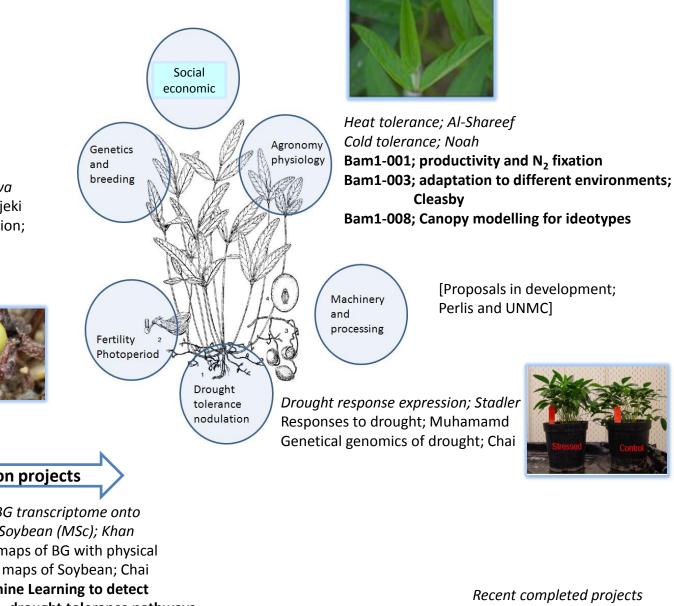


African genetic diversity; Stadler African genetic diversity; Molosiwa Indonesian genetic diversity; Redjeki Genetic mapping and domestication; Ahmad Bam1-006; MAGIC breeding

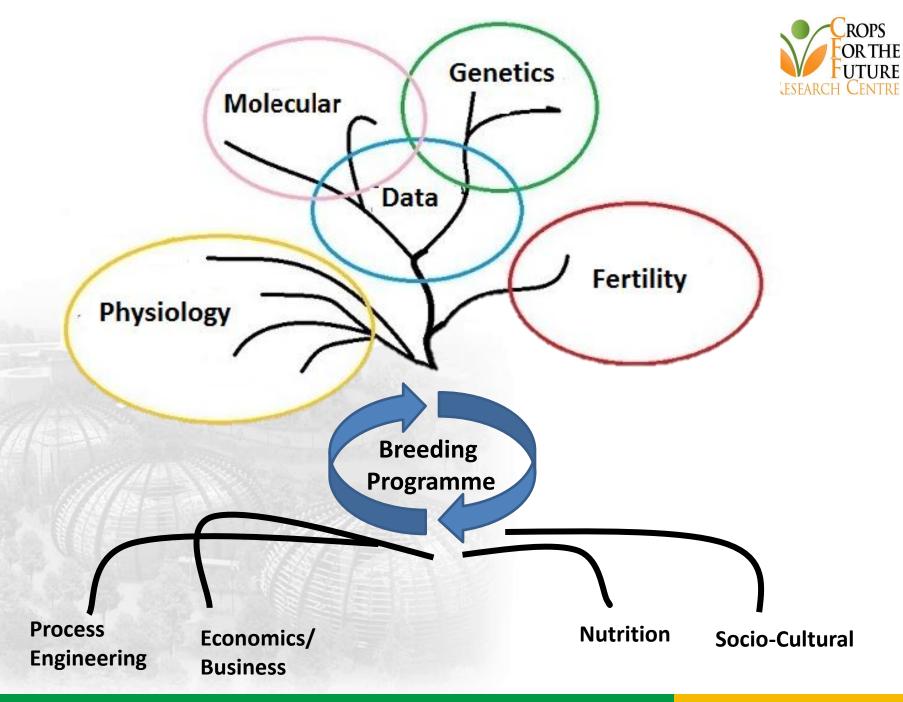
Photoperiod requirement; Kendabie Bam1-009 Fertility in bambara groundnut



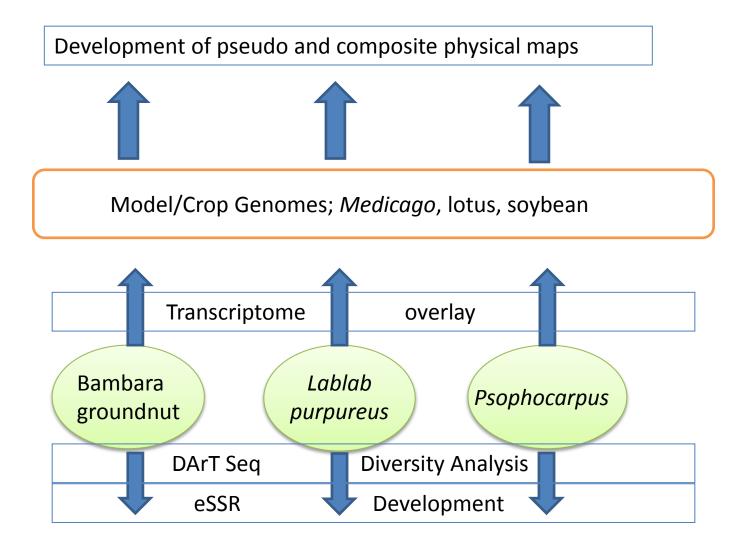
Recent completed projects Current on-going projects **BamYield agreed projects** 

**Translation projects** 

Overlaying the BG transcriptome onto Soybean (MSc); Khan Linking genetic maps of BG with physical maps of Soybean; Chai Bam1-004 Machine Learning to detect drought tolerance pathways Bam1-007 Combining multiple data types; Khan

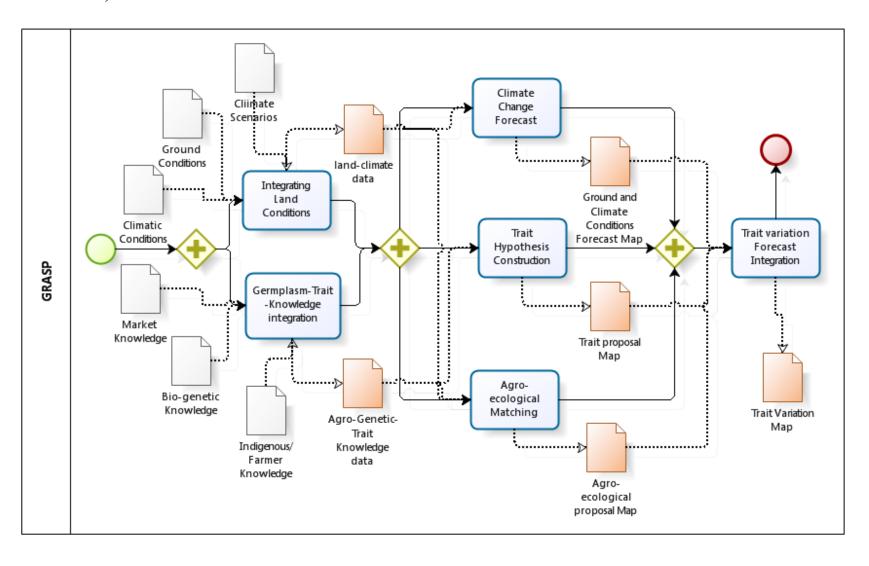


# Species cluster approach to working in underutilised crops



"Geospatial Resource for Agricultural Species and Pests with integrated workflow modelling to support Global Food Security (GRASP-GFS): a prototype"

BBSRC TRDF call 2 Support for Development of Bioinformatic Tools and Computational Approaches to the Biosciences; 18 months £127K



Bamyield; Bioinformatics, translation and data mining for minor crops;

Bam1-007

Combining Affymetrix, RNAseq and physical genome data - a multidimensional approach to transfer data from major crops to minor crops for Global Food Security. *PhD stipend and consumables only (Vice-Chancellor Fees Scholarship)* **Mr Faraz Khan** 

Bam1-004

Mining the gene networks in water-stress response in bambara groundnut: a machine learning approach to translating traits in model species to minor crops *PhD* 

### **Candidate identified**

Bam1-008 Mathematical Modelling of the bambara groundnut canopy *PhD 50% matching funding with Reading EPSRC DTC* **Advertised**  Bam1-007

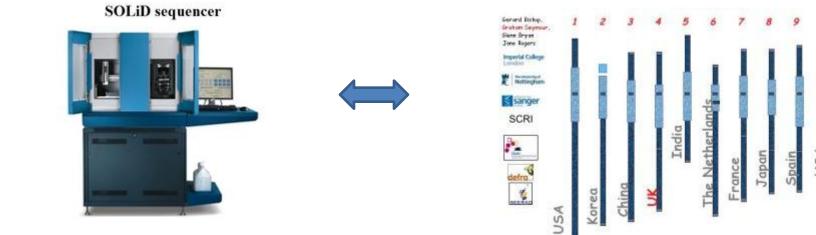
Combining Affymetrix, RNAseq and physical genome data - a multidimensional approach to transfer data from major crops to minor crops for Global Food Security.

PhD stipend and consumables only (Vice-Chancellor Fees Scholarship)

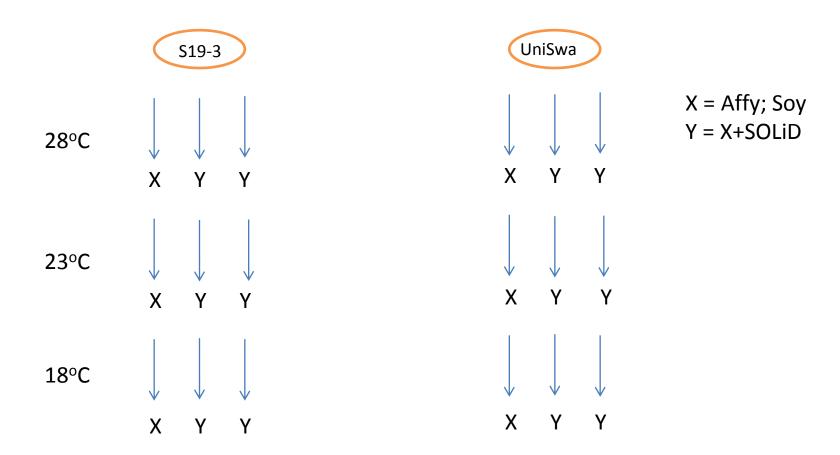
## Mr Faraz Khan







Temperature response in single genotypes of bambara groundnut by column transfer; *Noah, Al-Shareef, Mayes, Chai* 



1 plate 454 leaf transcriptome (16000 gene models), over layed on Soybean genome, with Soybean Affy oligos.

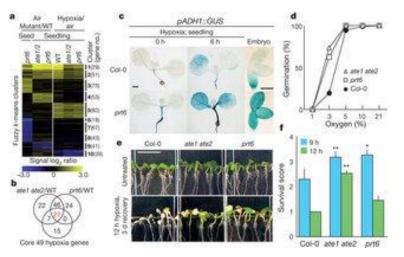
**Research questions:** 

- 1. How do the different data types relate?
- 2. How do we translate from models to subject crops?
- 3. What elements do we actually need to work in a crop and how can we do this efficiently and cheaply?
- 4. Can we work on 'clusters' of species at the genome level, rather than single species.
- 5. Can we develop/identify suitable software to make this work?

#### Bam1-004

Mining the gene networks in water-stress response in bambara groundnut: a machine learning approach to translating traits in model species to minor crops *PhD* 

# Candidate identified



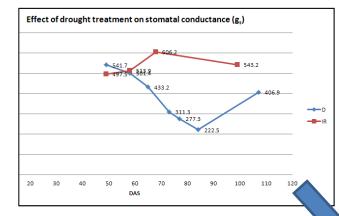
-Mike Holdsworth's group (UoN) have understood how plants sense anoxia stress -The sub1 mutant of rice is explained by the pathway they have uncovered and this offers an opportunity to see whether a similar pathway is active in Bambara groundnut Bambara groundnut landraces can survive
in rain levels between <400mm/year and</li>
>2500 mm/year

- Crosses have been made between S19-3 (Namibia) and Gresik (Indonesia)

- What are the mechanisms involved in such broad ranges of adaptation.

- Bambara groundnut is very sensitive to standing water





#### Compiled Storb-4

| 00<br>03<br>04<br>14<br>22<br>29<br>29<br>31 | 5060335<br>423050<br>6010885<br>423807<br>505038<br>506055<br>504508<br>506055<br>504508 |
|--|--|
| 19.8   | / P25b184AAG+AG4   |
| 27.9   | / P225297GCC+CA8   |
| 38.3 1                                       | 601401   |
| 38.8   | 598647   |
| 39.1   | 597687   |
| 39.2   | 601358   |
| 40.2   | y 423805   |
| 40.9   | 601656   |
| 41.8   | PRIMER98   |
| 42.4   | PRIMER2  |
| 43.0   | 422821   |
| 44.8   | 600972   |
| 44.9 %                                       | 601257   |
| 45.1 1                                       | 4601845 601050   |
| 47.0   | PRIMER45   |
| 49.5 1                                       | PRIMER95   |
| 50.1 //E                                     | P20E245ACG+CAG2  |
| 55.5 /                                       | \$ 597491  |
| 61.1   | PRIMER38   |
| 62.54  | P 601091   |
| 63.0   | 602174   |
| 63.14  | 598122   |
| 63.2   | 598598   |
| 63.44  | 598480 593939  |
| 71.0 /6                                      | 1595387  |
| 82.9   | P24b109AAC+AG4   |



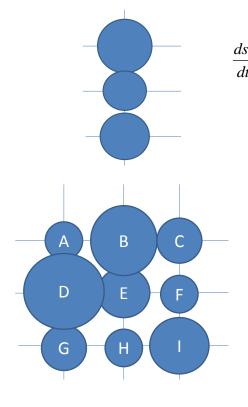


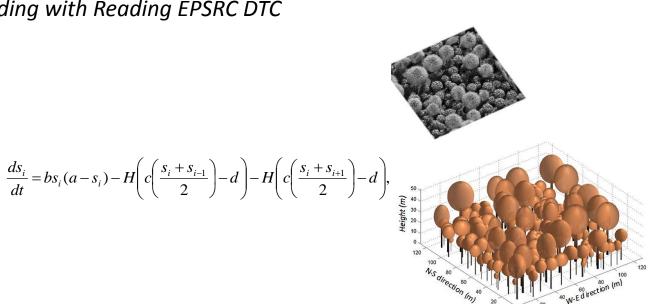
#### Machine Learning:

Do the same pathways exist as in models species? Can we mine datasets to find other important pathways?

Can we potentially address flooding sensitivity in bambara groundnut?

Bam1-008 Mathematical Modelling of the bambara groundnut canopy *PhD 50% matching funding with Reading EPSRC DTC* **Advertised** 





Mangrove modelling

- Analysis of single plants is not the same as growing a canopy
- These interaction between plants are one reason that translation from lab to field often works poorly
- If we can model these factors, we can potential use them to guide breeding and management practice and gain a deeper understanding of crop traits.