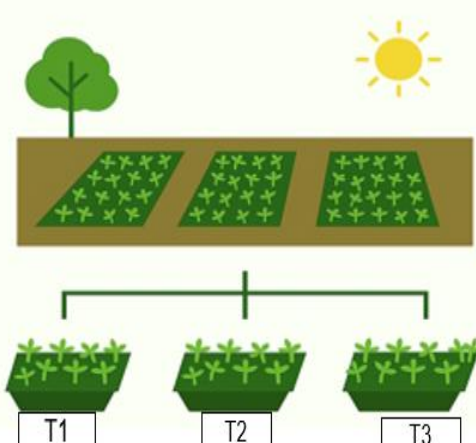


Biostimulant-Induced Modulation of Photosynthetic Efficiency in Alfalfa (*Medicago sativa* L.)Ebenezer Ayew Appiah<sup>1, 2</sup>, Erika Kutasy Tunde<sup>2</sup><sup>1</sup> Kálmán Kerpely Doctoral School of Crop Production and Horticultural Science, University of Debrecen, Böszörményi Str 138, H-4032 Debrecen, Hungary.<sup>2</sup> Institute of Crop Production, Breeding and Plant Technology, Faculty of Agricultural and Food Sciences and Environmental Management, University of Debrecen, Böszörményi Str. 138, H-4032 Debrecen, Hungary.

## INTRODUCTION &amp; AIM

Alfalfa (*Medicago sativa* L.) is a key forage crop valued globally for its high protein content and nitrogen-fixing capacity, making it essential in livestock feeding systems. However, its biomass yield is often constrained by reduced photosynthetic efficiency, particularly under stress conditions such as drought and nutrient imbalance. Biostimulants natural or synthetic substances including seaweed extracts, humic acids, and microbial or fungal inoculants have the potential to enhance plant metabolism, strengthen stress resilience, and mitigate these negative impacts. Therefore, this study aims to evaluate how two distinct biostimulant applications influence photosynthetic efficiency and overall productivity in alfalfa.

## METHOD



Study Design used

Code	Biostimulant treatments	Dosage
T1	Tricho Immun	21.6g/3L
T2	Biostimulant containing Si, MTU®, and pidolic acid	(2ml/3L)
T3	Control (no biostimulant)	

## Measurement

Measurements of (i) gas exchange parameters and (ii) chlorophyll fluorescence parameters were conducted under a photosynthetic photon flux density of 1500  $\mu\text{mol photons m}^{-2} \text{s}^{-1}$ .

Instrument used: Licor 6800



## RESULTS &amp; DISCUSSION

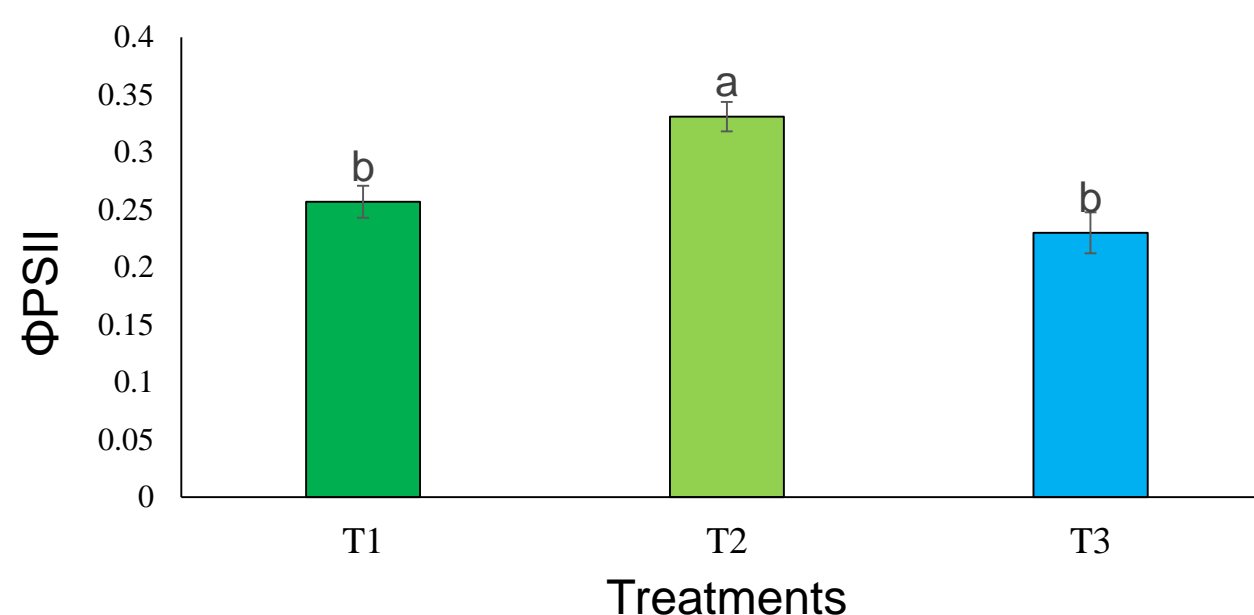
- Our findings show that biostimulant application significantly improved net photosynthetic rate, transpiration rate, intercellular CO<sub>2</sub> concentration, water use efficiency, and quantum yield of PSII, indicating improved photochemical efficiency.
- The result indicated that  $\Phi\text{PSII}$  was increased by 43.9%, net photosynthetic rate by 37.8%, and transpiration rate by 97.6% in T2 treatment compared to control.

## RESULTS &amp; DISCUSSION

**Table 2: Influence of biostimulants on gas exchange parameters of alfalfa (*Medicago Sativa* L.)**

Treatments	Ass ( $\mu\text{mol m}^{-2} \text{s}^{-1}$ )	Transpiration ( $\text{mmol m}^{-2} \text{s}^{-1}$ )	Intercellular CO <sub>2</sub> conc. ( $\mu\text{mol mol}^{-1}$ )	Ambient CO <sub>2</sub> conc. ( $\mu\text{mol mol}^{-1}$ )	WUE ( $\text{kg m}^{-3}$ )
T1	23.23±0.72 <sup>b</sup>	8.97±0.26 <sup>b</sup>	204.0±5.51 <sup>c</sup>	396.3±15.71 <sup>a</sup>	2.56±0.15 <sup>ab</sup>
T2	29.14±0.53 <sup>a</sup>	12.70±0.75 <sup>a</sup>	284.0±3.87 <sup>b</sup>	395.8±7.36 <sup>a</sup>	2.32±0.18 <sup>b</sup>
T3	21.14±0.48 <sup>b</sup>	6.43±0.26 <sup>c</sup>	720.0±10.52 <sup>a</sup>	396.6±3.82 <sup>a</sup>	3.33±0.26 <sup>a</sup>
Cv (%)	4.9	11.3	2.7	4.0	15.4
Lsd (5%)	2.71	2.39	24.85	35.72	0.95

Ass: photosynthesis rate, means with same letters are not different from each other

**Figure 1: Effect of biostimulant on Quantum yield of PSII of alfalfa (*Medicago Sativa* L.)**

- Similarly, Tricho Immun (T1) also increased  $\Phi\text{PSII}$ , net photosynthetic rate and transpiration rate by 11.7%, 9.8%, and 39.5% respectively, compared to the control.
- No significant effects were observed on ambient CO<sub>2</sub>. On the other hand, the control treatment tended to outperform the biostimulants in both water-use efficiency (WUE) and intercellular CO<sub>2</sub> concentration.

## CONCLUSION

- Biostimulant significantly enhance photosynthetic performance and resource-use efficiency in alfalfa
- T2 was most effective in improving photosynthetic parameters
- Overall, biostimulant application serve as a promising strategy to enhance forage crop productivity and sustainability in agriculture.

## FUTURE WORK / REFERENCES

Future research should assess changes in gene expression associated with photosynthesis and stress tolerance and apply transcriptomic approaches to identify the pathways activated by biostimulants.