



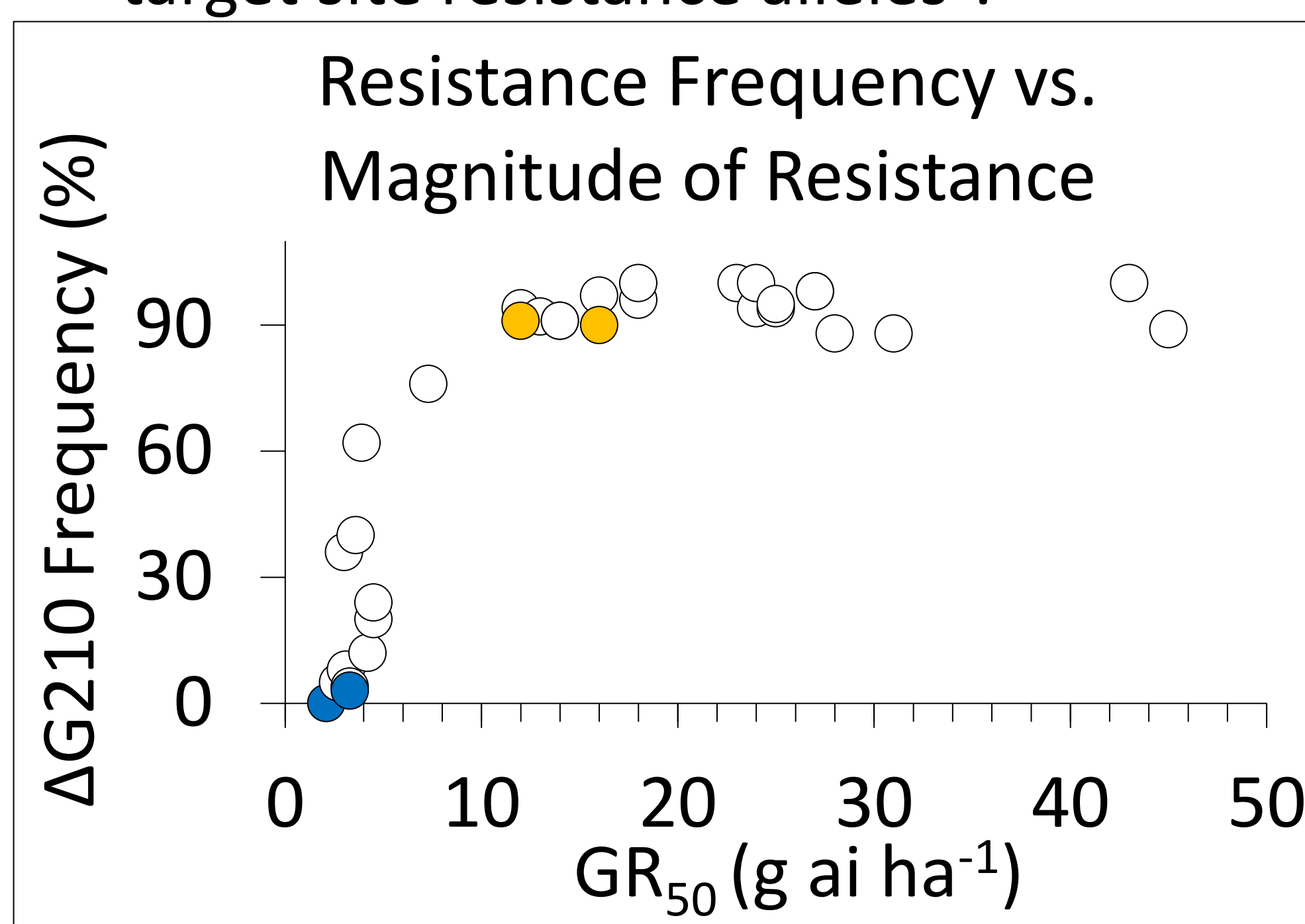
# Is Detoxification Contributing to PPO Inhibitor Resistance in Waterhemp (*Amaranthus tuberculatus*)?

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## Introduction

- PPO inhibitor resistance (PPO-R) in waterhemp threatens the long-term utility of PPO inhibiting herbicides<sup>1</sup>.
- A 2016 survey of PPO-R in the Midwest revealed a diversity of resistance responses from 4- to 17-fold that was unexplained by presence of R128 mutations, R allele zygosity, or frequency of resistant individuals (Figure 1).
- Non-target site resistance alleles can be a large contributor to overall resistance phenotype in the same individuals carrying target site resistance alleles<sup>2</sup>.



**Figure 1.** Resistance frequency of waterhemp populations as explained by  $GR_{50}$ . Resistance frequency accounts for a large portion of the change in  $GR_{50}$  until approximately  $10 \text{ g ai ha}^{-1}$  ( $R^2=0.52$ ). Blue dots indicate known S populations and yellow dots indicate known R populations.

- The compounds NBD-Cl and Malathion inhibit Glutathione S-transferases and Cytochrome P450s<sup>3</sup>, respectively.
- These Phase I and Phase II metabolism inhibitors have been used to demonstrate detoxification (detox) based herbicide resistance.<sup>4-7</sup>

## Hypothesis and Objective

**Hypothesis:** Enhanced fomesafen detoxification is contributing to the increased resistance response in two populations from the 2016 survey (IL-WAS and IN-DUB).

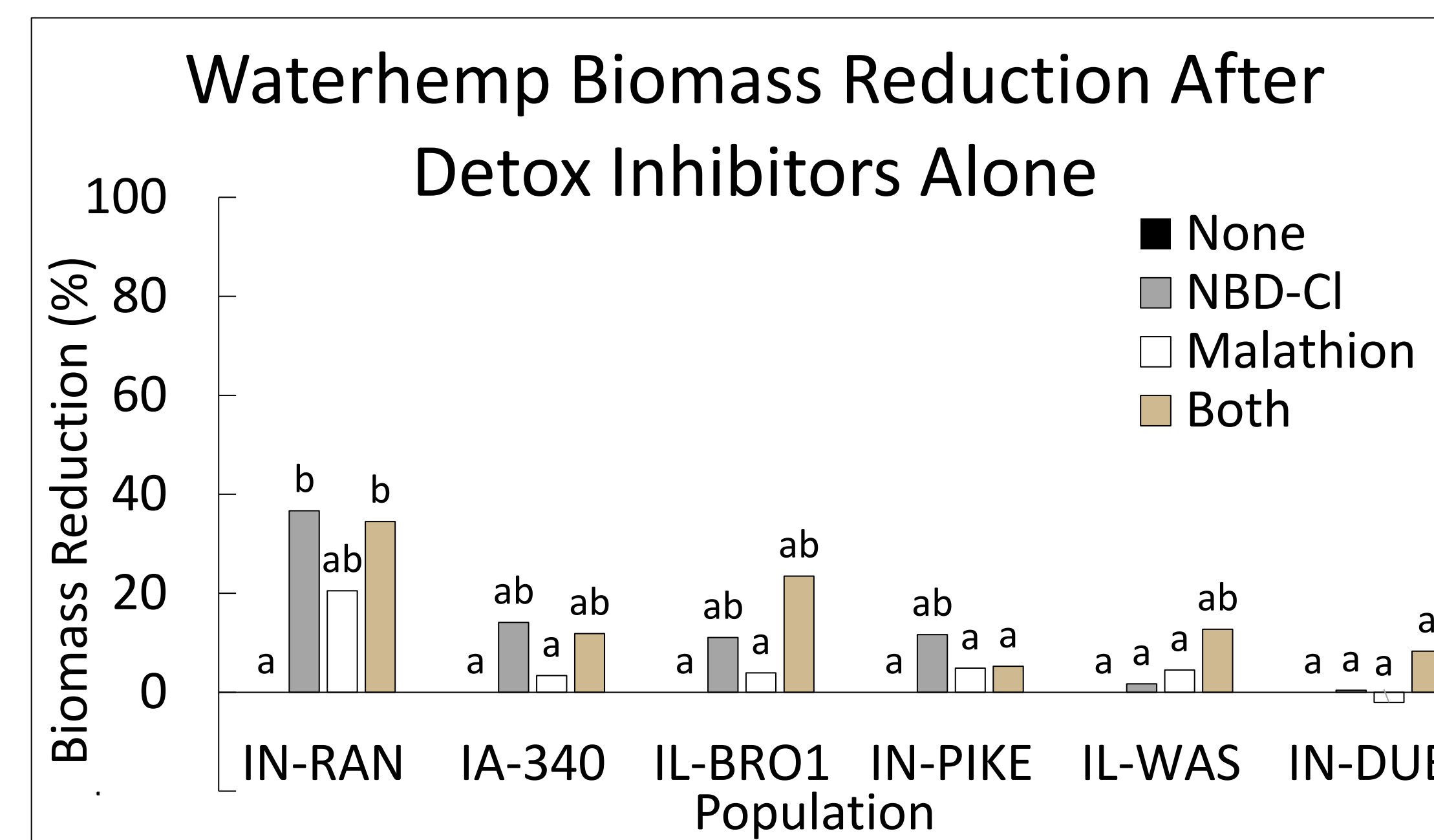
**Objective:** Determine the potential for co-occurrence of metabolic and target site resistance to PPO-inhibitors in two waterhemp populations.

## Materials and Methods

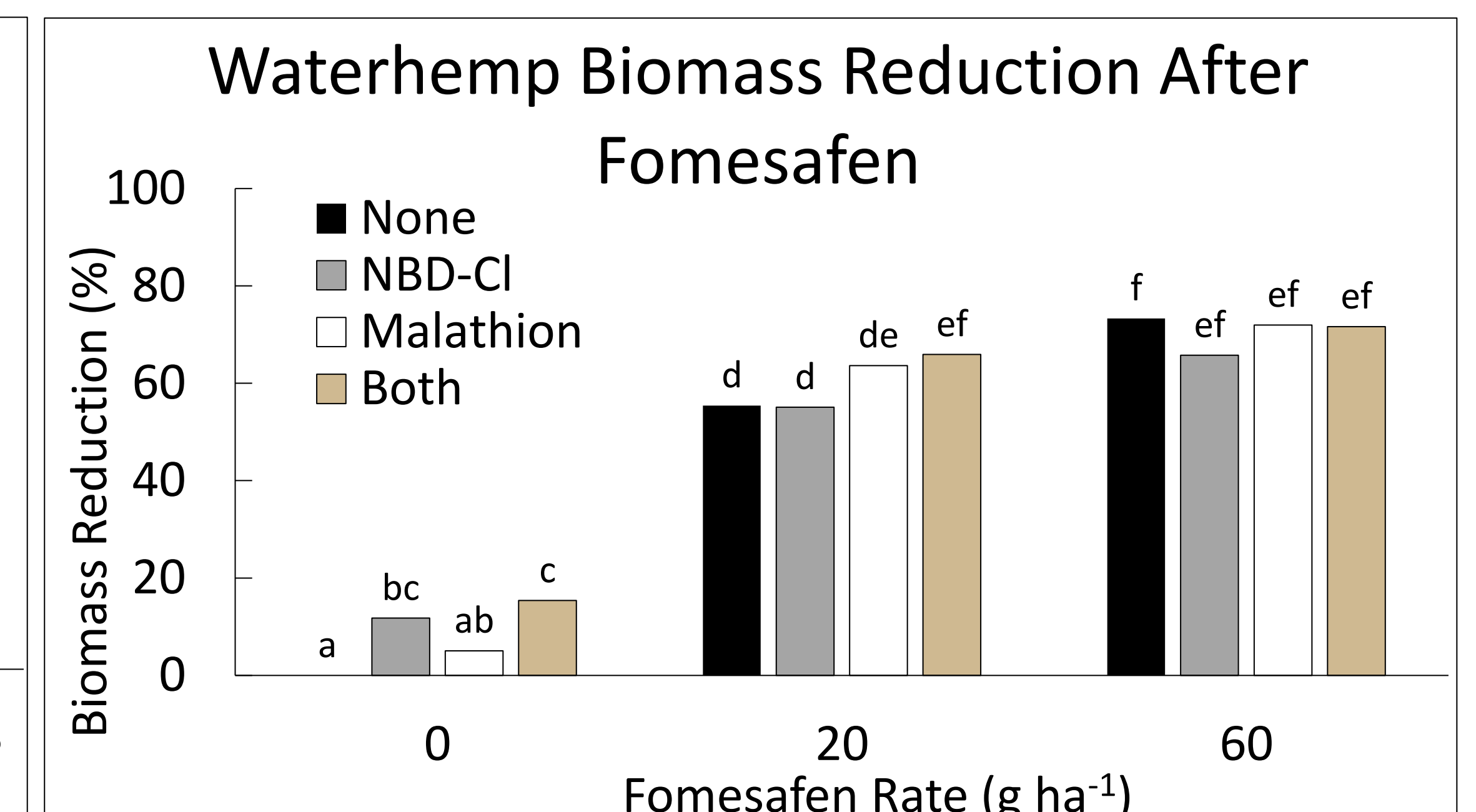
- Waterhemp from six populations was germinated in flats and then transplanted into  $3.8 \times 25 \text{ cm}$  tubes.
  - Susceptible (IN-RAN1 and IA-340)
  - Moderately resistant (IL-BRO1 and IN-PIKE1)
  - Highly resistant (IL-WAS and IN-DUB)
- Plants were sprayed with either NBD-CL ( $270 \text{ g ha}^{-1}$ ), malathion ( $1500 \text{ g ai ha}^{-1}$ ), NBD-Cl fb malathion, or blank DMSO solvent solution.
  - NBD-Cl and DMSO solvent solution applied 2d prior to fomesafen application.
  - Malathion applied 2h prior to fomesafen application.
- Plants were sprayed with rates of 0, 20, or  $60 \text{ g ai ha}^{-1}$  of fomesafen at the 5- to 7-leaf stage.
- Waterhemp control, height, and biomass data were collected at 7 and 14 Days after fomesafen treatment.
- Data were analyzed using PROC GLIMMIX in SAS 9.4 to conduct a three-way ANOVA.
- Means separated using Tukey's HSD at  $\alpha=0.05$ .

## Results and Discussion

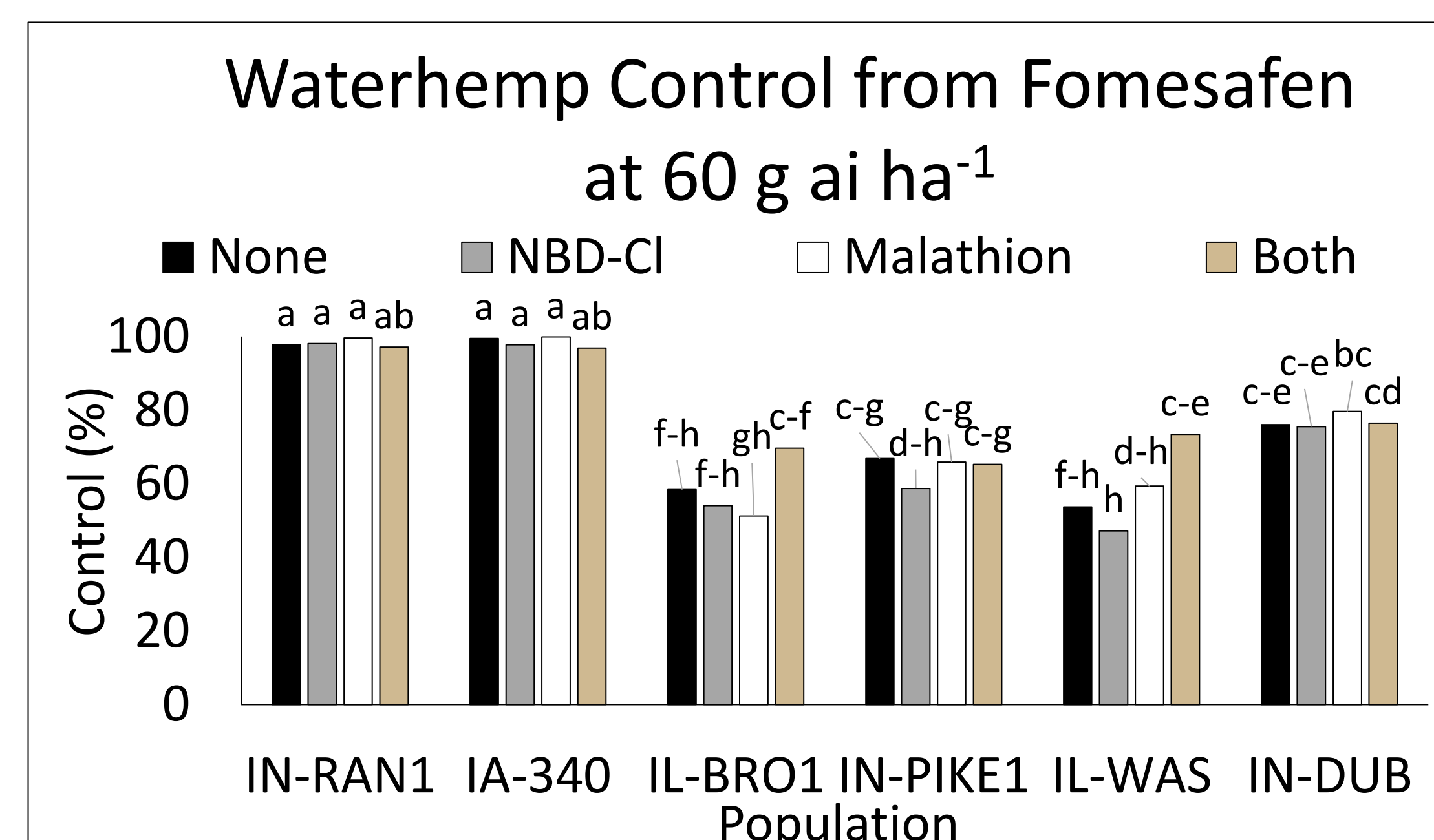
- Noticeable stunting was present for detox-inhibitor treatments similar to other groups<sup>6,7</sup> and was variable by population (Figure 2).
- At the 0 and  $20 \text{ g ha}^{-1}$  rate, NBD-Cl + Malathion increased fomesafen efficacy over all populations by 15 and 10 percentage points, respectively (Figure 3).
- At the  $60 \text{ g ha}^{-1}$  rate, NBD-Cl + Malathion increased control of IL-WAS by 20 to 26 percentage points compared to DMSO blank and NBD-Cl alone (Figure 4, Figure 5).



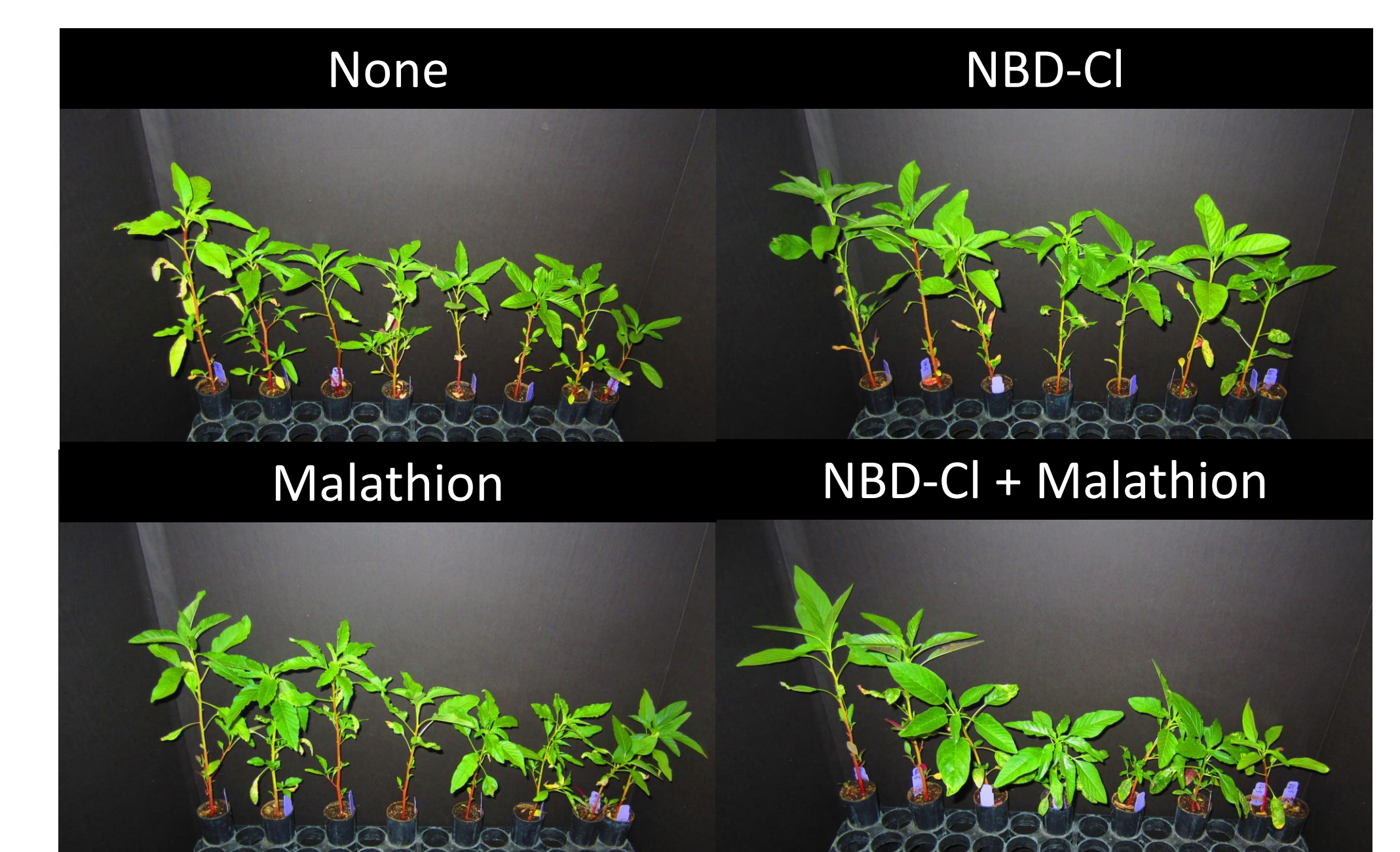
**Figure 2.** Biomass reduction 14 days after treatment with detox-inhibitors NBD-Cl and Malathion on waterhemp from six populations. Bars with different letters indicate significant difference between treatments.



**Figure 3.** Biomass reduction 14 days after treatment with detox-inhibitors in combination with fomesafen. Bars with different letters indicate significant difference between treatments.



**Figure 4.** Waterhemp control 14 days after application of  $60 \text{ g ai ha}^{-1}$  of fomesafen and detox-inhibitor treatment. Bars with different letters indicate significant difference between treatments.



**Figure 5.** IL-WAS sprayed with detox-inhibitor treatments followed by  $60 \text{ g ha}^{-1}$  of fomesafen.

## Conclusions

- Data were inconclusive with current methods and the hypothesis could not be fully tested.
- Evidence of detoxification based resistance should be pursued with additional methods such as LC-MS quantification of herbicide or detection of  $^{14}\text{C}$  labeled herbicide metabolites.

## Future Research

- Perform LC-MS analysis of fomesafen degradation through time in resistant populations.
- Investigate the variable response of different populations to malathion and NBD-Cl.

## References

- <sup>1</sup>Heap 2022 International Survey of Herbicide Resistant Weeds <http://www.weedscience.com>. Accessed November 1, 2022.
- <sup>2</sup>Kreiner JM, Tranel PJ, Weigel D, Stinchcombe JR, Wright SI (2021) The genetic architecture and population genomic signatures of glyphosate resistance in *Amaranthus tuberculatus*. *Mol Ecol*
- <sup>3</sup>Cummins I, Wortley DJ, Sabbadin F, He Z, Coxon CR, Straker HE, Sellars JD, Knight K, Edwards L, Hughes D, Kaundun SS, Hutchings SJ, Steel PG, Edwards R (2013) Key role for a glutathione transferase in multiple-herbicide resistance in grass weeds. *Proc Natl Acad Sci U S A* 110:5812–5817
- <sup>4</sup>Ma R, Kaundun SS, Tranel PJ, Riggins CW, McGinness DL, Hager AG, Hawkes T, Mc El, Riechers DE (2013) Distinct detoxification mechanisms confer resistance to mesotrione and atrazine in a population of waterhemp. *Plant Physiol* 163:363–377
- <sup>5</sup>Oliveira MC, Gaines TA, Dayan FE, Patterson EL, Jhala AJ, Knezevic SZ (2018) Reversing resistance to tembotrione in an *Amaranthus tuberculatus* (var. *rudis*) population from Nebraska, USA with cytochrome P450 inhibitors. *Pest Manag Sci* 74:2296–2305
- <sup>6</sup>Rangani G, Noguera M, Salas-Perez R, Benedetti L, Roma-Burgos N (2021) Mechanism of Resistance to S-metolachlor in Palmer amaranth. *Front Plant Sci* 0:392
- <sup>7</sup>Varanasi VK, Brabham C, Norsworthy JK (2018) Confirmation and Characterization of Non-target site Resistance to Fomesafen in Palmer amaranth (*Amaranthus palmeri*). *Weed Sci* 66:702–709