Mutation of Trp-574-Leu ALS gene confers resistance of radish biotypes to iodosulfuron and imazethapyr herbicides

Joanei Cechin¹*, Leandro Vargas², Dirceu Agostinetto¹, Fabiane Pinto Lamego³, Franciele Mariani⁴ and Taisa Dal Magro⁵

¹Faculdade de Agronomia Eliseu Maciel, Universidade Federal de Pelotas, Campus Universitário s/n., Cx. Postal 354, 96010-900, Pelotas, Rio Grande do Sul, Brazil. ²Empresa Brasileira de Pesquisa Agropecuária, Passo Fundo, Rio Grande do Sul, Brazil. ³Empresa Brasileira de Pesquisa Agropecuária, Embrapa Pecuária Sul, Bagé, Rio Grande do Sul, Brazil. ⁴Instituto Federal de Educação, Ciência e Tecnologia, Campus Sertão, Sertão, Rio Grande do Sul, Brazil. ⁵Universidade de Caxias do Sul, Vacaria, Rio Grande do Sul, Brazil. *Author for correspondence. E-mail: joaneicechin@yahoo.com.br

ABSTRACT. Acetolactate synthase inhibitors are the main group of herbicides used in winter crops in Southern Brazil where their intensive use has selected for herbicide-resistant biotypes of radish. The resistance affects the efficacy of herbicides, and identifying the resistance mechanism involved is important for defining management strategies. The aim of this study was to elucidate the resistance mechanism of radish biotypes by quantifying the enzyme activity, ALS gene sequencing and evaluating the response of biotypes to iodosulfuron and imazethapyr herbicide application after treatment with a cytochrome P 450 monooxygenase inhibitor. The susceptible (B₁) and resistant (B₄ and B₁₃) biotypes were from wheat fields in the Northwest of Rio Grande do Sul State. The results demonstrated that the enzyme affinity for the substrate (Kₘ) was not affected in biotypes B₄ and B₁₃ but that the Vₘₐₓ of the resistant biotypes was higher than that of biotype B₁. The resistant biotypes showed no differential metabolic response to iodosulfuron and imazethapyr herbicides when inhibited by malathion and piperonyl butoxide. However, gene sequencing of ALS showed a mutation at position 574, with an amino acid substitution of tryptophan for leucine (Trp-574-Leu) in resistant biotypes.

Keywords: Raphanus sativus, mechanism of resistance, ALS enzyme activity, gene mutation, metabolism.

Introduction

Raphanus sativus L. (radish) is a dicotyledonous weed species that is found in wheat, barley and canola fields of southern Brazil, where it causes yield reduction (Rigoli, Agostinetto, Schaedler, Dal Magro, & Tironi, 2008). Acetolactate synthase (ALS) is the most common enzyme in the branched-chain amino acid biosynthetic pathway and produces leucine, isoleucine and valine (McCourt & Duggleby, 2006). ALS inhibitor herbicides are essential for a variety of crops due to their selectivity, low effective dosage, reduced toxicity to animals and high potential for inhibiting the ALS enzyme (Yu, Han, & Vila-Aiub, 2010; Endo, Shimizu, Fujimori,
Yanagisawa, & Toki, 2013). Iodosulfuron and imazethapyr are the main herbicides used in wheat and canola where their intensive use has favored the selection of herbicide-resistant radish biotypes (Pandolfo, Presotto, Poverene, & Cantamutto, 2013).

The survival of biotypes can occur due to factors that may be related to the herbicide target site or non-target-site (Yuan, Tran, & Stewart, 2007). The occurrence of DNA mutations in the gene sequence and the overexpression of the ALS enzyme are possible factors resulting in reduced sensitivity to the herbicide due to insufficient or excessive levels of biosynthetic product (Duggleby, McCourt, & Guddat, 2008; Han et al., 2012). Mutations in the ALS gene can affect enzyme structure and function, thereby reducing the enzyme activity and herbicide affinity with the target site (Han et al., 2012). In Raphanus raphanistrum L., ALS gene mutations affecting the efficacy of ALS inhibitor herbicides were identified for proline (Pro197), aspartate (Asp756), tryptophan (Trp574) and alanine (Ala122); (Tan & Medd, 2002; Yu, Hashem, Walsh, & Powles, 2003; Yu, Han, Purba, Walsh, & Powles, 2012; Han et al., 2012).

Non-target-site herbicide resistance mechanisms can occur due to increased metabolism and compartmentalization, a reduction in sorption or the differential translocation of the herbicide molecule (Powles & Yu, 2010; Délye, Jasieniuk, & Le Corre, 2013). These mechanisms of resistance are also characterized by higher rates of herbicide detoxification due to increased glutathione-s-transferase, cytochrome P450 monooxygenase or glycosyltransferase activities (Délye et al., 2013). Nevertheless, both mechanisms affect herbicide efficacy and should be evaluated due to the possibility of their coexistence (Ahmad-Hamdani et al., 2013, Brosnan et al., 2016).

Therefore, elucidating the resistance mechanism in radish biotypes is important for determining alternative weed management strategies and for reducing the herbicide-resistance selective pressure. The aim of this study was to elucidate the resistance mechanism of radish biotypes by quantifying ALS enzyme activity, ALS gene sequencing and mechanism of radish biotypes is important for determining the zero activity control were subtracted from the enzymatic extraction method was adapted from methods described by Singh, Stidham, and Shomer (1988). Seven grams of young plant leaves was collected, frozen in liquid nitrogen (N2) and ground to a fine powder. Then, 70 mL (1:10 p/v) of 100 mM phosphate extraction buffer (pH = 7.5) containing 0.5 mM magnesium chloride (MgCl2), 10 mM sodium pyruvate, 0.5 mM thiamine pyrophosphate (TPP), 10 μM flavin adenine dinucleotide (FAD), 10% glycerol, 1 mM dithiothreitol and 5% polyvinylpyrrolidone (PVPP) was added. The material was homogenized for 20 minutes at 4°C, and the mixture was filtered to remove solid sediments. The liquid portion was centrifuged at 12,000 rpm for 20 minutes, the supernatant was collected and the solid residue was discarded.

The methodology used for the in vitro assay with the herbicide was adapted from the method of Gerwick (Gerwick, Mireles, & Eilers, 1993). The assay was performed in test tubes using three replicates with a factorial treatment design, where factor A was the different biotypes (B1, B4 and B13) and factor B consisted of different concentrations of iodosulfuron or imazethapyr (zero, 0.001, 0.01, 0.1, 1.0, 10, 100, and 1,000 μM). Each tube received 600 μL of enzyme solution, 100 μL of herbicide solution and 300 μL of 80 μM phosphate reaction buffer (pH = 7.0) containing 20 mM de magnesium chloride, 200 mM sodium pyruvate, 2 mM thiamine pyrophosphate and 20 μM flavin adenine dinucleotide. The assay had two standard treatments without herbicide to measure zero and 100% enzyme activity. The zero activity standard received 50 μL sulfuric acid (H2SO4 × 3 M) at the start of assay to prevent enzyme activity, and the 100% activity standard received 100 μL milli-Q water instead of the herbicide solution. The absorbance values for the zero activity control were subtracted from the

**Material and methods**

Seeds from radish plants that survived application of iodosulfuron herbicide were collected in wheat fields in the Northwest region of Rio Grande do Sul State. To screen for resistant biotypes, plants grown from these seeds were sprayed with 5 g a.i. ha⁻¹ of iodosulfuron and 106 g i.a. ha⁻¹ of imazethapyr; and dose response studies were then carried out for biotypes B1, B4, and B13. The B1 and B13 biotypes were from Três de Maio and Boa Vista do Cacedo, Rio Grande do Sul State municipalities, respectively, and demonstrated cross resistance to these herbicides and a high level of resistance to the iodosulfuron herbicide. Biotype B4, from Três de Maio, Rio Grande do Sul State, was susceptible to the iodosulfuron and imazethapyr herbicides (Cechin et al., 2016).

**ALS enzyme activity and in vitro assays with the herbicides**

The enzymatic extraction method was adapted from methods described by Singh, Stidham, and Shomer (1988). Seven grams of young plant leaves was collected, frozen in liquid nitrogen (N2) and ground to a fine powder. Then, 70 mL (1:10 p/v) of 100 mM phosphate extraction buffer (pH = 7.5) containing 0.5 mM magnesium chloride (MgCl2), 10 mM sodium pyruvate, 0.5 mM thiamine pyrophosphate (TPP), 10 μM flavin adenine dinucleotide (FAD), 10% glycerol, 1 mM dithiothreitol and 5% polyvinylpyrrolidone (PVPP) was added. The material was homogenized for 20 minutes at 4°C, and the mixture was filtered to remove solid sediments. The liquid portion was centrifuged at 12,000 rpm for 20 minutes, the supernatant was collected and the solid residue was discarded.
values read in other treatments. After preparation of the reaction, samples were incubated for 60 minutes at 30°C. The reactions were stopped with 50 μL of 3 M sulfuric acid for all treatments other than the zero activity control treatment, where the reaction had been stopped initially. Next, the tubes were incubated for 15 minutes at 60°C to create acetoin from the reaction of sulfuric acid with acetalactate. Then, 0.5% creatine (1,000 μL) and 0.5% 1-naphtol (1,000 μL) were prepared in 2.5 M sodium hydroxide (NaOH) and added to produce a colored complex. The final reaction was incubated for 15 minutes at 60°C, and the absorbance at 530 nm was read in a spectrophotometer. The ALS enzyme activity (μM acetoin min⁻¹ mL⁻¹) was determined by the amount of acetoin produced. For the standard curve, three replications were used; each tube received 1000 μL with different concentrations of solution acetoin (0, 10, 20, 40, 60, 80, 100, 200, and 400 μM). The colored complex was obtained by adding creatine, 1-naphtol and NaOH and incubated as described above.

The kinetic parameters of enzyme activity (K_M and V_max) were obtained using ten pyruvate concentrations (zero, 0.5, 1, 2, 4, 8, 16, 32, 64, and 100 mM). The substrate concentrations (pyruvate) were obtained by diluting phosphate reaction buffer (80 μM, pH = 7.0) with 100 mM pyruvate solution. The buffer contained 20 mM magnesium chloride, 2 mM thiamine pyrophosphate and 20 μM flavin adenine dinucleotide. The values of K_M and V_max were determined using the Michaelis-Menten equation: \( y = \frac{V_{max} \times X}{K_M + X} \) (Nelson & Cox, 2008), where \( y = \) ALS enzyme activity (μmol min⁻¹ mL⁻¹); \( V_{max} = \) maximum reaction velocity; \( X = \) substrate concentration (pyruvate); and \( K_M = \) substrate concentration, where the initial velocity is equal to half of the maximum reaction velocity. The data obtained were analyzed for normality (Shapiro-Wilk test) and submitted to analysis of variance (p ≤ 0.05), where the K_M and V_max values of biotypes were compared by a Duncan Test (p ≤ 0.05).

The absorbance values were corrected with the zero standard, and I_50 (amount of herbicide to inhibit 50% enzyme activity) was calculated using the logistic non-linear regression model: \( y = a / \left[ 1 + \left( x / x_{50} \right)^b \right] \) (Seefeldt, Jensen, & Fuerst, 1995), where \( y = \) ALS enzyme activity (%); \( a = \) maximum point; \( x = \) dose of iodosulfuron or imazethapyr (μM); \( x_{50} = \) dose of iodosulfuron or imazethapyr that corresponds to a 50% inhibition of the ALS enzyme; and \( b = \) curve declivity. The resistance factor (RF) was calculated dividing the I_50 of the resistant biotype by values from the susceptible biotype (Hall, Stromme, & Horsman, 1998). The level of total protein was obtained using the Bradford method (Bradford, 1976).

### ALS gene sequencing

RNA was extracted from leaf tissue (100 mg) in biotypes (B_1, B_4, and B_13) using 500 μL of extraction buffer (reagent Kit PureLink™ Plant RNA) according to the manufacturer's instructions. The quality and quantity of RNA were verified by electrophoresis gel and spectrophotometry, respectively. cDNA was synthesized from 2 μg RNA using the SuperScript™ III First-Strand Synthesis System for RT-PCR (Invitrogen™ - USA) according to the manufacturer's instructions. Polymerase chain reaction (PCR) products were obtained using the primers WR122F, WR653R, WR205R, WR376R, and WR574F (Han et al., 2012) and the ALS gene sequence of *R. raphanistrum* L. (AJ344986) that was deposited in GenBank of National Center for Biotechnology Information (NCBI).

PCR was conducted in a final volume of 25 μL containing 1.25 μL cDNA, 0.25 μM each of the forward (F) and reverse (R) primers, 12.5 μL 2x GoTaq™ Green Master Mix (Promega™) and nuclease-free water. cDNA denaturation was conducted at 94°C for 4 minutes, with 40 cycles of 94°C for 30 seconds. The annealing of primers occurred at 55°C for 30 seconds, and sequence extension was performed at 72°C for 30 to 120 seconds (Han et al., 2012). *Amplicons* were purified using the PCR Purification Combo Kit (Invitrogen™ - USA) and sequenced using the ABI-PRISM 3100 Genetic Analyzer (Applied Biosystems). Sequences from the various biotypes were aligned using the Bioedit version 7.2.5 software and compared with the sequence of *R. raphanistrum* L. (AJ344986) deposited at GenBank (www.ncbi.nlm.nih.gov/genbank).

### Metabolization

The metabolization of radish biotypes was analyzed in two additional experiments that were performed in a greenhouse with a completely randomized design and four replications. Seeds were sown in plastic trays, and two days after emergence, each seedling was transplanted to plastic pots with volume capacities of 0.75 L that contained soil and PlantMax substrate at a 2:1 ratio.

The treatments were arranged in a factorial design, where factor A was the radish biotypes (B_1, B_4, and B_13) and factor B were the cytochrome P_450 monoxygenase inhibitors malathion and piperonyl butoxide. Spraying occurred when plants were at the three to four leaf stage by using a CO₂ backpack.
sprayer calibrated to deliver 120 L ha\textsuperscript{-1}. The metabolism inhibitors were sprayed 30 minutes before herbicide application at a dose of 500 g a.i. ha\textsuperscript{-1} for malathion and 525 g a.i. ha\textsuperscript{-1} for piperonyl butoxide (PBO). The doses of iodosulfuron and imazethapyr herbicides sprayed were 3.5 and 106 g a.i. ha\textsuperscript{-1}, respectively.

The control and shoot dry matter (SDM) were evaluated at 28 days after application (DAA). A percentage scale for control was adopted, in which zero (0) and one hundred (100) corresponded to the absence of damage and complete death of the plants, respectively (Frans & Crowley, 1986). The SDM was determined by drying the vegetable material in a kiln with circulated forced air at 60°C for 72 hours and expressed as grams per plant.

The obtained data were analyzed for normality (Shapiro-Wilk test), which did not require data transformation, and were then submitted to analysis of variance (p ≤ 0.05). When statistical significance was observed, the data were submitted to the Duncan test (p ≤ 0.05).

Results and discussion

ALS enzyme activity and in vitro assay with the herbicides

The K\textsubscript{M} values (pyruvate) of resistant biotypes (B\textsubscript{4} and B\textsubscript{13}) did not show statistically significant differences in enzyme affinity for the substrate, and their V\textsubscript{max} was higher than that of the susceptible biotype (Figure 1). However, the K\textsubscript{M} values of the B\textsubscript{4} and B\textsubscript{13} biotypes were 20 and 25% higher, respectively, than that of the susceptible biotype (Table 1). Further, the V\textsubscript{max} values were 7.78, 5.73, and 1.87 μM acetoin mg\textsuperscript{-1} protein h\textsuperscript{-1} for B\textsubscript{4}, B\textsubscript{13} and B\textsubscript{1} biotypes, respectively (Table 1). Other studies have demonstrated that biotypes that are resistant to ALS inhibitor herbicides do not present changes in the kinetic parameters (K\textsubscript{M} and V\textsubscript{max}) compared to susceptible biotypes (Ashigh & Tardif, 2007; Dal Magro et al., 2010), which suggests that resistance does not affect pyruvate binding. Similar results were found in Cyperus difformis L. biotypes resistant to pirazosulfuron-ethyl herbicide, in which the affinity of enzyme was not affected and the V\textsubscript{max} was 52% greater (Dal Magro et al., 2010).

The ALS inhibitor herbicides act by blocking the channel that leads to the active site of the enzyme; therefore, any changes in this channel can impede the binding of the herbicide while maintaining the conformation and function of the enzyme (McCourt, Panf, King-Scott, Guddat, & Duggleby, 2006). In Lolium rigidum (Gaudin) biotypes resistant to sulfometuron and imazapyr herbicides, a mutation of tryptophan to leucine in position 574 (Trp\textsubscript{574-Leu}) did not change the K\textsubscript{M} and V\textsubscript{max} was 287% greater in resistant biotypes (Yu et al., 2010). The results obtained for in vitro assays with herbicide treatment demonstrated that 0.043 μM iodosulfuron and 3.2 μM imazethapyr inhibited 50% of the ALS enzyme activity (I\textsubscript{50}) in the susceptible biotype. For the resistant biotypes, the I\textsubscript{50} was 0.65 and 0.82 μM for iodosulfuron and 718 and 425 μM for imazethapyr in B\textsubscript{4} and B\textsubscript{13} biotypes, respectively (Figure 2, Table 2).

**Figure 1.** ALS activity (μM acetoin mg\textsuperscript{-1} protein h\textsuperscript{-1}) of susceptible (B\textsubscript{1}) and resistant (B\textsubscript{4} and B\textsubscript{13}) radish biotypes to iodosulfuron and imazethapyr herbicides subjected to differential pyruvate concentrations (mM). Points represent the mean values and bars represent least significant difference (p < 0.05).

**Table 1.** Kinetic parameters K\textsubscript{M} (mM) and V\textsubscript{max} (μM acetoin mg\textsuperscript{-1} protein h\textsuperscript{-1}) of susceptible (B\textsubscript{1}) and resistant (B\textsubscript{4} and B\textsubscript{13}) radish biotypes in response to iodosulfuron and imazethapyr herbicide treatment.

<table>
<thead>
<tr>
<th>Biotypes</th>
<th>K\textsubscript{M} (mM)</th>
<th>V\textsubscript{max} (μM acetoin mg\textsuperscript{-1} protein h\textsuperscript{-1})</th>
</tr>
</thead>
<tbody>
<tr>
<td>B\textsubscript{1} (Susceptible)</td>
<td>20.02\textsuperscript{*}</td>
<td>1.87 C</td>
</tr>
<tr>
<td>B\textsubscript{4} (Resistant)</td>
<td>16.70</td>
<td>7.78 A</td>
</tr>
<tr>
<td>B\textsubscript{13} (Resistant)</td>
<td>16.10</td>
<td>5.73 B</td>
</tr>
</tbody>
</table>

*V.C. (%) 6.44

*mean followed by the same uppercase letter (column) do not differ by Duncan’s test (p ≤ 0.05). \textsuperscript{*} = not significant (p > 0.05).

These values of I\textsubscript{50} for biotypes B\textsubscript{4} and B\textsubscript{13} resulted in a resistance factor (RF) of 15 and 19 to iodosulfuron herbicide and a RF of 224 and 133 to imazethapyr herbicide, respectively (Figure 2, Table 2). High enzyme inhibition was observed in R. raphanistrum L. where the I\textsubscript{50} value in resistant and susceptible biotypes to clorosulfuron herbicide was 1.55 and 0.009 μM, respectively (Yu et al., 2012). The results demonstrated that B\textsubscript{4} and B\textsubscript{13} biotypes present high levels of resistance to the herbicide imazethapyr (Table 2). Similar results were reported for radish biotypes resistant to ALS inhibitor herbicides, for which the I\textsubscript{50} was higher for imazethapyr than for flumetsulam and metoxuron-methyl herbicides (Yu et al., 2012; Pandolfo et al.,...
Different levels of resistance to herbicides may be related to the position of the mutation sites in relation to the herbicide-coupling site (Yu et al., 2012; Pandolfo et al., 2016). Different levels of resistance to herbicides may be related to the position of the mutation sites in relation to the herbicide-coupling site (Yu et al., 2012; Pandolfo et al., 2016).

This mutation was identified in different weeds resistant to ALS inhibitor herbicides, including a recent discovery in biotypes of \textit{R. sativus} L. (Pandolfo et al., 2016). Mutation of the ALS gene can compromise the herbicide coupling to the target site of the enzyme and affect the weed control with ALS inhibitor herbicides (McCourt et al., 2006). The Trp-574-Leu substitution is considered the most relevant mutation because it confers resistance to all five chemical groups of the ALS inhibitor herbicides (Pandolfo et al., 2016). However, this mutation has not been reported in resistant biotypes of \textit{R. sativus} L. in Brazil; this can be considered the first identified case. In biotypes of \textit{R. raphanistrum} L., the most frequent mutations involve a proline at amino acid position 197 (Pro197), which confers resistance to the chemical groups of sulfonylureas and triazolopyrimidines (Tan & Medd, 2002; Yu et al., 2003). However, the levels of cross-resistance will depend on the position where the mutation occurred and the modified amino acid (Yu et al., 2012; Han et al., 2012).

A 1758 bp fragment of the ALS gene with five conserved regions was sequenced from the cDNA of susceptible (B1) and resistant (B4 and B13) radish biotypes. This region includes all domains with mutation points that were previously identified in biotypes of \textit{R. raphanistrum} L. resistant to ALS inhibitor herbicides (Tan & Medd, 2002; Yu et al., 2003; 2012; Han et al., 2012). The partial sequence presented a single nucleotide change of TGG to TTG, which led to a Trp-574-Leu substitution in resistant biotypes (Figure 3).

Figures 2 and 3 show the in vitro inhibition of ALS activity (%) of susceptible (B1) and resistant (B4 and B13) radish biotypes subjected to different concentrations of iodosulfuron (A) and imazethapyr (B) herbicides (µM). Points represent the mean values of repetitions in each biotype.

<p>| Table 2. | I50 values with confidence intervals (95% CI) and resistance factors (RF) of susceptible (B1) and resistant (B4 and B13) radish biotypes subjected to different concentrations of iodosulfuron and imazethapyr herbicides. |</p>
<table>
<thead>
<tr>
<th>Biotypes</th>
<th>I50</th>
<th>95 CI</th>
<th>Resistance factor</th>
</tr>
</thead>
<tbody>
<tr>
<td>Iodosulfuron (B1)</td>
<td>0.043</td>
<td>0.09 - 0.004</td>
<td>15</td>
</tr>
<tr>
<td>Iodosulfuron (B4)</td>
<td>0.65</td>
<td>1.05 - 0.25</td>
<td>19</td>
</tr>
<tr>
<td>Iodosulfuron (B13)</td>
<td>0.82</td>
<td>1.00 - 0.64</td>
<td>133</td>
</tr>
<tr>
<td>Imazethapyr (B1)</td>
<td>3.2</td>
<td>5.8 - 0.6</td>
<td>224</td>
</tr>
<tr>
<td>Imazethapyr (B4)</td>
<td>718</td>
<td>763.04 - 672.96</td>
<td>32,14</td>
</tr>
<tr>
<td>Imazethapyr (B13)</td>
<td>425</td>
<td>646.94 - 203.06</td>
<td>133</td>
</tr>
</tbody>
</table>

1I50 = dose required to inhibit 50% of ALS enzyme activity.

2RF obtained by division I50 of the resistant biotype by I50 of the susceptible biotype.

### ALS gene sequencing

A 1758 bp fragment of the ALS gene with five conserved regions was sequenced from the cDNA of susceptible (B1) and resistant (B4 and B13) radish biotypes. The partial sequence presented a single nucleotide change of TGG to TTG, which led to a Trp-574-Leu substitution in resistant biotypes (Figure 3).
Furthermore, all biotypes presented silent mutations at the 122 position that contains an alanine (Ala122), where the codon GCC was substituted for GCT (data not shown) without resulting in a modification to the amino acid sequence. Amino acid substitutions at Ala-122 are more unlikely due to the necessity of two nucleotide alterations to modify the amino acid (Yu et al., 2012). However, in *R. raphanistrum* biotypes, the substitution of Alanine 122 to Tyrosine (Ala-122-Tyr) conferred resistance to three chemical groups of ALS inhibitors herbicides (Han et al., 2012).

The mutation diagnosed here (Trp-574-Leu) is one of the most important because it confers resistance to all chemical groups of ALS inhibitor herbicides, thus reducing the control options for resistant biotypes (Pandolfo et al., 2016; Cechin et al., 2016).

**Metabolization**

The results demonstrated that the application of cytochrome P450 monooxygenase inhibitors followed by the application of iodosulfuron or imazethapyr herbicides did not efficiently control the resistant biotypes (Table 3). The isolated spraying of cytochrome P450 monooxygenase inhibitors did not cause phytotoxicity in any of the radish biotypes (data not shown).

**Table 3.** Control (%) in susceptible (B1) and resistant (B4 and B13) radish biotypes subjected to the application of iodosulfuron or imazethapyr herbicides alone or preceded by cytochrome P450 monooxygenase inhibitors at 28 days after application (DAA).

<table>
<thead>
<tr>
<th>Treatment</th>
<th>B1</th>
<th>B4</th>
<th>B13</th>
</tr>
</thead>
<tbody>
<tr>
<td>Iodosulfuron</td>
<td>99 a</td>
<td>3.0</td>
<td>0.0</td>
</tr>
<tr>
<td>Malathion + iodosulfuron</td>
<td>100 aB</td>
<td>4.0</td>
<td>21.0</td>
</tr>
<tr>
<td>PBO + iodosulfuron</td>
<td>99 a</td>
<td>2.0</td>
<td>4.0</td>
</tr>
<tr>
<td>Control</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
</tr>
</tbody>
</table>

V.C. (%) 10.00

<table>
<thead>
<tr>
<th>Treatment</th>
<th>B1</th>
<th>B4</th>
<th>B13</th>
</tr>
</thead>
<tbody>
<tr>
<td>Imazethapyr</td>
<td>97 a</td>
<td>1.0</td>
<td>1.0</td>
</tr>
<tr>
<td>Malathion + imazethapyr</td>
<td>97 a</td>
<td>0.0</td>
<td>0.0</td>
</tr>
<tr>
<td>PBO + imazethapyr</td>
<td>98 a</td>
<td>2.0</td>
<td>1.0</td>
</tr>
<tr>
<td>Control</td>
<td>0.0</td>
<td>0.0</td>
<td>0.0</td>
</tr>
</tbody>
</table>

V.C. (%) 6.95

*Means followed by the same lowercase letter (line) and the same uppercase letter (column) do not differ by Duncan’s test (p ≤ 0.05). "a" = not significant (p > 0.05).

However, in an experiment with ALS herbicide-resistant biotypes of *Poa annua* L., an increase in control by bispyribac-sodium herbicide was observed when malathion was used as an inhibitor of P450 monoxygenase. However, these results were not observed for other ALS enzyme inhibitors, indicating that the mechanism of resistance involved cannot be attributed to differential metabolism by P450 monoxygenase (Brosnan et al., 2016).

The differences found among the biotypes for control and SDM when malathion or PBO inhibitors where used indicate cytochrome P450 may be involved in metabolic resistance, which may be specific to a given herbicide.

Therefore, the results of control and SDM observed in B4 and B13 biotypes did not indicate the involvement of cytochrome P450 enzyme in resistance of radish to iodosulfuron and imazethapyr herbicides.
Conclusion

The kinetic parameters of ALS enzyme were not affected in resistant biotypes. A mutation was detected at position 574 of the ALS gene, which resulted in a substitution of tryptophan to leucine (Trp-574-Leu) in resistant radish biotypes. The results demonstrated that there was no differential metabolism of iodosulfuron and imazethapyr herbicides in resistant biotypes when cytochrome P450 was inhibited by malathion and piperonyl butoxide.

Acknowledgements

We thank the coordination of the Higher Education Personnel Training (CAPES) for the scholarship to the first author and also the Embrapa/Monsanto Partnership.

References


Received on July 2, 2016.
Accepted on October 18, 2016.

License information: This is an open-access article distributed under the terms of the Creative Commons Attribution License, which permits unrestricted use, distribution, and reproduction in any medium, provided the original work is properly cited.