSn-techniques.

11.2.6. Microscopy (SEM, TEM, AFM, Light Microscopy) Studies of Silicon in Plants

Silicon deposition can be observed by fluorescent microscopy techniques. Shimizu et al. [200] discussed the fluorescent dye of silicon deposition 2-(4-pyridyl)-5-((4-(2-dimethylaminoethylaminocarbamoyl)methoxy)-phenyl)oxazole (PDMPO), whereby silicon can be visualized at a concentration higher than 3.2 mM. Dabney et al. [201] developed a method to employ autofluorescence to study silica size and distribution in plants. The three types of silica bodies or silica body related to mineral structures were examined in both abaxial and adaxial epidermis of *K. macrantha* leaves. Sizes of these species, density, and distribution patterns varied.

Blecher et al. [202] compared three methods (SEM, light microscopy, and Raman imaging) to observe the silicon deposition in *Equisetum hyemale, Carex pendula*, and *Miscanthus sinensis*. They discussed the pros and cons of these solutions, although the combination of EDX and vCD (low-voltage-high-contrast) offered the best possibility for identifying the localization of silica insertions [202].

SEM imaging gives an opportunity to visualize silicon. Alvarez et al. [203] showed large amounts, in comparison to the control, of amorphous silica in the flag leaf blades of rice plants treated with soluble silicate and nanosilica with the comparable density of particles from these two sources. Głazowska et al. [204] showed that low silicon availability in the mutant defective in silicon uptake called low-silicon 1 Brachypodium distachyon promotes deposition of Si in the amorphous form or bound to cell wall polymers rather than as silicified structures. He et al. [96] showed, using AFM microscopy that silicon is covalently bound to hemicellulose. In roots, stegmata cells containing Si aggregates were positioned on the outer surface of the sclerenchyma bundles, while silicon was undetected in the epidermis, nor was it found in association with the cuticle [99]. Kido et al. [100] showed that Si was localized in the marginal areas of leaf blades of wild-type rice plants. Hodson and Sangster [205] showed that silicon was localized primarily in the endodermis associated with the inner tangential wall. Zexer and Elbaum [206] detected putative silica aggregation loci identified in the roots of Sorghum bicolor. An aggregate form of silicon was present in the roots of oat plants (268–366 nm in size), and nanoparticle silicon was also present in the cell walls of leaves [207]. The silica aggregates were lamellar in structure [164]. In contrast, Perry et al. [208] showed that the shape of silica deposited in the lemma of

Phalaris canariensis L. changed from sheet-like (days 0–11 post-emergence) and globular (days 12–32) to fibrillar (days 33–40) [208,209].

The coupling TEM or SEM with an EDX detector enables the detection of other elements in plant material, whereas the information about these elements is lost after the digestion of solid plant material [205]. For example, the preparation of a sample using digestion with HF excludes the detection of calcium in the sample due to the precipitation of calcium fluoride from the sample. Gu et al. [169] used SEM in a study on rice plants and showed that the highest Si concentration was around the endodermis; the second most abundant sites were around the exodermis and sclerenchyma in roots, while in leaf sheaths, the preferential localization of Si was around the epidermis and sclerenchyma. The impact of iron on the distribution of silicon in the roots and leaves of rice plants was examined. The results displayed the trend for silicon accumulation in the root epidermis and vascular cylinder, glandular trichomes, the epidermis, and the vascular cylinder of leaves of the rice plants grown in a medium with silicon supplementation and absence of iron [210].

11.2.7. Structural Analysis (IR and Raman) of Silicon in Plants

IR spectrometry is a technique to identify compounds owing to very specific absorption of chemical groups of compounds on IR radiation. The presence of silicon can be detected in the range of "the fingerprint region" in a spectrum. Soukup et al. [164] stated that only small differences were observed in the spectra between silicon-supplemented and non-supplemented plants. The differences were visible in regions of the C=O stretching band (1670–1760 cm⁻¹), to -OCH₃ and -OH groups conjugated to aromatic rings (1140–1190 cm⁻¹) and to phenolic compounds (1550–1670 cm⁻¹) [164]. Bokor et al. [99] showed a broad band in the region 400–490 cm⁻¹ and assigned the bond-rocking vibration to Si-O-Si, underlining the amorphous nature of silica. The broad and asymmetrical band around 800 cm⁻¹ visible in all three spectra was attributed to symmetric Si-O-Si stretching vibrations [99].

11.3. Analytical Techniques for the Separation and Isolation of Biologically Active Compounds

Depending on a plant species, Si source, Si amount, and the plant's ability to take up Si, the plant Si content can range from 0.1% to 10% (dry weight) [18,211]. The high Si deposition in plant tissues improves their strength and rigidity [14]. On the other hand, phytoliths are found in silica cells in the leaves of Si-accumulating species, such as those belonging to the Poaceae family (rice and wheat), and they vary with the species and age of a plant [212]. In addition, it was reported that most monocots, for example, plants of the Poaceae family such as *T. aestivum* (wheat), *Oryza sativa* (rice), *Lolium perenne* (ryegrass), *Hordeum vulgare* (barley), and *Zea mays* (maize), are high Si accumulators [212,213]; rice accumulates up to 10% Si in shoots [213]. Further, some dicots (Fabaceae family), for instance, fava bean (*Vicia faba*), are found to incorporate Si into their roots [211]. Si doses play a very important role in the accumulation of plant bioactive compounds and the antioxidant capacity, where positive effects of Si are closely related to the accumulation of Si in different plant tissues [214].

Once plant materials are extracted with different techniques, such as maceration, ASE, SFE, MAE, etc., the extracts are subjected to a separation technique. The chromatographic techniques most frequently applied in the phytochemical analysis are thin-layer chromatography (TLC), high-performance liquid chromatography (HPLC), and gas chromatography (GC); these analytical techniques could also be used in the identification and isolation of individual components from composite mixtures (preparative and micro-preparative scale) [11] (Figure 4).

11.3.1. Thin-Layer Chromatography (TLC)

Thin-layer chromatography (TLC), also called planar chromatography, is a technique for the qualitative and quantitative analysis of organic or inorganic compounds, isolation of individual compounds from multicomponent mixtures, and preparative-scale isolation; it is also a principal separation technique in plant chemistry research [11,215]. A classical thin layer has a thickness of 250 μ m, a particle size of 10–12 μ m, and allows an analysis of approx. 10 samples in 0.1 to 1000 μ L amount at a max. 15 cm development distance simultaneously; thus, the thickness of the layer (stationary phase) and the particle size distribution based on the properties of the analyte are important for the quality and efficiency of separation [215].



Figure 4. Determination and identification of Si-based biologically active compounds in *Medicago sativa* L., *T. aestivum* L., and *P. sativum* L.

For planar chromatography with high-performance layers, detection, and data acquisition, high-performance thin-layer chromatography (HPTLC) could be used [8], offering more rapid separation with thinner layers, smaller particles, and a narrower particle-size distribution [216]. HPLTC uses pre-coated plates coated with a sorbent of particle size 5–7 microns and a layer thickness of 150–200 microns [8]. Additionally, HPTLC appears to be more sensitive and reproducible than classical TLC [216].

After the biologically active compounds present in a sample are separated using preparative TLC or HPTLC, those compounds can be identified by employing GC-MS and NMR techniques [217], LC-MS [218], LA-ICP-MS [219,220], SEM-EDS, and ICP-OES [221].

The advantages of TLC are particularly important in research on plant extracts, which are very complex mixtures, as it allows the identification of known and unknown compounds, as well as the selection of biologically active compounds [11]. With a minimal sample, the sample preparation can be applied for the separation, quantification, and structural elucidation of secondary metabolites from crude plant extracts [216]. Often, such extracts contain polar (e.g., phenols and tannins) and nonpolar (lipids, chlorophylls, and waxes) ballasts, as well as a fraction of active substances very important for phytochemistry [11].

The study by Rodrigues et al. [222] provided evidence that *Oryza sativa* L. (rice–Poaceae family) produces several phenolic compounds along with phytoalexins, and the use of TLC and HPLC led to the observation that silicon enhances the accumulation of diterpenoid phytoalexins in rice, a potential mechanism for blast resistance. Oksana Sytar et al. [223] studied the content of anthocyanins (a subgroup of flavonoids) in grains and sprouts of *T. aestivum* L. (wheat), such as cyanidin 3-O-rutinoside, cyanidin 3-O-glucoside, cyanidin 3-O-galactoside, and cyanidin 3-O-galactopyranosyl-rhamnoside, using the HPTLC technique. These scholars reported that cyanidin was two- to three-fold more abundant in the grains than in the sprouts.

11.3.2. High Performance Liquid Chromatography (HPLC)

High-performance liquid chromatography (HPLC) is a popular, versatile, robust, and widely used technique for the analysis and isolation of bioactive natural products [11,12]. HPLC allows the separation (or partition) between a mobile phase (eluent) and a stationary phase (the column packing); hence, depending on the nature of the stationary phase, the separation process can be (i) based on repeated adsorption-desorption steps, (ii) based on the separation between the mobile and the stationary phase, (iii) the stationary phase is made up of an ionic surface of the opposite charge to that of the sample, and (iv) size exclusion chromatography, where the sample is separated according to the molecular size of its constituents [224]. The mobile phases generally employed are water/acetonitrile or water/methanol, which may be run in the gradient elution mode, or in the isocratic elution mode [11]. Natural products are frequently isolated following the evaluation of a relatively crude extract in order to characterize the biologically active entity, which is often present only as a minor component in the extract; the HPLC resolution is suitable for fast processing on an analytical and preparative scale of multicomponent samples [12]. For the phytochemical analyses, HPLC is used for (i) the isolation and purification of secondary metabolites from a complex mixture present in an extract or its fractions, (ii) metabolomic study or principal component analysis, and (iii) quantitative and qualitative analysis [225].

HPLC can be used with a different detector as HPLC-ICP-MS; this technique coupled to ICP-MS is the most important hyphenated technique for the multi-elemental speciation analysis of biological and environmental samples, which requires liquid samples [226]; and LC coupled with a UV–Vis detector is a powerful tool for the separation and identification of anthraquinones, flavonoids, carotenoids, anthocyanins, or indigoids [227].

On the other hand, ultra-performance liquid chromatography (UPLC) is a modern chromatographic system, which is simply the modification of HPLC [228] rooted in the same basic principle, where the main difference between the UPLC and HPLC is in the particle size of the sorbent of the column, namely UPLC is applied in assays of samples with smaller particles <2 μ m in diameter; it also operates at high pressure (6000–15,000 psi) and uses 1.7- μ m reverse-phase packing material, while conventional HPLCs use 3–5 μ m packing material and are set at different pressure (2000 and 4000 psi) [229]. Even more, to improve the UPLC efficiency, a high temperature is set to help reduce the viscosity of the mobile phase, and if the flow rate is high, mass transfer is increased by increasing the diffusivity of the analytes on the column, thereby reducing the back pressure significantly [230,231]. Thus, UPLC has better resolution, speed, and sensitivity compared with HPLC [228]. The application of UPLC in the phytochemical analysis is similar to that of HPLC, namely the identification and quantification of components in a complex mixture; it is the technique's principal application, i.e., the identification and detection of all possible metabolites in a mixture, in a crude extract or in fractions [225].

Vega et al. [231] explored the impact of silicon on phenolic compounds found in shoot extracts of *Hordeum vulgare* (barley–Poaceae family) under aluminum stress conditions. They discovered that silicon influenced the production of a particular group of flavonoids known as flavone-glucosides. With the help of HPLC, they identified several flavone-glucosides, including isoorientin-7-O-glucoside (lutonarin), apigenin-pentoxide-hexoside isomers 1 and 2, and isoorientin and isovitexin derivatives containing sinapoyl and feruloyl moieties.

In another study, Troszyńska et al. [109] analyzed the seed coat extracts of *Pisum sativum* (pea–Fabaceae family) to determine the presence of various phenolic acids, such as benzoic (p-OH benzoic acid, protocatechuic acid, vanillic acid) and cinnamic acids and their derivatives (*p*-coumaric acid trans, *p*-coumaric derivatives, ferulic derivatives). They also detected flavones (apigenin-7-glucoside, apigenin-8 C-glucoside, luteolin-7-glucoside) and flavonol glycosides (kaempferol-3-glucoside, quercetin-3-rutinoside, quercetin-3-ramnoside) through HPLC assays, emphasizing the potent antioxidant activity of the analyzed extracts.

Kabir et al. [72], on the other hand, investigated the role of silicon in alleviating cadmium toxicity in *Medicago sativa* L. (alfalfa–Fabaceae family). They analyzed plant

metabolites, including glutathione, cysteine, methionine, and proline, via HPLC. These metabolites are crucial in combatting oxidative stress induced by cadmium in alfalfa plants. Figure 5 illustrates the chemical structures of selected phenolic compounds, focusing on the flavonoids and phenolic acids described in the literature [72,109,232].



Figure 5. Chemical structures of selected phenolic compounds: (**a**) isoorientin-7-O-glucoside; (**b**) isoorientin 6^{''}-O-glucoside; (**c**) isoorientin 7-O-(6^{'''}-O-(E)-feruloyl)glucoside; (**d**) apigenin-7-O-beta-D-glucopyranoside; (**e**) 4-Hydroxybenzoic acid; (**f**) 3,4-dihydroxybenzoic acid; (**g**) vanillic acid; (**h**) 4-hydroxycinnamic acid; (**i**) ferulic acid; (**j**) apigenin 7-glucoside; (**k**) luteolin-7-glucoside; (**l**) kaempferol 7-O-glucoside; (**m**) quercetin-3-rutinoside; (**n**) quercetin-3-glucoside.

Researchers have suggested the presence of several potential links between silicon and phenol metabolism: (i) silicon could interact with the OH groups of phenols by condensing with Si(OH) in biological systems, (ii) silicon may be involved with lignin–carbohydrate

complexes present in cell walls, and (iii) silicon might play a role in signal transduction pathways responsible for inducing lignin production [193,233,234]. These potential mechanisms offer insight into the complex relationship between silicon and the production of secondary metabolites in plants. Silicon has been reported to enhance the production of secondary metabolites derived from the shikimate pathway, which is responsible for the synthesis of aromatic amino acids and phenolic compounds. Studies have suggested that silicon may indirectly modulate this pathway by increasing the activity of key enzymes, such as 3-deoxy-D-arabino-heptulosonate 7-phosphate (DAHP) synthase, shikimate kinase, and chorismate synthase. Enhanced production of secondary metabolites through the shikimate pathway has been associated with improved plant resistance to both biotic and abiotic stresses, as many of these compounds possess antioxidant and antimicrobial properties [5,14,235]. The malonate pathway plays a crucial role in the biosynthesis of various polyketide secondary metabolites, including flavonoids and stilbenes. Silicon may influence the malonate pathway by regulating the expression of genes encoding key enzymes, such as chalcone synthase and stilbene synthase. Additionally, silicon might modulate the activity of these enzymes through post-translational modifications or by affecting substrate availability. As a result, silicon could enhance the production of specific secondary metabolites that contribute to plant defense mechanisms and overall health [5,14,235]. The mevalonate pathway is responsible for the biosynthesis of isoprenoids, a diverse group of secondary metabolites that include terpenoids, carotenoids, and steroids. Silicon has been reported to stimulate the activity of key enzymes in this pathway, such as hydroxymethylglutaryl-CoA reductase (HMGR) and isopentenyl pyrophosphate (IPP) isomerase. The exact mechanism by which silicon affects these enzymes is not yet fully understood, but it may involve changes in gene expression, enzyme activity, or substrate availability. The enhanced production of isoprenoids can lead to increased stress tolerance in plants, as many of these compounds have antioxidant, allelopathic, and signaling properties [5,14,235]. It is worth highlighting the potential role of silicon in modulating the shikimate, malonate, and mevalonate pathways, leading to the enhanced production of secondary metabolites that contribute to plant defense and overall health. Although our understanding of the specific mechanisms by which silicon influences these pathways remains incomplete, our findings contribute to the growing body of evidence suggesting that silicon plays a crucial role in regulating a plant's secondary metabolism.

11.3.3. Gas Chromatography (GC)

Gas chromatography (GC) is used for assays of volatile and nonpolar (hydrophobe) compounds [11]. Gas chromatography is usually coupled with mass spectrometry (GC-MS), allowing the measurement of the molecular weight of a compound, and once a molecular ion has been identified, it is possible to measure this ion accurately to ascertain the exact number of hydrogens, carbons, oxygens, and any other elements that might be present in the molecule [11]. A conventional GC-MS is composed of an injection port, a detector, and a narrow, coiled column; the mobile phase is often carrier gas, such as helium, nitrogen, or argon; thus, a sample is injected, vaporized, and propelled into a column with the carrier gas [225]. For an analysis of secondary metabolites, volatile and nonpolar ones, GC-MS is a valuable analytical technique, e.g., in assays of stable and volatile alkaloids; nevertheless, flavonoids, isoflavonoids, and hydroxylated alkaloids are converted into derivatives for analysis with GC-MS [225].

Karpagasundar and Kulothungan [236] reported a study on *Physalis minima* (Solanaceae family), where 31 bioactive compounds detected in leaves were evaluated using the GC-MS technique; it was concluded that the prevailing compounds were heneicosanoic acid, bicyclo [4.1.0] hepta-2, 4-dien, octadecanoic acid (CAS), stearic acid, and octadeca-9, 12-dienoic acid. In addition, some studies on plants secondary metabolites analyzed using GC-MS were reported in *Origanum vulgare* L. (arvacrol, thymol, γ-terpinene and linalyl acetate), *Ziziphora tenuior* L. (monoterpenes: pulegone, p-menth-3-en-8-ol, isomenthone, and 8-hydroxymenthone), *Clerodendrum viscosum* (steroids, triterpenoids, alkaloids,

saponins, flavonoids, and tannins) and in *Zingiber roseum* (terpenoids: α -pinene, β -pinene, p-cymene, limonene, β -cubebene, transcaryophyllene, terpinen-4-ol, α -terpeneol, epi-cubebol, caryophyllene oxide, and verticiole) [225]. However, no identification studies of silica-based bioactive compounds in *Medicago sativa* L., *T. aestivum* L., and *P. sativum* L. have been reported to use the GC-MS technique.

12. Antimicrobial Properties of Medicago spp., P. sativum, and Triticum spp.

Medicinal plants, owing to the rich composition of biologically active compounds, are currently considered to be a promising source of substances with a broad range of antiseptic activity. One such popular plant is *Medicago* spp., commonly known as lucerne or alfalfa, which synthesizes many substances possessing biological properties, such as isoflavones, saponins, coumarins, tannins, terpenoids, naphthoquinones, and alkaloids [112,113,237]. Many researchers indicate the high antibacterial potential of extracts obtained from this plant [117,237]. Khaledi et al. [117] indicate that the extract obtained with the maceration of this plant inhibited the growth of *Enterococcus faecalis* at a concentration of 512 µg/mL and effectively killed this strain of bacteria at a concentration of 1024 µg/mL. The authors suggest that the content of phenolic compounds in the raw material may be responsible for this effect.

In addition, Chavan et al. [237] showed that the alfalfa leaf methanol extract exhibits the minimum inhibitory concentration (MIC) at 37 μ g/mL, 12.03 μ g/mL, and 111 μ g/mL against Escherichia coli, Pseudomonas aeruginosa, and Staphylococcus aureus, respectively. Avato et al. [131] investigated the antimicrobial properties of extracted saponins from roots and aerial parts of alfalfa. The results confirmed the ability of the plant extract to inhibit the development of Bacillus subtilis, Bacillus cereus, E. feacalis, S. aureus, and Saccharomyces cerevisiae. The authors conclude that the antimicrobial activity of M. sativa is related to the content of medicagenic acid. Chegini et al. [119] demonstrated the antibacterial effect of alfalfa root extract on the common bacteria in bronchitis and sinusitis. The results showed that the MIC value of the tested extract was 125 mg/mL against Haemophilus influenza, Streptococcus pneumoniae, and Moraxella catarrhalis. However, it was noted that the extract did not affect S. aureus. Rodrigues et al. [122] examined water-alcohol extracts obtained from seven *Medicago* species for their antimicrobial activity. The results of the research allowed the researchers to establish the lowest values of the minimum inhibitory concentrations of extracts at 31.3 µg/mL against K. pneumoniae, S. aureus, and *Staphylococcus epidermidis*, and 125 µg/mL against *E. coli*.

On the other hand, the extracts obtained from any of the studied *Medicago* species were unable to inhibit the growth of *P. aeruginosa* and *Candida albicans*. Antifungal properties of *M. sativa* were reported by Sadowska et al. [112]. They used *Medicago*-derived saponin fractions to counteract the development of *C. albicans*—one of the most common fungal pathogens of animals and humans. The research results showed that the tested phytochemicals exhibit a direct fungicidal/fungistatic activity. The saponin fraction of *M. sativa* proved to be able to inhibit *C. albicans* germ tube formation and eradicate the mature Candida biofilm. In turn, Abbruscato et al. [132] investigated the antifungal properties of saponin mixtures from alfalfa tops and roots against *Pyricularia oryzae*, a fungus that causes rice blight. They confirm that the best antifungal effects on different cultivars of rice are exhibited by a prosapogenin mixture from *M. sativa* tops. Saniewska et al. [238] compared the antifungal activity of saponins isolated from the roots of the plant evince better fungicidal properties. The saponins isolated from *M. sativa* roots at the concentration of 0.1% could totally inhibit the linear growth of *Phoma narcissi*.

Moreover, the linear growth of *Botrytis cinerea* was limited to about 74%, *Botrytis tulipae* to about 68%, *Alternaria zinniae* to about 67%, *Rhizoctonia solani* to about 74%, and *Phoma poolensis* to about 38%. It was revealed that saponins obtained from alfalfa could also be effective against phytoparasites. The activity of alfalfa against parasites has also been confirmed. D'Addabbo et al. [239] studied five different *Medicago* species to determine

their nematicidal properties. The application of saponin extract solutions at concentrations of 1000 µg/mL after 24 h of the treatment led to almost complete mortality of nematodes in the case of the three tested *Medicago* species. Therefore, it can be concluded that extracts obtained from *Medicago* spp. may represent an alternative to chemical plant protection products. Further, some alfalfa proteins can exhibit antibacterial properties. Another interesting plant with a wide spectrum of antiseptic properties is *P. sativum* (garden pea). Saeed and Tariq [240] investigated the antibacterial activity of juice obtained from seeds and skin of *P. sativum* in relation to 56 isolates belonging to 11 species of Gram-negative bacilli. The results demonstrate good antibacterial properties of both seeds and skin of *P. sativum* juices, with average inhibition zones of 16.30 ± 2.02 mm and 16.39 ± 3.16 mm, respectively. Hadrich et al. [124] investigated the antibacterial and antifungal properties of extracts obtained by the maceration of pea skin. The best properties were reported for the ethyl acetate extract, while the water extract did not exhibit antibacterial properties. The inhibition zone diameter ranged from 13.0 ± 1.0 mm for Aspergillus niger to 19.0 ± 1.0 mm for E. coli. Nair et al. [127] examined the antibacterial properties of lectin extracts obtained from *P. sativum*. They showed good properties (MIC value was 1 mg/mL) of the tested extracts in relation to S. aureus, B. subtilis, E. coli, and P. aeruginosa. However, the study did not show the killing effect of the examined extracts on the tested bacteria strains.

In turn, Wang et al. [241] investigated the antifungal properties of protein isolated from pea seeds with a molecular mass of 11 kDa and a lysine-rich N-terminal sequence. The isolated protein exhibits good antifungal properties against *Physalospora piricola*, Fusarium oxysporum, and Mycosphaerella arachidicola. Ye and Ng [242] isolated another protein from *P. sativum* with antifungal properties with a novel N-terminal sequence and a molecular mass of 31 kDa. The protein, designated pisumin, shows good antifungal properties against *Pleurotus ostreatus* and *Coprinus comatus* and much weaker action against R. solani and F. oxysporum. Another interesting study on the antimicrobial action of *P. sativum* compounds was conducted by Rehman and Khanum [243]. They isolated and characterized the antimicrobial peptides from the seed/pod of P. sativum against Micrococcus luteus, S. typhi, S. aureus, S. epidermidis, Klebsiella pneumoniae, E. coli, P. aeruginosa, Proteus vulgaris, and Pasterurella multocida. Observations of the inhibition zones and MIC revealed that the two active peptides from the pod, i.e., P7 (~10 kDa), P8 (~11 kDa), and from the seed, i.e., S4 (~19 kDa), S5 (~22 kDa) showed good antibacterial properties, and the most sensitive strain was S. aureus. The research on the antimicrobial properties of pea peptides was also conducted by Golla et al. [120], who showed that germinated seeds had the potential to accumulate peptides with antimicrobial activity. The peptides isolated from peas were effective against E. coli, P. aeruginosa, and S. aureus. Moreover, the extracts from *P. sativum* germinated seeds showed the highest inhibition activity among all extracts obtained in this study. The inhibition zone size obtained in this experiment was 8.58 ± 0.03 mm in relation to *E. coli*, 9.35 ± 0.05 mm in relation to *P. aeruginosa*, and 22.16 \pm 0.04 mm in relation to S. aureus. Erecevit and Kırbağ [244] investigated the antimicrobial properties of fatty acids, vitamins, and flavonoids extracted from peas against E. coli, B. megaterium, K. pneumoniae, S. aureus, C. glabrata, C. albicans, Epidermophyton sp., and *Trichophyton* sp. While the analyzed fatty acids show the best antimicrobial effects, the flavonoids did not inhibit the growth of any of the tested strains except *B. megaterium*.

T. aestivum, known as common wheat, also deserves attention owing to its rich composition of biologically active substances. As early as 1972, the antibacterial activity of peptides isolated from this plant against *Xanthomonas campestris*, *Pseudomonas solanacearum*, and *Corynebacterium michiganense* was confirmed [245]. Twenty-four years later, Caruso et al. [246] purified and characterized other proteins from the wheat kernel which exhibited antifungal activity against *B. cinerea*, *Fusarium culmorum*, and *Fusarium graminearum*. In 1988, another study demonstrated the antifungal potential of chitinases isolated from embryos of wheat [247]. Recent studies conducted by Narendhirakannan et al. [128] show that ethanolic and aqueous extracts of *T. aestivum* inhibit the growth of *E. coli*, *K. pneumoniae*, *E. faecalis*, *S. aureus*, and *P. aeruginosa* at a concentration level of 12.5 µg/mL. In addition,

Pagnussatt et al. [121] proved the antifungal activity of wheat crude protein extracts against *F. graminearum*.

Saha et al. [248] investigated the antimicrobial potential of the methanol and ethanol extracts obtained from leaf, seed, awn, stem, root, and whole plants of two wheat varieties (Kheri and Pavon76). The results indicated that the methanol and ethanol extracts of Pavon76 seed and awn, as well as the whole plant extract, resulted in the highest diameter of inhibition zones in relation to *E. coli* and *S. aureus* in comparison to Kheri extracts. With respect to the root extract, the methanol extract of both varieties turned out to be more effective in relation to the tested strains than the ethanol extract. The authors attribute the antibacterial properties of the tested extracts to the presence of phytochemicals in wheat.

Rajpurohit et al. [249] tested the antimicrobial activity of wheatgrass extract obtained with the cold extract method against some Gram-positive bacteria strains. The MIC values were found to be 1.25% of the extract against *Lactobacillus* spp. and 5% against *S. mutans*. Sundaresan et al. [125] evaluated the antimicrobial activity of wheat grass extracts obtained using five different solvents (water, hexane, methanol, ethanol, and ethyl acetate). The results reveal that all the tested extracts exhibit antibacterial potential against seven foodborne pathogens. Among the investigated extracts, the hexane extracts from seven-day-old wheat grass exhibit the highest antibacterial action, especially against Listeria monocytogenes and Yersinia enterocolitica. The study conducted by Sangwan et al. [123] assessed the antimicrobial effects of the simultaneous application of 10% T. aestivium and 5% Triticum bellirica extracts against two fungal and eight bacterial strains tested in the agar cup plate method. The best results were obtained for the aqueous extracts against M. luteus, E. coli, P. vulgaris, *P. aeruginosa*, and *A. niger*, where the inhibition zone diameter was in the range of 21–23 mm. Furthermore, the antimicrobial activity was also satisfactory with respect to the other strains (inhibition zone diameter -16-18 mm). Schalchli et al. [250] evaluated the allelopathic effect of root exudate extracts from wheat on Gaeumannomyces graminis var. tritici growth. The results established the MIC value for the most effective extract variant at 0.36 mg/mL.

In turn, Amber et al. [118] investigated the antimicrobial potential of phytochemicals (alkaloids, flavonoids, and saponins) and crude methanolic extracts of wheat against S. aureus, K. pneumonia, and E. coli. The MIC values for all the tested strains ranged between 25–50 mg/mL. Sharma et al. [251] determined the antiseptic potential of anthocyanins extracted from three colored wheat varieties of *T. aestivum* viz., as well as uncolored white wheat. The highest antimicrobial activity was noted for the black flour extract, which has the highest content of anthocyanin. The determined minimum microbicidal concentration of this variant against E. coli, C. albicans, S. aureus, and P. aeruginosa was established at 200, 200, 100, and 150 mg/mL, respectively. In addition, black wheatgrass juice extracts were also the most effective among all of the tested wheatgrass juice extracts and showed an MIC value in the range of 100–150 mg/mL in relation to all pathogens. Kim et al. [126] showed that the extract of 2,6-dimethoxy-1,4-benzoquinone included in the wheat germ strongly inhibits S. aureus and B. cereus growth. The Rajoria research team [252] undertook to identify the major bioactive compound that exhibited antimicrobial properties in various organic extracts of *T. aestivum* L. grass. All the extracts obtained revealed the qualitative presence of the most important phytochemicals, such as steroids and cardiac glycosides, tannins, alkaloids, flavonoids, and carbohydrates.

Moreover, the conducted chromatographic analyses revealed that the presence of bioactive compounds in the extracts, e.g., chlorogenic acid, rutin, chlorogenic acid, and tocopherol, were responsible for the maximum noted antimicrobial activity of wheat grass against *S. aureus*, *Salmonella typhi*, and *Vibrio cholerae*. However, growth inhibition was not observed in the case of *Flavobacterium* sp., *E. coli*, *P. stringii*, *B. subtilis*, and *S. faecalis*. Despite this, the authors conclude that the plant submitted to their research, containing many of the medicinally important bioactive compounds, has great potential to be used in medicine for the treatment of diseases caused by pathogenic bacteria.

Besides traditional techniques for identifying plant biologically active compounds with antimicrobial activity, more innovative tools are becoming available. One such interesting method is thin-layer chromatography-direct bioautography (TLC-DB). This technique combines the separation of biologically active compounds on an adsorbent layer with direct biological tests. For analysis, a TLC plate is immersed in the microbial suspension and incubated in appropriate conditions. Then, after the separation of the active compounds, the inhibition zones of microbial growth can be observed on the TLC plate. Therefore, rapid detection of antibacterial compounds in complex plant extracts is possible in both screening and semi-quantitative tests.

Moreover, it is not only the classic TLC technique but also high-performance thin-layer chromatography (HPTLC), planar electro-chromatography (PEC), and overpressured-layer chromatography (OPLC) that can be easily combined with bioautography [253]. After separation and the evaluation of antimicrobial properties, individual compounds can be directly identified on the plate using spectrometric methods. A technique often used for this purpose is matrix-assisted laser desorption/ionization mass spectrometry (MALDI MS). TLC/MALDI enables the analysis of complex organic mixtures [253,254]. Previous reports indicate that this approach is effective in the identification of active compounds of plant origin, such as lipids or flavonoids [255,256]. The use of "direct analysis in real-time" (DART) in conjunction with TLC also seems to be an interesting approach. This method employs the principles of Penning ionization, in which excited gas atoms interact with air molecules, resulting in the production of secondary ions able to desorb/ionize the compounds present on the surface of the tested plate [254]. The simultaneous application of TLC with derivatization, bioautography, and DART-MS was successfully applied for the analysis of plant-origin substances by Bañuelos-Hernández et al. [257].

Besides their well-known antibacterial properties, plant extracts from *Medicago* spp., *P. sativum*, and *Triticum* spp. can also counteract biofilm formation, making them an important area of focus in this chapter. Biofilms, which are complex communities of microorganisms embedded within a self-produced extracellular matrix, can be challenging to treat due to their increased resistance to antimicrobial agents. Consequently, the discovery of plant extracts' antibiofilm properties could lead to novel therapeutic approaches against biofilm-related infections.

Abouzeid et al. [258] explored the antibiofilm properties of alfalfa exudate and its identified components. The findings suggested that the primary active compounds influencing biofilm formation are canavanine and hyperoside. These compounds reduced the number of *E. coli* isolates with a moderate and strong biofilm-forming potential from 68.42% to 31.58% and 21.05%, respectively. Moreover, research conducted by Chamachar et al. [259] demonstrated that alfalfa extracts have the potential to inhibit E. coli biofilm formation by decreasing the expression of the rcsA and papC genes. Another group of active compounds with antibiofilm properties found in Medicago spp. are saponins, which possess significant antifungal properties. Sadowska et al. [112] showed that the saponin fraction of *M. sativa* was capable of inhibiting *Candida albicans* germ tube formation and eradicating mature Candida biofilms. In the case of Candida biofilm formation, Psd1 protein isolated from peas [260] and lectins isolated from wheat [261] also proved to be effective. It was demonstrated that lectins could inhibit Candida biofilm formation at a concentration of 50 µg/mL with an efficiency of up to 44%. González-Ortiz et al. [262] also investigated the effect of wheat on biofilm formation. In their study, a 0.5% concentration of soluble wheat-bran extract exhibited antibiofilm potential against Staphylococcus aureus.

13. The Effect of Medicago spp., P. sativum, and Triticum spp. on Eukaryotic Cells

The bioactive compounds present in plants can exhibit cytotoxic properties that have a dual nature in terms of their impact. While their cytotoxic effects on normal, healthy cells may pose limitations to potential applications of plant extracts, the same toxic effects on cancerous or abnormal cells can be highly beneficial. These cytotoxic properties can be harnessed and integrated into existing anti-cancer treatment strategies, potentially improving their effectiveness and offering alternative therapeutic options (Figure 6). This underscores the importance of studying and understanding the cytotoxicity of plant extracts



and their active components to maximize their potential in the development of novel anticancer treatments.

Figure 6. The effect of Medicago spp., P. sativum, and Triticum spp. on eukaryotic cells.

13.1. The Antitumor Activity of Medicago spp., P. sativum, and Triticum spp.

Many studies indicate that extracts and bioactive compounds present in *Medicago* exhibit cytotoxic effects, especially on tumor cells. Gatouillat et al. [263] proved that extract obtained from *M. sativa* leaf could induce apoptosis of sensitive and multidrug-resistant tumor cells. Furthermore, it was determined that terpene derivatives and flavonoids present in the extract were responsible for this effect. The cytotoxic properties were confirmed for chrysoeriol, tricin, (-)-medicarpin, (-)-melilotocarpan E, and millepurpan in relation to the mouse leukemia P388 cell line and its doxorubicin-resistant counterpart (P388/DOX). Further research confirmed that two isoflavonoids: millepurpan and medicarpin, isolated from alfalfa leaves, show a cytotoxic effect on these cell lines. The application of millepurpan and medicarpin leads to the apoptosis of both sensitive and resistant P388 cells. The estimated IC50 values were 54 μ M for P388 and 69 μ M for P388/DOX after the application of millepurpan and about 90 μ M for P388 as well as for P388/DOX cells after the application of medicarpin [264].

Another group of bioactive substances derived from *Medicago* plants that demonstrate antitumor properties are saponins. Research has revealed that mixtures of saponins from various alfalfa species can effectively inhibit the growth of breast cancer (MCF-7) and cervical cancer (HeLa) cell lines [265]. Furthermore, studies conducted on the same cancer cell lines have indicated that a trypsin inhibitor (a low molecular weight protein) sourced from *Medicago* plants could also diminish their clonogenic survival [266]. This evidence underscores the potential value of these biologically active compounds in the development of novel therapeutic strategies for combating different types of cancer.

The active compounds present in *P. sativum* may also exhibit cytotoxic properties. Studies conducted by El-Feky et al. [267] show that the ethyl acetate fraction obtained from peels of *P. sativum* shows a high cytotoxic activity in relation to human breast carcinoma MCF-7 cell line at 73.6% and low cytotoxic action in relation to the colon carcinoma HCT-

116 cell line at 21% efficacy. On the other hand, n-hexane extract activity was established at 26.1% and 28.3% for the MCF-7 and HCT-116 cell lines, respectively. More detailed studies allowed researchers to determine the IC50 value of the most effective extracts, which was calculated as 73.4 on the MCF-7 cell line. Furthermore, the antitumor activity of compounds isolated from peas was determined. The most effective compound in relation to the MCF-7 cell line turned out to be apigenin. In addition, this compound was not toxic to normal human skin cell lines. Quercetin turned out to be the least active in relation to the HCT-116 cell line. In another study, it was found that aqueous extracts from the seed coat of dark-colored varieties of *P. sativum* exerted concentration-dependent toxic abilities on human breast carcinoma MDA-MB-453, human colon adenocarcinoma LS174, human lung carcinoma A594, and myelogenous leukemia K562 malignant cell lines. Moreover, the correlation analysis demonstrated that the intensities of cytotoxic properties of the extracts. It turned out that the tested extracts were probably able to activate not only caspase-3-dependent apoptosis but also other cell death modalities [268].

The activity against HepG-2 cancer cell lines is also exerted by the sulfated alkaline soluble and insoluble extracts from pea peel. In addition, the sulfated acidic insoluble, neutral soluble, and alkaline soluble extracts from the pea peel exhibit activity in relation to MCF-7 cancer cell lines. Their effectiveness was 55.8% and 52.8% (HepG-2), as well as 73.4%, 72.4%, and 72.8% (MCF-7), respectively. For comparison, the effectiveness of Doxorubicin[®] was 52.6% and 72% for HepG-2 and MCF-7 cells, respectively [269]. It was also proved that the purified asparaginase extracted from peas exhibited a toxic effect on the L20B tumor cell line. A Neutral Red assay was used in this study, which is a cell survival/viability test based on the binding of dye by viable cells. The results indicated that the tested plant-derived enzyme had a significant cytotoxicity effect on tumor cells in the concentration range from 150 μ g/mL to 1.17 μ g/mL. Moreover, the inhibition of the growth of the L20B cell line was gradually increasing with the increase in the used asparaginase concentration [270]. Another promising compound derived from peas with anti-cancer properties are lectins. El-Aassar et al. [271] demonstrated the cytotoxic properties of lectins extracted and purified from *P. sativum* seeds in relation to hepatocellular carcinoma HepG2 cells. The application of the recommended lectin dose (5 mg/100 μ L) results in a decrease in the HepG2 cell proliferation rate by 60.77%. The cytotoxic effect of pea lectins on neoplastic cells was also confirmed by Patel [272]. Some protease inhibitors from peas can also exhibit anti-cancer properties. Clemente et al. [273] reported a significant and dose-dependent decrease in the proliferation of human colorectal adenocarcinoma HT29 cells after the application of the Bowman-Birk trypsin-chymotrypsin inhibitor from peas. The rTI1B had the highest cytotoxic activity in this study (IC50 = 46μ M).

Triticum spp. was also found as a potential source of anti-cancer substances. Prolamins obtained from *T. aestivum* spp. *spelta* were found to possess cytotoxic properties against human colon cancer Caco-2/TC7 cells and to agglutinate human myelogenous leukemia K562(S) cells [274]. Barisone et al. [211] investigated the anti-tumor properties of fermented extracts of wheat sprouts (FWGE) and the 10-200 kDa protein fraction (FWGP) obtained from this extract. The lymphomacidal activity of FWGE against NHL cell lines was determined by the evaluation of apoptosis and cell cycle. The determined IC50 was 120, 250, and 275 µg/mL in relation to Jurkat, Ramos, and Raji cell lines, respectively. In addition, the fermented extract was much less toxic against normal human primary B cells $(IC50 = 582 \ \mu g/mL)$. However, the results showed that the IC50 value for FWGP was lower than in the case of FWGE. The protein fraction showed significant cytotoxicity to malignant NHL cell lines. Moreover, the cytotoxic action in relation to lung carcinoma (H1650, A549) and hepatic carcinoma (HepG2) cell lines was also shown. The protein extract was only ineffective against breast cancer (MCF-7) cells. Another study assessed the antiproliferative potential of chloroform extract from the dried shoots of *T. aestivum*. The tested extract at the concentration of 250 μ g/mL was able to inhibit the growth of 87.23% of the HepG2 human hepatocellular carcinoma cancer cell line [275]. The crude aqueous extract from

T. aestivum leaves was also tested against the HeLa cell line. The antiproliferative potential of the extract was assayed using the MTT method. The extract exhibited dose-dependent cytotoxic action against the cancer cell line, and, at the same time, it did not exhibit toxicity to the normal VERO cell line [276]. Later studies by the same author confirmed the cytotoxic effect of this extract against the colorectal carcinoma HCT-15 cell line. The application of the MTT method proved the antiproliferative properties of the tested extract, and the IC50 value was estimated at 258.8 μg/mL [277].

13.2. The Influence of Medicago spp., P. sativum, and Triticum spp. on Normal Eukaryotic Cells

The widespread application of plant extracts in medicine requires them to fulfill two fundamental criteria. First, they must exhibit cytotoxic properties against cancer cells, and second, they should not have any negative effects on normal, healthy cells. Consequently, it is crucial to study the impact of plant extracts on normal cell lines.

Research on the HaCaT and HFF-1 skin cell lines revealed that hydro-alcoholic extracts from seven different alfalfa species exhibited low cytotoxicity [122]. Additionally, saponinrich fractions derived from *Medicago sativa* did not demonstrate significant cytotoxic effects on the mouse fibroblast cell line L929 [112]. Another investigation showed that alfalfa leaf extracts obtained using enzyme-assisted supercritical fluid extraction did not reduce the viability of the L929 cell lines up to 0.5 mg/mL, with an IC50 value of 1.36 mg/mL [142]. Furthermore, it was found that plant extracts could also positively influence human cells. *Medicago sativa* extracts obtained through ultrasound-assisted extraction increased the metabolism and proliferation of keratinocyte and fibroblast cell lines. This effect is likely attributed to the high antioxidant activity of these extracts, suggesting their potential use in the cosmetics industry [278]. Additionally, apigenin derived from *Pisum sativum* demonstrated no toxic activity against normal human skin cell lines [267].

In another study, researchers investigated the effects of lectin isolated from peas at a concentration of 5 mg/100 μ L, which demonstrated a toxic impact on tumor cells while not causing any adverse effects on peripheral blood mononuclear cells (PBMCs). Interestingly, the study also revealed that using lower concentrations of lectins actually promoted increased proliferation of these cells [271]. Further research by Clemente et al. [279] showed that pea proteins did not hinder the growth of normal, non-malignant colonic fibroblast CCD-18Co cells yet displayed toxic effects on cancer cell lines.

Similarly, the crude aqueous extract derived from *Triticum aestivum* did not exhibit any toxic potential toward the normal VERO cell line. In contrast, it displayed cytotoxic effects on cancer cells [276]. Subsequent studies by the same author confirmed that the *Triticum aestivum* aqueous extract did not exhibit significant toxic properties against fibroblasts (3T6 cell line) at concentrations ranging from 4 to 10 mg/mL. However, the extract did show notable cytotoxicity at a concentration of 2 mg/mL. Remarkably, this concentration also demonstrated greater efficacy in wound healing [280].

14. Conclusions

This review shows that silicon has been recognized as one of the most beneficial mineral elements for plants, performing an array of functions, for instance, fortifying plants' tolerance to abiotic and biotic stresses. Hence, the action of silicon in three plants belonging to the Fabaceae (especially *Pisum sativum* L. and *Medicago sativa* L.) and Poaceae (particularly *Triticum aestivum* L.) families regarding their physiology, strength, and utility for phytoremediation activities has been described in detail. Moreover, the methods of silicon determination and speciation are presented in this paper. Additionally, the review covers current analytical techniques, instruments, and methodologies used to investigate the influence of silicon on plant development. Finally, the manuscript focuses on questions of the isolation and characterization of bioactive compounds in plant material with antimicrobial properties and cytotoxic effects. This review highlights the potential role of silicon in modulating the biochemical and metabolic pathways, leading to the enhanced production of secondary metabolites that contribute to plant defense and overall

human health. Although our understanding of the specific mechanisms by which silicon influences these pathways remains incomplete, the literature findings contribute to the growing body of evidence suggesting that silicon plays a crucial role in regulating a plant's secondary metabolism.

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Abbreviations

AFM	Atomic Force Microscopy
ASE	Accelerated Solvent Extraction
AID	Autoclave Induced Digestion
BEC	Background Equivalent Concentration
DART	Direct Analysis in Real-Time
ETV	Electrothermal Vaporization
FTIR	Fourier Transform Infrared Spectroscopy
GAE	Gallic Acid Equivalents
HPTLC	High-Performance Thin-Layer Chromatography
HRE	Heat Reflux Extraction
HSAB	Hard and Soft Acids and Base Theory
ICP	Inductively Coupled Plasma
IC50	half maximal inhibitory concentration
LA	Laser Ablation
LIBS	Laser-Induced Breakdown Spectrometry
MIC	Minimum Inhibitory Concentration
MALDI MS	Matrix-Assisted Laser Desorption/Ionization Mass Spectrometry
MBC	Molybdenum Blue Colorimetry
MAE	Microwave Assisted Extraction
NMR	Nuclear magnetic resonance
NIRS	Near Infrared Reflectance Spectroscopy
OPLC	Overpressure-Layer Chromatography
OES	Optical Emission Spectrometry
OID	Oven Induced Digestion
PEC	Planar Electro-Chromatography
RF	Radio Frequency
SFE	Supercritical Fluid Extraction
SEM-EDX	Scanning Electron Microscopy with Energy Dispersive X-ray Detector
SEC	Size Exclusion Chromatography
TLC-DB	Thin-Layer Chromatography-Direct Bioautography
TEM	Transmission Electron Microscopy
UAE	Ultrasound-Assisted Extraction
XRF	X-ray Fluorescence

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