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The β -cyanoalanine pathway is involved in the response to water deficit in Arabidopsis thaliana.

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1	TITLE: The β -cyanoalanine pathway is involved in the response to water deficit in
2	Arabidopsis thaliana
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4	Running head: The β -cyanoalanine pathway and water deficit in Arabidopsis
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12 Abstract

The β -cyanoalanine pathway is primarily responsible for detoxification of excess cyanide 13 produced by plants. Recent evidence suggests that cyanide detoxification via this pathway may 14 be involved in the response and tolerance to water deficit in plants. The aim of this study was to 15 explore this role in Arabidopsis thaliana in greater detail. The first objective was to establish 16 responsiveness of the pathway to the magnitude and duration of water deficit. The second 17 objective was to examine how interruption of single genes (AtCysA1, AtCysC1 and AtNIT4) 18 encoding enzymes of the pathway influenced the ability to metabolize cyanide and withstand 19 20 water deficit. Arabidopsis plants were exposed to conditions which emulated acute and chronic water deficit, followed by measurement of tissue cyanide concentration, activity of enzymes, and 21 physiological parameters. The results for wild-type Arabidopsis demonstrated a transient 22 increase in cyanide concentration and β -cyanoalanine synthase activity, followed by a decrease 23 in both. The increase in enzyme activity was localized to the tissue in direct proximity to the 24 stress. The knockdown AtCysA1 mutant did not differ from wild-type while AtCysC1 mutants 25 were slightly more sensitive to water deficit. The AtNIT4 mutant was the most sensitive showing 26 decreased growth along with altered chlorophyll content under water deficit as compared to 27 28 wild-type. Collectively, the results indicated that the pathway is responsive to water deficit although the severity of stress did not alter the nature of the response, implying that the capacity 29 to remove cyanide generated during water deficit may contribute to tolerance to this stress in 30 31 Arabidopsis.

Author keywords Arabidopsis thaliana; Cyanide; Cyanoalanine synthase; Nitrilase; Water
 deficit

34

1. Introduction

Cyanide is produced by higher plants via multiple metabolic pathways. The two most 35 prevalent sources of endogenous cyanide are the turnover of cyanogenic glycosides or 36 cyanolipids [1] and ethylene biosynthesis [2, 3]. While formation of cyanogenic glycosides and 37 cyanolipids is limited to ~2,500 plant species, ethylene biosynthesis is the ubiquitous source of 38 endogenous cyanide in plants. Ethylene in higher plants is synthesized by oxidation of 1-amino-39 cyclopropane-1-carboxylic acid (ACC) by ACC oxidase (ACO)(EC 1.14.17.4), releasing 40 hydrogen cyanide and carbon dioxide as co-products [2]. The β -cyanoalanine synthase pathway 41 42 is the principle pathway for cyanide detoxification in plants. In the first step of the pathway catalyzed by β -cyanoalanine synthase (EC 4.4.1.9), cyanide reacts with cysteine to form β -43 cyanoalanine, releasing bisulphide [4]. The second step is mediated by a dual function nitrile 44 hydratase/nitrilase (EC 3.5.5.1) designated in Arabidopsis as nitrilase 4 (NIT4). The NIT4 45 enzyme catalyzes reactions that convert cyanoalanine to either asparagine or aspartate in 46 conjunction with ammonia, respectively [5]. 47 Recent evidence suggests a role of the β -CAS pathway in plant response and acclimation 48 to abiotic stress. For example, when birch plants were exposed to ozone, there was an increase in 49 50 ethylene production and increased β -CAS transcript abundance [6]. Tissue cyanide concentrations and β-CAS activity showed concomitant increases when tobacco (Nicotiana 51 tabacum L.) plants were subjected to water deficit [7]. Cyanide concentration decreased after 52 two days of stress due to the action of β -CAS enzyme, even though ethylene production was still 53 high. The conclusion offered by the authors was that water deficit induced cyanide production, 54

and that removal of cyanide by the β -CAS enzyme contributed to the response, if not also the tolerance, to water deficit.

The overarching goal of the work here was to further investigate the contribution of the 57 β-cyanoalanine synthase pathway to the response to water deficit. Arabidopsis thaliana (Col-0) 58 59 and three SALK T-DNA insertion lines for the three genes directly associated with the pathway 60 [5, 8], namely O-acetylserine(thiol) lyase (OASTL) (EC 2.5.1.47) (AtCysAI), β -cyanoalanine synthase (AtCysC1), and nitrilase 4 (AtNIT4), were used. There were two main objectives for 61 this study. The first was to examine the response of the pathway in wild type Arabidopsis Col-0 62 63 to the magnitude and duration of water deficit. Since ethylene production depends on the severity and duration of stress [9], the concomitant production of cyanide should also be variable. 64 Activity of β-CAS may also vary in order to maintain cyanide at a steady state concentration 65 below that potentially inhibitory for metabolism [3]. The second objective was to examine how 66 interruption of single genes encoding enzymes of the pathway influenced the ability of 67 Arabidopsis to metabolize cyanide and respond to water deficit. Given that β -CAS (AtCysC1, 68 Bsas 3;1) and one OASTL (AtCysA1, Bsas 1;1) are the only enzymes with β -cyanoalanine 69 synthase activity under physiological conditions [8], the efforts here sought to determine if both 70 71 of these genes were required for the function of the pathway *in vivo* or if one gene alone was sufficient to maintain cyanide assimilation under water deficit conditions. As a single gene in 72 Arabidopsis [5], the use of the nitrilase 4 SALK line allows for an examination of the impact of 73 74 water deficit when the second step of the pathway is specifically interrupted.

75

77 **2. Results**

2.1 Response of the β -cyanoalanine pathway to the magnitude and duration of water deficit 78 Acute water deficit significantly increased the relative abundance of AtCysC1 transcript, 79 but by less than two-fold overall in whole Arabidopsis seedlings (Figure 1A). A two-fold 80 increase in β-CAS-like enzyme activity was also observed in response to acute water deficit 81 (Figure 1B). While there was a significant increase in expression of AtCysC1 transcript and β -82 CAS-like enzyme activity, there was no significant difference in tissue cyanide content between 83 control plants and plants exposed to water deficit treatments (data not shown). No significant 84 85 difference was observed also in expression of AtNIT4 transcript in response to water deficit as compared to control plants (data not shown). 86 The acute water deficit treatment was repeated with plants removed from the growing 87 medium and allowed to air dry for 20 min intervals up to 60 min. As expected, RWC of the 88 tissues fell significantly over the course of 1 h, from 96% to ~85% after 20 min and to ~64% by 89 60 min. When the tissues were analyzed for cyanide concentration and β -CAS-like activity, both 90 were observed to follow similar, transient changes (Figure 2). Both tissue cyanide and β -CAS-91 like activity increased at 20 min by 2.5- and 10-fold higher than the basal levels, respectively, 92 although these changes were not significant. After 20 min, the values for each decreased so that 93 94 at 60 min, both tissue cyanide concentration and β -CAS-like activity were significantly lower than the peak at 20 min. To confirm that the patterns in the data represented specific changes in 95 tissue cyanide and β-CAS-like activity and not changes driven by differences in fresh weight 96 biomass or total protein content, respectively, the data were recalculated to express the former in 97

pmol cyanide and the latter as nmol product h⁻¹ (Figure S1) and the same general trends were
obtained although the difference for each parameter as a function of time was not significant.

One approach to apply a more environmentally-relevant, chronic water deficit was to 100 101 simply withhold watering and monitor the water status of the growing medium. Plants were first harvested for measurement four days after watering ceased when the volumetric water content of 102 the medium fell below 50%. Additional harvests were carried out on the consecutive fifth and 103 sixth days after watering ceased. Tissue cyanide concentration was significantly higher in plants 104 that had been deprived of water for four and five days as compared to plants prior to the onset of 105 106 water deficit treatment (Figure 3). After six days without water, tissue cyanide concentrations were not significantly different from untreated plants. A significant difference in β-CAS-like 107 activity was observed between plants deprived of water and plants prior to onset of treatment. 108 109 There was however no significant difference in β -CAS activity between sampling dates. To determine the response of the pathway to a longer duration of water deficit, the experiment was 110 essentially repeated except that plants were harvested after 5, 10, and 15 d without watering, at 111 112 which point there was an obvious loss of turgor in the leaves of all treated plants. The RWC of the treated tissues decreased significantly from 84% to 64% during the 15 d period, while the 113 well watered control plants maintained a RWC of 84-88%. Compared to control plants, tissue 114 cyanide concentration was significantly higher after 5 d without watering but significantly lower 115 than untreated controls after 10 and 15 d (Figure 4). There was no significant difference in β -116 117 CAS-like activity between well watered control plants and plants deprived of water for 5 or 10 days. However, enzyme activity decreased significantly after 15 d without water. Overall, the 118 activity of β-CAS pathway to water deficit was generally similar regardless of the magnitude and 119

duration of the stress. As for the acute water deficit exposure, recalculating the results from the
two chronic water deficit exposures as pmol cyanide and as nmol product h⁻¹ (Figure S1), the
same general trends were again obtained. The differences for each parameter as a function of
time did show significant differences in this case with the same transient patterns as those
observed in Figs. 3-4.

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- 126

5 2.2 The pathway response is localized to stressed tissue

To demonstrate that the response of the β -cyanoalanine synthase pathway to water deficit 127 128 stress was a phenomenon localized to tissues experiencing the water deficit, a split-root 129 technique was employed. As anticipated, discontinuing watering led to a significant decrease in the volumetric soil water content, reaching a value of 20% in root compartment 2 as compared to 130 131 58% in the watered root compartment 1. The volumetric soil water content of the two root compartments for control plants were each >80%. These differences in soil water between the 132 two equally-watered compartments for control plants and the watered side of the pots receiving a 133 134 water deficit treatment suggested that water was drawn more heavily from the watered pots opposite the water deficit pots, allowing the plant to compensate for the deficit in the dry 135 136 compartment. Despite these differences in soil hydration, the RWC of the shoots of all plants was unaffected, with no significant difference between the treated and control plants (88 and 137 85%, respectively). At harvest, there was no significant difference in the root tissue cyanide 138 139 concentration between the dry and wet compartments or in comparison to the two wet compartments of the control split-root. The tissue cyanide concentration was significantly higher 140 in shoots of plants that had a dry root compartment as compared to control plants with both root 141

142	compartments well watered (Figure 5). While there was no significant difference in β -CAS-like
143	activity in the shoots between treatments, enzyme activity in the roots experiencing water deficit
144	was significantly greater than well-watered roots (Figure 6). These trends in the data did not
145	change when the data were recalculated as former in pmol cyanide or as nmol product h^{-1} (Figure
146	S2).
147	
148	2.3 Response of the β -cyanoalanine synthase pathway mutants to water deficit
149	Consistent with studies by others [8], the established SALK lines for β -CAS (<i>AtCysC1</i>)
150	and the cytosolic OASTL (AtCysA1) were shown to be knockdowns for enzyme activity (Figure
151	7A). The β -CAS-like activity decreased by 22% for <i>AtCysA1</i> and 42% for <i>AtCysC1</i> lines as
152	compared to Col-0 plants. The magnitude of the decrease for each line was similar to the 33%
153	

154 observation made here and elsewhere [15, 16] was that the interruption of *AtNIT4* had a

155 concomitant influence on β -CAS-like activity, with a decrease in activity of 71% in comparison

to Col-0 plants (Figure 7A). Analysis of AtNIT4 enzyme activity for the *AtCysA1* and *AtCysC1*

157 lines was not significantly different from Col-0 (Figure 7B). The *AtNIT4* mutant line did display

nitrile hydratase activity although at a rate significantly lower (~20%) than for Col-0 and the

159 other two lines (Figure 8A). There was however no significant difference in nitrilase activity

under control conditions (4.1 and 3.6 nKat mg⁻¹ protein h⁻¹ respectively for Col-0 and *AtNIT4*) or

161 in response to KCN treatment (3.9 and 4.0 nKat mg^{-1} protein h^{-1} , respectively). Activity of this

162 enzyme was not cyanide inducible in either line (Figure 8B), indicating that the presence of

163 cyanide produced during water deficit did not produce the results in Figure 8A due to differential164 induction of nitrile hydratase activity in the two Arabidopsis lines.

Growth parameters (e.g., leaf area and biomass) were similar for the Col-0 plants and the 165 AtCysAl mutant line under nominal conditions. Leaf area and fresh weight biomass for AtCysCl 166 were significantly different from Col-0 while the dry weight biomass was not different. There 167 were no significant differences in these parameters between the AtCysA1 and AtCysC1 lines 168 under well-watered conditions. The AtNIT4 plants showed significantly less leaf area and 169 biomass than the Col-0 and AtCysAlline under nominal growth conditions but none of the values 170 171 were significantly different from the AtCysCl line (Figure 9). For the AtNIT4 line, the plants also displayed a smaller rosette diameter and leaf size, but not a decrease in leaf number 172 compared to the other lines (Figure S3). On imposition of water deficit there was a statistically 173 significant decrease in RWC for all lines and the magnitude of the decrease was the same for 174 each line. Well-watered controls across all four lines had a RWC of 83% while the plants 175 deprived of water had a RWC of 71%. Although the magnitude of the decrease in RWC was the 176 177 same for all the lines, the effect of this mild water deficit in terms of the subsequent effect on growth varied for each of the mutants. There was a significant decrease in leaf surface area and 178 179 fresh weight biomass after 15 d of water deficit for each line as compared to corresponding wellwatered plants of the same line (Figure 9). The magnitude of the decrease was not significantly 180 different between the AtCysA1 mutant line and the Col-0 plants (Table 1). The AtCysA1 mutant 181 182 had a 35% decrease in leaf area. The AtCysCl mutant had 50% reductions in leaf area and significantly different from Col-0. The AtNIT4 mutant was the most sensitive, with the same 183 parameter decreasing by 66%. For dry weight biomass, the AtCysAl showed a small decrease 184

while the *AtCysCl* mutant plants were slightly larger (Table 1). The *AtNIT4* plants decreased
significantly in dry weight biomass (25.4%) relative to well watered plants of the same line. The
magnitude of decrease in dry weight biomass was however greatest for the Col-0 plants at 42%
(Table 1).

The relative chlorophyll content of Col-0, the AtCysA1 and AtCysC1 lines all showed a 189 significant increase with time as compared to well-watered plants of the same line (Figure 10). 190 The measured values for these lines were higher than the control values, representing relative 191 chlorophyll values greater than the well watered plants of the same line. The AtCysA1 and 192 193 AtCysCl mutants were however not different from Col-0 except at the final time point where the AtCysAl line showed the greatest relative chlorophyll content. Contrary to the above, the 194 relative chlorophyll content of the AtNIT4 mutant showed an initial decline in the first 5 d after 195 196 initiation of water deficit and stabilized thereafter at 90% of the untreated plants of the same line. At the end of the water deficit period (day 15), the only plants showing an appreciable increase 197 in tissue cyanide was the AtNIT4 mutant under water deficit (Figure 11). There was no difference 198 199 in β -CAS-like activity between the four lines or between treatments (data not shown). As 200 compared to previous experiments with Col-0 only, the magnitude of the values for β -CAS 201 activity for all lines (Col-0 and the mutants) was lower, in these plants which seemed reasonable since the cyanide content had stabilized at low levels. 202

203

3. Discussion

205 Water deficit is a limiting factor for plant growth and development. Plants utilize a 206 number of mechanisms at the cellular level to stave off the negative effects of water deficit, 207 including osmotic adjustment, synthesis of compatible solutes, increased synthesis of abscisic 208 acid, and upregulated expression of DREB transcription factors and late embryogenesis proteins [17]. Plants also need energy under stress and the relative importance of the mitochondria as the 209 210 principal organelle for ATP production increases [18]. While other work has shown that under intense water stress, mitochondria activity decreases [19], some have reported an increase in 211 212 mitochondrial function under moderate, slow onset stress [20]. Water stress generates ethylene, and would also therefore generate cyanide. Since a principle target of cyanide inhibition is the 213 terminal oxidase of the mitochondria, cyanide detoxification by the β -CAS pathway is likely 214 215 necessary to help offset the negative effects of cyanide on the mitochondria [21], and therefore 216 contribute to the response to water deficit.

217 The first objective for this study was to establish how responsive the β -CAS pathway is 218 to the magnitude and duration of water deficit. In Arabidopsis the AtCysCl gene shows high, constitutive expression that is not responsive to cyanide exposure or stress [22], hence the 219 220 significant yet minor difference in transcript abundance even in response to an acute exposure to 221 water deficit was not wholly unexpected (Figure 1). The high protein expression is perhaps necessary to compensate for the fact that the cysteine synthesis activity displayed by β-CAS 222 represents a competing reaction for cyanide assimilation [23]. The generally parallel changes in 223 tissue cyanide and β-CAS-like enzyme activity observed here in response to water deficit 224 225 (Figures 2-4) are consistent with reports from a similar water deficit study with tobacco [7]. 226 Both studies showed a transient increase in β-CAS-like enzyme activity coinciding with a water deficit-induced spike in tissue cyanide. It was interesting to note that the pattern displayed for 227 228 cyanide concentration and β-CAS-like activity was similar for each combination of water deficit intensity and duration imposed on the Arabidopsis seedlings (Figures 2-4) suggestive of a coordinated response that includes metabolic components beyond the β -CAS pathway. With little evidence for an induction of *AtCysC1*gene, the results suggests that innate activity of the existing pool of enzymes in Arabidopsis may be sufficient to assimilate pulses of cyanide elicited during stress-induced cyanide synthesis. The same may be true for the NIT4 enzyme and may therefore explain the lack of a response at the level of transcription.

The split-root experiment further demonstrated that the response of the β -CAS pathway 235 to water deficit was localized predominantly to the tissue directly subjected to the stress as 236 237 illustrated by the significant increase in β -CAS-like activity in roots from the dry compartment as compared to the well watered compartment (Figure 6). Within the dry compartment, the 238 imposition of water deficit would presumably promote a localized increase in stress ethylene 239 which would in turn be expected to increase local activity of β -CAS. The tissue cyanide 240 concentration from the split root experiment was ~two-fold higher as compared to the other 241 drying techniques (Figure 5). Increased ethylene production was also reported in tomato plants 242 treated as split-roots [24] suggesting that the splitting of the roots imposed an additional stress, 243 perhaps mechanical. It was however, unexpected that the shoots of treated plants had higher 244 245 cyanide concentrations as compared to the controls and this accumulation of cyanide was not accompanied by an increase in β -CAS activity (Figure 6). As reported in other split-root studies 246 [e.g., 25] the RWC result here did not indicate water deficit stress in the shoots. The cyanide 247 248 detected in that tissue would not immediately seem to arise as a collateral result of the root-level treatment since the shoots of the treated plants maintained their hydration by drawing the 249 necessary water from the well-watered compartment. Two hypotheses may explain the source of 250

251 the cyanide detected in the shoots. The first is that cyanide was produced in the drying roots and 252 moved by diffusion into the dead xylem cells resulting in its translocation to shoots via the transpiration stream. However, considering the volatile nature of the HCN molecule and the 253 widespread detoxification system in all living tissues [26], an alternate explanation is that 254 cyanide was synthesized in the shoots in response to chemical signals from roots. Studies have 255 demonstrated chemical signaling between tissues upon perception of stress [e.g. 24]. In that 256 study increased abscisic acid (ABA) and ethylene were reported in the xylem sap of split-root 257 treated plants. It is also well established that production of ACC, the ethylene precursor is 258 259 increased in response to stress and may be translocated in the xylem [27]. An increase in ACC 260 production in the roots from the dry root compartment followed by translocation to the shoots may have thus resulted in increased ethylene production in shoots, and therefore cyanide 261 262 production. Simultaneous analysis of ethylene and cyanide would be necessary in future studies to provide confirmation of this supposition. 263

The efforts to evaluate how the β -CAS mutants responded to the imposition of water 264 deficit were complicated by the inherent differences in growth characteristics for the lines 265 (Figure 9) and the specific response of each to the mild water deficit imposed (Table 1, Figures 266 267 9-11). Collectively the decrease in RWC, dry weight biomass, and leaf area (Figure 9) all indicate that withholding water did affect the water status of all four lines. The progressively 268 greater decrease in these values from the *AtCysA1* to *AtCysC1* and *AtNIT4* lines (Table 1) 269 270 appears inversely related to the apparent importance of these genes and their encoded enzymes for the β -CAS pathway in that AtCysC1 is responsible for a larger proportion of β -CAS enzyme 271 activity than AtCysA1 while AtNIT4 acts at a critical bottleneck on the pathway. This 272

relationship is not fully supported though by the data for tissue cyanide and β-CAS enzyme 273 274 activity. With the exception of the AtNIT4 line, this mild water deficit did not produce an increase in β -CAS enzyme activity or an increase in tissue cyanide for the AtCysA1 and AtCysC1 275 276 lines (Figure 11). Even in the absence of an increase in cyanide in those compartments, the imposition of mild water deficit must have interacted in a different way with these two mutant 277 lines to influence the change in leaf area and fresh weight biomass. The decrease in dry weight 278 279 biomass for the Col-0 line further indicates that the change in water status did have a concomitant effect on growth of that line. The AtCysAl and AtCysCl lines did not show a 280 281 comparable decrease in dry weight biomass, suggesting that each line was more robust than Col-0 in growth following the imposition of the mild water deficit. An increase in rosette size at the 282 seedling stage was reported previously for the AtCysA1 mutant [28], although for this study the 283 growth enhancement was reflected through to maturity at 10 weeks. No statistically significant 284 evidence of this was observed here. 285

While these two lines and Col-0 showed an increase in relative chlorophyll content under 286 water deficit as compared to well-watered plants of the same line, these results may not 287 specifically indicate a change in chlorophyll synthesis. The apparent change in chlorophyll 288 289 could be due to the reduced leaf area since changes in chloroplast size, number, and/or chlorophyll content may be independent of changes in leaf area [29]. The divergence at Day 15 290 (Figure 10) with a sustained increased demonstrated by the AtCysA1 line as compared to a sharp 291 decrease for Col-0 and AtCysCl suggest the possible onset of more distinct differences between 292 the two mutant lines in response to the mild water stress imposed. 293

294 In the absence of additional data, one speculative explanation for the differences in response of the AtCysA1 and AtCysC1 mutant may involve the interaction of the β -CAS pathway 295 with signaling molecules such as ethylene, ABA, reactive oxygen species, and hydrogen sulfide. 296 297 The interaction between cyanide and ethylene synthesis is well documented [3, 30], with cyanide enhancing expression of genes such as ACS6 associated with ethylene synthesis [31] and 298 displaying crosstalk with elements of the ethylene signaling pathway during germination of 299 300 sunflower seed [32]. There is a growing body of evidence indicating interactions between ethylene and ABA in the control of growth under water stress [33]. Increased production of 301 302 reactive oxygen species (ROS) under water deficit has been demonstrated [34] and these 303 molecules are both sources of cellular damage and cellular signals. The ROS are thought to be important for regulating activity of the cytochrome and alternative oxidase pathways of 304 mitochondrial electron transport. Cyanide also influences these two pathways via its inhibition 305 of the terminal oxidase of the cytochrome pathway and may have direct roles in modulating 306 production of ROS. There is emerging evidence of signaling roles of hydrogen sulfide in plant 307 308 cells. As enzymes with dual function in cysteine synthesis and β -CAS enzyme activity, insertional mutations of either AtCysA1 or AtCysC1 could alter the regulation of both cyanide 309 and hydrogen sulfide concentration. Given the numerous potential interactions of these two β-310 CAS pathway enzymes in this landscape of signaling molecules, a complex network of 311 interactions could emerge that could influence growth in the presence of absence of mild water 312 313 stress, stomatal closure, and by extension photosynthetic activity and photochemical quenching for PS II protection. Obviously a possible disruption to this signaling network and/or a shift in 314 cellular carbon balance resulting from one of the two specific insertional mutations would 315

require extensive additional work to verify, but these two possibilities offer some explanation forthe results obtained for these two lines.

The results of this work demonstrated that the insertional mutation of AtNIT4 has the 318 most drastic effect on growth under both nominal and water limiting conditions as compared to 319 AtCysA1 or AtCysC1. The results from Figure 8B illustrate that the latter is not due to a lack of 320 enzyme inducibility resulting from the t-DNA insertion as neither the Col-0 nor AtNIT4 line 321 showed increased activity in response to cyanide exposure. The phenotypic difference in nitrile 322 hydratase activity would be expected then to remain the same between the Col-0 and AtNIT4 323 324 lines in the presence of water deficit-elicited cyanide. While sulfurtransferases in plants have been identified and do show metabolism of cyanide to thiocyanate in vitro [35], the data here on 325 hypersensitivity of the AtNIT4 mutant indirectly supports prior studies which demonstrated that 326 327 sulfurtransferases show no specific role in cyanide homeostasis *in planta*. The results offer additional evidence that the β -CAS pathway is the primary pathway for cyanide detoxification in 328 Arabidopsis. The *AtNIT4* mutant plants are not only knockdowns for nitrile hydratase activity but 329 330 also knockdowns for β -CAS activity (Figure 7 and 8). The knockdown in β -CAS activity may be indicative of feedback control of the pathway. With decreased activity of both enzymes in the 331 pathway, there would be an accumulation of β -cyanoalanine and decreased assimilation of 332 cyanide. Accumulation of both cyanide and β -cyanoalanine would be potentially detrimental to 333 the plants [8], especially in the absence of any activity from the sulfurtransferase pathway for 334 335 cyanide assimilation. Although ostensibly a t-DNA insertional mutant, the modest decrease in nitrile hydratase and lack of a difference in nitrilase activity suggest that this line may be leaky. 336 While AtNIT4 is a single gene in Arabidopsis which shows high specificity in vitro for β -337

338 cyanoalanine as a substrate [5], there seem to be no published studies with AtNIT4 mutants in 339 Arabidopsis to demonstrate that a t-DNA insertion in this gene abolishes enzymatic activity. The nitrilase/nitrile hydratase from maize (Zea mays), sorghum (Sorghum bicolor) and tobacco are 340 341 heterodimers [36]. Loss of one protein of the heterodimer may not completely abolish the enzyme activity but may decrease the overall rate and/or efficiency of enzymatic activity [36]. 342 Nonetheless, the sensitivity of AtNIT4 plants is reflected in the magnitude of the reduction in leaf 343 area and fresh weight biomass (Figure 9), and relative chlorophyll content (Figure 10). The small 344 phenotype is comparable to what might be expected for an auxin-deficient mutant, especially 345 346 since nitrilases are also involved in auxin biosynthesis [37]. However, given the purported substrate specificity of the AtNIT4 protein for β -cyanoalanine [5], this argues against AtNIT4 as 347 an auxin-deficient mutant. If the AtNIT4 is indeed a single gene in Arabidopsis responsible for 348 hydrolysis of β-cyanoalanine, knocking down this gene should cause a build-up of cyanide 349 (Figure 11) as well as β -cyanoalanine. β -cyanoalanine can be toxic to plants [8] via inhibition of 350 Asparagine:tRNA synthatase [38] and inhibition of root growth [37, 39]. These effects could 351 352 then be additive to or synergistic with the hypothesized cellular changes resulting from the 353 decreased β -CAS activity in the *AtCysA1* or *AtCysC1* lines.

In conclusion, the study has shown that the β -CAS pathway in Arabidopsis Col-0 shows a transient response to water deficit stress. The pattern of response is similar even with differences in the magnitude and duration of stress. The results have shown that the response of the β -CAS enzyme is specific to the tissue directly experiencing the stress. This study has also shown enzymatic functional redundancy between the AtCysA1 and AtCysC1 proteins and supports prior results [8] demonstrating that AtCysC1 makes a greater contribution to β -CAS activity than 360 AtCvsA1. Insertional mutation of one of the corresponding genes does not appear to affect cyanide homeostasis under water deficit but does give rise to distinct differences in growth 361 parameters associated with water status (i.e., leaf surface area and fresh weight biomass). Such 362 results suggest a complex metabolic network involving the proteins encoded by AtCysAl and 363 AtCysC1. The results from the AtNIT4 line demonstrate the innate differences in growth of this 364 line under both normal and water limited conditions. A functional sulfurtransferase pathway, 365 which should theoretically be able to remove excess cyanide during water deficit, did not 366 function as such in the AtNIT4 insertional mutants. Given the modest reduction in nitrile 367 368 hydratase activity in this line, further investigation will be needed to examine the contribution of the encoded enzyme to nitrilase and nitrile hydratase activity and potential *in vivo* redundancies 369 that may exist. The degree to which β -CAS activity contributes to abiotic stress tolerance will 370 also require further examination to establish the extent of cyanide production under other water 371 deficit scenarios and in response to other abiotic stresses. The results obtained are relevant to 372 both the fundamental study of abiotic stress in plants and to the herbicide industry in that the 373 374 mode of action of auxinic herbicides is induction of ethylene biosynthesis and cyanide autotoxicity [40]. Tolerance to such herbicides has been linked to CAS activity suggesting both 375 376 stress and herbicide tolerance may be provided in plants in part by the β -CAS pathway. An intriguing question for future study would be to examine how plants respond to simultaneous 377 exposure to water deficit in cyanide contaminated soils. As tolerance to both stresses may 378 379 require action of the β -CAS pathway, such studies could provide information on the capacity of the pathway to respond under more extreme conditions in the presence of multiple stresses. The 380 complexity of the results obtained here and the interaction of the β -CAS pathway with 381

mitochondrial function and several important cellular signaling molecules (e.g., ethylene, ROS,
hydrogen sulfide) also imply a broader contribution to cellular function than simple cyanide
detoxification.

385

4. Materials and Methods

387 *4.1 Plant culture*

Seeds of wild type Arabidopsis thaliana (Col-0) and the SALK t-DNA insertional 388 mutants for cysteine synthase (AtCysA1, At4g14880, SALK 72213), β -cyanoalanine synthase 389 (*AtCysC1*, At3g61440, SALK 22479) and nitrilase 4 (*AtNIT4*, At5g22300, SALK 016289C) 390 were obtained from TAIR (www.Arabidopsis.org). For some experiments, seeds were surface 391 sterilized by fumigation and germinated on plates containing half-strength MS medium with 1% 392 agar and 0.5% sucrose. After a three day vernalization period at 4°C in the dark the plates were 393 placed at a 30° angle in a Percival growth chamber (Model E-36 L, Des Moines, IA, USA) at 394 ambient humidity with an 8 h photoperiod at a light intensity of ~150 μ M m⁻² s⁻¹. The day/night 395 396 temperatures were 22° and 18°C respectively, and plants were left to grow for three weeks. Plants were removed from plates and transferred to sterile perlite and vermiculite (1:1) and 397 398 grown for an additional three weeks. For one experiment, plants were transferred from plates directly to hydroponic solution with the following composition: 6 mM KNO₃, 4 mM Ca(NO₃)₂, 399 0.1 mM NH₄H₂PO₄, 1 mM MgSO₄, 50 µM KCl, 12.5 µM H₃BO₃, 1 µM MnSO₄, 1 µM ZnSO₄, 400 0.5 µM CuSO₄, 0.1 µM NiSO₄, and 0.016 µM (NH₄)₆Mo₇O₂₄ [10]. The solution was buffered 401 with 1 mM n-morpholinoethanesulfonic acid (MES) titrated to pH 6.0 with MES-TRIS. Iron 402 was provided as 10 µM Fe-EDTA, the solution was continuously aerated, and the growth 403

404 conditions were as indicated above. After three weeks of growth plants were transferred from the hydroponic culture to sterile sand for the split-root experiment described below. Surface 405 sterilized seeds of wild type (Col-0) for one additional experiment were also germinated directly 406 407 in sterile potting mix using the same vernalization and growth conditions as above. These plants were watered with 50 mL of the same nutrient solution as above and 100 mL water per week. 408 The 100 mL of water was split into two applications of 50 mL. The plants were grown for six 409 weeks under the same growth conditions and photoperiod as above to establish biomass and were 410 then subjected to the desired treatments. 411

412

4.2 Response of the β -cyanoalanine synthase pathway to magnitude and duration of water deficit 413 Two different strategies were used to impose water deficit on Arabidopsis Col-0 plants to 414 observe the response of the β -cyanoalanine synthase pathway. For the first, the goal was to 415 examine the response to a severe, short-term (i.e., acute) water deficit. Three week old plants 416 grown on MS plates were subjected to this acute water deficit by removing the plants from the 417 plate and exposing the whole seedlings to air for 20 min. This method of creating water deficit 418 was used previously in a study that examined global changes in gene expression in Arabidopsis 419 420 in response to abiotic stresses [11]. There were three replicates each for the acute water deficit treatment and the control. Whole plants were harvested, snap-frozen, and stored at -80°C. 421 Additional plants, grown three weeks in MS plates and then three weeks in potting mix were left 422 423 unwatered for seven days prior. An acute water deficit treatment was imposed by gently removing the seedlings from the potting mix, removing adhering soil particles, and exposing the 424 roots to air for 45 min. Following the treatments, whole plants (three replicates each of control or 425

treated plants) were snap-frozen in liquid nitrogen for later analysis of whole plant tissue cyanide concentration and β -CAS-like activity. Additional plants were germinated and grown in perlite and vermiculite up to the 10-12 leaf stage. These plants were pulled out of the medium and air dried for 60 min to impose acute water deficit. Plants were sacrificed at 20 min intervals for 60 min and snap-frozen in liquid nitrogen. Replicate samples of treated and control plants (*n*=3) were analyzed for relative water content, tissue cyanide and β -CAS-like activity in the whole plant.

The second strategy for imposing water deficit was more physiologically and 433 434 environmentally relevant (i.e. chronic water deficit). For one experiment, watering of Col-0 plants grown in perlite and vermiculite was withdrawn and the volumetric water content of the 435 medium was monitored using an Echo EC-5 soil moisture sensor (Decagon Devices Inc, Pullman 436 WA USA). After 4 days, when the volumetric water content of the medium decreased to 50%, 437 plants were harvested at one day intervals for three days. These samples (n=3) were analyzed for 438 relative water content, tissue cyanide and β -CAS-like activity. In a second experiment, watering 439 440 was discontinued for 15 days. Plants (n=4) were harvested every five days for 15 days, at which point all treated plants were showing a significant loss of turgor. Relative water content of the 441 442 shoots, tissue cyanide and β -CAS-like activity were determined.

To examine whether the response was localized to tissues in contact with dry soil, a splitroot experiment was performed using six-week-old Col-0 plants grown for three weeks in hydroponics. The plants were grown hydroponically to reduce the damage associated with their transfer and to allow for an easier separation of the root system into two roughly comparable components. The two components of the root systems were each established in separate pots

containing sterile sand. The plants were provided one week to acclimate to the new growing 448 medium. During this recovery period, there were no signs of stress and new growth was evident. 449 After the recovery period, the split-root systems were assigned to one of two treatment groups, 450 control and treated. For the control plants, the two pots containing the split-root systems were 451 each watered with 50 mL nutrient solution once per week and two applications of 50 mL of 452 453 water twice per week. For the treated plants, water was completely withheld from one compartment, while the other compartment received the same watering pattern as used for the 454 control plants. Plants were harvested on day 7 and separated into shoots and the roots from each 455 456 pot. The relative water content of shoots, tissue cyanide concentration in roots and shoots, and β -CAS-like activity in roots and shoots were determined for these samples. There were four 457 replicates of each treatment in this experiment with a replicate represented by a single split-root 458 459 plant.

460

461 4.3 Response of the β -cyanoalanine synthase pathway mutants to water deficit

The decrease in β -cyanoalanine synthase-like activity and/or nitrilase/nitrile hydratase 462 activity in the insertional mutants was demonstrated using the procedures described below. 463 464 Plants were germinated and grown in plates under the same growth conditions as above. After three weeks, seedlings were transferred from MS plates to hydroponic culture and allowed to 465 grow for two additional weeks. Whole plants were harvested, rinsed, snap-frozen in liquid 466 nitrogen and stored at -80°C prior to analysis. The assay for β -cyanoalanine synthase-like 467 activity was conducted on whole seedlings and each line was replicated four times. Similarly, 468 nitrilase/nitrile hydratase activity was determined in Col-0 and the AtNIT4 mutants. 469

470 Nitrilase/nitrile hydratase activity was determined in untreated plants of each line and in plants 471 treated for 36 h with 1 mM cyanide (as KCN). Three week old seedlings were transferred from MS plates to hydroponic culture as above, and grown for three weeks. The 36 h cyanide 472 473 treatment was then imposed on half the Col-0 and AtNIT4 plants. The composition of the hydroponic cyanide treatment solution was similar to the basal nutrient solution except that the 474 phosphate concentration was lowered to 0.02 mM, MnSO₄ concentration was lowered to 1.6 µM, 475 and iron was omitted to preclude formation of metal cyanide solids [10]. Aeration of the nutrient 476 solution was discontinued to reduce cyanide volatilization [10]. After 36 h of exposure, whole 477 478 seedlings were harvested, rinsed, snap-frozen in liquid nitrogen, and stored at -80°C until the nitrile hydratase/nitrilase assays were performed. Each line and treatment had four replicates. 479 To examine how the putative lines for β -cyanoalanine synthase (AtCysC1), the cytosolic 480 OASTL (AtCysA1), and the nitrile hydratase/nitrilase (AtNIT4) responded to water deficit, seeds 481 of each line were germination and grown for three weeks on MS plates and then three weeks in 482 1:1 sterile perlite and vermiculite. A subset of plants of each line was harvested and the whole 483 plants were snap-frozen in liquid nitrogen for determination of basal β -CAS-like activity. For 484 the remaining plants, watering was withheld for 15 days and the physiological status of the plants 485 486 was monitored by measuring relative chlorophyll content. The plants were harvested and leaf area was measured (Li-COR leaf area meter, model LI-3000 A, Lincoln NE USA). The tissues 487 were then subdivided into two subsamples. The first was used for the determination of relative 488 489 water content. Using the dry weight and the relative water content of this subsample, the total dry weight of the tissue at harvest was calculated. The second tissue subsample was rinsed with 490

491 deionized water to remove adhering soil particles, and snap-frozen for storage as described above

492 for determination of tissue cyanide concentration and β -CAS activity.

493

494 *4.4 Biochemical and physiological measurements*

495 <u>4.4.1 Quantitative RT-PCR</u>

To assess expression of genes of interest, total RNA was extracted from whole seedlings using 496 the E.Z.N.A Plant RNA Kit (OMEGA Bio-tek, Norcross, GA, USA) according to the 497 manufacturer's instructions. The RNA concentration was determined using a Nanodrop 498 spectrophotometer, (NanoDrop ND-1000, Wilmington, DE, USA). First strand cDNA was 499 synthesized from 1 µg of RNA using the SuperScriptTM III First Strand Synthesis System for RT-500 PCR (Invitrogen, CA, USA) according to the manufacturer's instructions. Real-time PCR was 501 conducted using gene specific primers and the Chromo4 real-time PCR system (Bio-Rad 502 Laboratories, Hercules, CA, USA). Quantification of transcript abundance was achieved using a 503 standard curve which related DNA concentration to the Ct value [12]. Expression of genes was 504 505 normalized to the ubiquitin 10 gene (At4g05320) and melting curves were used to confirm that only a single transcript was being amplified. 506

507

508 <u>4.4.2 Measurement of plant tissue cyanide</u>

Tissue cyanide content was determined using a microdiffusion technique [13] to consist of a 50 mL conical centrifuge tube and a 5 mL glass tube as a gas trap. Plants were ground under liquid nitrogen and then ground again in an extraction buffer (50 mM of K₂HPO₄, pH 8.9) using 10 mL of buffer for each gram of fresh tissue. The slurry was transferred to a 15 mL centrifuge

tube and centrifuged for 10 min at 3,000 g at 4°C. The supernatant was transferred to the 50 mL 513 514 Falcon tube. The smaller inner tube, which contained 1.5 mL of 5 mM K₂HPO₄ and 0.5 mL of phenolphthalin reagent, was inserted into the 50 mL centrifuge tube. To force the cyanide into 515 the volatile phase, $18 \text{ M} \text{ H}_2\text{SO}_4$ was added to the supernatant in the 50 mL tube to a final 516 concentration of 0.41 M and swirled gently to mix. The Falcon tube was sealed and incubated at 517 25°C for 1 h. After the incubation, the smaller gas trap was removed and 0.5 mL of 0.1% KOH 518 was added. After a period of color development, the absorption of the trap solution was then 519 read at 550 nm wavelength with a Cary 50 UV-Visible spectrophotometer (Varian Inc., 520 521 Mulgrave, Australia), and compared to a standard curve constructed from known concentrations

of KCN.

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522

524 <u>4.4.3 DMPDA assay for β -cyanoalanine synthase activity</u>

To determine activity of β -CAS in the seedlings, whole plants or individual tissues were 525 ground under liquid nitrogen and then under a buffer consisting of 2 mM EDTA-Na₂, 10 mM 526 527 cysteine, 0.1 M Tris-HCl, pH 9.5. The homogenate was centrifuged at 4,000 g for 10 min at 4° C and then split into three subsamples. One subsample was used for determination of the total 528 529 protein content using the Pierce BCA assay Kit (Thermo Scientific) according to the manufacturer's instructions. The other two samples were used for the determination of β -CAS 530 activity. One of these two tubes was boiled for 10 min to provide a heat-killed control to correct 531 532 for background absorbance. The activity of β -CAS was assayed using the DMPDA assay [7]. The crude protein supernatant was mixed 1:1 (v/v) with the substrate (10 mM cysteine, 3 mM 533 KCN, 160 mM 2-amino-2-methyl-1-propanol, pH 9.8). The reaction mixture was incubated in 534

sealed vials at 26° C for 20 min. The reaction was terminated with the addition of 0.5 mL of
acidic dye precursor reagent (15 mM N,N-dimethyl-1,4-phenylenediamine dihydrochloride, 3
mM ferric chloride, 4.2 mM HCl). After a 20 min period of color development, methylene blue
formation was measured spectrophotometrically at 745 nm. Enzyme activity data was
normalized to the protein concentration.

540

541 <u>4.4.4 Colorimetric determination of nitrilase and nitrile hydratase activity</u>

Nitrilase and nitrile hydratase activity was assayed via determination of ammonia 542 according to Piotrowski et al. [5]. Briefly, 1 g (FW) of tissue was ground in liquid nitrogen and 543 then in 3 ml of 100 mM KH₂PO₄ buffer, pH 8.9. The homogenate was centrifuged at 4000 g for 544 15 min at 4°C. The supernatant was decanted and centrifuged again for 20 min to obtain a crude 545 extract. Total protein in the extract was determined as indicated above. For determination of 546 background ammonia in the extract, an aliquot containing 100 µg of total protein was heat 547 denatured in boiling water for 10 min and then incubated with substrate (3 mM β -cyanoalanine 548 549 in 0.05 M potassium phosphate buffer, pH 8.9) in a total volume of 1 mL. Nessler's reagent (1.33 mL) was added and after 10 min of color development the absorption was read at 480 nm. 550 A second aliquot of the crude extract containing 100 µg of total protein was incubated with the 551 same substrate in a total volume of 1 mL for 30 min at 30°C after which the sample was boiled at 552 100°C for 10 min to stop the reaction. For parallel determination of nitrilase and hydratase 553 554 activity, two aliquots, each 0.1 mL were taken from the reaction mixture and each was diluted to 1 mL with deionized water. To one sample, 1.33 mL Nessler's reagent was added for 555 determination of ammonia resulting from the nitrilase activity. The second sample was incubated 556

with 0.25 units of asparaginase at 37°C for 30 min, allowing for determination of ammonia
resulting from both nitrilase and nitrile hydratase activity. After incubation, 1.33 mL Nessler's
reagent was added and both samples were read at 480 nm. Nitrile hydratase activity expressed as
the difference between ammonia produced from the combined activity of the two enzymes and
the activity of nitrilase [5].

562

563 <u>4.4.5 Relative water content</u>

Samples were immediately weighed after harvest to determine fresh weight (FW). The samples were placed in the dark overnight in vials containing DI water at 4°C and then weighed to determine turgid weight (TW). The samples were oven dried to constant mass at 55°C and the dry weight (DW) was obtained. Relative water content (RWC) was determined according to Barr and Weatherley [14]:

569
$$RWC (\%) = \frac{FW - DW}{TW - DW} * 100$$

570

571 <u>4.4.6 Relative chlorophyll content</u>

572 Relative chlorophyll content of leaves was determined using a SPAD 502+ Chlorophyll Meter
573 (Konica Minolta Sensing Inc., Osaka, Japan).

574

575 *4.5 Statistical analyses*

576 Data from the experiments using only Col-0 plants were analyzed using SAS package Version

- 577 9.1 as one-way ANOVA with Tukey's test used for post hoc analysis. Data from experiments
- 578 with Col-0 and the three mutant lines was first analyzed as a two-way ANOVA. Where no

significant interactions between the main effects for a parameter were observed, the main effects
were analyzed separately by one-way ANOVA with Tukey's test used for post hoc analysis.
When a significant interaction between main effects did occur, the data were reanalyzed by oneway ANOVA with each interaction mean (line x treatment) representing an individual treatment.
Tukey's test was again used for the post hoc analysis.

584

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