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Genetic Variability of the 5-Enolpyruvylshikimate-3-Phosphate (EPSP) Synthase Gene in Round-Up™ Resistant Soil Bacteria

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Genetic Variability of the 5-Enolpyruvylshikimate-3-Phosphate (EPSP) Synthase Gene in Round-Up™ Resistant Soil Bacteria

Abstract

Global use of Round-up™ has had a profound effect on agricultural systems. While there is a clear challenge with the emergence of resistant weeds, less emphasis has been placed on its impacts regarding the soil ecosystem. In this study, we sought to identify glyphosate resistance in two specific soil bacteria, *Bacillus subtilis* and *Pseudomonas syringae* pv. *glycinea*. To accomplish this purpose, selective media and colony PCR were used to isolate and identify colonies with glyphosate resistance. The presence of glyphosate resistance in soil bacteria was verified, and continuing work is being done to analyze sequence data that may lead to a better biochemical understanding of how resistance to glyphosate is achieved in various soil microorganisms.

Keywords

glyphosate, chemical resistance, Round-Up™, bacteria, genes, organisms, microbial mutation, enzymes

Disciplines

Soil Science

Comments

Poster presented at the University of Iowa Better FUTURES for Iowans Symposium in Iowa City, Iowa, on July 31, 2015.



Genetic variability of the 5-enolpyruvylshikimate-3-phosphate (EPSP) synthase gene in Round-up™ resistant soil bacteria

Dr. Jeffrey Ploegstra, Seth Steenwyk, Hannah Van Maanen
July 31, 2015
University of Iowa, Better Futures for Iowans



ABSTRACT

Global use of Round-up™ has had a profound effect on agricultural systems. While there is a clear challenge with the emergence of resistant weeds, less emphasis has been placed on its impacts regarding the soil ecosystem. In this study, we sought to identify glyphosate resistance in two specific soil bacteria, *Bacillus subtilis* and *Pseudomonas syringae* pv. *glycinea*. To accomplish this purpose, selective media and colony PCR were used to isolate and identify colonies with glyphosate resistance. The presence of glyphosate resistance in soil bacteria was verified, and continuing work is being done to analyze sequence data that may lead to a better biochemical understanding of how resistance to glyphosate is achieved in various soil microorganisms.

BACKGROUND

Round-up™ ready technology has been highly successful. As such, application of Round-up™ is increasing. Over 1.4 billion pounds are applied annually in 160 countries around the world.[†] Genetically modified Round-up™ Ready crops are protected from the herbicide while the susceptible weeds are killed.

However, glyphosate is also an antibiotic. Research has already looked into Round-up™ resistant weeds, but less research has been conducted on how glyphosate application affects soil microbe populations.

Glyphosate, the active ingredient, targets the EPSP synthase enzyme in the Shikimate pathway. Because animals and humans do not utilize this metabolic route, Round-up™ has a low toxicity for them. However, this pathway is common to plants, bacteria, and fungi.

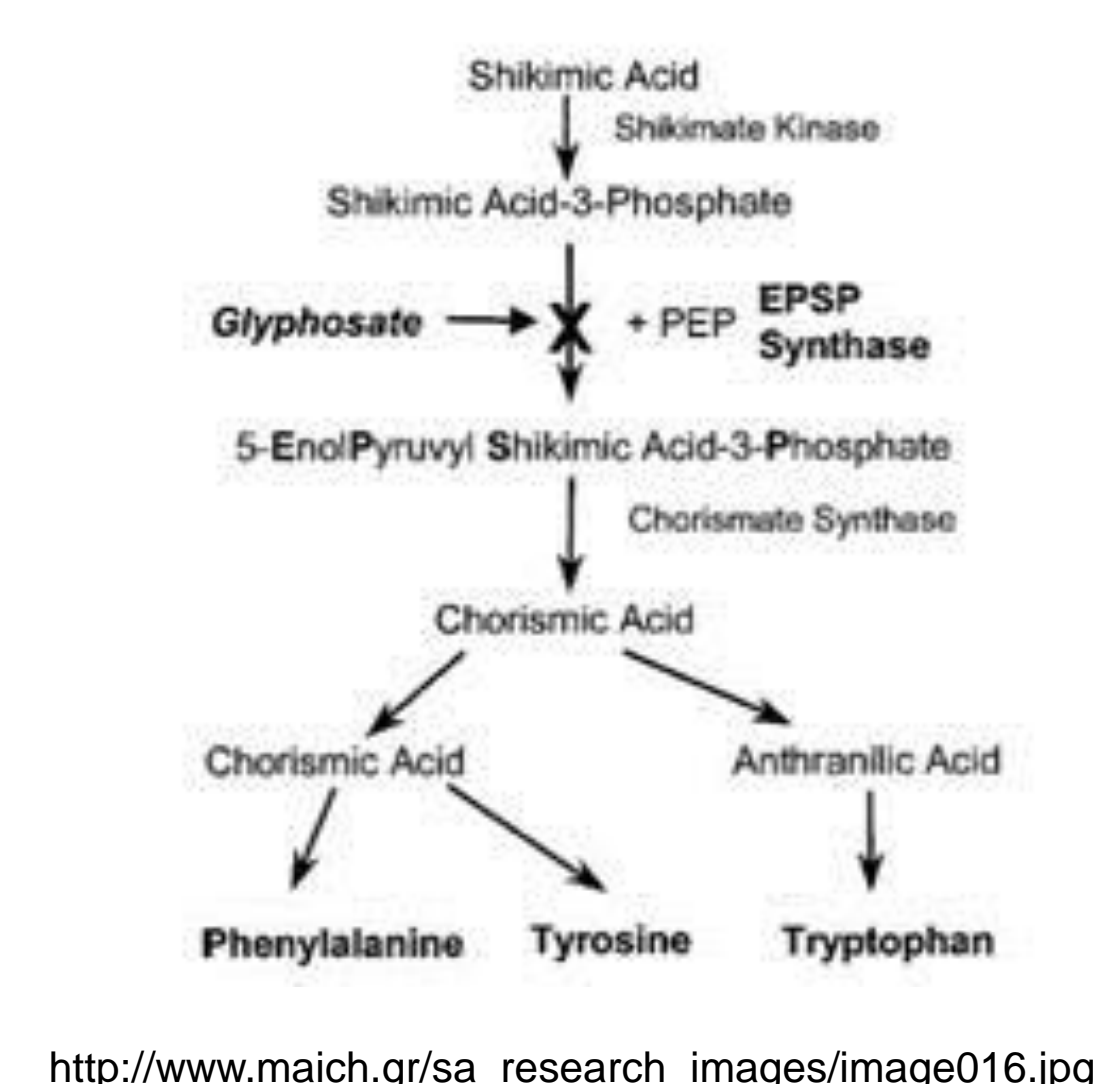
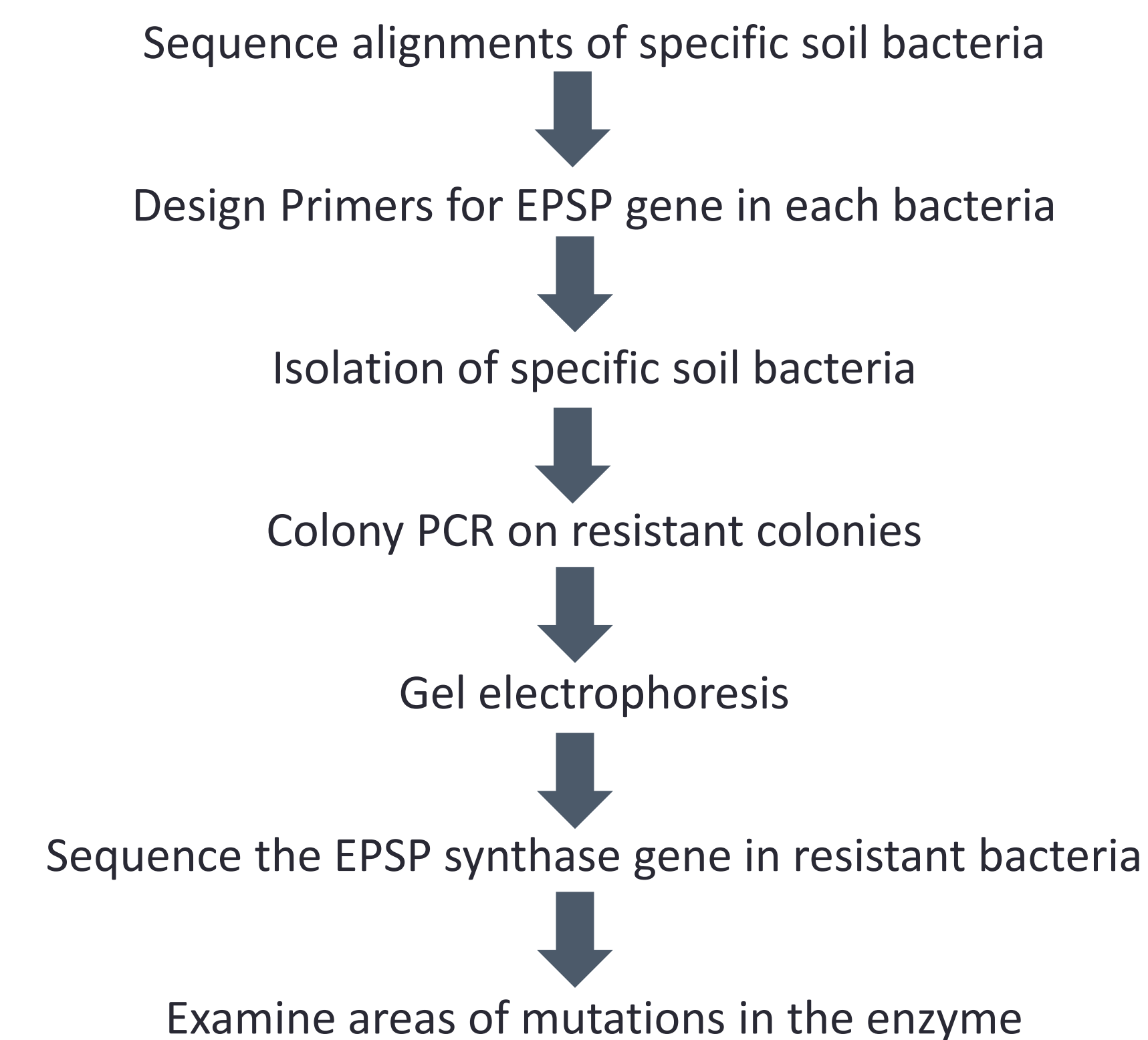
Soil microbes are an essential part of all ecosystems. The implications of alterations in the soil microbial communities in row crops is poorly understood. This research represents an opportunity to broaden the scope of assessment for the impact of agricultural interventions on overall ecosystem quality.

[†]Grossman, 2015: <http://news.nationalgeographic.com/2015/04/150422-glyphosate-roundup-herbicide-weeds/>

OBJECTIVES

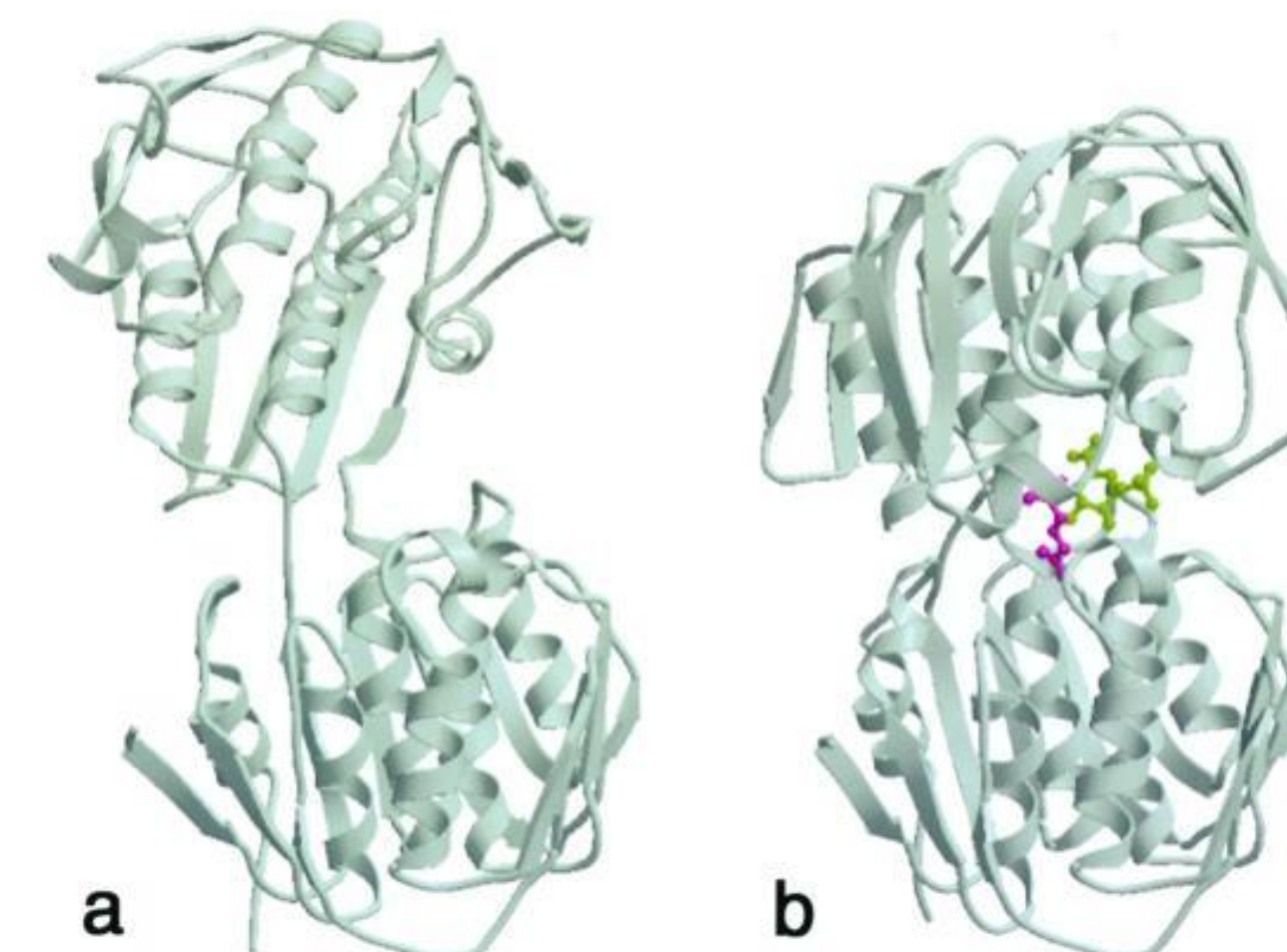
- Better understand Round-up™ resistance in beneficial and pathogenic microbes
 - Isolate and identify target soil organisms based on morphology and resistance to glyphosate
 - Create “tool kit” for assessing the EPSP synthase gene
 - Identify changes in the EPSP synthase gene
 - Relating genetic changes to soil history and glyphosate resistance

METHODS



http://www.maich.gr/sa_research_images/image016.jpg

Figure 1: Shikimate pathway with the EPSP synthase inhibited by glyphosate.

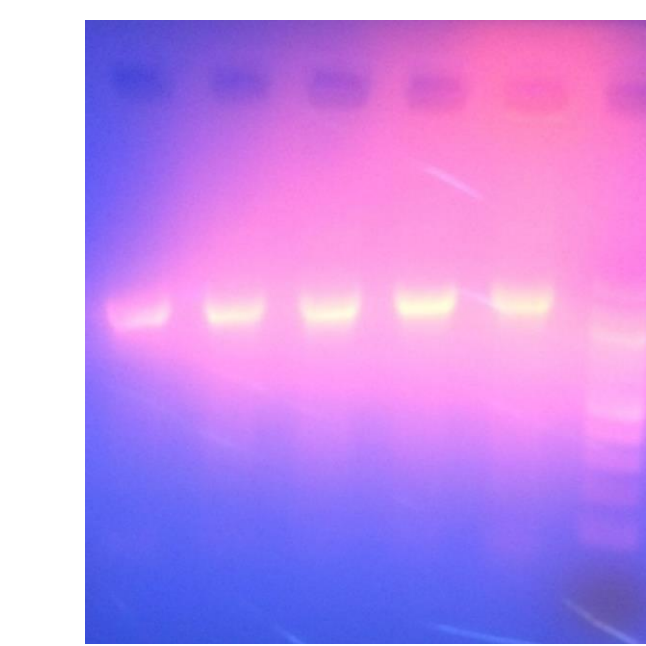


<http://www.ncbi.nlm.nih.gov/pmc/articles/PMC29264/>

Figure 2: Shows the EPSP gene without ligands (a) and with ligands glyphosate and S3P (b).

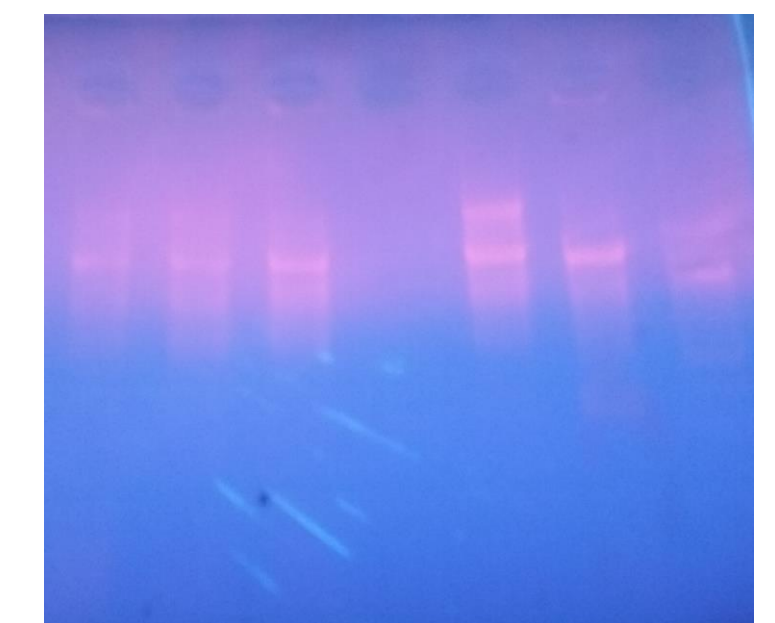
RESULTS

Bacillus subtilis

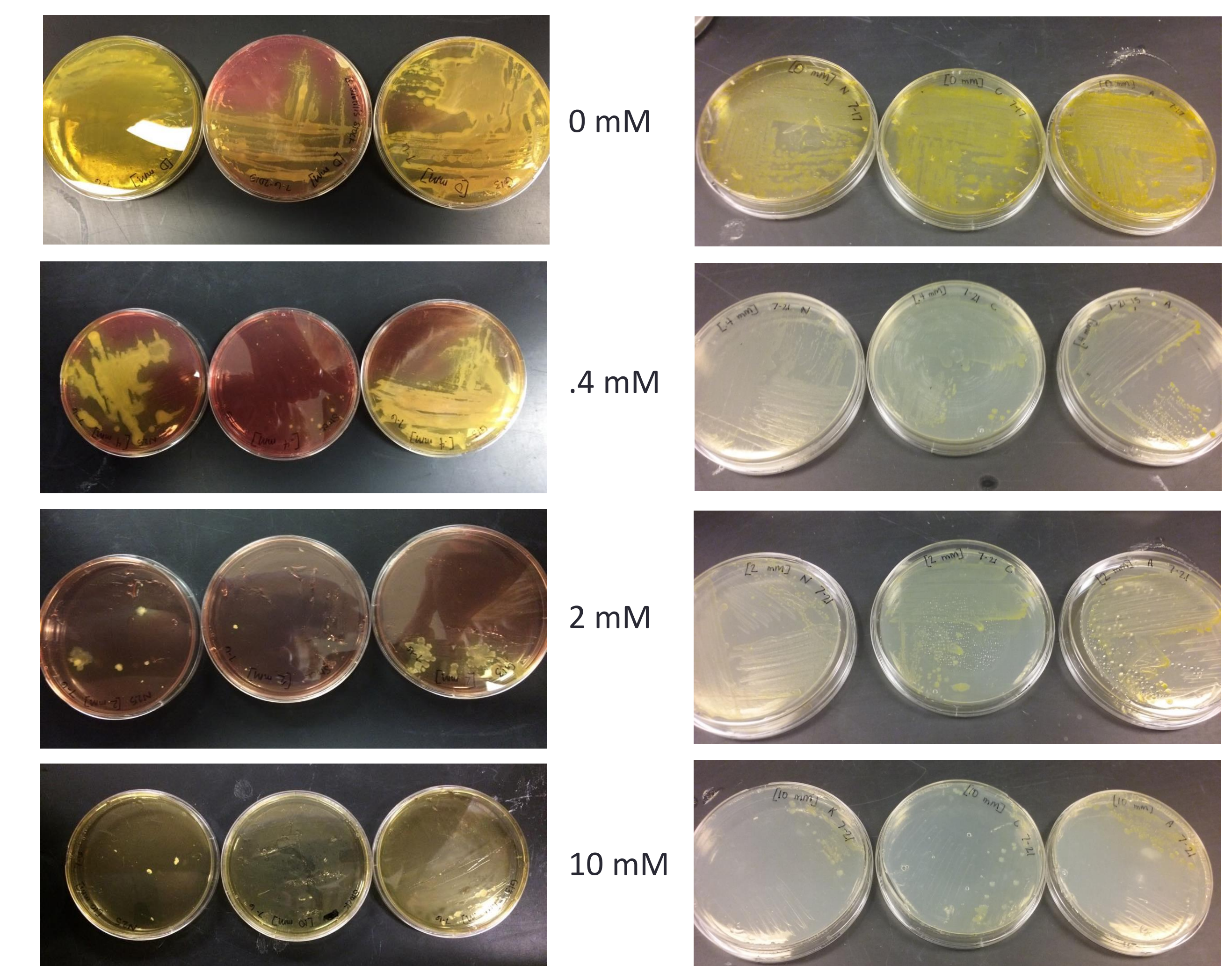


Confirmation of effective B. subtilis primers

Pseudomonas syringae pv. *glycinea*



Confirmation of effective P. syringae primers



MSA and Nutrient Agar plates containing increasing concentrations of glyphosate

RESULTS

Nucleotide Mutations

T219G A258G G264T A297G C471T C474T C510A T789A

Amino Acid Alignment with Synonymous Mutations Highlighted

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B_sub  MLNNIENNNDGPFENSTIQHIFKEIFKAGLELQEEEDHSHKALLVSRKKKPEDTIVDIKGEK 60
G13N  MLNNIENNNDGPFENSTIQHIFKEIFKAGLELQEEEDHSHKALLVSRKKKPEDTIVDIKGEK 60
*****

B_sub  IGDGQQRFTVIGPCAVESYEQVAEVAARAKKQKIKILRGSAFKPRTSPYDFQGLGVEGLQI 120
G13N  IGDGQQRFTVIGPCAVESYEQVAEVAARAKKQKIKILRGSAFKPRTSPYDFQGLGVEGLQI 120
*****

B_sub  LKRVADEFDLAVISEIVTPAHIEEALDYIDVIQIGARNMQNFELLKAAGAVKKPVLLKRG 180
G13N  LKRVADEFDLAVISEIVTPAHIEEALDYIDVIQIGARNMQNFELLKAAGAVKKPVLLKRG 180
*****

B_sub  LAATISEFINAAEYIMSGQNDQIILCERGIRTYETATRNLDISAVPILKQETHLPVFVD 240
G13N  LAATISEFINAAEYIMSGQNDQIILCERGIRTYETATRNLDISAVPILKQETHLPVFVD 240
*****

B_sub  VHTSGRRDLLPTAKAALAIGDGVMAEVHPDPSVALSDSAQMAIPEFEKWLNLKPM 300
G13N  VHTSGRRDLLPTAKAALAIGDGVMAEVHPDPSVALSDSAQMAIPEFEKWLNLKPM 300
*****

B_sub  VKYNA 305
G13N  VKYNA 305
*****
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CONCLUSIONS & NEXT STEPS

- Conclusions
 - Glyphosate resistance is prevalent among soil bacteria.
 - We have designed primers capable of amplifying the EPSP synthase gene
 - Sequence variations exist between resistant strains and organisms.
- Next Steps
 - Include more organisms in the study
 - Annotate areas of common mutations
 - Predict structural effects of mutations on the EPSP synthase enzyme
 - Look at differential gene expression of resistant isolates with and without glyphosate pressure