Recent Discovery of *Amaranthus palmeri* S. Watson in Italy: Characterization of ALS-Resistant Populations and Sensitivity to Alternative Herbicides

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Abstract: *Amaranthus palmeri* S. Watson (Amaranthaceae Juss.) is a dioecious noxious weed, native to the Americas, which infests summer crops. It causes high crop losses, and rapidly evolves resistance to herbicides. In Europe, *A. palmeri* was recorded mostly as a casual alien, but in 2018 it was reported infesting a soybean field in Italy, and the next year two more populations were found in the same area. Experiments were conducted on these three populations to evaluate the resistance to ALS-inhibiting herbicides, to determine the main resistance mechanisms involved and assess the efficacy of alternative herbicides with different sites of action than ALS. The three populations were confirmed cross-resistant to ALS-inhibiting herbicides (thifensulfuron-methyl and imazamox). Gene sequencing identified a Trp to Leu substitution at position 574 of ALS gene in resistant plants, proving that the main resistance mechanism for the three populations is target-site related. The presence of other resistance mechanisms cannot be excluded. Metobromuron, metribuzin and glyphosate are still effective on these populations.

Keywords: herbicide resistance; palmer amaranth; alternative herbicides; soybean

1. Introduction

*Amaranthus palmeri* S. Watson (Amaranthaceae Juss.), commonly known as palmer amaranth or careless weed, is a dioecious noxious weed naturally distributed from Northern Mexico to southwest North America [1]. Outside its native range, *A. palmeri* occurs as alien species in all continents. In Europe it is considered as a casual alien in most countries and naturalized in a few (Belarus, Greece, Italy, Lithuania, Spain and Sweden), where it grows on human-made habitats, especially along roads [2]. In Italy, *A. palmeri* was found for the first time in 2014 in the Emilia Romagna region (NE-Italy) and recorded as casual along a major route [3]. The species was also recorded in Piedmont region (NW-Italy) on a riverbank and under a railway bridge [4]. In 2018, *A. palmeri* was found in a soybean field in the Veneto region (NE-Italy) [5]. In 2019, two more populations were found, 10 and 22 km far from that site, again in soybean. In 2020, the species was scouted for the first time in Lombardy, in a harvested corn field [6].

A wide range of pre-emergence and post-emergence herbicides are available to control *A. palmeri* in soybean. However, worldwide, *A. palmeri* has evolved resistance to eight sites of action (SoA) [7] and a population from Kansas has evolved multiple resistance to up to six SoA [8]. All records of herbicide resistant *A. palmeri* were found in the United States until 2008, when it was first reported in Israel [9], then in Brazil [10], Argentina [11] and, more recently, in Spain, where the introduction of already resistant biotypes was suspected [12].
The most common mechanism endowing ALS inhibitors resistance in *A. palmeri* is target-site mediated and involves several point mutations at the ALS gene [13], but a few cases of non-target-site mechanism have recently been reported [8,14]. Due to both economic and environmental factors, post-emergence application of ALS-inhibiting herbicides is the most common weed control strategy in soybean in Italy, and it is likely one of the driving factors of the recent spreading of ALS-resistant amaranths [15]. Several herbicides with alternative sites of action, such as inhibitors of photosystem (PS) I and II, 5-enolpyruvylshikimate-3-phosphate (EPSP) synthase and carotenoid biosynthesis can provide effective control of ALS-resistant populations of *A. palmeri* [16,17]. However, given the ability of this species to evolve resistance against herbicides with different sites of action [18], the efficacy of potential alternative herbicides should be assessed at local level before recommending their use.

The aims of this work were (i) to verify the resistance status to ALS-inhibiting herbicides of three Italian *A. palmeri* populations; (ii) to determine the main resistance mechanisms involved; and (iii) to evaluate the efficacy of alternative herbicides with different sites of action than ALS.

2. Materials and Methods

2.1. Plant Material

Three *A. palmeri* populations were collected in north-eastern Italy in soybean fields where poor herbicide control was suspected. For each population, seeds were collected from at least 30 female plants, randomly distributed across the field. Population 18-100 was collected in 2018 and a voucher herbarium specimen was deposited at La Sapienza University of Rome (Herbarium Flaminio, HFLA) [5]. Populations 19-174 and 19-177 were collected in 2019. In addition, an American susceptible *A. palmeri* check (kindly provided by Prof. T. A. Gaines [19]) was used as susceptible reference population (20-179).

2.2. Herbicide Bioassays

Bioassays with post-emergence and pre-emergence herbicides were conducted twice, in a greenhouse located in north-eastern Italy (45°21' N, 11°58' E). Greenhouse temperatures varied between 15 and 20 °C and from 25 to 34 °C, during the night and day, respectively.

For experiments with herbicides applied in post-emergence, seed germination, seedlings growth and herbicide treatments were done as previously described [20]. In brief, seeds were sown in transparent plastic boxes (diameter 12 cm, height 5 cm) containing 50 mL of 0.6% agar; after 1 week at 4 °C (dark), the boxes were transferred into a germination cabinet at alternating light and temperature to stimulate germination: 12 h light, 28 °C and 12 h dark, 18 °C. After 5 days, seedlings were transplanted into plastic trays (325 × 265 × 95 mm³, 20 plants each) with a standard potting mix (60% clay loam soil, 15% sand, 15% perlite and 10% peat) and watered as required. Just prior to the treatment, plants in each tray were counted. Three- to four-leaf stage plants, corresponding to growth stage 13–14 of the BBCH scale [21], were treated with ALS inhibitors at the recommended field rates (1×) and three times that (3×): thifensulfuron-methyl 6 and 18 g a.i. ha⁻¹ (Harmony 50 SX, DuPont™, 500 g a.i. 1 kg⁻¹), imazamox 40 and 120 g a.i. ha⁻¹ (Tuareg®, DuPont™, 40 g a.i. L⁻¹). Glyphosate was applied at the field rate: 480 g a.i. ha⁻¹ (Roundup Platinum®, Monsanto, 480 g a.i. L⁻¹). Agrochemicals were applied using a precision bench sprayer delivering 300 L ha⁻¹ at a pressure of 215 kPa and speed of about 0.75 m s⁻¹, with a boom equipped with three flat-fan (extended range) hydraulic nozzles (Teejet, 11002). The experimental layout was a complete randomized design with two replicates (trays).

Four weeks after herbicide application, the number of surviving plants and visual estimation of their biomass (VEB) were assessed. The VEB scores, ranging from 100 (for plants not affected by the herbicide, equal to the untreated control) to 0 (when the plants were clearly dead), were given to each treated tray. Standard error (SE) was calculated for each data mean.
For experiments with herbicides applied in pre-emergence, the same kind of trays used for post-emergence experiments were filled with a potting mix with no peat. A layer of sterilized clay loam soil was added on top (3 days at 105 °C, 2–3 cm, organic matter 1.1 mg kg⁻¹ of dry matter). Seeds were soaked in a transparent plastic box with a wet paper filter for 3 days at 4 °C (dark), then transferred into a germination cabinet for 24 h, 12 h light at 28 °C and 12 h dark at 18 °C to stimulate germination. Seeds were then gently wiped with paper and sown in trays (50 seeds per tray). Seeds were then covered with a layer of 1–2 mm of the sterilized soil. An aluminum foil pan was placed at the base of each tray and water was distributed both by sprinkler irrigation and by filling the pan to maintain soil moisture. The experimental layout was a complete randomized design with 3 replicates (trays) per treated thesis and 4 replicates per non-treated thesis. Twenty-four hours after sowing, pre-emergence herbicides were applied at the field rates: metribuzin 245 g a.i. ha⁻¹ (Feinzin® 70 DF, Adama, 70 g a.i. 100 g⁻¹), metobromuron 1500 g a.i. ha⁻¹ (Proman Flow®, Belchim Crop Protection, 500 g a.i. 1000 g⁻¹), clomazone 144 g a.i. ha⁻¹ (Sirtaki®, Sipcam, 360 g a.i. 1000 mL⁻¹). Pre-emergence herbicide application was done with the same precision bench sprayer and same settings as for the post-emergence application. Two weeks after herbicide application, the number of seedlings and shoot fresh weight were recorded. The percentage of plant survival was calculated as the ratio between the emerged seedlings of treated trays with respect to those of non-treated trays. The relative fresh weight was calculated as the ratio between the fresh weight of treated trays with respect to that of non-treated trays. Standard error (SE) was calculated for each data mean.

2.3. DNA Extraction and Detection of Mutated ALS Alleles

DNA was extracted from 15 plants that survived the field dose of thifensulfuron-methyl treatment, following a CTAB protocol [22]. The domain B of ALS was amplified using the PCR protocol previously described for Amaranthus retroflexus [19], with primers AMA-2F (5′-TCCCGGGTTAAAATCATGCTC-3′) and AMA-2R (5′-CTTCTTCCATCACCCCTCTGT-3′). PCR were performed using GoTaq® G2 Hot Start Polymerase (Promega) in a 25 µL mixture including 5 µL of 5× Green GoTaq Flexi Buffer, dNTPs mix (0.2 mM each), MgCl2 (1.5 mM), forward and reverse primers (0.2 µM), 0.125 µL GoTaq DNA Polymerase and 50 ng DNA. Amplification conditions: 2 min at 95 °C; 35 cycles of 30 s at 95 °C, 30 s at 58 °C, 40 s at 72 °C; 5 min at 72 °C. PCR products were purified using NucleoSpin® Gel and PCR Clean-up kit (Macherey-Nagel GmbH & Co., Duren, Germany) following manufacturer’s instructions. Once purified, both strands of the PCR products obtained from each plant were Sanger-sequenced by BMR Genomics (Padova, Italy) and edited with FinchTV 1.4.0. Sanger sequencing data (ab1 files) were deposited in a dedicated repository [23].

3. Results

3.1. Resistance to ALS Inhibitors

The percentage of plant survival and the visual estimation of biomass (VEB) between the two experiments were comparable (considering the SE); therefore, the data were pooled, and mean values were reported. The susceptible check (20-179) was completely controlled by both thifensulfuron-methyl (SU) and imazamox (IMI) at the field rate (Figure 1). Conversely, for populations 18-100, 19-174 and 19-177, a high percentage of plants survived both thifensulfuron-methyl and imazamox. At the recommended field rate, plant survival ranged from 77% to 86% for imazamox and from 83% to 100% for thifensulfuron-methyl. The VEB of the ALS-treated plants was also high, with values ranging from 78% to 94% and from 86% to 93% for imazamox and thifensulfuron-methyl, respectively, indicating a negligible effect of both herbicides on biomass production. Similar survival rates and VEB were recorded when the plants of the three populations were treated at three times the field rates of both ALS inhibitors (Figure 1).
Figure 1. Response of A. palmeri populations treated with imazamox (IMI) and thifensulfuron-methyl (SU) at the field rate (1×) and three-times that (3×). Blue bars refer to the percentage of plant survival and red bars to the visual estimation of biomass (VEB). Vertical thin bars represent the standard error.

3.2. Response to Non-ALS Herbicides

All populations were completely controlled by glyphosate and no plants survived (data not shown).

In the experiments with pre-emergence herbicides, the emergence of seedlings in non-treated trays started two days after sowing. The percentages of emerged seedlings were significantly different between the two experiments (Table 1). For all populations, the efficacy of metobromuron and metribuzin was near 100% in both experiments. Instead, the percentage of plant survival and fresh weight of clomazone-treated plants varied significantly between experiments. Therefore, the data were not pooled (Figure 2, Table 2). The percentage of control for the susceptible A. palmeri population 20-179 ranged between 63 and 72%, while for the ALS-resistant populations the percentage of control ranged from (approximately) 1 to 74% (Figure 2). The relative fresh weight of the plants of population 20-179 that survived to clomazone ranged between 3% and 22%, while the fresh weight of the ALS-resistant populations (18-100, 19-174 and 19-177) was generally higher, with values ranging from 17 to 100% (Table 2). At the field dose, if the efficacy is lower than the 80% or the fresh weight of survivors exceed 20% of the fresh weight of non-treated controls, the chemical control is incomplete a population might be considered at risk of evolving resistance.

Table 1. Percentage of emerged seedlings of the four A. palmeri populations in non-treated conditions 14 days after sowing. Standard error in brackets.

| Population   | 20-179 | 18-100 | 19-174 | 19-177 |
|--------------|--------|--------|--------|--------|
| Experiment 1 | 69 (±3) | 49 (±5) | 32 (±4) | 35 (±1) |
| Experiment 2 | 98 (±7) | 83 (±4) | 23 (±1) | 59 (±4) |

Table 2. Relative fresh weight of A. palmeri plants that survived clomazone, calculated as the ratio between the fresh weight of treated trays with respect to non-treated trays, in the first and second experiment. Standard error in brackets.

| Population   | 20-179 | 18-100 | 19-174 | 19-177 |
|--------------|--------|--------|--------|--------|
| Experiment 1 | 3 (±1)  | 51 (±7) | 17 (±7) | 100 (±7) |
| Experiment 2 | 22 (±14)| 24 (±7) | 65 (±15)| 39 (±13) |
were heterozygous, four out of fifteen plants had no mutations at this locus, and three had no mutations at this locus. Nine out of fifteen plants of population 19-177 were homozygous, seven out of fifteen plants were heterozygous, while only two had no mutations at this locus. Five out of fifteen plants of population 19-174 were homozygous, seven out of fifteen plants were heterozygous, while three had no mutations at this locus. Nine out of fifteen plants of population 19-177 were heterozygous for the mutation, while only two had no mutations at this locus. Thirteen out of fifteen plants of population 18-100 were heterozygous for the ALS gene mutation, while only two had no mutations at this locus. Five out of fifteen plants of population 18-100 were heterozygous for the ALS gene mutation, while only two had no mutations at this locus.

### 3.3. ALS Gene Mutation

A point mutation (TGG to TTG), which caused an amino acid change from tryptophan to leucine, was found at codon 574 of ALS in most plants of all resistant populations (Figure 3). Thirteen out of fifteen plants of population 18-100 were heterozygous for the mutation, while only two had no mutations at this locus. Five out of fifteen plants of population 19-174 were homozygous, seven out of fifteen plants were heterozygous, while three had no mutations at this locus. Nine out of fifteen plants of population 19-177 were heterozygous, four out of fifteen plants had no mutations at this locus, and sequencing failed for two plants. Sanger sequencing data (ab1 files) are freely accessible through a dedicated repository [23].

![Chromatogram detail of ALS nucleotide sequences obtained from population 17-174, showing different point mutation at codon 574 (highlighted by the orange frame). TGG codon encodes for the amino acid tryptophan, while TTG for leucine. The double peak indicates that the plant is heterozygous.](image)

**Figure 3.** Chromatogram detail of ALS nucleotide sequences obtained from population 17-174, showing different point mutation at codon 574 (highlighted by the orange frame). TGG codon encodes for the amino acid tryptophan, while TTG for leucine. The double peak indicates that the plant is heterozygous.

### 4. Discussion

Three populations of *A. palmeri* were confirmed to be cross-resistant to thifensulfuronmethyl and imazamox. The negligible effect of these herbicides on plant biomass and the presence of the Trp574Leu substitution at ALS, known to confer broad spectrum resistance to ALS inhibitors in various weed species [12], proved that the main resistance mechanism is target-site related. Some plants had no mutations at position 574: the presence of other mutations or other resistance mechanisms cannot be excluded. However, target-site resis-
tance is the main mechanism that has been reported in other *A. palmeri* populations. In Arkansas, a study conducted in 13 counties revealed that plants frequently had this mutation conferring a broad cross-resistance to all ALS herbicide families [24]. The Trp574Leu substitution has also been found in Brazilian *A. palmeri* populations [25]. Population 18-100 was the third record of *A. palmeri* in Italy and the first for the Veneto Region [5]. Finding it already resistant to ALS herbicides was surprising and alarming. A similar situation has recently been observed in Italy with the congeneric weed species *Amaranthus tuberculatus* (Moq.) J.D. Sauer, that has spread rather fast since appearing in agronomic habitats. Numerous ALS-resistant populations were reported in several locations [15], very likely after the introduction of ALS-resistant biotypes from outside Italy [26]. It is worth mentioning that *A. tuberculatus* had been present in Italy since the 1980s; therefore, the evolution of herbicide resistance from resident unnoticed populations was still possible (although unlikely). Instead, in the case of these three *A. palmeri* populations, the evolution of herbicide resistance from naturally occurring populations is very unlikely; rather, a very recent introduction from outside Italy appears to be the most plausible hypothesis. The presence of the same resistance-endowing mutation in most resistant individuals of the three populations clearly indicates that field-applied ALS inhibitors exerted a strong selective pressure. ALS-resistant populations of *A. palmeri* have also been described in Spain [12]. In that case, the populations were found along roadsides and field borders, while no persistent field infestations were reported. In those Spanish populations, the low frequency of several different mutations at ALS locus, conferring resistance with different patterns may reflect a weak selective pressure due to ALS inhibitors.

The recommended field rates of glyphosate, metribuzin and metobromuron completely controlled the three ALS-resistant *A. palmeri* populations. Glyphosate and metribuzin are known to control *A. palmeri* [27,28], whereas little information is available on the efficacy of metobromuron applied alone. Our results show that all three herbicides should be considered valuable options to control these *A. palmeri* populations. Instead, the efficacy of clomazone was weak, with variable responses between the ALS-resistant populations and an incomplete control of the susceptible check. This is in keeping with the results from other studies [29,30] and with the indication that *Amaranthus* spp. is a “medium susceptible species” reported on the label of most herbicides containing clomazone as single active ingredient. The use of this active ingredient should be evaluated only if required to control other weed species (such as velvetleaf, *Abutilon theophrasti* Medik) and always in mixture with other active ingredients.

The early detection of *A. palmeri* in fields is crucial, especially when herbicide resistant. The recognition of *A. palmeri* might be easier if compared with other congeneric species, because it is morphologically well characterized [1]. Most of the *Amaranthus* species that are widely diffused in agricultural environments are monoecious while *A. palmeri*, together with *Amaranthus tuberculatus* (Moq.) J.D. Sauer, are dioecious species, thus male and female plants should be observed in the field. Flowers of female plants of *A. palmeri* normally have five tepals, while flowers of female plants of *A. tuberculatus* normally have 0–3 [31]. A simplified botanical key was previously developed to recognize some of the most common *Amaranthus* species in agronomic habitats [15].

Given the high fertility and dispersion ability of *A. palmeri*, a zero-tolerance strategy should be adopted to avoid the spread of seeds. An experiment conducted in a glyphosate-resistant cotton field in the USA demonstrated that 20,000 seeds distributed in a square meter (simulating a single glyphosate resistant female palmer amaranth seed output), can infest 95–100% of the field area, causing a complete harvest loss in three years [32]. At crop harvest, infested patches should be avoided by harvest machinery [33]. Crop and herbicide rotation, mechanical weeding, hand weeding (whenever possible) and accurate cleaning of agricultural machinery are crucial to avoid further evolution of resistance. Innovative tools and tactics are progressively becoming available for farmers. For example, the adoption of harvest weed seed control (HWSC) systems could significantly contribute to the control of *A. palmeri* given its high seed retention until crop harvest [34,35]. With 1 million tons
(Mt) produced in 2019, Italy is the third soybean producer in Europe (after Ukraine, 3.6 Mt, and the Russian Federation, 4.3 Mt), and the first in the European Union (before Serbia, 0.7 Mt; Romania, 0.4 Mt; and France, 0.4 Mt) [36]. Since we are, very likely, at the initial stage of the invasion and the disrupting potential of this weed is well known, every effort to eradicate it is profitable.

5. Conclusions

Amaranthus palmeri had not been recorded in Italy before 2015. The detection of three ALS-resistant populations of A. palmeri in soybean is an alarming event that needs to be adequately followed up. These A. palmeri populations are cross-resistant to both thifensulfuron-methyl and imazamox. The main resistance mechanism is due to the well-known Trp to Leu substitution at position 574 of ALS. The presence of more than one resistance mechanism cannot be excluded. Nevertheless, herbicides with different sites of action than ALS (glyphosate, metribuzin and metobromuron) still control these populations.

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