

**Enhancing pulse productivity on problem
soils by smallholder farmers:
Challenges and opportunities**

Workshop held August 14-18, 2011 at Penn
State

Goal:

“... formulate recommendations on future **research foci** to achieve **major increases** in pulse productivity under **edaphic and abiotic stress** conditions in small-holder farm settings.”

Bob Herdt, Cornell (facilitator)

Jim Beaver, U Puerto Rico

Steve Beebe, CIAT

Jonathan Lynch, Penn State

Mark Westgate, Iowa State

I. Rao, CIAT

Jeff White, USDA/ARS

Sieg Snapp, MSU

45 participants

US PLGUs	19	(11 universities)
CG IRRI)	8	(CIAT, ICRISAT, IITA,
USDA/ARS	6	
African NARS	6	
USAID	4	
USDA	1	
McKnight	1	

working groups:

- genetic improvement
- phenomics
- BNF/plant microbe interactions
- systems analysis and management

each group developed

- goals and potential
- actions and outputs
- outcomes

Genetic Improvement

Goals and potentials:

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought
- heat

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought
- heat
- low soil fertility (low P, N, K, toxicities of Al and Mn)

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought
- heat
- low soil fertility (low P, N, K, toxicities of Al and Mn)
- grain filling under stress

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought
- heat
- low soil fertility (low P, N, K, toxicities of Al and Mn)
- grain filling under stress

Data will

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought
- heat
- low soil fertility (low P, N, K, toxicities of Al and Mn)
- grain filling under stress

Data will

- aid breeding

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought
- heat
- low soil fertility (low P, N, K, toxicities of Al and Mn)
- grain filling under stress

Data will

- aid breeding
- help physiologists understand utility of specific traits

Genetic Improvement

Goals and potentials:

Biology now stands at the threshold of the ultimate genetic tool: whole genome sequencing of individual organisms to reveal the genetic basis of useful traits.

Our goal is to lay the genetic groundwork to address the edaphic and climatic constraints that limit legume yields in the developing world to a fraction of their genetic potential.

Reveal the genes underlying phenotypic for tolerance of

- drought
- heat
- low soil fertility (low P, N, K, toxicities of Al and Mn)
- grain filling under stress

Data will

- aid breeding
- help physiologists understand utility of specific traits
- raw material for soil microbiologists, agronomists, modeling

Genetic Improvement

Actions and outputs:

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;
- Share, access and interpret information;

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;
- Share, access and interpret information;
- Convert this information into selection strategies.

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;
- Share, access and interpret information;
- Convert this information into selection strategies.

Outputs:

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;
- Share, access and interpret information;
- Convert this information into selection strategies.

Outputs:

- Sources and markers for traits to improve yield under stress;

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;
- Share, access and interpret information;
- Convert this information into selection strategies.

Outputs:

- Sources and markers for traits to improve yield under stress;
- Greater stress tolerance in elite lines;

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;
- Share, access and interpret information;
- Convert this information into selection strategies.

Outputs:

- Sources and markers for traits to improve yield under stress;
- Greater stress tolerance in elite lines;
- Better understanding of how genotype affects phenotype;

Genetic Improvement

Actions and outputs:

To realize the potential of genome sequencing, legume communities must develop breeding platforms that:

- Create networks to evaluate many lines for multiple traits, across multiple stressful environments;
- Outsource genome sequencing of germplasm sets;
- Share, access and interpret information;
- Convert this information into selection strategies.

Outputs:

- Sources and markers for traits to improve yield under stress;
- Greater stress tolerance in elite lines;
- Better understanding of how genotype affects phenotype;
- A platform incorporating genomic tools for legume breeding.

Genetic Improvement

Outcomes:

Genetic Improvement

Outcomes:

- Improved breeding efficiency for stress tolerance.

Genetic Improvement

Outcomes:

- Improved breeding efficiency for stress tolerance.
- Improved germplasm for farmers (especially women) with greater, more stable yields under smallholder conditions.

Genetic Improvement

Outcomes:

- Improved breeding efficiency for stress tolerance.
- Improved germplasm for farmers (especially women) with greater, more stable yields under smallholder conditions.
- Better understanding of stress tolerance, useful for guided breeding and modeling system impacts/future climates.

Genetic Improvement

Outcomes:

- Improved breeding efficiency for stress tolerance.
- Improved germplasm for farmers (especially women) with greater, more stable yields under smallholder conditions.
- Better understanding of stress tolerance, useful for guided breeding and modeling system impacts/future climates.
- Genetic information will benefit other breeding objectives, including biotic stress resistance and biofortification.

Phenomics

Goals and potentials:

Phenomics

Goals and potentials:

Recent research advances provide the foundation for breakthroughs in the development of more drought-tolerant, nutrient-efficient lines, including advances in our understanding of root biology, yield formation, and new phenotyping platforms.

Phenomics

Goals and potentials:

Recent research advances provide the foundation for breakthroughs in the development of more drought-tolerant, nutrient-efficient lines, including advances in our understanding of root biology, yield formation, and new phenotyping platforms.

Our goal is to integrate modern phenomics and genomics to develop pulse genotypes with substantially improved performance in stressful environments.

Phenomics

Goals and potentials:

Recent research advances provide the foundation for breakthroughs in the development of more drought-tolerant, nutrient-efficient lines, including advances in our understanding of root biology, yield formation, and new phenotyping platforms.

Our goal is to integrate modern phenomics and genomics to develop pulse genotypes with substantially improved performance in stressful environments.

This activity seeks to define the mechanistic basis of legume adaption to low soil fertility, drought, and heat. This information is needed to

Phenomics

Goals and potentials:

Recent research advances provide the foundation for breakthroughs in the development of more drought-tolerant, nutrient-efficient lines, including advances in our understanding of root biology, yield formation, and new phenotyping platforms.

Our goal is to integrate modern phenomics and genomics to develop pulse genotypes with substantially improved performance in stressful environments.

This activity seeks to define the mechanistic basis of legume adaption to low soil fertility, drought, and heat. This information is needed to

- guide focused breeding strategies

Phenomics

Goals and potentials:

Recent research advances provide the foundation for breakthroughs in the development of more drought-tolerant, nutrient-efficient lines, including advances in our understanding of root biology, yield formation, and new phenotyping platforms.

Our goal is to integrate modern phenomics and genomics to develop pulse genotypes with substantially improved performance in stressful environments.

This activity seeks to define the mechanistic basis of legume adaption to low soil fertility, drought, and heat. This information is needed to

- guide focused breeding strategies
- develop high-throughput phenotypic screens to identify adapted germplasm and molecular markers.

Phenomics

Goals and potentials:

Recent research advances provide the foundation for breakthroughs in the development of more drought-tolerant, nutrient-efficient lines, including advances in our understanding of root biology, yield formation, and new phenotyping platforms.

Our goal is to integrate modern phenomics and genomics to develop pulse genotypes with substantially improved performance in stressful environments.

This activity seeks to define the mechanistic basis of legume adaption to low soil fertility, drought, and heat. This information is needed to

- guide focused breeding strategies
- develop high-throughput phenotypic screens to identify adapted germplasm and molecular markers.
- Develop integrated crop management strategies that achieve synergies between genetic and agroecological technologies.

Phenomics

Goals and potentials:

Recent research advances provide the foundation for breakthroughs in the development of more drought-tolerant, nutrient-efficient lines, including advances in our understanding of root biology, yield formation, and new phenotyping platforms.

Our goal is to integrate modern phenomics and genomics to develop pulse genotypes with substantially improved performance in stressful environments.

This activity seeks to define the mechanistic basis of legume adaption to low soil fertility, drought, and heat. This information is needed to

- guide focused breeding strategies
- develop high-throughput phenotypic screens to identify adapted germplasm and molecular markers.
- Develop integrated crop management strategies that achieve synergies between genetic and agroecological technologies.

Phenomics

Actions and outputs:

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.
- Understanding the basis of adaptation to low availability of soil K, Ca, and Mg.

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.
- Understanding the basis of adaptation to low availability of soil K, Ca, and Mg.

For **drought and heat**, research priorities include:

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.
- Understanding the basis of adaptation to low availability of soil K, Ca, and Mg.

For **drought and heat**, research priorities include:

- Root traits enhancing water acquisition under drought.

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.
- Understanding the basis of adaptation to low availability of soil K, Ca, and Mg.

For **drought and heat**, research priorities include:

- Root traits enhancing water acquisition under drought.
- Processes controlling reproduction under stress.

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.
- Understanding the basis of adaptation to low availability of soil K, Ca, and Mg.

For **drought and heat**, research priorities include:

- Root traits enhancing water acquisition under drought.
- Processes controlling reproduction under stress.
- Understanding the role of phenology in stress avoidance.

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.
- Understanding the basis of adaptation to low availability of soil K, Ca, and Mg.

For **drought and heat**, research priorities include:

- Root traits enhancing water acquisition under drought.
- Processes controlling reproduction under stress.
- Understanding the role of phenology in stress avoidance.

Two cross-cutting themes are 1) **root traits** for soil resource acquisition, and 2) **resource remobilization into seeds**.

Phenomics

Actions and outputs:

Discover key traits, characterize their utility, and phenotypically profile pulse germplasm. For **low soil fertility** research priorities include:

- Root traits enhancing acquisition of P and mineral N.
- Processes enhancing utilization of P and N in the plant.
- Understanding the basis of adaptation to low availability of soil K, Ca, and Mg.

For **drought and heat**, research priorities include:

- Root traits enhancing water acquisition under drought.
- Processes controlling reproduction under stress.
- Understanding the role of phenology in stress avoidance.

Two cross-cutting themes are 1) **root traits** for soil resource acquisition, and 2) **resource remobilization into seeds**.

Substantial impact can be realized by deploying traits of known value such as **root hairs**, **Al resistance** and **Mn tolerance** in breeding programs.

Phenomics

Actions and outputs (b):

Phenomics

Actions and outputs (b):

This will lead to the following outputs:

Phenomics

Actions and outputs (b):

This will lead to the following outputs:

- validated ideotypes for focused breeding;

Phenomics

Actions and outputs (b):

This will lead to the following outputs:

- validated ideotypes for focused breeding;
- high-throughput phenotyping platforms;

Phenomics

Actions and outputs (b):

This will lead to the following outputs:

- validated ideotypes for focused breeding;
- high-throughput phenotyping platforms;
- sources of key traits;

Phenomics

Actions and outputs (b):

This will lead to the following outputs:

- validated ideotypes for focused breeding;
- high-throughput phenotyping platforms;
- sources of key traits;
- greater stress tolerance in elite lines;

Phenomics

Actions and outputs (b):

This will lead to the following outputs:

- validated ideotypes for focused breeding;
- high-throughput phenotyping platforms;
- sources of key traits;
- greater stress tolerance in elite lines;
- integration of genetic and management technologies for improved Integrated Crop Management.

Phenomics

Outcomes:

Phenomics

Outcomes:

Outcomes include all of those listed for Genetic Improvement, including improved efficiency of breeding programs, improved germplasm, and improved understanding of relevant processes that will guide model development and Integrated Crop Management strategies.

BNF/plant microbe interactions

Goals and potentials:

BNF/plant microbe interactions

Goals and potentials:

Despite considerable investment in improving BNF, legumes grown by smallholders on degraded soils produce only a fraction of their genetic potential.

BNF/plant microbe interactions

Goals and potentials:

Despite considerable investment in improving BNF, legumes grown by smallholders on degraded soils produce only a fraction of their genetic potential.

Our overall goal is to identify and capitalize on favorable microbial – plant interactions to improve legume productivity in degraded soils common to priority cropping systems in Sub-Saharan Africa.

BNF/plant microbe interactions

Goals and potentials:

Despite considerable investment in improving BNF, legumes grown by smallholders on degraded soils produce only a fraction of their genetic potential.

Our overall goal is to identify and capitalize on favorable microbial – plant interactions to improve legume productivity in degraded soils common to priority cropping systems in Sub-Saharan Africa.

Potential for BNF is determined by multiple factors including rhizobial genotype, plant genotype, environment, and management system. As such, improving BNF requires:

BNF/plant microbe interactions

Goals and potentials:

Despite considerable investment in improving BNF, legumes grown by smallholders on degraded soils produce only a fraction of their genetic potential.

Our overall goal is to identify and capitalize on favorable microbial – plant interactions to improve legume productivity in degraded soils common to priority cropping systems in Sub-Saharan Africa.

Potential for BNF is determined by multiple factors including rhizobial genotype, plant genotype, environment, and management system. As such, improving BNF requires:

- integrating greater knowledge of microbial symbiont populations with understanding of the impact of environmental and management conditions on those populations;

BNF/plant microbe interactions

Goals and potentials:

Despite considerable investment in improving BNF, legumes grown by smallholders on degraded soils produce only a fraction of their genetic potential.

Our overall goal is to identify and capitalize on favorable microbial – plant interactions to improve legume productivity in degraded soils common to priority cropping systems in Sub-Saharan Africa.

Potential for BNF is determined by multiple factors including rhizobial genotype, plant genotype, environment, and management system. As such, improving BNF requires:

- integrating greater knowledge of microbial symbiont populations with understanding of the impact of environmental and management conditions on those populations;
- a systematic breeding methodology to identify superior legumes and rhizobial strains; and

BNF/plant microbe interactions

Goals and potentials:

Despite considerable investment in improving BNF, legumes grown by smallholders on degraded soils produce only a fraction of their genetic potential.

Our overall goal is to identify and capitalize on favorable microbial – plant interactions to improve legume productivity in degraded soils common to priority cropping systems in Sub-Saharan Africa.

Potential for BNF is determined by multiple factors including rhizobial genotype, plant genotype, environment, and management system. As such, improving BNF requires:

- integrating greater knowledge of microbial symbiont populations with understanding of the impact of environmental and management conditions on those populations;
- a systematic breeding methodology to identify superior legumes and rhizobial strains; and
- capacity building encompassing training, germplasm curation, and inoculum standardization through a regional Rhizobium Resource Center.

BNF/plant microbe interactions

Actions and outputs:

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

- identification of beneficial microbes in soils of priority cropping systems;

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

- identification of beneficial microbes in soils of priority cropping systems;
- relation of current microbial population structures to soil conditions and crop management;

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

- identification of beneficial microbes in soils of priority cropping systems;
- relation of current microbial population structures to soil conditions and crop management;
- identification of soil biological factors that vary with microbial population structure, soil conditions, and cropping systems and their potential impact on plant growth and yield;

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

- identification of beneficial microbes in soils of priority cropping systems;
- relation of current microbial population structures to soil conditions and crop management;
- identification of soil biological factors that vary with microbial population structure, soil conditions, and cropping systems and their potential impact on plant growth and yield;
- identification of genetic markers for persistent rhizobia in adverse soils that can be used to select superior strains;

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

- identification of beneficial microbes in soils of priority cropping systems;
- relation of current microbial population structures to soil conditions and crop management;
- identification of soil biological factors that vary with microbial population structure, soil conditions, and cropping systems and their potential impact on plant growth and yield;
- identification of genetic markers for persistent rhizobia in adverse soils that can be used to select superior strains;
- development of models to predict inoculant success;

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

- identification of beneficial microbes in soils of priority cropping systems;
- relation of current microbial population structures to soil conditions and crop management;
- identification of soil biological factors that vary with microbial population structure, soil conditions, and cropping systems and their potential impact on plant growth and yield;
- identification of genetic markers for persistent rhizobia in adverse soils that can be used to select superior strains;
- development of models to predict inoculant success;
- development of a practical decision tool for farmers on profitable use of inoculants; and

BNF/plant microbe interactions

Actions and outputs:

Actions/Outputs related to predicting rhizobia populations in degraded soils include:

- identification of beneficial microbes in soils of priority cropping systems;
- relation of current microbial population structures to soil conditions and crop management;
- identification of soil biological factors that vary with microbial population structure, soil conditions, and cropping systems and their potential impact on plant growth and yield;
- identification of genetic markers for persistent rhizobia in adverse soils that can be used to select superior strains;
- development of models to predict inoculant success;
- development of a practical decision tool for farmers on profitable use of inoculants; and
- identification of enabling interactions between a broader array of soil microbes and legumes that promote favorable plant growth and suppress soil-borne diseases.

BNF/plant microbe interactions

Actions and outputs (b):

BNF/plant microbe interactions

Actions and outputs (b):

Actions/Outputs related to identifying yield enhancing interactions between microbes and plants:

BNF/plant microbe interactions

Actions and outputs (b):

Actions/Outputs related to identifying yield enhancing interactions between microbes and plants:

- creation of a set of “Differential Bean Genotypes” to screen candidate rhizobia strains;

BNF/plant microbe interactions

Actions and outputs (b):

Actions/Outputs related to identifying yield enhancing interactions between microbes and plants:

- creation of a set of “Differential Bean Genotypes” to screen candidate rhizobia strains;
- development of a set of promiscuous inoculants;

BNF/plant microbe interactions

Actions and outputs (b):

Actions/Outputs related to identifying yield enhancing interactions between microbes and plants:

- creation of a set of “Differential Bean Genotypes” to screen candidate rhizobia strains;
- development of a set of promiscuous inoculants;
- determination of how plant phenotypes affect BNF;

BNF/plant microbe interactions

Actions and outputs (b):

Actions/Outputs related to identifying yield enhancing interactions between microbes and plants:

- creation of a set of “Differential Bean Genotypes” to screen candidate rhizobia strains;
- development of a set of promiscuous inoculants;
- determination of how plant phenotypes affect BNF;
- quantification of effects of plant removal (e.g. for animal feed, green manure) on soil N and microbial dynamics;

BNF/plant microbe interactions

Actions and outputs (b):

Actions/Outputs related to identifying yield enhancing interactions between microbes and plants:

- creation of a set of “Differential Bean Genotypes” to screen candidate rhizobia strains;
- development of a set of promiscuous inoculants;
- determination of how plant phenotypes affect BNF;
- quantification of effects of plant removal (e.g. for animal feed, green manure) on soil N and microbial dynamics;
- characterization of host response with and without environmental stress, to explain a “BNF-gap” under stress;

BNF/plant microbe interactions

Actions and outputs (b):

Actions/Outputs related to identifying yield enhancing interactions between microbes and plants:

- creation of a set of “Differential Bean Genotypes” to screen candidate rhizobia strains;
- development of a set of promiscuous inoculants;
- determination of how plant phenotypes affect BNF;
- quantification of effects of plant removal (e.g. for animal feed, green manure) on soil N and microbial dynamics;
- characterization of host response with and without environmental stress, to explain a “BNF-gap” under stress;
- cross-legume comparisons to reveal host traits explaining differences in BNF efficiency.

BNF/plant microbe interactions

Actions and outputs (c):

BNF/plant microbe interactions

Actions and outputs (c):

Actions/Outputs related to establishing a Regional Rhizobia Resource Center:

BNF/plant microbe interactions

Actions and outputs (c):

Actions/Outputs related to establishing a Regional Rhizobia Resource Center:

- a germplasm resource for rhizobia and favorable soil microbe collection and curation;

BNF/plant microbe interactions

Actions and outputs (c):

Actions/Outputs related to establishing a Regional Rhizobia Resource Center:

- a germplasm resource for rhizobia and favorable soil microbe collection and curation;
- training for small business owners producing and distributing inoculant, students, lab managers, farmers, and extensionists;

BNF/plant microbe interactions

Actions and outputs (c):

Actions/Outputs related to establishing a Regional Rhizobia Resource Center:

- a germplasm resource for rhizobia and favorable soil microbe collection and curation;
- training for small business owners producing and distributing inoculant, students, lab managers, farmers, and extensionists;
- mini-grant program to develop skills in inoculant preparation and management;

BNF/plant microbe interactions

Actions and outputs (c):

Actions/Outputs related to establishing a Regional Rhizobia Resource Center:

- a germplasm resource for rhizobia and favorable soil microbe collection and curation;
- training for small business owners producing and distributing inoculant, students, lab managers, farmers, and extensionists;
- mini-grant program to develop skills in inoculant preparation and management;
- industry standards for inoculant quality control, quality control detection, and inoculant production at large scale and on-farm.

BNF/plant microbe interactions

Outcomes:

- Increased pulse production by smallholder farmers.
- New knowledge about the effects of crop management and soil conditions on soil microbial dynamics will inform needed changes in cropping systems management.
- Establishing the genetic basis for superior plant-microbe interactions will speed development and deployment of improved varieties with greater yields in degraded soils.
- Creation of the Regional Rhizobium Resource Center will provide training, germplasm curation, and inoculum production, which are critical for improving BNF on a regional scale.

Systems analysis and management

Goals and potentials:

Systems analysis and management

Goals and potentials:

Improved crop and soil management are fundamental to ensuring that gains through genetic improvement attain their full potential and that such gains are truly sustainable.

Systems analysis and management

Goals and potentials:

Improved crop and soil management are fundamental to ensuring that gains through genetic improvement attain their full potential and that such gains are truly sustainable.

Our goal is to identify and promote promising crop management options for pulse agroecosystems that emphasize more efficient use of existing or readily available resources and that ultimately improve the productivity, profitability and sustainability of smallholder agriculture.

Systems analysis and management

Goals and potentials:

Improved crop and soil management are fundamental to ensuring that gains through genetic improvement attain their full potential and that such gains are truly sustainable.

Our goal is to identify and promote promising crop management options for pulse agroecosystems that emphasize more efficient use of existing or readily available resources and that ultimately improve the productivity, profitability and sustainability of smallholder agriculture.

Four advances and innovations can support the required transformation of legume systems agronomy in Africa:

Systems analysis and management

Goals and potentials:

Improved crop and soil management are fundamental to ensuring that gains through genetic improvement attain their full potential and that such gains are truly sustainable.

Our goal is to identify and promote promising crop management options for pulse agroecosystems that emphasize more efficient use of existing or readily available resources and that ultimately improve the productivity, profitability and sustainability of smallholder agriculture.

Four advances and innovations can support the required transformation of legume systems agronomy in Africa:

- Simple, robust ecophysiological models combined with geospatial tools can simulate pulse productivity as affected by crop and soil management scenarios in Africa. The cost and time required to generate a similar set of data using conventional agronomic trials is prohibitive.

Systems analysis and management

Goals and potentials:

Improved crop and soil management are fundamental to ensuring that gains through genetic improvement attain their full potential and that such gains are truly sustainable.

Our goal is to identify and promote promising crop management options for pulse agroecosystems that emphasize more efficient use of existing or readily available resources and that ultimately improve the productivity, profitability and sustainability of smallholder agriculture.

Four advances and innovations can support the required transformation of legume systems agronomy in Africa:

- Simple, robust ecophysiological models combined with geospatial tools can simulate pulse productivity as affected by crop and soil management scenarios in Africa. The cost and time required to generate a similar set of data using conventional agronomic trials is prohibitive.
- Advances in fundamental understanding of soil processes.

Systems analysis and management

Goals and potentials:

Improved crop and soil management are fundamental to ensuring that gains through genetic improvement attain their full potential and that such gains are truly sustainable.

Our goal is to identify and promote promising crop management options for pulse agroecosystems that emphasize more efficient use of existing or readily available resources and that ultimately improve the productivity, profitability and sustainability of smallholder agriculture.

Four advances and innovations can support the required transformation of legume systems agronomy in Africa:

- Simple, robust ecophysiological models combined with geospatial tools can simulate pulse productivity as affected by crop and soil management scenarios in Africa. The cost and time required to generate a similar set of data using conventional agronomic trials is prohibitive.
- Advances in fundamental understanding of soil processes.
- Widespread recognition and acceptance of farmer participatory research.

Systems analysis and management

Goals and potentials:

Improved crop and soil management are fundamental to ensuring that gains through genetic improvement attain their full potential and that such gains are truly sustainable.

Our goal is to identify and promote promising crop management options for pulse agroecosystems that emphasize more efficient use of existing or readily available resources and that ultimately improve the productivity, profitability and sustainability of smallholder agriculture.

Four advances and innovations can support the required transformation of legume systems agronomy in Africa:

- Simple, robust ecophysiological models combined with geospatial tools can simulate pulse productivity as affected by crop and soil management scenarios in Africa. The cost and time required to generate a similar set of data using conventional agronomic trials is prohibitive.
- Advances in fundamental understanding of soil processes.
- Widespread recognition and acceptance of farmer participatory research.
- Networking tools can ensure that producers have access to information in a timely and location-relevant manner and can provide feedback to the research community in a cost-effective manner.

Systems analysis and management

Actions and outputs:

Systems analysis and management

Actions and outputs:

Quantitative systems analysis (QSA), which combines simulations with other geospatial analyses, will be used to predict stress effects on crops and cropping systems across time and space. This approach will

Systems analysis and management

Actions and outputs:

Quantitative systems analysis (QSA), which combines simulations with other geospatial analyses, will be used to predict stress effects on crops and cropping systems across time and space. This approach will

- evaluate the potential benefits of different pulse crops, plant traits, and management alternatives in target environments;

Systems analysis and management

Actions and outputs:

Quantitative systems analysis (QSA), which combines simulations with other geospatial analyses, will be used to predict stress effects on crops and cropping systems across time and space. This approach will

- evaluate the potential benefits of different pulse crops, plant traits, and management alternatives in target environments;
- Identify ways to enhance the efficiency of resource utilization, for example, through improved access to nutrients in deeper soil layers or through increased capture of precipitation by reducing runoff.

Systems analysis and management

Actions and outputs:

Quantitative systems analysis (QSA), which combines simulations with other geospatial analyses, will be used to predict stress effects on crops and cropping systems across time and space. This approach will

- evaluate the potential benefits of different pulse crops, plant traits, and management alternatives in target environments;
- Identify ways to enhance the efficiency of resource utilization, for example, through improved access to nutrients in deeper soil layers or through increased capture of precipitation by reducing runoff.

Participatory research teams will evaluate strategies identified by QSA in the target environments.

Systems analysis and management

Actions and outputs:

Quantitative systems analysis (QSA), which combines simulations with other geospatial analyses, will be used to predict stress effects on crops and cropping systems across time and space. This approach will

- evaluate the potential benefits of different pulse crops, plant traits, and management alternatives in target environments;
- Identify ways to enhance the efficiency of resource utilization, for example, through improved access to nutrients in deeper soil layers or through increased capture of precipitation by reducing runoff.

Participatory research teams will evaluate strategies identified by QSA in the target environments.

- Diagnostic and decision tool kits for farmers, extensionists and NGOs can be developed based on feedback from these participatory studies.

Systems analysis and management

Actions and outputs:

Quantitative systems analysis (QSA), which combines simulations with other geospatial analyses, will be used to predict stress effects on crops and cropping systems across time and space. This approach will

- evaluate the potential benefits of different pulse crops, plant traits, and management alternatives in target environments;
- Identify ways to enhance the efficiency of resource utilization, for example, through improved access to nutrients in deeper soil layers or through increased capture of precipitation by reducing runoff.

Participatory research teams will evaluate strategies identified by QSA in the target environments.

- Diagnostic and decision tool kits for farmers, extensionists and NGOs can be developed based on feedback from these participatory studies.
- *Ex-ante* and *ex-post* socio-economic analyses will be conducted to evaluate the impact of promising strategies.

Systems analysis and management

Outcomes:

QSA in coordination with participatory research will accelerate locally-appropriate adaptations to African pulse agroecosystems that are more diverse, productive and resilient to edaphic and climatic constraints. This research approach also permits the development agronomic practices that are more attuned to the needs and limitations of specific target environments.

closing thoughts